

```
%-- 11/16/20, 12:10 AM --%
ft_read_cifti_mod✓
('/Users/dianaperez/Box/Quest_Backup/HCP_analyses/from_Ben/HCP_variants/spCorr/100206_all
Runs_mean_vs_120_avg_corr_LR_corr.dtseries.nii')
help MRIread
%-- 11/16/20, 1:04 AM --%
ft_read_cifti_mod✓
('/Users/dianaperez/Box/DATA/HCP/CIFTI/bold_data/100206_rfMRI_REST1_LR_surf_subcort_norma
lwall.dtseries.nii')
unique(ans.brainstructure)
cifti = ft_read_cifti_mod✓
('/Users/dianaperez/Box/DATA/HCP/CIFTI/bold_data/100206_rfMRI_REST1_LR_surf_subcort_norma
lwall.dtseries.nii')
ft_read_cifti_mod✓
('/Users/dianaperez/Box/Quest_Backup/HCP_analyses/from_Ben/HCP_variants/spCorr/100206_all
Runs_mean_vs_120_avg_corr_LR_corr.dtseries.nii')
unique(ans.brainstructure)
%-- 11/16/20, 1:33 AM --%
help MRIread
ft_read_cifti_mod✓
('/Users/dianaperez/Box/Quest_Backup/HCP_analyses/from_Ben/HCP_variants/spCorr/100206_all
Runs_mean_vs_120_avg_corr_LR_corr.dtseries.nii')
addpath '/Users/dianaperez/Box/Dependencies/cifti-matlab-master'
ft_read_cifti_mod✓
('/Users/dianaperez/Box/Quest_Backup/HCP_analyses/from_Ben/HCP_variants/spCorr/100206_all
Runs_mean_vs_120_avg_corr_LR_corr.dtseries.nii')
cifti = ans
unique(cifti.brainstructure)
lhem = cifti.data(cifti.brainstructure==1);
rhem = cifti.data(cifti.brainstructure==2);
find(cifti.brainstructure==2)
find(cifti.brainstructure==2)
rhem = cifti.data(cifti.brainstructure==2);
roi_data = load_untouch_nii_wrapper✓
('/Users/dianaperez/Box/Quest_Backup/Atlases/Seitzman300/Seitzman300_MNI_res02_allROIs.
nii.gz')
addpath '/Users/dianaperez/Box/Dependencies/cifti-matlab-master'
roi_data = load_untouch_nii_wrapper✓
('/Users/dianaperez/Box/Quest_Backup/Atlases/Seitzman300/Seitzman300_MNI_res02_allROIs.
nii.gz')
addpath '/Users/dianaperez/Box/Quest_Backup/Scripts'
roi_data = load_untouch_nii_wrapper✓
('/Users/dianaperez/Box/Quest_Backup/Atlases/Seitzman300/Seitzman300_MNI_res02_allROIs.
nii.gz')
cd '/Users/dianaperez/Documents/GitHub/GrattonLab-General-Repo/FCProcess'
roi_data = load_untouch_nii_wrapper✓
('/Users/dianaperez/Box/Quest_Backup/Atlases/Seitzman300/Seitzman300_MNI_res02_allROIs.
nii.gz')
unique(roi_data)
brain = cifti.brainstructure(cifti.brainstructure>0)
lhem = cifti.data(brain==1);
rhem = cifti.data(brain==2);
unique(lhem)
map = ft_read_cifti_mod✓
('/Users/dianaperez/Box/Research/lateralizationVariants/variantMaps/100206_ThresholdedVar
iantMap_SNRExclude_7.5.dtseries.nii')
brain = map.data(map.brainstructure>0)
brain = map.brainstructure(map.brainstructure>0)
lhem = map.data(brain==1);
unique(lhem)
```

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unique(map.data)
map = ft_read_cifti_mod✓
('/Users/dianaperez/Box/Research/lateralizationVariants/variantMaps/111514_ThresholdedVar✓
iantMap_SNRExclude_7.5.dtseries.nii')
unique(map.data)
map = ft_read_cifti_mod✓
('/Users/dianaperez/Box/Research/lateralizationVariants/variantMaps/temp/101006_RH_Varian✓
t_Size_50_TempVariantMap_7.5.dtseries.nii')
unique(map.data)
CreateVariantFiles
file = [dataLoc fileName num2str(subject)];
file = [dataLoc num2str(subject) fileName ];
CreateVariantFiles
workbenchdir = '/Users/dianaperez/Applications/workbench/bin_macosx64/';
CreateVariantFiles
map = ft_read_cifti_mod✓
('/Users/dianaperez/Box/Research/lateralizationVariants/variantMaps/100206_ThresholdedVar✓
iantMap_SNRExclude_7.5.dtseries.nii')
unique(map.data)
CreateVariantFiles
map = ft_read_cifti_mod✓
('/Users/dianaperez/Box/Research/lateralizationVariants/variantMaps/100206_ThresholdedVar✓
iantMap_SNRExclude_7.5.dtseries.nii')
unique(map.data)
CreateVariantFiles
unique(cifti.data)
CreateVariantFiles
size(vars_sizes,1)
CreateVariantFiles
map = ft_read_cifti_mod✓
('/Users/dianaperez/Box/Research/lateralizationVariants/variantMaps/100206_ThresholdedVar✓
iantMap_SNRExclude_7.5.dtseries.nii')
unique(map.data)
CreateVariantFiles
clear all
map = ft_read_cifti_mod✓
('/Users/dianaperez/Box/Research/lateralizationVariants/variantMaps/100206_ThresholdedVar✓
iantMap_SNRExclude_7.5.dtseries.nii')
brain = map.brainstructure(map.brainstructure>0);
lhem = map.data(brain==1);
unique(lhem)
rhem = map.data(brain==2);
unique(rhem)
unique(L_hem>0)
unique(lhem>0)
unique(lhem)
unique(lhem)>0
sum(unique(lhem)>0)
L_hem_varIDs = unique(L_hem);
L_hem_varIDs = unique(lhem);
varcount = {LHvars, RHvars, MHvars};
varcount = {}
varcount{1} = {}
varcount{2} = 4
varcount{5} = 4
varcount = {}
varcount{3,1} = 1
varcount{3,2} = 2
varcount{3(1)} = 2
varcount{3[1]} = 2

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```
varcount{3{1}} = 2
countVarsxHem
file = [dataLoc num2str(subs(x)) fileName];
file = [dataLoc num2str(subs(1)) fileName];
countVarsxHem
subs{1,1}
subs{1,1,1}
countVarsxHem
files(n)
ft_read_cifti_mod(files(n));
countVarsxHem
subsxgroup(1)
subsxgroup{1}
countVarsxHem
groups{g}
countVarsxHem
clear all
load varcount.mat
LH = varcount{1,1};
RH = varcount{1,2};
MH = varcount{1,3};
x = LH(:,2);
y = LH(:,3);
b1 = x/y;
x\y
b1 = x\y;
yCalc1 = b1*x;
scatter(x,y)
hold on
plot(x,yCalc1)
xlabel('Handedness Score')
ylabel('Number of Left Hemisphere Variants')
title('Linear Regression Handedness & Number of Lhem Variants')
X = [ones(length(x),1) x];
b = X\y
yCalc2 = X*b;
plot(x,yCalc2,'--')
legend('Data','Slope','Slope & Intercept','Location','best');
Rsqr1 = 1 - sum((y - yCalc1).^2)/sum((y - mean(y)).^2)
Rsqr2 = 1 - sum((y - yCalc2).^2)/sum((y - mean(y)).^2)
LH_sorted = sort(LH(:,2),'descend');
[LH_sorted, I] = sort(LH(:,2),'descend');
LH_sorted2 = LH(I)
LH_sorted2 = LH(I,:);
LH = LH_sorted2;
x1 = LH(1:14,2);
x1 = LH(1:24,2);
x2 = LH(25:48,2);
y1 = LH(1:24,3);
y2 = LH(25:48,3);
scatter(x1,y1, 'filled')
clear yCalc1
clear plot
xLH = LH(:,2);
clear all
load('varcount.mat')
LH = varcount{1,1};
RH = varcount{1,2};
MH = varcount{1,3};
xLH = LH(:,2)
```

```
y1LH = LH(:,3);
y2LH = LH(:,4);
scatter(xLH,y1LH)
hold on
scatter(xLH, y2LH)
scatter(xLH, y2LH, 'filled')
X = [ones(length(xLH),1) xLH];
b1 = X\y
b1 = X\y1LH
yCalc1 = X*b1
plot(xLH, yCalc1, '--')
b1 = X\y2LH
yCalc2 = X*b1
plot(xLH, yCalc1)
countVarsxHem
count(Lhem>0)
help count
Lhem(Lhem>0)
Lhem>0
countVarsxHem
clear all
countVarsxHem
clear
load('varcount.mat')
load('vertcount.mat')
LHvars = varcount{1,1};
LHverts = vertcount{1,1};
LH = [LHvars LHverts(:,2:3)];
LH = [LHvars LHverts(:,3:4)];
x=LH(:,2);
y=LH(:,5);
scatter(x,y)
hold on
y2=LH(:,6);
scatter(x,y2,'filled')
X = [ones(length(x),1) x];
b = X\y;
b2 = X\y2;
yCalc = X*b;
yCalc2 = X*b2;
plot(x,yCalc)
hold on
plot(x,yCalc2, '--')
legend('Lhem','Rhem','Lhem slope', 'Rhem slope', 'Location', 'best')
xlabel('Handedness')
ylabel('Number of Variant Vertices')
corr(x,y)
corr(x,y2)
[r p] = corrcoef(x,y2)
[r p] = corrcoef(x,y)
help repmat
x=LH(:,2);
y=LH(:,3);
y2=LH(:,4);
scatter(x,y)
hold on
scatter(x,y2,'filled')
X = [ones(length(x),1) x];
b = X\y;
b2 = X\y2;
```

```
yCalc = X*b;
yCalc2 = X*b2;
plot(x,yCalc)
plot(x,yCalc2, '--')
legend('Lhem','Rhem','Lhem slope', 'Rhem slope', 'Location', 'best')
xlabel('Handedness')
ylabel('Number of Variant Clusters')
[r p] = corrcoef(x,y2)
[r p] = corrcoef(x,y)
RHverts = vertcount{1,2};
RHvars = varcount{1,2};
RH = [RHvars RHverts(:,3:4)];
handedness = RH(:,2);
lhemvars = RH(:,3);
rhemvars = RH(:,4);
lhemverts = RH(:,5);
rhemverts = RH(:,6);
scatter(handedness,lhemvars)
scatter(handedness,rhemvars, 'filled')
hold on
X = [ones(length(handedness),1) handedness];
b = X\lhemvars;
b2 = X\rhemvars;
yCalc = X*b;
yCalc2 = X*b2;
plot(x,yCalc)
plot(x,yCalc2, '--')
legend('Lhem','Rhem','Lhem slope', 'Rhem slope', 'Location', 'best')
xlabel('Handedness')
ylabel('Number of Variant Clusters')
[r_lhem p_lhem] = corrcoef(handedness,lhemvars)
[r_rhem p_rhem] = corrcoef(handedness,rhemvars)
RHverts = vertcount{1,2};
RHvars = varcount{1,2};
RH = [RHvars RHverts(:,3:4)];
clear all
load('varcount.mat')
load('vertcount.mat')
RHverts = vertcount{1,2};
RHvars = varcount{1,2};
RH = [RHvars RHverts(:,3:4)];
clear RHvars
clear RHverts
handedness = RH(:,2);
lhemvars = RH(:,3);
rhemvars = RH(:,4);
lhemverts = RH(:,5);
rhemverts = RH(:,6);
scatter(handedness,lhemvars)
scatter(handedness,rhemvars, 'filled')
scatter(handedness,lhemvars)
hold on
scatter(handedness,rhemvars, 'filled')
X = [ones(length(handedness),1) handedness];
b = X\lhemvars;
b2 = X\rhemvars;
yCalc = X*b;
yCalc2 = X*b2;
plot(x,yCalc)
plot(handedness,yCalc)
```

```
hold on
plot(handedness,yCalc2, '--')
legend('Lhem','Rhem','Lhem slope', 'Rhem slope', 'Location', 'best')
xlabel('Handedness')
ylabel('Number of Variant Clusters')
[r_lhem p_lhem] = corrcoef(handedness,lhemvars)
[r_rhem p_rhem] = corrcoef(handedness,rhemvars)
scatter(handedness,lhemverts)
hold on
scatter(handedness,rhemverts, 'filled')
X = [ones(length(handedness),1) handedness];
b = X\lhemverts;
b2 = X\rhemverts;
yCalc = X*b;
yCalc2 = X*b2;
plot(handedness,yCalc)
hold on
plot(handedness,yCalc2, '--')
legend('Lhem','Rhem','Lhem slope', 'Rhem slope', 'Location', 'best')
xlabel('Handedness')
ylabel('Number of Variant Vertices')
[r_lhem p_lhem] = corrcoef(handedness,lhemverts)
[r_rhem p_rhem] = corrcoef(handedness,rhemverts)
clear all
load('varcount.mat')
load('vertcount.mat')
%% Middle Handers
MHverts = vertcount{1,3};
MHvars = varcount{1,2};
MH = [MHvars MHverts(:,3:4)];
handedness = MH(:,2);
lhemvars = MH(:,3);
rhemsvars = MH(:,4);
lhemverts = MH(:,5);
rhemsverts = MH(:,6);
scatter(handedness,lhemvars)
hold on
scatter(handedness,rhemsvars, 'filled')
X = [ones(length(handedness),1) handedness];
b = X\lhemvars;
b2 = X\rhemsvars;
yCalc = X*b;
yCalc2 = X*b2;
plot(handedness,yCalc)
hold on
plot(handedness,yCalc2, '--')
legend('Lhem','Rhem','Lhem slope', 'Rhem slope', 'Location', 'best')
xlabel('Handedness')
ylabel('Number of Variant Clusters')
[r_lhem p_lhem] = corrcoef(handedness,lhemvars)
[r_rhem p_rhem] = corrcoef(handedness,rhemsvars)
MHverts = vertcount{1,3};
MHvars = varcount{1,2};
MH = [MHvars MHverts(:,3:4)];
MHverts = vertcount{1,3};
MHvars = varcount{1,3};
MH = [MHvars MHverts(:,3:4)];
handedness = MH(:,2);
lhemvars = MH(:,3);
rhemsvars = MH(:,4);
```

```
lhemverts = MH(:,5);
rhemverts = MH(:,6);
scatter(handedness,lhemvars)
hold on
scatter(handedness,rhemvars, 'filled')
X = [ones(length(handedness),1) handedness];
b = X\lhempvars;
b2 = X\rhemvars;
yCalc = X*b;
yCalc2 = X*b2;
plot(handedness,yCalc)
hold on
plot(handedness,yCalc2, '--')
legend('Lhem','Rhem','Lhem slope', 'Rhem slope', 'Location', 'best')
xlabel('Handedness')
ylabel('Number of Variant Clusters')
[r_lhem p_lhem] = corrcoef(handedness,lhemvars)
[r_rhem p_rhem] = corrcoef(handedness,rhemvars)
scatter(handedness,lhemverts)
hold on
scatter(handedness,rhemverts, 'filled')
X = [ones(length(handedness),1) handedness];
b = X\lhempverts;
b2 = X\rhemverts;
yCalc = X*b;
yCalc2 = X*b2;
plot(handedness,yCalc)
hold on
plot(handedness,yCalc2, '--')
legend('Lhem','Rhem','Lhem slope', 'Rhem slope', 'Location', 'best')
xlabel('Handedness')
ylabel('Number of Variant Vertices')
[r_lhem p_lhem] = corrcoef(handedness,lhemverts)
[r_rhem p_rhem] = corrcoef(handedness,rhemverts)
LHvars = varcount{1,1};
RHvars = varcount{1,2};
ALL_vars = [LHvars;RHvars;MHvars];
clear LHvars
clear RHvars
clear b
clear b2
clear handedness
clear lhempvars
clear lhempvars
clear lhempverts
clear MH
clear MHvars
clear MHverts
clear p_lhem
clear p_rhem
clear X
clear yCalc
clear yCalc2
clear r_lhem
clear r_rhem
clear rhempcars
clear rhempverts
clear rhempvars
LH = vertcount{1,1};
RH = vertcount{1,2};
```

```
MH = vertcount{1,3};
ALL_verts = [LH;RH;MH];
clear LH
clear RH
clear MH
ALL = [ALL_vars ALL_verts(:,3:4)];
clear ALL_vars
clear ALL_verts
handedness = ALL(:,2);
hand = handedness;
clear handedness
lhemvars = ALL(:,3);
rhemvars = ALL(:,4);
scatter(hand,lhemvars)
hold on
scatter(hand,rhemvars,'filled')
X = [ones(length(hand),1) hand];
b = X\lhemvars;
b2 = X\rhemvars;
yCalc = X*b;
yCalc2 = X*b2;
plot(hand,yCalc)
hold on
plot(hand,yCalc2, '--')
legend('Lhem','Rhem','Lhem slope', 'Rhem slope', 'Location', 'best')
xlabel('Handedness')
ylabel('Number of Variant Clusters')
[r_lhem p_lhem] = corrcoef(hand,lhemvars)
[r_rhem p_rhem] = corrcoef(hand,rhemvars)
handedness = hand;
clear hand
scatter(handedness,lhemverts)
hold on
lhemverts = ALL(:,5);
rhemverts = ALL(:,6);
scatter(handedness,lhemverts)
X = [ones(length(handedness),1) handedness];
b = X\lhemverts;
yCalc = X*b;
plot(handedness,yCalc)
scatter(handedness,lhemverts)
hold on
X = [ones(length(handedness),1) handedness];
b = X\lhemverts;
yCalc = X*b;
plot(handedness,yCalc)
hold on
[r_lhem p_lhem] = corrcoef(handedness,lhemverts)
scatter(handedness,rhemverts, 'filled')
b2 = X\rhemverts;
yCalc2 = X*b2;
hold on
plot(handedness,yCalc2, '--')
[r_rhem p_rhem] = corrcoef(handedness,rhemverts)
legend('Lhem','Rhem','Lhem slope', 'Rhem slope', 'Location', 'best')
xlabel('Handedness')
ylabel('Number of Variant Vertices')
legend('Lhem','Lhem slope','Rhem', 'Rhem slope', 'Location', 'best')
%-- 11/22/20, 2:01 PM --%
usrpath
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userpath
%-- 11/23/20, 12:26 PM --%
reliability
addpath(genpath('/Users/dianaperez/Box/Quest_Backup/Scripts'))
reliability
%-- 11/29/20, 6:19 PM --%
help pause
%-- 11/30/20, 12:06 AM --%
checkisfloat
%-- 11/30/20, 7:28 PM --%
overlapmap
%-- 12/8/20, 8:23 PM --%
uiopen('/Users/dianaperez/Desktop/100206_Testing_surface_to_metric_leftCTX.func.gii',1)
load('/Users/dianaperez/Desktop/100206_Testing_surface_to_metric_leftCTX.func.gii')
%-- 12/9/20, 8:12 PM --%
help niftiread
%-- 1/4/21, 10:37 PM --%
flip_cifti_ind
%-- 1/5/21, 12:01 PM --%
flip_cifti_ind
hems = cifti.brainstructure(cifti.brainstructure>0);
L_hem = data(hems==1);
L_hem = cifti.data(hems==1);
L_hem = cifti.data(hems==1);
R_hem = cifti.data(hems==2);
for l = 1:length(L_hem)
L_hem(l) == l;
end
for l = 1:length(L_hem)
L_hem(1,l) == l;
end
for l = 1:length(L_hem)
L_hem(l,1) == l;
end
L_hem(1,1) == 1
L_hem(1,1) = 1
for l = 1:length(L_hem)
L_hem(l,1) = l;
end
find(hems==1)
L_hem_ind = find(hem==1);
L_hem_ind = find(hems==1);
R_hem_ind = find(hems==2);
for l = 1:length(L_hem_ind)
cifti.data(l) = l;
end
for r = 1:length(R_hem_ind)
cifti.data(r) = r;
end
for r = 1:length(R_hem_ind)
cifti.data(r) = r;
end
for r = 1:length(R_hem_ind)
cifti.data(r) = R_hem_ind(r);
end
for r = 1:length(R_hem_ind); cifti.data(R_hem_ind(r)) = R_hem_ind(r); end
for l = 1:length(L_hem_ind)
cifti.data(L_hem_ind(l)) = L_hem_ind(l); end
for r = 1:length(R_hem_ind); cifti.data(R_hem_ind(r)) = r; end
ft_write_cifti_mod('/Users/dianaperez/Desktop/testing_symmetry.dtseries.nii', cifti)
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```
test_cifti = cifti;
test_point = 21200;
test_cifti.data = zeros(length(test_cifti.data));
test_cifti.data = zeros(size(test_cifti.data));
test_cifti.data(test_point) = 1;
test_cifti.data(R_hem_ind(test_point)) = 1;
R_hem_ind(test_point)
ft_write_cifti_mod('/Users/dianaperez/Desktop/testing_symm_of_point.dtseries.nii', ↵
test_cifti)
test_cifti.data = zeros(size(test_cifti.data));
test_point = 1558;
test_cifti.data(test_point) = 1;
test_cifti.data(R_hem_ind(test_point)) = 1;
ft_write_cifti_mod('/Users/dianaperez/Desktop/testing_symm_of_point_2.dtseries.nii', ↵
test_cifti)
test_point = 15578;
test_cifti.data(test_point) = 1;
test_cifti.data(R_hem_ind(test_point)) = 1;
ft_write_cifti_mod('/Users/dianaperez/Desktop/testing_symm_of_point_2.dtseries.nii', ↵
test_cifti)
R_hem_ind(test_point)
%-- 1/6/21, 4:00 PM --%
ft_write_cifti_mod('/Users/dianaperez/Desktop/testing_symmetry.dtseries.nii', cifti)
flip_cifti_ind
test_cifti = cifti;
test_point = 15578;
test_cifti.data = zeros(size(test_cifti.data));
test_cifti.data(test_point) = 1;
hems = cifti.brainstructure(cifti.brainstructure>0);
L_hem_ind = find(hem==1);
L_hem_ind = find(hems==1);
R_hem_ind = find(hems==2);
test_cifti.data(R_hem_ind(test_point)) = 1;
test_cifti.data(44942) = 2;
ft_write_cifti_mod('/Users/dianaperez/Desktop/testing_symmetry_new_script.dtseries.nii', ↵
test_cifti)
test_cifti.data(test_point) = 1;
ft_write_cifti_mod('/Users/dianaperez/Desktop/testing_symmetry_new_script.dtseries.nii', ↵
test_cifti)
%-- 1/10/21, 3:18 PM --%
filename = ↵
'/Users/dianaperez/Box/Research/lateralizationVariants/variantMaps/100206_ThresholdedVari ↵
antMap_SNRExclude_7.5.dtseries.nii';
cifti = ft_read_cifti_mod(filename)
data = zeros(1:length(cifti.data));
data = zeros(1:length(cifti.data));
data = zeros(size(cifti.data));
data(43342) = 1 ;
data(13869) = 1 ;
cifti.data = data;
ft_write_cifti_mod('/Users/dianaperez/Desktop/100206_flipped_variant_map.dtseries.nii', ↵
flip_cifti)
ft_write_cifti_mod('/Users/dianaperez/Desktop/test_homotopic_pair.dtseries.nii', cifti)
x1 = 52.08;
x2 = -11.79;
z1 = 23.13;
x2 = 50.17;
y2 = 8.72;
z2 = -1.73;
distance = sqrt(((x2-x1)^2)+((y2-y1)^2)+((z2-z1)^2))
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y1 = -11.79
distance = sqrt(((x2-x1)^2)+((y2-y1)^2)+((z2-z1)^2))
x2 = 51.48;
y2 = -10.39
z2 = 23.76
distance = sqrt(((x2-x1)^2)+((y2-y1)^2)+((z2-z1)^2))
x2 = 59.26
y2 = -10.39
z2 = 23.76
distance = sqrt(((x2-x1)^2)+((y2-y1)^2)+((z2-z1)^2))
%-- 1/12/21, 12:41 PM --%
homotopic_overlap
addpath(genpath('/Users/dianaperez/Documents/GitHub/Lateralization_Variants'))
homotopic_overlap
brain = cifti.brainstructure(cifti.brainstructure>0);
hems = brain; clear brain
left_hem = cifti.data(hems==1);
right_hem = cifti.data(hems==2);
left_right_hems = [left_hem right_hem];
left_right_hems = [left_hem right_hem(1:length(left_hem))];
left_right_hems = [left_hem right_hem(1:length(left_hem))];
for i = 1:length(left_right_hems)
if left_right_hems(1) == 1 && left_right_hems(2) == 1
end
if left_right_hems(i,1) == 1 && left_right_hems(i,2) == 1
left_right_hems(i,3) == 1;
else
left_right_hems(i,3) == 0;
end
end
ht_variants = [];
for i = 1:length(left_right_hems)
if left_right_hems(i,1) == 1 && left_right_hems(i,2) == 1
ht_variants = [];
else
end
end
for i = 1:length(left_right_hems)
if left_right_hems(i,1) == 1 && left_right_hems(i,2) == 1
ht_variants = 1;
else
ht_variants = 0;
end
end
for i = 1:length(left_right_hems)
if left_right_hems(i,1) == 1 && left_right_hems(i,2) == 1
ht_variants(i) = 1;
else
ht_variants(i) = 0;
end
end
ht_variants = ht_variants';
sum(ht_variants)
homotopic_overlap
L = template.brainstructure(1:32492);
R = template.brainstructure(32493:end);
unique(L)
unique(R)
for v = 1:64984
if cifti.brainstructure(v) == -1

```

```
ht_variants = 0;
ht_variants(v) = 0;
end
end
for v = 1:64984
if cifti.brainstructure(v) == -1
ht_variants(v) = 0;
elseif cifti.brainstructure(v) == 1
ht_variants(v) =
end
end
end
homotopic_overlap
ht_variants = ht_variants';
unique(ht_variants)
left_hem = ht_variants(1:32492);
right_hem = ht_variants(32492:end);
homotopic_overlap
unique(left_hem)
unique(right_hem)
sum(ht_variants)
sum(allsubs)
homotopic_overlap
groupmap2 = groupmap;
groupmap = [groupmap; groupmap2];
homotopic_overlap
for v = 1:64984
end
ind = [];
for i = 1:32492
ind(i,1) = i
end
for i = 1:32492
ind(i,1) = i;
end
ind = [ind;ind];
ind(35000)
template2 = template;
template2.data = ind;
ft_write_cifti_mod('/Users/dianaperez/Desktop/vertex_equals_index_inc_noncortical.
dtseries.nii', template2)
cif = ind(template.brainstructure>0);
template2.data = cif;
ft_write_cifti_mod('/Users/dianaperez/Desktop/vertex_equals_index_inc_noncortical.
dtseries.nii', template2)
load('brain_ind.mat')
brain_ind(15578)
find(cif==17288)
cif2 = zeros(size(cif));
cif2(15578) = 1;
cif2(45292) = 1;
template2.data = cif2;
ft_write_cifti_mod('/Users/dianaperez/Desktop/testing_symmetry_points_11578_45292.
dtseries.nii', template2)
ft_write_cifti_mod('testing_symmetry_new_script.dtseries.nii')
ft_read_cifti_mod('testing_symmetry_new_script.dtseries.nii')
cifti = ans;
find(cifti.data==1)
find(cif2.data==1)
find(cif2==1)
```

```
cif2(45274) =1;
cif2(54306) = 3;
cif2(45274) =2;
cif2(45274) = 3;
cif2(15578) = 1;
cif2(45292) = 2;
cif2(54306) = 4;
find(cif2>0)
template2.data = cif2;
ft_write_cifti_mod('/Users/dianaperez/Desktop/testing_symmetry_three_methods.dtseries.
nii', template2)
find(brain_ind==26389)
find(brain_ind==49731)
cif2(54306) = 0;
cif2(45243) = 4;
ft_write_cifti_mod('/Users/dianaperez/Desktop/testing_symmetry_three_methods.dtseries.
nii', template2)
template2.data = cif2;
ft_write_cifti_mod('/Users/dianaperez/Desktop/testing_symmetry_three_methods.dtseries.
nii', template2)
cif2(45292) = 0;
cif2(45274) = 0;
cif2(45243) = 1;
template2.data = cif2;
find(cif2>0)
ft_write_cifti_mod('/Users/dianaperez/Desktop/testing_symmetry_15578_45243.dtseries.nii',
template2)
ind = [];
for i = 1:32492
ind(i,1) = i;
end
cif = [ind;ind];
cif = cif(template.brainstructure>0);
homotopic_test_points
clear all
homotopic_test_points
brain_indices = brain_indices';
cifti_indices(21200)
64984/2
23001 + 32492
Left_hem = brain_indices(1:32492);
Right_hem = brain_indices(32493:end);
Right_hem(23001)
find(cifti_indices==55493)
homotopic_test_points
unique(cifti2.data)
cifti2.data(LeftCortex) = 1;
cifti2.data(RightCortex) = 1;
homotopic_test_points
left_ctx = cifti.brainstructure(1:32492);
right_ctx = cifti.brainstructure(32493:end);
for i = 1:length(left_ctx)
if left_ctx(i) > 0
left_ctx(i) = Left_hem(i);
end
end
left_cortex = left_ctx>0;
left_cortex = left_ctx(left_ctx>0);
for i = 1:length(right_ctx)
if right_ctx(i) > 0
```

```
right_ctx(i) = Right_hem(i);
end
end
right_cortex = right_ctx(right_ctx>0);
clear ans
clear count
clear i
clear LeftCortex
clear new_cifti
clear pair
clear RightCortex
clear subject
clear target_point
clear test_point
clear threshold
find(left_cortex==15578)
find(right_cortex==15578)
right_ctx(15578)
Right_hem(15578)
cifti.brainstructure(15578+32492)
right_ctx(15578)
find(right_cortex==right_ctx(15578))
LeftCortex = 13868;
right_cortex(13868)
RightCortex = 48070;
cifti2.data = zeros(size(cifti2.data));
cifti2.data(LeftCortex) = 1;
cifti2.data(RightCortex) = 1;
outfile = [outfile_dir 'testing_homotopic_pairs_' num2str(LeftCortex) '_' num2str(RightCortex) '.dtseries.nii'];
ft_write_cifti_mod(outfile, cifti2);
clear all
homotopic_test_points
clear dataLoc
clear filename
clear subject
clear threshold
clear test_point
x = find(cifti_indices == 69)
ind = cifti.brainstructure;
for i = 1:length(ind)
if ind(i) > 0
ind(i) = find(cifti.brainstructure == i );
end
end
for i = 1:length(ind)
if ind(i) > 0
ind(i) = find(cifti.brainstructure == ind(i));
end
end
ind(i) = find(cifti_indices == ind(i));
for i = 1:length(ind)
if ind(i) > 0
ind(i) = find(cifti_indices == ind(i));
end
end
x = find(cifti_indices == 69)
for i = 1:length(ind)
if ind(i) > 0
ind(i) = find(cifti_indices == i);
```

```

end
end
LH = ind(1:(64984/2));
RH = ind((64984/2)+1:end);
clear ind
ind = [LH RH];
ind(15578)
ind(15578,:)
find(ind == 15578)
ind(17288)
ind(17288,:)
i = find(ind == 15578)
homotopic_test_points
%-- 1/18/21, 3:27 PM --%
help niftiread
cd /Volumes/GRATTONLAB/Lifespan/BIDS/Nifti
niftiinfo('sub-LS02_ses-1_task-rest_run-1_space-MNI152NLin6Asym_res-2_desc-preproc_bold.
nii.gz')
%-- 1/19/21, 10:07 AM --%
diff_map_homotopic_variants
new_cifti = new_cifti';
find(right_hem>0)
unique(right_hem)
diff_map_homotopic_variants
homotopic_overlap
outfile = '/Users/dianaperez/Desktop/HCP_allSubs_diffMap_394GoodSubs_FlippedHems_thresh7.5.dtseries.nii'
cifti1 = HCP_allSubs_OverlapMap_384GoodSubs_threshold_7.5.dtseries.nii
cifti1 = 'HCP_allSubs_OverlapMap_384GoodSubs_threshold_7.5.dtseries.nii'
cifti2 = 'HCP_allSubs_FlippedOverlapMap_384GoodSubs_threshold_7.5.dtseries.nii'
makeDiffMap(cifti1, cifti2, outfile)
cifti1 = '/Users/dianaperez/Desktop/HCP_allSubs_OverlapMap_384GoodSubs_threshold_7.5.
dtseries.nii'
cifti2 =
'/Users/dianaperez/Desktop/HCP_allSubs_FlippedOverlapMap_384GoodSubs_threshold_7.5.
dtseries.nii'
makeDiffMap(cifti1, cifti2, outfile)
cifti2 = '/Users/dianaperez/Desktop/HCP_LH_FlippedOverlapMap_384GoodSubs_threshold_7.5.
dtseries.nii'
cifti1 = '/Users/dianaperez/Desktop/HCP_LH_OverlapMap_384GoodSubs_threshold_7.5.dtseries.
nii'
outfile = '/Users/dianaperez/Desktop/HCP_LH_diffMap_394GoodSubs_FlippedHems_thresh7.5.
dtseries.nii'
makeDiffMap(cifti1, cifti2, outfile)
cifti2 = '/Users/dianaperez/Desktop/HCP_MH_FlippedOverlapMap_384GoodSubs_threshold_7.5.
dtseries.nii'
cifti1 = '/Users/dianaperez/Desktop/HCP_MH_OverlapMap_384GoodSubs_threshold_7.5.dtseries.
nii'
outfile = '/Users/dianaperez/Desktop/HCP_MH_diffMap_394GoodSubs_FlippedHems_thresh7.5.
dtseries.nii'
makeDiffMap(cifti1, cifti2, outfile)
cifti2 = '/Users/dianaperez/Desktop/HCP_RH_FlippedOverlapMap_384GoodSubs_threshold_7.5.
dtseries.nii'
cifti1 = '/Users/dianaperez/Desktop/HCP_RH_OverlapMap_384GoodSubs_threshold_7.5.dtseries.
nii'
outfile = '/Users/dianaperez/Desktop/HCP_RH_diffMap_394GoodSubs_FlippedHems_thresh7.5.
dtseries.nii'
makeDiffMap(cifti1, cifti2, outfile)
%-- 1/19/21, 4:16 PM --%
load('indices.mat')

```

```
%-- 1/20/21, 12:24 PM --%
permute_diffMaps_flippedVars
clear all
flip_switch = zeros(384,1);
flip_switch(1:length(flip_switch)/2) = 1;
permute_diffMaps_flippedVars
help randperm
ind = randperm(size(flip_switch));
ind = randperm(length(flip_switch));
ind = randperm(length(flip_switch),1);
rand_flip_switch = flip_switch(ind');
help rand
permute_diffMaps_flippedVars
disp(['Permutation #' num2str(p) ': The maximum number of subjects that overlap in the
flipped left hemisphere is ' num2str(max(flip_left_sum))])
permute_diffMaps_flippedVars
permute_diffMaps
rand_files(s,1) = allSubs_files(ind(s));
allSubs_files(294)
rand_files={}
rand_files{s} = allSubs_files(ind(s));
rand_files{2} = allSubs_files(ind(49));
rand_files{2,1} = allSubs_files(ind(49));
clear all
permute_diffMaps
cifti = ft_read_cifti_mod(files(x));
files(x)
files{x}
cifti = ft_read_cifti_mod(files(x));
cifti = ft_read_cifti_mod(files{x});
permute_diffMaps
pseudo_RH_files = rand_files(end-length(RH_files):end);
pseudo_RH_files = rand_files(end-length(RH_files)+1:end);
pseudo_RH_files = rand_files{end-length(RH_files)+1:end};
pseudo_RH_files = rand_files(end-length(RH_files)+1:end);
permute_diffMaps
allSubs_files{2}
allSubs_files(2)
allSubs_files{ind(33)}
permute_diffMaps
files(1)
files{1}
ifti = ft_read_cifti_mod(files{x});
permute_diffMaps
%-- 1/23/21, 11:06 AM --%
overlapmap
outfile = ['/Users/dianaperez/Desktop/HCP_' groups{g}
'_OverlapMap_752GoodSubs_matchedGroups_threshold_' num2str(threshold(t)) '.dtseries.
nii'];
overlapmap
outfile = ['/Users/dianaperez/Desktop/HCP_brainMask.dtseries.nii'];
[overlap_map] = makemap('allSubs', allSubs_files, template);
mask = [];
for v = 1:length(overlap_map)
end
for v = 1:length(overlap_map.data)
if overlap_map.data > 0
end
if overlap_map.data(v) > 0
brain_mask(v) = 1;
```



```

else
brain_mask(v) = 0;
end
end
sum(overlap_map.data>0)
sum(brain_mask)
brainmask = brainmask';
brainmask = brain_mask';
ft_write_cifti_mod(outfile, brainmask);
template.data = brainmask;
ft_write_cifti_mod(outfile, template);
sum(template.brainstructure==1)
permute_diffMaps_flippedVars
left_masked = left(mask==1,:);
flip_left_masked = flip_left(mask==1,:);
left_sum = sum(left_masked,2);
disp(['Permutation #' num2str(p) ': The maximum number of subjects that overlap in the
left hemisphere is ' num2str(max(left_sum))])
flip_left_sum = sum(flip_left_masked,2);
disp(['Permutation #' num2str(p) ': The maximum number of subjects that overlap in the
flipped left hemisphere is ' num2str(max(flip_left_sum))])
overlap_left = sum(left,2)/length(files);
disp(['Permutation #' num2str(p) ': The maximum proportion of subjects that overlap in
the left hemisphere is ' num2str(max(overlap_left))])
overlap_flip_left = sum(flip_left,2)/length(files);
disp(['Permutation #' num2str(p) ': The maximum proportion of subjects that overlap in
the flipped left hemisphere is ' num2str(max(overlap_flip_left))])
spCorrs(p,t) = corr(overlap_left, overlap_flip_left);
permute_diffMaps_flippedVars
make_SNR_mask
read_4dfpimg_HCP_good
make_SNR_mask
etype = 'bigendian';
MeanSNR = mean(inputdata,2);
make_SNR_mask
find(MeanSNR==Nan)
find(MeanSNR==nan)
find(MeanSNR==NaN)
overlapmap
%-- 1/24/21, 9:35 PM --%
permute_diffMaps_flippedVars
clear all
load
( '/Users/dianaperez/Box/Research/Lateralization_Variants/Permutation_Tests/VariantsvsFlip
pedVariants_spCorr_300permutations_5.mat')
dist = spCorrs;
load
( '/Users/dianaperez/Box/Research/Lateralization_Variants/Permutation_Tests/VariantsvsFlip
pedVariants_spCorr_300permutations_4.mat')
dist = [dist; spCorrs];
load
( '/Users/dianaperez/Box/Research/Lateralization_Variants/Permutation_Tests/VariantsvsFlip
pedVariants_spCorr_100-200permutations_2.mat')
dist = [dist; spCorrs];
load
( '/Users/dianaperez/Box/Research/Lateralization_Variants/Permutation_Tests/VariantsvsFlip
pedVariants_spCorr_1-100permutations.mat')
dist = [dist; spCorrs];
min(dist)
max(dist)

```

```
%-- 1/27/21, 10:00 AM --%
permute_diffMaps_flippedVars
clear all
load
(' /Users/dianaperez/Box/Research/Lateralization_Variants/Permutation_Tests/VariantsvsFlip
pedVariants_spCorr_300permutations_5.mat')
dist = spCorrs;
load
(' /Users/dianaperez/Box/Research/Lateralization_Variants/Permutation_Tests/VariantsvsFlip
pedVariants_spCorr_300permutations_4.mat')
dist = [dist; spCorrs];
load
(' /Users/dianaperez/Box/Research/Lateralization_Variants/Permutation_Tests/VariantsvsFlip
pedVariants_spCorr_200permutations_3.mat')
dist = [dist; spCorrs];
load
(' /Users/dianaperez/Box/Research/Lateralization_Variants/Permutation_Tests/VariantsvsFlip
pedVariants_spCorr_1-100permutations.mat')
dist = [dist; spCorrs];
load
(' /Users/dianaperez/Box/Research/Lateralization_Variants/Permutation_Tests/VariantsvsFlip
pedVariants_spCorr_100-200permutations_2.mat')
dist = [dist; spCorrs];
min(dist)
max(dist)
plot(dist)
help plot
plot(dist, 1:1000)
plot(dist, 'circle')
plot(dist, 'o')
pseudo_maps = dist;
clear dist
load
(' /Users/dianaperez/Box/Research/Lateralization_Variants/Permutation_Tests/VariantsvsFlip
pedVariants_spCorr_1permutations.mat')
true_map = spCorrs;
clear spCorrs
hold on
plot(true_map, 'o')
hold on
plot(pseudo_maps, 'o')
hold on
plot(true_map, 500, 'o')
plot(500, true_map, 'o')
plot(pseudo_maps, 'o')
hold on
plot(500, true_map, 'o')
p = plot(500, true_map, 'o', pseudo_maps, 'o')
p = plot(500, true_map, 'o', 1:1000, pseudo_maps, 'o')
clear p
p = plot(500, true_map, 'o', 1:1000, pseudo_maps, 'o')
p = plot(500, true_map, 'o', 'filled', 1:1000, pseudo_maps, 'o', 'filled')
p = plot(500, true_map, 'ro', 1:1000, pseudo_maps, 'ro')
p = plot(500, true_map, 'ro', 'MarkerFaceColor', 'r', 1:1000, pseudo_maps, 'bo',
'MarkerFaceColor', 'b')
help plot
plot(500, true_map, 'ro', 'MarkerFaceColor', 'r')
hold on
plot(1:1000, pseudo_maps, 'bo', 'MarkerFaceColor', 'b')
plot(1:1000, pseudo_maps, 'bo')
```

```
hold on
plot(500, true_map, 'ro', 'MarkerFaceColor', 'r')
%-- 1/27/21, 1:16 PM --%
overlapmap
g = {'allSubs'};
datafiles = {allSubs_files};
outfile =
'/Users/dianaperez/Box/Research/Lateralization_Variants/OverlapMaps/OverlapMaps-
Jan19_2021/HCP_allSubs_diffMap_394GoodSubs_FlippedHems_thresh7.5.dtseries.nii';
overlapmap
for y = 1:nfiles
cifti = ft_read_cifti_mod(files{x});
new_cifti = [];
count = 1;
for i = 1:length(cifti.brainstructure)
if cifti.brainstructure(i) > 0
if cifti.data(count) > 0
new_cifti(i,1) = 1;
else new_cifti(i,1) = 0;
end
count = count + 1;
else
new_cifti(i,1) = cifti.brainstructure(i);
end
end
left_hem = new_cifti(1:(length(new_cifti)/2));
right_hem = new_cifti((length(new_cifti)/2)+1:end);
flipped_hems = [right_hem; left_hem];
flipped_cifti = flipped_hems(cifti.brainstructure>0);
for q = 1:length(flipped_cifti)
if flipped_cifti(q) > 0
allSubs(q,y) = 1;
elseif flipped_cifti(q) == 0
allSubs(q,y) = 0;
end
end
end
flipped_groupsum = sum(allsubs,2);
disp(['The maximum number of subjects that overlap in the ' group ' group is ' num2str
(max(groupsum))])
flipped_groupmap = sum(allsubs,2)/nfiles;
disp(['The maximum proportion of subjects that overlap in the ' group ' group is '
num2str(max(groupmap))])
diff = groupmap - flipped_groupmap;
template.data = diff
outfile =
'/Users/dianaperez/Box/Research/Lateralization_Variants/OverlapMaps/OverlapMaps-
Jan19_2021/HCP_allSubs_diffMap_394GoodSubs_FlippedHems_thresh7.5.dtseries.nii';
ft_write_cifti_mod(outfile, template)
dir = '/Users/dianaperez/Box/Research/Lateralization_Variants/OverlapMaps/OverlapMaps-
Jan19_2021/';
makeDiffMap([dir 'HCP_allSubs_OverlapMap_384GoodSubs_threshold_7.5.dtseries.nii'], [dir
'HCP_allSubs_FlippedOverlapMap_384GoodSubs_threshold_7.5.dtseries.nii'], [[dir
'HCP_allSubs_diffMap_384GoodSubs_FlippedHems_thresh7.5.dtseries.nii']]);
makeDiffMap([dir 'HCP_allSubs_OverlapMap_384GoodSubs_threshold_7.5.dtseries.nii'], [dir
'HCP_allSubs_FlippedOverlapMap_384GoodSubs_threshold_7.5.dtseries.nii'], [[dir
'HCP_allSubs_diffMap_384GoodSubs_FlippedHems_thresh7.5.dtseries.nii']]);
permute_diffMaps_flippedVars
permute_left_right_handlers
clear all
```

```
load
('/Users/dianaperez/Box/Research/Lateralization_Variants/Permutation_Tests/VariantsvsFlippedVariants_LH_spCorr_1_thresh_7.5_permutations.mat')
true_map = spCorrs;
load
('/Users/dianaperez/Box/Research/Lateralization_Variants/Permutation_Tests/VariantsvsFlippedVariants_LH_spCorr_1000_thresh_7.5_permutations.mat')
pseudo_maps = spCorrs;
plot(1:1000, pseudo_maps, 'bo', 'MarkerFaceColor', 'b')
hold on
plot(500, true_map, 'ro', 'MarkerFaceColor', 'r')
permute_left_right_handers
clear all
load
('/Users/dianaperez/Box/Research/Lateralization_Variants/Permutation_Tests/VariantsvsFlippedVariants_MH_spCorr_1_thresh_7.5_permutations.mat')
true_map = spCorrs;
load
('/Users/dianaperez/Box/Research/Lateralization_Variants/Permutation_Tests/VariantsvsFlippedVariants_MH_spCorr_1000_thresh_7.5_permutations.mat')
pseudo_maps = spCorrs;
plot(1:1000, pseudo_maps, 'bo')
hold on
plot(500, true_map, 'ro', 'MarkerFaceColor', 'r')
ylim([0 1]);
ylim([0.5 1]);
ylabel('r value');
xlabel('Permutations');
m = findobj(gca, 'Type', 'marker');
m = findobj(gca);
m = findobj(gca, 'Type', 'line');
hleg1 = legend(m(1:2), 'randomized maps', 'true map', 'Location', 'SouthWest');
hleg1 = legend(m(1:2), 'true map', 'randomized maps', 'Location', 'SouthWest');
hleg1.FontSize = 14;
ax = gca;
ax.FontSize = 14;
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.5, 0.7]);
print(gcf, [outputDir 'VariantsvsFlippedVariants_' groups{g} '_spCorr_' num2str(threshold
(t)) '_thresh_' num2str(numperms) '_permutations_results.jpg'], '-dpng', '-r300');
rootDir = '/Users/dianaperez/Box/Research/Lateralization_Variants/';
outputDir = [rootDir 'Permutation_Tests/'];
print(gcf, [outputDir 'VariantsvsFlippedVariants_' groups{g} '_spCorr_' num2str(threshold
(t)) '_thresh_' num2str(numperms) '_permutations_results.jpg'], '-dpng', '-r300');
print(gcf, [outputDir 'VariantsvsFlippedVariants_MH_spCorr_7.5_
5_thresh_1000_permutations_results.jpg'], '-dpng', '-r300');
permute_left_right_handers
figure;
plot(1:numperms, spCorrs, 'bo')
ylim([.4 1]);
hold on
plot(round(numperms/2), true_spCorr, 'ro', 'MarkerFaceColor', 'r')
ylabel('r value');
xlabel('Permutations');
title(['Left Hem vs Pseudo Left Hem Variant Maps - ' groups{g}]);
m = findobj(gca, 'Type', 'line');
hleg1 = legend(m(1:2), 'randomized maps', 'true map', 'Location', 'SouthWest');
hleg1.FontSize = 14;
ax = gca;
ax.FontSize = 14;
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.5, 0.7]);
```

```

print(gcf,[outputDir 'VariantsvsFlippedVariants_' groups{g} '_spCorr_' num2str(threshold<
(t)) '_thresh_' num2str(numperms) '_permutations_results.jpg'],'-dpng','-r300');
close gcf
permute_left_right_handlers
load<
('/Users/dianaperez/Box/Research/Lateralization_Variants/Permutation_Tests/VariantsvsFlip<
pedVariants_RH_spCorr_1000_thresh_7.5_permutations.mat')
mean(spCorrs)
load<
('/Users/dianaperez/Box/Research/Lateralization_Variants/Permutation_Tests/VariantsvsFlip<
pedVariants_LH_spCorr_1000_thresh_7.5_permutations.mat')
mean(spCorrs)
load<
('/Users/dianaperez/Box/Research/Lateralization_Variants/Permutation_Tests/VariantsvsFlip<
pedVariants_MH_spCorr_1000_thresh_7.5_permutations.mat')
mean(spCorrs)
load<
('/Users/dianaperez/Box/Research/Lateralization_Variants/Permutation_Tests/VariantsvsFlip<
pedVariants_randomizedMaps_spCorr_1000permutations.mat')
mean(spCorrs)
load<
('/Users/dianaperez/Box/Research/Lateralization_Variants/Permutation_Tests/VariantsvsFlip<
pedVariants_randomizedMaps_spCorr_1000permutations.mat')
clear all
load<
('/Users/dianaperez/Box/Research/Lateralization_Variants/Permutation_Tests/VariantsvsFlip<
pedVariants_randomizedMaps_spCorr_1000permutations.mat')
mean(pseudo_maps)
%-- 1/28/21, 5:10 PM --%
permute_left_right_handlers
unique(pseudoLH)
unique(pseudoRH);
permute_left_right_handlers
%-- 2/3/21, 10:43 AM --%
permute_flippedVars
help niftiwrite
permutations_cluster_correction
load('/Users/dianaperez/Box/Quest_Backup/Scripts/CIFTI_RELATED/Resources/Conte69_atlas-<
v2.LR.32k_fs_LR.wb/Cifti_surf_neighbors_LR_normalwall.mat')
permutations_cluster_correction
clear all
permutations_cluster_correction
permute_flippedVars
permutations_cluster_correction
clear all
permutations_cluster_correction
unique(diffmat_bin)
permutations_cluster_correction
unique(diffmat_bin)
load<
('/Users/dianaperez/Box/Research/Lateralization_Variants/Permutation_Tests/PseudoRightvsP<
seudoLeftHanders_spCorr_1000_thresh_7.5_permutations_allRH.mat')
load<
('/Users/dianaperez/Box/Research/Lateralization_Variants/Permutation_Tests/RightvsLeftHan<
ders_true_spCorr_1000_thresh_7.5.mat')
p = spCorrs<true_spCorr;
count(p)
p = count(spCorrs<true_spCorr);
p = sum(spCorrs<true_spCorr);
p = 23/1000;

```

```

mean(spCorrs)
permute_left_right_handlers
load
('/Users/dianaperez/Box/Research/Lateralization_Variants/Permutation_Tests/RightvsLeftHanders_true_spCorr_1000_thresh_7.5.mat')
load
('/Users/dianaperez/Box/Research/Lateralization_Variants/Permutation_Tests/PseudoRightvsPseudoLeftHanders_spCorr_1000_thresh_7.5_permutations_allRH.mat')
p = count(spCorrs<true_spCorr);
p = sum(spCorrs<true_spCorr);
clear all
load
('/Users/dianaperez/Box/Research/Lateralization_Variants/Permutation_Tests/PseudoRightvsPseudoLeftHanders_spCorr_1000_thresh_7.5_permutations_allRH.mat')
load
('/Users/dianaperez/Box/Research/Lateralization_Variants/Permutation_Tests/RightvsLeftHanders_true_spCorr_1000_thresh_7.5.mat')
permute_left_right_handlers
p = sum(spCorrs<true_spCorr);
mean(spCorrs)
load
('/Users/dianaperez/Box/Research/Lateralization_Variants/Permutation_Tests/PseudoRightvsPseudoLeftHanders_spCorr_1000_thresh_7.5_permutations_allRH.mat')
mean(spCorrs)
p = sum(spCorrs<true_spCorr);
figure;
plot(1:numperms, spCorrs, 'bo')
ylim([.4 1]);
hold on
plot(round(numperms/2), true_spCorr, 'ro', 'MarkerFaceColor', 'r')
ylabel('r value');
xlabel('Permutations');
title('Left Handers vs Right Handers Variant Maps');
m = findobj(gca, 'Type', 'line');
hleg1 = legend(m(1:2), 'randomized maps', 'true map', 'Location', 'SouthWest');
hleg1.FontSize = 14;
ax = gca;
ax.FontSize = 14;
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.5, 0.7]);
load
('/Users/dianaperez/Box/Research/Lateralization_Variants/Permutation_Tests/PseudoRightvsPseudoLeftHanders_spCorr_1000_thresh_7.5_permutations.mat')
mean(spCorrs)
p = sum(spCorrs<true_spCorr);
390/1000
true_spCorr = .71
p = sum(spCorrs<true_spCorr);
p/1000
true_spCorr = .73
p = sum(spCorrs<true_spCorr);
p/1000
%-- 2/5/21, 9:34 PM --%
permute_flippedVars
mean(diffmats)
mean(diffmats);
mean(ans);
permutations_cluster_correction
mean(diffmats);
mean(ans);
permutations_cluster_correction

```

```

help makenifti
help writenifti
niftiwrite
help niftiwrite
outlier_voxels
filename = [outdir '/sub-' sub '_ses-' num2str(ses(s)) '_run-' num2str(r)
'_outlier_voxels.nii.gz'];
niftiwrite(new_nii, filename)
niftiinfo(filename)
info = niftiinfo(filename)
clear all
filename = [outdir '/sub-' sub '_ses-' num2str(ses(s)) '_run-' num2str(r)
'_outlier_voxels.nii.gz'];
outdir = '/Users/dianaperez/Desktop/Research/Lifespan/';
filename = [outdir '/sub-' sub '_ses-' num2str(ses(s)) '_run-' num2str(r)
'_outlier_voxels.nii.gz'];
sub = 'LS05';
filename = [outdir '/sub-' sub '_ses-1_run-1_outlier_voxels.nii.gz'];
info = niftiinfo(filename)
info = niftiread(filename)
%-- 2/8/21, 11:17 AM --%
addpath(genpath('/Users/dianaperez/Box/Quest_Backup/Scripts/'))
CreateVariantFiles_LS03specific
addpath(genpath('/Users/dianaperez/Box/Dependencies/cifti-matlab-master'))
CreateVariantFiles_LS03specific
[cifti_rest.data] = ExcludeVariantSize(cifti_rest.data, subject, threshold(x), 50);
[cifti_rest.data] = ExcludeVariantbySize(cifti_rest.data, subject, threshold(x), 50);
CreateVariantFiles_LS03specific
subsampling
CreateVariantFiles_LS03specific
map_1 = ft_read_cifti_mod
('/Users/dianaperez/Desktop/Research/Lifespan/LS03_variants_sizeExcluded_first-
half_thresh-5_smooth_2.55.dtseries.nii');
map_2 = ft_read_cifti_mod
('/Users/dianaperez/Desktop/Research/Lifespan/LS03_variants_sizeExcluded_second-
half_thresh-5_smooth_2.55.dtseries.nii');
spCorr = corr(map_1, map_2);
spCorr = corr(map_1.data, map_2.data);
map_1 = ft_read_cifti_mod
('/Users/dianaperez/Desktop/Research/Lifespan/LS03_variants_sizeExcluded_first-
half_thresh-10_smooth_2.55.dtseries.nii');
map_2 = ft_read_cifti_mod
('/Users/dianaperez/Desktop/Research/Lifespan/LS03_variants_sizeExcluded_second-
half_thresh-10_smooth_2.55.dtseries.nii');
spCorr = corr(map_1.data, map_2.data);
clear all
load
('/Users/dianaperez/Box/Research/Lateralization_Variants/Permutation_Tests/PseudoRightvsP
seudoLeftHanders_spCorr_1_thresh_7.5_permutations.mat')
load
('/Users/dianaperez/Box/Research/Lateralization_Variants/Permutation_Tests/PseudoRightvsP
seudoLeftHanders_spCorr_1000_thresh_7.5_permutations_allRH.mat')
mean(spCorrs)
load
('/Users/dianaperez/Box/Research/Lateralization_Variants/Permutation_Tests/PseudoRightvsP
seudoLeftHanders_spCorr_1_thresh_7.5_permutations.mat')
load
('/Users/dianaperez/Box/Research/Lateralization_Variants/Permutation_Tests/RightvsLeftHan
ders_true_spCorr_1_thresh_7.5.mat')
load

```

```
( '/Users/dianaperez/Box/Research/Lateralization_Variants/Permutation_Tests/PseudoRightvsPseudoLeftHanders_spCorr_1000_thresh_7.5_permutations_allRH.mat')
p = sum(spCorrs<true_spCorr)
p = p/length(spCorrs)
permute_left_right_handers
trueCorrs = true_spCorr;
permute_left_right_handers
load
( '/Users/dianaperez/Box/Research/Lateralization_Variants/Permutation_Tests/PseudoRightvsPseudoLeftHanders_spCorr_1000_thresh_7.5_permutations.mat')
mean(spCorrs)
p = sum(spCorrs<true_spCorr)
p = p/length(spCorrs)
true_spCorr = 0.73
p = sum(spCorrs<true_spCorr)
p = p/length(spCorrs)
true_spCorr = [.73, .75, .76];
figure;
plot(1:numperms, spCorrs, 'bo')
ylim([.4 1]);
hold on
plot(round(numperms/2), true_spCorr, 'ro', 'MarkerFaceColor', 'r')
plot([498, 500, 502], true_spCorr, 'ro', 'MarkerFaceColor', 'r')
figure;
plot(1:numperms, spCorrs, 'bo')
ylim([.4 1]);
hold on
plot([498, 500, 502], true_spCorr, 'ro', 'MarkerFaceColor', 'r')
plot([400, 500, 600], true_spCorr, 'ro', 'MarkerFaceColor', 'r')
figure;
plot(1:numperms, spCorrs, 'bo')
ylim([.4 1]);
hold on
plot([400, 500, 600], true_spCorr, 'ro', 'MarkerFaceColor', 'r')
ylabel('r value');
xlabel('Permutations');
title('Left Handers vs Right Handers Variant Maps');
m = findobj(gca, 'Type', 'line');
hleg1 = legend(m(1:2), 'true maps', 'randomized map', 'Location', 'SouthWest');
hleg1.FontSize = 14;
ax = gca;
ax.FontSize = 14;
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.5, 0.7]);
print(gcf,[outputDir 'RightvsLeftHanders_spCorr_' num2str(threshold(t)) '_thresh_'
num2str(numperms) '_permutations_results_all_RH.jpg'],'-dpng','-r300');
close gcf
clear all
map_1 = ft_read_cifti_mod
( '/Users/dianaperez/Desktop/Research/Lifespan/LS03_variants_sizeExcluded_first-half_thresh-10_smooth_2.55.dtseries.nii');
map_2 = ft_read_cifti_mod
( '/Users/dianaperez/Desktop/Research/Lifespan/LS03_variants_sizeExcluded_second-half_thresh-10_smooth_2.55.dtseries.nii');
diff=map_1.data - map_2.data;
ft_write_cifti_mod('/Users/dianaperez/Desktop/diff_variants_LS03_thresh_10.dtseries.nii',
diff)
template = map_1;
template.data = diff;
ft_write_cifti_mod('/Users/dianaperez/Desktop/diff_variants_LS03_thresh_10.dtseries.nii',
template)
```



```
map_1 = ft_read_cifti_mod(
('/Users/dianaperez/Desktop/Research/Lifespan/LS03_variants_sizeExcluded_first-
half_thresh-5_smooth_2.55.dtseries.nii');
map_2 = ft_read_cifti_mod(
('/Users/dianaperez/Desktop/Research/Lifespan/LS03_variants_sizeExcluded_second-
half_thresh-5_smooth_2.55.dtseries.nii');
diff=map_1.data - map_2.data;
template.data = diff;
ft_write_cifti_mod('/Users/dianaperez/Desktop/diff_variants_LS03_thresh_5.dtseries.nii',
template)
map_1 = ft_read_cifti_mod('/Users/dianaperez/Desktop/sub-LS03_spCorr_first-
half_try1_vs_120_allsubs_corr_cortex_corr.dtseries.nii');
map_2 = ft_read_cifti_mod('/Users/dianaperez/Desktop/sub-LS03_spCorr_second-
half_try1_vs_120_allsubs_corr_cortex_corr.dtseries.nii');
% correlation between the two
spCorr = corr(map_1.data, map_2.data);
%-- 2/17/21, 2:05 PM --%
corr_to_corr
clear LS02_1 LS02_2 LS02_3
clear LS03_1 LS03_2 LS03_3 LS03_4 LS03_5
clear LS05_1 LS05_2 LS05_3 LS05_4 LS05_5
clear LS07_1 LS07_2
corr_to_corr
for s = 1:4 %different subjects, avg over sessions
figure_corrmat_GrattonLab(squeeze(group_corrmat(s,:,:,:),atlas_params,-0.4,1);
title(['AllSessAvg, Subject ' subs{s}]);
colormap('jet');
saveas(gcf, [outdir 'Corrmat_' subs{s} '_sessmean.tiff'],'tiff');
end
close('all');
maskmat = ones(atlas_params.num_rois);
maskmat = logical(triu(maskmat,1));
%not averaged across sessions
tmp = corrmat_2(1,:,:,:);
corrln(1,:) = tmp(maskmat);
tmp = corrmat_2(2,:,:,:);
corrln(2,:) = tmp(maskmat);
tmp = corrmat_2(3,:,:,:);
corrln(3,:) = tmp(maskmat);
tmp = corrmat_3(1,:,:,:);
corrln(4,:) = tmp(maskmat);
tmp = corrmat_3(2,:,:,:);
corrln(5,:) = tmp(maskmat);
tmp = corrmat_3(3,:,:,:);
corrln(6,:) = tmp(maskmat);
tmp = corrmat_3(4,:,:,:);
corrln(7,:) = tmp(maskmat);
tmp = corrmat_3(5,:,:,:);
corrln(8,:) = tmp(maskmat);
% tmp = mean_LS04;
% corrln(9,:) = tmp(maskmat);
tmp = corrmat_5(1,:,:,:);
corrln(9,:) = tmp(maskmat);
tmp = corrmat_5(2,:,:,:);
corrln(10,:) = tmp(maskmat);
tmp = corrmat_5(3,:,:,:);
corrln(11,:) = tmp(maskmat);
tmp = corrmat_5(4,:,:,:);
corrln(12,:) = tmp(maskmat);
tmp = corrmat_5(5,:,:,:);
```

```
corrln(13,:) = tmp(maskmat);
tmp = corrmats_7(1,:,:,:);
corrln(14,:) = tmp(maskmat);
tmp = corrmats_7(2,:,:,:);
corrln(15,:) = tmp(maskmat);
% tmp = mean_LS10;
% corrln(17,:) = tmp(maskmat);
simmat = corr(corrln');
figure('Position',[1 1 1000 800]);
imagesc(simmat,[0 1]); colormap('jet');
hline_new([3,8,13,15]+0.5,'k',2);
vline_new([3,8,13,15]+0.5,'k',2);
set(gca,'XTick',[2,6,11,14.5], 'YTick', [2,6,11,14.5], 'XTickLabel',...
{'LS02', 'LS03', 'LS05', 'LS07'}, 'YTickLabel', {'LS02', 'LS03', 'LS05', 'LS07'});
axis square;
colorbar;
title('Correlation Matrix Similarity');
saveas(gcf,[outdir 'SimilarityMat_rest.tiff'],'tiff');
maskmat = ones(atlas_params.num_rois);
maskmat = logical(triu(maskmat,1));
count = 1;
for s = 1:6
tmp = mean_corrmats(s,:,:,:);
corrln(count,:) = tmp(maskmat);
count = count+1;
end
% end
simmat = corr(corrln');
figure('Position',[1 1 1000 800]);
imagesc(simmat,[0 1]); colormap('jet');
hline_new([3,8,9,14,16]+0.5,'k',2);
vline_new([3,8,9,14,16]+0.5,'k',2);
set(gca,'XTick',[2,6,9,12,15.5,17], 'YTick', [2,6,9,12,15.5,17], 'XTickLabel',...
{'LS02', 'LS03', 'LS04', 'LS05', 'LS07', 'LS10'}, 'YTickLabel', {'LS02', 'LS03', 'LS04',
'LS05', 'LS07', 'LS10'});
axis square;
colorbar;
title('Correlation Matrix Similarity');
saveas(gcf,[outdir 'SimilarityMat_averaged.tiff'],'tiff');
clear corrln
tmp = corrmats_2(1,:,:,:);
corrln(1,:) = tmp(maskmat);
tmp = corrmats_2(2,:,:,:);
corrln(2,:) = tmp(maskmat);
tmp = corrmats_2(3,:,:,:);
corrln(3,:) = tmp(maskmat);
simmat = corr(corrln');
figure('Position',[1 1 1000 800]);
imagesc(simmat,[0 1]); colormap('jet');
hline_new([3,8,13,15]+0.5,'k',2);
vline_new([3,8,13,15]+0.5,'k',2);
axis square;
colorbar;
clear corrln
tmp = corrmats_3(1,:,:,:);
corrln(1,:) = tmp(maskmat);
tmp = corrmats_3(2,:,:,:);
corrln(2,:) = tmp(maskmat);
tmp = corrmats_3(3,:,:,:);
corrln(3,:) = tmp(maskmat);
```

```
tmp = corrmat_3(4,:,:,:);
corrln(4,:) = tmp(maskmat);
tmp = corrmat_3(5,:,:,:);
corrln(5,:) = tmp(maskmat);
simmat = corr(corrln');
figure('Position',[1 1 1000 800]);
imagesc(simmat,[0 1]); colormap('jet');
hline_new([1,2,3,4,5]+0.5,'k',2);
vline_new([1,2,3,4,5]+0.5,'k',2);
set(gca, 'XTick', [1,2,3,4,5], 'YTick', [1:5], 'XTickLabel', {'ses-1', 'ses-2', 'ses-3', 'ses-4', 'ses-5'}, 'YTickLabel', {'ses-1', 'ses-2', 'ses-3', 'ses-4', 'ses-5'});
axis square;
colorbar;
title('Similarity Across Sessions for sub-LS03')
saveas(gcf, [outdir 'simmat_sub-LS03.tiff'], 'tiff');
tmp = corrmat_5(1,:,:,:);
corrln(1,:) = tmp(maskmat);
tmp = corrmat_5(2,:,:,:);
corrln(2,:) = tmp(maskmat);
tmp = corrmat_5(3,:,:,:);
corrln(3,:) = tmp(maskmat);
tmp = corrmat_5(4,:,:,:);
corrln(4,:) = tmp(maskmat);
tmp = corrmat_5(5,:,:,:);
corrln(5,:) = tmp(maskmat);
simmat = corr(corrln');
figure('Position',[1 1 1000 800]);
imagesc(simmat,[0 1]); colormap('jet');
hline_new([1,2,3,4,5]+0.5,'k',2);
vline_new([1,2,3,4,5]+0.5,'k',2);
set(gca, 'XTick', [1,2,3,4,5], 'YTick', [1:5], 'XTickLabel', {'ses-1', 'ses-2', 'ses-3', 'ses-4', 'ses-5'}, 'YTickLabel', {'ses-1', 'ses-2', 'ses-3', 'ses-4', 'ses-5'});
axis square;
colorbar;
title('Similarity Across Sessions for sub-LS05')
saveas(gcf, [outdir 'simmat_sub-LS05.tiff'], 'tiff');
mean_corr = mean(triu(simmat))
maskmat = logical(triu(simmat,1));
mean_corr = mean(simmat(maskmat))
tmp = corrmat_3(1,:,:,:);
corrln(1,:) = tmp(maskmat);
tmp = corrmat_3(2,:,:,:);
corrln(2,:) = tmp(maskmat);
tmp = corrmat_3(3,:,:,:);
corrln(3,:) = tmp(maskmat);
tmp = corrmat_3(4,:,:,:);
corrln(4,:) = tmp(maskmat);
tmp = corrmat_3(5,:,:,:);
corrln(5,:) = tmp(maskmat);
simmat = corr(corrln');
maskmat = ones(atlas_params.num_rois);
maskmat = logical(triu(maskmat,1));
tmp = corrmat_3(1,:,:,:);
corrln(1,:) = tmp(maskmat);
tmp = corrmat_3(2,:,:,:);
corrln(2,:) = tmp(maskmat);
tmp = corrmat_3(3,:,:,:);
corrln(3,:) = tmp(maskmat);
tmp = corrmat_3(4,:,:,:);
corrln(4,:) = tmp(maskmat);
```

```
tmp = corrmat_3(5,:,:,:);
corrln(5,:) = tmp(maskmat);
simmat = corr(corrln');
maskmat2 = logical(triu(simmat,1));
mean_corr = mean(simmat(maskmat2))
clear corrln
tmp = corrmat_2(1,:,:,:);
corrln(1,:) = tmp(maskmat);
tmp = corrmat_2(2,:,:,:);
corrln(2,:) = tmp(maskmat);
tmp = corrmat_2(3,:,:,:);
corrln(3,:) = tmp(maskmat);
simmat = corr(corrln');
maskmat2 = logical(triu(simmat,1));
mean_corr = mean(simmat(maskmat2))
tmp = corrmat_7(1,:,:,:);
clear corrln
tmp = corrmat_7(1,:,:,:);
corrln(1,:) = tmp(maskmat);
tmp = corrmat_2(2,:,:,:);
corrln(2,:) = tmp(maskmat);
simmat = corr(corrln');
maskmat2 = logical(triu(simmat,1));
mean_corr = mean(simmat(maskmat2))
tmp = corrmat_7(2,:,:,:);
corrln(2,:) = tmp(maskmat);
simmat = corr(corrln');
mean_corr = mean(simmat(maskmat2))
tmp = corrmat_2(1,:,:,:);
corrln(1,:) = tmp(maskmat);
tmp = corrmat_2(2,:,:,:);
corrln(2,:) = tmp(maskmat);
tmp = corrmat_2(3,:,:,:);
corrln(3,:) = tmp(maskmat);
tmp = corrmat_3(1,:,:,:);
corrln(4,:) = tmp(maskmat);
tmp = corrmat_3(2,:,:,:);
corrln(5,:) = tmp(maskmat);
tmp = corrmat_3(3,:,:,:);
corrln(6,:) = tmp(maskmat);
tmp = corrmat_3(4,:,:,:);
corrln(7,:) = tmp(maskmat);
tmp = corrmat_3(5,:,:,:);
corrln(8,:) = tmp(maskmat);
% tmp = mean_LS04;
% corrln(9,:) = tmp(maskmat);
tmp = corrmat_5(1,:,:,:);
corrln(9,:) = tmp(maskmat);
tmp = corrmat_5(2,:,:,:);
corrln(10,:) = tmp(maskmat);
tmp = corrmat_5(3,:,:,:);
corrln(11,:) = tmp(maskmat);
tmp = corrmat_5(4,:,:,:);
corrln(12,:) = tmp(maskmat);
tmp = corrmat_5(5,:,:,:);
corrln(13,:) = tmp(maskmat);
tmp = corrmat_7(1,:,:,:);
corrln(14,:) = tmp(maskmat);
tmp = corrmat_7(2,:,:,:);
corrln(15,:) = tmp(maskmat);
```

```
simmat = corr(corrln');
figure('Position',[1 1 1000 800]);
imagesc(simmat,[0 1]); colormap('jet');
help squeeze
clear corrln
corrln(1,:) = mean_LS02(maskmat);
corrln(2,:) = mean_LS03(maskmat);
corrln(3,:) = mean_LS05(maskmat);
corrln(4,:) = mean_LS07(maskmat);
simmat = corr(corrln');
figure('Position',[1 1 1000 800]);
imagesc(simmat,[0 1]); colormap('jet');
maskmat2 = logical(triu(simmat,1));
mean_corr = mean(simmat(maskmat2))
hline_new([0, 1,2,3,4]+0.5,'k',2);
vline_new([0, 1,2,3,4]+0.5,'k',2);
set(gca,'XTick',[1,2,3,4], 'YTick', [1,2,3,4], 'XTickLabel', {'LS02', 'LS03', 'LS05', 'LS07'}, 'YTickLabel', {'LS02', 'LS03', 'LS05', 'LS07'});
set(gca,'XTick',[1,2,3,4], 'YTick', [1,2,3,4], 'XTickLabel', {'LS02', 'LS03', 'LS05', 'LS07'}, 'YTickLabel', {'LS02', 'LS03', 'LS05', 'LS07'});
a = get(gca, 'XTickLabel');
set(gca,'XTickLabel',a,'fontsize',15,'FontWeight','bold')
set(gca,'XTickLabel',a,'fontsize',20,'FontWeight','bold')
axis square;
colorbar;
title('Correlation Matrix Similarity');
title('fontsize', 30)
title('Correlation Matrix Similarity');
saveas(gcf,[outDir 'SimilarityMat_averaged.tiff'],'tiff');
corr_to_corr
title('Similarity Across Subjects (r)');
clear corrln
for s = 1:numSubs
    corrln(s,:) = group_corrmat(s,maskmat);
end
clear corrln
if multi_ses_only
    corrln(1,:) = mean_LS02(maskmat);
    corrln(2,:) = mean_LS03(maskmat);
    corrln(3,:) = mean_LS05(maskmat);
    corrln(4,:) = mean_LS07(maskmat);
else
    corrln(1,:) = mean_LS02(maskmat);
    corrln(2,:) = mean_LS03(maskmat);
    corrln(3,:) = mean_LS04(maskmat);
    corrln(4,:) = mean_LS05(maskmat);
    corrln(5,:) = mean_LS07(maskmat);
    corrln(6,:) = mean_LS10(maskmat);
end
corr_to_corr
hline_new([0,3,8,13,15]+0.5,'k',2);
vline_new([0,3,8,13,15]+0.5,'k',2);
title('Across Subject Similarity');
corr_to_corr
sub_mats = [];
sub_mats = [sub_mats corrmat];
sub_mats = [];
sub_mats = [sub_mats corrmat];
sub_mats = [sub_mats; corrmat];
corr_to_corr
```

```
two_mean = sub_means;
sub_means = [];
sub_means_2(1,:,:,:) = two_mean;
sub_means_2(sub,:,:,:) = mean_corrmat;
sub_means = sub_means_2;
clear sub_means_2
clear two_mean
for sub = 1:numSubs
    for ses = 1:sessions(sub)
        tmp = sub_mats(count,:,:,:);
        corrlin(count,:) = tmp(maskmat);
        count = count + 1;
    end
end
sessions(sub)
for sub = 1:numSubs
    for ses = 1:sessions(sub)
        tmp = sub_mats(count,:,:,:);
        corrlin(count,:) = tmp(maskmat);
        count = count + 1;
    end
end
corr_to_corr
ones(ses,ses)
x = [0:5]
corr_to_corr
tick_labels = {}
tick_labels = {tick}
tick_labels = {tick, tick}
tick_labels = {tick_labels, tick}
tick_labels = {tick_labels tick}
tick_labels = {tick}
tick_labels = {tick_labels tick}
tick_labels = {tick}
tick_labels{end+1} = {tick}
tick_labels{end+1} = tick
corr_to_corr
permutations_cluster_correction
subsampling
addpath '/Users/dianaperez/Documents/GitHub/Lifespan-Analysis'
subsampling
plot(times(1:5),corr(1,:), 'Color',[1, 0.5, 0], 'LineWidth', 3)
ylim([0 1]);
plot(corr(1,:))
plot(corr(1,1:5))
plot(times(1:5),corr(1,1:5))
plot(times(1:5),corr(1,5), 'Color',[1, 0.5, 0], 'LineWidth', 3)
ylim([0 1]);
hold on
plot(times, corr(2,:), 'Color', [1,1,0], 'LineWidth', 3)
ylim([0 1]);
hold on
plot(times(1:18),corr(3,1:18), 'Color',[0, 0, 1], 'LineWidth', 3)
hold on
plot(times(1:5),corr(1,5), 'Color',[1, 0.5, 0], 'LineWidth', 3)
ylim([0 1]);
hold on
plot(times, corr(2,1:12), 'Color', [1,1,0], 'LineWidth', 3)
ylim([0 1]);
hold on
```

```
plot(times(1:18),corr(3,1:18),'Color',[0, 0, 1],'LineWidth', 3)
hold on
plot(times(1:5),corr(1,5),'Color',[1, 0.5, 0],'LineWidth', 3)
ylim([0 1]);
hold on
plot(times, corr(2,1:12), 'Color', [1,1,0], 'LineWidth', 3)
ylim([0 1]);
hold on
plot(times(1:18),corr(3,1:8),'Color',[0, 0, 1],'LineWidth', 3)
hold on
plot(times(1:5),corr(1,5),'Color',[1, 0.5, 0],'LineWidth', 3)
ylim([0 1]);
hold on
plot(times, corr(2,1:12), 'Color', [1,1,0], 'LineWidth', 3)
ylim([0 1]);
hold on
plot(times(1:8),corr(3,1:8),'Color',[0, 0, 1],'LineWidth', 3)
hold on
plot(times(1:5),corr(1,5),'Color',[1, 0.5, 0],'LineWidth', 3)
ylim([0 1]);
hold on
plot(times(1:12), corr(2,:), 'Color', [1,1,0], 'LineWidth', 3)
ylim([0 1]);
hold on
plot(times(1:8),corr(3,1:8),'Color',[0, 0, 1],'LineWidth', 3)
hold on
plot(times(1:5),corr(1,5),'Color',[1, 0.5, 0],'LineWidth', 3)
ylim([0 1]);
hold on
plot(times(1:12), corr(2,:), 'Color', [1,1,1], 'LineWidth', 3)
ylim([0 1]);
hold on
plot(times(1:8),corr(3,1:8),'Color',[0, 0, 1],'LineWidth', 3)
hold on
plot(times(1:5),corr(1,5),'Color',[1, 0.5, 0],'LineWidth', 3)
ylim([0 1]);
hold on
plot(times(1:12), corr(2,:), 'Color', [1,1,1], 'LineWidth', 3)
ylim([0 1]);
hold on
plot(times(1:12), corr(2,:), 'Color', [0,0,0], 'LineWidth', 3)
ylim([0 1]);
hold on
plot(times(1:8),corr(3,1:8),'Color',[0, 0, 1],'LineWidth', 3)
hold on
plot(times(1:5),corr(1,5),'Color',[1, 0.5, 1],'LineWidth', 3)
ylim([0 1]);
hold on
figure;
plot(times(1:5),corr(1,5),'Color',[1, 0.5, 1],'LineWidth', 3)
ylim([0 1]);
```

```
hold on
figure;
plot(times(1:5),corr(1,1:5),'Color',[1, 0.5, 0],'LineWidth', 3)
ylim([0 1]);
hold on
plot(times(1:12), corr(2,:), 'Color', [0,0,0], 'LineWidth', 3)
ylim([0 1]);
hold on
plot(times(1:8),corr(3,1:8),'Color',[0, 0, 1],'LineWidth', 3)
hold on
CreateVariantFiles_LS03specific
varmap_first = ft_read_cifti_mod
('/Users/dianaperez/Box/Research/Lifespan/LS03_variants_sizeExcluded_first-half_thresh-
5_smooth_2.55.dtseries.nii');
varmap_first = ft_read_cifti_mod
('/Users/dianaperez/Box/Research/Lifespan/LS03_variants_sizeExcluded_second-half_thresh-
5_smooth_2.55.dtseries.nii');
varmap_first = ft_read_cifti_mod
('/Users/dianaperez/Box/Research/Lifespan/LS03_variants_sizeExcluded_first-half_thresh-
5_smooth_2.55.dtseries.nii');
varmap_second = ft_read_cifti_mod
('/Users/dianaperez/Box/Research/Lifespan/LS03_variants_sizeExcluded_second-half_thresh-
5_smooth_2.55.dtseries.nii');
clear all
varmap_first = ft_read_cifti_mod
('/Users/dianaperez/Box/Research/Lifespan/LS03_variants_sizeExcluded_first-half_thresh-
5_smooth_2.55.dtseries.nii');
varmap_second = ft_read_cifti_mod
('/Users/dianaperez/Box/Research/Lifespan/LS03_variants_sizeExcluded_second-half_thresh-
5_smooth_2.55.dtseries.nii');
data_first = varmap_first.data;
data_second = varmap_first.data;
new_cifti = zeros(data_first);
new_cifti = zeros(data_first,1);
new_cifti = zeros(size(data_first));
for v = 1:length(new_cifti)
if data_first(v,1) == 1 && data_second(v,1) == 1
new_cifti(v,1) == 3;
elseif data_first(v,1) == 1 && data_second(v,1) == 0
new_cifti(v,1) == 1;
elseif data_first(v,1) == 0 && data_second(v,1) == 1
new_cifti(v,1) == 2;
else
disp('Check Variant IDs');
end
end
unique(data_first)
new_cifti = zeros(size(data_first));
for v = 1:length(new_cifti)
if data_first(v,1) == 1 && data_second(v,1) == 1
new_cifti(v,1) == 3;
elseif data_first(v,1) == 1 && data_second(v,1) == 0
new_cifti(v,1) == 1;
elseif data_first(v,1) == 0 && data_second(v,1) == 1
new_cifti(v,1) == 2;
elseif data_first(v,1) == 0 && data_second(v,1) == 0
new_cifti(v,1) == 0;
else
disp('Check Variant IDs');
end
end
```



```

end
unique(new_cifti)
plot_variant_overlap_splithalf
find(data_first>0)
find(data_second>0)
plot_variant_overlap_splithalf
unique(new_cifti)
plot_variant_overlap_splithalf
data_first(598,1)
data_second(598,1)
plot_variant_overlap_splithalf
unique(new_cifti)
if data_first(598,1) == 1 && data_second(598,1) == 1
disp('hope it works')
end
plot_variant_overlap_splithalf
new_cifti(v)
plot_variant_overlap_splithalf
new_cifti(v)
plot_variant_overlap_splithalf
unique(new_cifti)
varmap_first.data = new_cifti;
ft_write_cifti_mod('/Users/dianaperez/Desktop/sub-LS03_split-half_variant-overlap.
dtseries.nii', data_first)
ft_write_cifti_mod('/Users/dianaperez/Desktop/sub-LS03_split-half_variant-overlap.
dtseries.nii', varmap_first)
CreateVariantFiles_LS03specific
plot_variant_overlap_splithalf
permutations_make_zmap
CreateVariantFiles_LS03specific
help contains
addpath(genpath('/Users/dianaperez/Documents/GitHub/GrattonLab-General-Repo/FCProcess'))
addpath(genpath('/Users/dianaperez/Documents/GitHub/GrattonLab-General-Repo/FCProcess/'))
subsampling
addpath '/Users/dianaperez/Documents/GitHub/Lifespan-Analysis/Load_nii'
subsampling
figure;
% plot(times(1:5),corr(1,1:5),'Color',[1, 0.5, 0],'LineWidth', 3)
% ylim([0 1]);
% hold on
plot(times(1:12), corr(1,:), 'Color', [1,0.5,0], 'LineWidth', 3)
ylim([0 1]);
hold on
plot(times(1:8),corr(2,1:8),'Color',[0, 0, 1],'LineWidth', 3)
hold on
ylabel('Pearson Correlation (r)');
xlabel('Time (Minutes)');
m = findobj(gca,'Type','line');
hleg1 = legend(m(1:2), 'LS05', 'LS03', 'Location', 'SouthEast');
hleg1.FontSize = 14;
ax = gca;
ax.FontSize = 17;
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.5, 0.7]);
print(gcf,['/projects/p31161/ReliabilityLifespanRestDatatruhalf' num2str(pts2sample) '.
jpg'],'-dpng','-r300');
figure
plot(times(1:12), corr(1,:), 'Color', [1,0.5,0], 'LineWidth', 3)
plot(times, corr(1,:), 'Color', [1,0.5,0], 'LineWidth', 3)
ylim([0 1]);
hold on

```

```

plot(times(1:8),corr(2,1:8),'Color',[0, 0, 1],'LineWidth', 3)
hold on
ylabel('Pearson Correlation (r)');
xlabel('Time (Minutes)');
m = findobj(gca,'Type','line');
hleg1 = legend(m(1:2), 'LS05', 'LS03', 'Location', 'SouthEast');
hleg1.FontSize = 14;
ax = gca;
ax.FontSize = 17;
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.5, 0.7]);
print(gcf,['/projects/p31161/ReliabilityLifespanRestDatatruhalf' num2str(pts2sample) '.\
jpg'],'-dpng','-r300');
print(gcf,[output_dir '/ReliabilityLifespanRestDatatruhalf' num2str(pts2sample) '.\
jpg'],'-dpng','-r300');
times = [5:5:100];
figure;
% plot(times(1:5),corr(1,1:5),'Color',[1, 0.5, 0],'LineWidth', 3)
% ylim([0 1]);
% hold on
plot(times, corr(1,:), 'Color', [1,0.5,0], 'LineWidth', 3)
ylim([0 1]);
hold on
plot(times(1:18),corr(2,1:18),'Color',[0, 0, 1],'LineWidth', 3)
hold on
ylabel('Pearson Correlation (r)');
xlabel('Time (Minutes)');
m = findobj(gca,'Type','line');
hleg1 = legend(m(1:2), 'LS05', 'LS03', 'Location', 'SouthEast');
hleg1.FontSize = 14;
ax = gca;
ax.FontSize = 17;
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.5, 0.7]);
print(gcf,[output_dir '/ReliabilityLifespanRestDatatruhalf' num2str(pts2sample) '.\
jpg'],'-dpng','-r300');
corr_to_corr
tmp = mean_LS02;
corrln(1,:) = tmp(maskmat);
tmp = mean_LS03;
corrln(2,:) = tmp(maskmat);
tmp = mean_LS04;
corrln(3,:) = tmp(maskmat);
tmp = mean_LS05;
corrln(4,:) = tmp(maskmat);
tmp = mean_LS07;
corrln(5,:) = tmp(maskmat);
tmp = mean_LS10;
corrln(6,:) = tmp(maskmat);
simmat = corr(corrln');
clear corr
simmat = corr(corrln');
figure('Position',[1 1 1000 800]);
imagesc(simmat,[0 1]); colormap('jet');
hline_new([0,1,2,3,4,5,6]+0.5,'k',2);
vline_new([0,1,2,3,4,5,6]+0.5,'k',2);
set(gca,'XTick',[1,2,3,4,5,6], 'YTick', [1,2,3,4,5,6], 'XTickLabel',...
{'LS02', 'LS03', 'LS04', 'LS05', 'LS07', 'LS10'}, 'YTickLabel', {'LS02', 'LS03', 'LS05',\
'LS07'});
axis square;
colorbar;
set(gca,'XTick',[1,2,3,4,5,6], 'YTick', [1,2,3,4,5,6], 'XTickLabel',...

```

```

{'LS02', 'LS03', 'LS04', 'LS05', 'LS07', 'LS10'}, 'YTickLabel', {'LS02', 'LS03', 'LS05', 'LS07', 'LS10'});
set(gca, 'XTick', [1,2,3,4,5,6], 'YTick', [1,2,3,4,5,6], 'XTickLabel', {'LS02', 'LS03', 'LS04', 'LS05', 'LS07', 'LS10'}, 'YTickLabel', {'LS02', 'LS03', 'LS05', 'LS07', 'LS10'});
set(gca, 'XTick', [1,2,3,4,5,6], 'YTick', [1,2,3,4,5,6], 'XTickLabel', {'LS02', 'LS03', 'LS04', 'LS05', 'LS07', 'LS10'}, 'YTickLabel', {'LS02', 'LS03', 'LS05', 'LS07', 'LS10'});
title('Similarity Across Subjects')
axis square;
colorbar;
saveas(gcf, [outdir 'SimilarityMat_AcrossSubs.tiff'], 'tiff');
maskmat2 = ones(simmat);
maskmat2 = ones(size(simmat));
maskmat2 = logical(triu(maskmat2,1));
sim = simmat(maskmat);
sim = simmat(maskmat2);
mean(sim)
subsampling
%-- 2/28/21, 7:08 PM --%
permutations_cluster_correction
permute_flippedVars
permutations_cluster_correction
clear all
permutations_cluster_correction
permute_flippedVars
permute_flippedVars
permute_flippedVars
template = ft_read_cifti_mod(LH_files{1});
new_cifti = [];
count = 1;
left_hem = length(template.brainstructure)/2;
for i = 1:left_hem
if template.brainstructure(i) == 1
homotopic_overlap
sum(allSubs)
sum(allsubs)
homotopic_overlap
permute_flippedVars
new_cifti = [];
count = 1;
template = ft_read_cifti_mod(LH_files{1});
for i = 1:length(template.brainstructure)
if template.brainstructure(i) > 0
new_cifti(i,1) = trueDiff;
count = count + 1;
else
new_cifti(i,1) = template.brainstructure(i);
end
end
for i = 1:length(template.brainstructure)
if template.brainstructure(i) > 0
new_cifti(i,1) = trueDiff(count);
count = count + 1;
else
new_cifti(i,1) = template.brainstructure(i);
end
end
for i = 1:(length(template.brainstructure)/2)
if template.brainstructure(i) > 0
new_cifti(i,1) = trueDiff;
count = count + 1;

```

```
else
new_cifti(i,1) = template.brainstructure(i);
end
end
left_hem = template.brainstructure(1:(length(template.brainstructure)/2));
for i = 1:(length(left_hem))
if left_hem(i) > 0
new_cifti(i,1) = trueDiff;
count = count + 1;
else
new_cifti(i,1) = left_hem(i);
end
end
for i = 1:length(left_hem)
if left_hem(i) > 0
new_cifti(i,1) = trueDiff;
count = count + 1;
else
new_cifti(i,1) = left_hem(i);
end
end
count = 1;
cifti = [];
for i = 1:length(left_hem)
if left_hem(i) > 0
new_cifti(i,1) = trueDiff(count);
count = count + 1;
else
new_cifti(i,1) = left_hem(i);
end
end
new_cifti = [new_cifti; new_cifti];
new_cifti = new_cifti(template.brainstructure>0);
ft_write_cifti_mod('/Users/dianaperez/Desktop/LH_diffmap.dtseries.nii', new_cifti);
template.data = new_cifti;
ft_write_cifti_mod('/Users/dianaperez/Desktop/LH_diffmap.dtseries.nii', template);
template = ft_read_cifti_mod(LH_files{1});
count = 1;
cifti = [];
for i = 1:length(left_hem)
if left_hem(i) > 0
new_cifti(i,1) = trueDiff(count);
count = count + 1;
else
new_cifti(i,1) = left_hem(i);
end
end
new_cifti = [new_cifti; new_cifti];
new_cifti = new_cifti(template.brainstructure>0);
template.data = new_cifti;
ft_write_cifti_mod('/Users/dianaperez/Desktop/MH_diffmap.dtseries.nii', template);
template = ft_read_cifti_mod(LH_files{1});
count = 1;
cifti = [];
for i = 1:length(left_hem)
if left_hem(i) > 0
new_cifti(i,1) = trueDiff(count);
count = count + 1;
else
new_cifti(i,1) = left_hem(i);
```

```

end
end
new_cifti = [new_cifti; new_cifti];
new_cifti = new_cifti(template.brainstructure>0);
template.data = new_cifti;
ft_write_cifti_mod('/Users/dianaperez/Desktop/RH_diffmap.dtseries.nii', template);
template = ft_read_cifti_mod(LH_files{1});
count = 1;
cifti = [];
for i = 1:length(left_hem)
if left_hem(i) > 0
new_cifti(i,1) = trueDiff(count);
count = count + 1;
else
new_cifti(i,1) = left_hem(i);
end
end
new_cifti = [new_cifti; new_cifti];
new_cifti = new_cifti(template.brainstructure>0);
template.data = new_cifti;
ft_write_cifti_mod('/Users/dianaperez/Desktop/allSubs_diffmap.dtseries.nii', template);
permute_left_right_handlers
open make_zmap
template = ft_read_cifti_mod(RH_files{1});
template.data = trueDiff;
ft_write_cifti_mod('/Users/dianaperez/Desktop/permutations_RH-LH_trueDiffMap.dtseries.
nii', template);
HCP_assignNetworksToVariants
clear all
HCP_assignNetworksToVariants
size(subs,1)
HCP_assignNetworksToVariants
addpath '/Users/dianaperez/Documents/GitHub/Lifespan-Analysis'
HCP_assignNetworksToVariants
cifti_geo.distances = cifti_geo.distances(1:59412,1:59412);
HCP_assignNetworksToVariants
bold_data = cifti.data;
concat = [bold_data bold_data];
HCP_assignNetworksToVariants
%-- 3/17/21, 1:46 PM --%
permute_flippedVars
get_rid_of_this
64984/2
get_rid_of_this
ft_write_cifti_mod('/Users/dianaperez/Desktop/allSubs_diffmap.dtseries.nii', trueDiff)
template = LH_files(1)/
template = LH_files(1);
template.data = trueDiff;
ft_read_cifti_mod(template)
template = ft_read_cifti_mod(LH_files(1))
template = ft_read_cifti_mod(LH_files{1})
template.data = trueDiff;
ft_write_cifti_mod('/Users/dianaperez/Desktop/allSubs_diffmap.dtseries.nii', template)
load('brain_ind.mat')
find(brain_ind==18330)
find(brain_ind==18330+(64984/2))
find(brain_ind==18988)
find(brain_ind==19070+(64984/2))
neigh = load
('/Users/dianaperez/Box/Quest_Backup/Scripts/CIFTI_RELATED/Resources/Conte69_atlas-v2.LR.

```

```
32k_fs_LR.wb/Cifti_surf_neighbors_LR_normalwall.mat');
neigh = neigh.neighbors;
min_subject_threshold= .04;
clear all
fname = '
'/Volumes/GRATTONLAB/HCP/Variants/networkAssignments/Templatematch_spCorr_bysubject_IDs_r
eassigned.dtseries.nii';
diffmap = ft_read_cifti_mod('/Users/dianaperez/Desktop/allSubs_diffmap.dtseries.nii');
netvars = ft_read_cifti_mod(fname);
neigh = load
('/Users/dianaperez/Box/Quest_Backup/Scripts/CIFTI_RELATED/Resources/Conte69_atlas-v2.LR.
32k_fs_LR.wb/Cifti_surf_neighbors_LR_normalwall.mat');
neigh = neigh.neighbors;
min_subject_threshold= .04;
diffmat_bin= logical(abs(diffmap.data)>= min_subject_threshold);
VOI = 46334;
mask = VOI;
neigh(VOI)
neigh(VOI,:)
for n = 1:7
if diffmat_bin(neigh(VOI,n))
mask = [mask; neigh(VOI,n)];
end
end
mask = VOI;
for n = 1:7
if diffmat_bin(neigh(VOI,n))
mask = [mask; neigh(VOI,n)];
end
end
mask = unique(mask)
for m = 1:length(mask)
for n = 1:7
if diffmat_bin(neigh(m,n))
mask = [mask; neigh(m,n)];
end
end
mask = unique(mask);
end
length(mask)
neigh(m,n)
for m = 1:length(mask)
for n = 1:7
if diffmat_bin(neigh(mask(m),n))
mask = [mask; neigh(mask(m),n)];
end
end
mask = unique(mask);
unique(mask)
for m = 1:length(mask)
for n = 1:7
if diffmat_bin(neigh(mask(m),n))
mask = [mask; neigh(mask(m),n)];
end
end
mask = unique(mask);
end
diffmat_bin(mask)
for m = 1:length(mask)
```

```
for n = 1:7
if diffmat_bin(neigh(mask(m),n))
mask = [mask; neigh(mask(m),n)];
end
end
mask = unique(mask);
end
new_cifti = zeros(59412,1);
new_cifti(mask) == 1;
template = diffmap;
template.data = new_cifti;
ft_write_cifti_mod('/Users/dianaperez/Desktop/HCP_operculum_variant_cluster_mask.
dtseries.nii', template);
unique(new_cifti)
new_cifti(mask) = 1;
unique(new_cifti)
template.data = new_cifti;
ft_write_cifti_mod('/Users/dianaperez/Desktop/HCP_operculum_variant_cluster_mask.
dtseries.nii', template);
clear all
fname =
'/Volumes/GRATTONLAB/HCP/Variants/networkAssignments/Templatematch_spCorr_bysubject_IDs_r
eassigned.dtseries.nii';
maskname = '/Users/dianaperez/Desktop/HCP_operculum_variant_cluster_mask.dtseries.nii';
netvars = ft_read_cifti_mod(fname);
ROI_mask = ft_read_cifti_mod(maskname);
uiopen('/Users/dianaperez/Desktop/HCPsublist.txt',1)
load('/Users/dianaperez/Box/Research/Lateralization_Variants/subjectData/goodSubs384.
mat')
load('/Users/dianaperez/Box/Research/Lateralization_Variants/subjectData/goodSubs752.
mat')
goodSubs384 = goodSubs384(:,1);
goodSubs752 = goodSubs752(:,1);
help sort
goodSubs384 = sort(goodSubs384,'descend');
goodSubs752 = sort(goodSubs752,'descend');
goodSubs384 = sort(goodSubs384,'ascend');
goodSubs752 = sort(goodSubs752,'ascend');
netvars382 = [];
help ismember
for s = 1:length(goodSubs752)
if ismember(goodSubs752(s),goodSubs384)
netvars382 = [netvars382 netvars.data(:,s)];
end
end
variants_of_interest = netvars382(ROI_mask, :);
variants_of_interest = netvars382(ROI_mask.data, :);
verts_of_interest = ROI_mask.data(ROI_mask.data==1);
variants_of_interest = netvars382(verts_of_interest, :);
get_network_IDs = unique(variants_of_interest);
verts_of_interest = find(ROI_mask.data==1);
variants_of_interest = netvars382(verts_of_interest, :);
get_network_IDs = unique(variants_of_interest);
c = 2;
num_verts_per_net(c,1) = get_networks_IDs(c);
num_verts_per_net(c,1) = get_network_IDs(c);
num_verts_per_net(c,1) = get_networks_IDs(c);
num_verts_per_net(c,1) = get_network_IDs(c);
num_verts = find(verts_of_interest==get_network_IDs(c));
find(verts_of_interest=get_network_IDs(c))
```

```
find(verts_of_interest==get_network_IDs(c))
num_verts = find(variants_of_interest==get_network_IDs(c));
num_verts_per_net(c,2) = length(num_verts);
for c = 1:length(get_network_IDs)
num_verts_per_net(c,1) = get_network_IDs(c);
num_verts = find(variants_of_interest==get_network_IDs(c));
num_verts_per_net(c,2) = length(num_verts);
end
num_verts_per_net = num_verts_per_net(2:end,:);
pie(num_verts_per_net)
find(variants_of_interest>0);
clear all
fname = ✓
'/Volumes/GRATTONLAB/HCP/Variants/networkAssignments/Templatematch_spCorr_bysubject_IDs_r✓
eassigned.dtseries.nii';
% diffmap = ft_read_cifti_mod('/Users/dianaperez/Desktop/allSubs_diffmap.dtseries.nii');
diffmap = ft_read_cifti_mod('/Users/dianaperez/Desktop/HCP_RH-✓
LH_diffMap_384GoodSubs_thresh7.5.dtseries.nii');
netvars = ft_read_cifti_mod(fname);
neigh = load✓
('/Users/dianaperez/Box/Quest_Backup/Scripts/CIFTI_RELATED/Resources/Conte69_atlas-v2.LR.✓
32k_fs_LR.wb/Cifti_surf_neighbors_LR_normalwall.mat');
neigh = neigh.neighbors;
min_subject_threshold= .04;
diffmat_bin= logical(abs(diffmap.data)>= min_subject_threshold);
VOI = 16620;
mask = VOI;
for n = 1:7
if diffmat_bin(neigh(VOI,n))
mask = [mask; neigh(VOI,n)];
end
end
for m = 1:length(mask)
for n = 1:7
if diffmat_bin(neigh(mask(m),n))
mask = [mask; neigh(mask(m),n)];
end
end
mask = unique(mask);
end
for n = 1:7
if diffmat_bin(neigh(VOI,n))
mask = [mask; neigh(VOI,n)];
end
end
mask = unique(mask);
neigh(VOI,1)
diffmat_bin(neigh(VOI,1))
diffmat_bin(neigh(VOI,2))
diffmat_bin(neigh(VOI,3))
diffmat_bin(neigh(VOI,4))
diffmat_bin(neigh(VOI,5))
diffmat_bin(neigh(VOI,6))
diffmat_bin(neigh(VOI,7))
VOI = 16487;
for n = 1:7
if diffmat_bin(neigh(VOI,n))
mask = [mask; neigh(VOI,n)];
end
end
end
```



```
for m = 1:length(mask)
for n = 1:7
if diffmat_bin(neigh(mask(m),n))
mask = [mask; neigh(mask(m),n)];
end
end
mask = unique(mask);
end
new_cifti = zeros(59412,1);
new_cifti(mask) = 1;
template = diffmap;
template.data = new_cifti;
ft_write_cifti_mod('/Users/dianaperez/Desktop/HCP_operculum_Lefthem_variant_cluster_mask.
dtseries.nii', template);
VOI = 17278;
mask = VOI;
for n = 1:7
if diffmat_bin(neigh(VOI,n))
mask = [mask; neigh(VOI,n)];
end
end
mask = unique(mask);
for m = 1:length(mask)
for n = 1:7
if diffmat_bin(neigh(mask(m),n))
mask = [mask; neigh(mask(m),n)];
end
end
mask = unique(mask);
end
new_cifti = zeros(59412,1);
new_cifti(mask) = 1;
template = diffmap;
template.data = new_cifti;
ft_write_cifti_mod(
('/Users/dianaperez/Desktop/HCP_operculum_Lefthem_variant_cluster_mask_RHvsLH.dtseries.
nii', template);
VOI = 47074;
mask = VOI;
for n = 1:7
if diffmat_bin(neigh(VOI,n))
mask = [mask; neigh(VOI,n)];
end
end
mask = unique(mask);
for m = 1:length(mask)
for n = 1:7
if diffmat_bin(neigh(mask(m),n))
mask = [mask; neigh(mask(m),n)];
end
end
mask = unique(mask);
end
xfor m = 1:length(mask)
for n = 1:7
if diffmat_bin(neigh(mask(m),n))
mask = [mask; neigh(mask(m),n)];
end
end
mask = unique(mask);
```

```
end
for m = 1:length(mask)
for n = 1:7
if diffmat_bin(neigh(mask(m),n))
mask = [mask; neigh(mask(m),n)];
end
end
mask = unique(mask);
end
help isnan
for m = 1:length(mask)
neighbors = neigh(mask(m),n);
neigh_ind = isnan(neighbors);
neighbors = neighbors(neigh_ind==0);
for n = 1:length(neighbors)
if diffmat_bin(neighbors(m))
mask = [mask; neighbors(m)];
end
end
mask = unique(mask);
end
for m = 1:length(mask)
neighbors = neigh(mask(m),n);
neigh_ind = isnan(neighbors);
neighbors = neighbors(neigh_ind==0);
for n = 1:length(neighbors)
if diffmat_bin(neighbors(m))
mask = [mask; neighbors(m)];
end
end
mask = unique(mask);
end
min_subject_threshold= -.04;
%diffmat_bin= logical(abs(diffmap.data)>= min_subject_threshold);
diffmat_bin= logical(diffmap.data<= min_subject_threshold);
VOI = 47074;
mask = VOI;
for n = 1:7
if diffmat_bin(neigh(VOI,n))
mask = [mask; neigh(VOI,n)];
end
end
for m = 1:length(mask)
neighbors = neigh(mask(m),n);
neigh_ind = isnan(neighbors);
neighbors = neighbors(neigh_ind==0);
for n = 1:length(neighbors)
if diffmat_bin(neighbors(m))
mask = [mask; neighbors(m)];
end
end
mask = unique(mask);
end
make_operculum_mask
for m = 1:length(mask)
neighbors = neigh(mask(m),:);
neigh_ind = isnan(neighbors);
neighbors = neighbors(neigh_ind==0);
for n = 1:length(neighbors)
if diffmat_bin(neighbors(n))
```

```
mask = [mask; neighbors(n)];
end
end
mask = unique(mask);
end
new_cifti = zeros(59412,1);
new_cifti(mask) = 1;
template = diffmap;
template.data = new_cifti;
ft_write_cifti_mod(
    ('/Users/dianaperez/Desktop/HCP_operculum_Righthem_variant_cluster_mask_RHvsLH.dtseries.
nii', template);
VOI = 17278;
mask = VOI;
for n = 1:7
    if diffmat_bin(neigh(VOI,n))
        mask = [mask; neigh(VOI,n)];
    end
end
for m = 1:length(mask)
    neighbors = neigh(mask(m),:);
    neigh_ind = isnan(neighbors);
    neighbors = neighbors(neigh_ind==0);
    for n = 1:length(neighbors)
        if diffmat_bin(neighbors(n))
            mask = [mask; neighbors(n)];
        end
    end
end
mask = unique(mask);
end
for m = 1:length(mask)
    neighbors = neigh(mask(m),:);
    neigh_ind = isnan(neighbors);
    neighbors = neighbors(neigh_ind==0);
    for n = 1:length(neighbors)
        if diffmat_bin(neighbors(n))
            mask = [mask; neighbors(n)];
        end
    end
end
mask = unique(mask);
end
new_cifti = zeros(59412,1);
new_cifti(mask) = 1;
template = diffmap;
template.data = new_cifti;
ft_write_cifti_mod(
    ('/Users/dianaperez/Desktop/HCP_operculum_Lefthem_variant_cluster_mask_RHvsLH.dtseries.
nii', template);
diffmap = ft_read_cifti_mod('/Users/dianaperez/Desktop/allSubs_diffmap.dtseries.nii');
min_subject_threshold= .04;
diffmat_bin= logical(abs(diffmap.data)>= min_subject_threshold);
%diffmat_bin= logical(diffmap.data<= min_subject_threshold);
VOI = 16487;
mask = VOI;
for n = 1:7
    if diffmat_bin(neigh(VOI,n))
        mask = [mask; neigh(VOI,n)];
    end
end
end
for m = 1:length(mask)
```

```
neighbors = neigh(mask(m),:);
neigh_ind = isnan(neighbors);
neighbors = neighbors(neigh_ind==0);
for n = 1:length(neighbors)
if diffmat_bin(neighbors(n))
mask = [mask; neighbors(n)];
end
end
mask = unique(mask);
end
for n = 1:7
if diffmat_bin(neigh(VOI,n))
mask = [mask; neigh(VOI,n)];
end
end
VOI = 16566;
mask = VOI;
for n = 1:7
if diffmat_bin(neigh(VOI,n))
mask = [mask; neigh(VOI,n)];
end
end
for m = 1:length(mask)
neighbors = neigh(mask(m),:);
neigh_ind = isnan(neighbors);
neighbors = neighbors(neigh_ind==0);
for n = 1:length(neighbors)
if diffmat_bin(neighbors(n))
mask = [mask; neighbors(n)];
end
end
mask = unique(mask);
end
new_cifti = zeros(59412,1);
new_cifti(mask) = 1;
template = diffmap;
template.data = new_cifti;
ft_write_cifti_mod('/Users/dianaperez/Desktop/HCP_operculum_Lefthem_variant_cluster_mask.
dtseries.nii', template);
VOI = 46334;
mask = VOI;
for n = 1:7
if diffmat_bin(neigh(VOI,n))
mask = [mask; neigh(VOI,n)];
end
end
for m = 1:length(mask)
neighbors = neigh(mask(m),:);
neigh_ind = isnan(neighbors);
neighbors = neighbors(neigh_ind==0);
for n = 1:length(neighbors)
if diffmat_bin(neighbors(n))
mask = [mask; neighbors(n)];
end
end
mask = unique(mask);
end
new_cifti = zeros(59412,1);
new_cifti(mask) = 1;
template = diffmap;
```

```
template.data = new_cifti;
ft_write_cifti_mod
('/Users/dianaperez/Desktop/HCP_operculum_Righthem_variant_cluster_mask.dtseries.nii',
template);
maskname = '/Users/dianaperez/Desktop/HCP_operculum_Righthem_variant_cluster_mask.
dtseries.nii';
ROI_mask = ft_read_cifti_mod(maskname);
load('/Users/dianaperez/Box/Research/Lateralization_Variants/subjectData/goodSubs384.
mat')
load('/Users/dianaperez/Box/Research/Lateralization_Variants/subjectData/goodSubs752.
mat')
goodSubs384 = goodSubs384(:,1);
goodSubs752 = goodSubs752(:,1);
goodSubs384 = sort(goodSubs384,'ascend');
goodSubs752 = sort(goodSubs752,'ascend');
netvars382 = [];
for s = 1:length(goodSubs752)
if ismember(goodSubs752(s),goodSubs384)
netvars382 = [netvars382 netvars.data(:,s)];
end
end
verts_of_interest = find(ROI_mask.data==1);
variants_of_interest = netvars382(verts_of_interest, :);
network_IDs = unique(variants_of_interest);
num_verts_per_net = [];
for c = 1:length(network_IDs)
num_verts_per_net(c,1) = network_IDs(c);
num_verts = find(variants_of_interest==network_IDs(c));
num_verts_per_net(c,2) = length(num_verts);
end
num_verts_per_net = num_verts_per_net(2:end,:);
maskname = '/Users/dianaperez/Desktop/HCP_operculum_Lefthem_variant_cluster_mask.
dtseries.nii';
ROI_mask = ft_read_cifti_mod(maskname);
clear verts_of_interest
clear variants_of_interest
clear networks_IDs
verts_of_interest = find(ROI_mask.data==1);
variants_of_interest = netvars382(verts_of_interest, :);
network_IDs = unique(variants_of_interest);
num_verts_per_net = [];
for c = 1:length(network_IDs)
num_verts_per_net(c,1) = network_IDs(c);
num_verts = find(variants_of_interest==network_IDs(c));
num_verts_per_net(c,2) = length(num_verts);
end
num_verts_per_net = num_verts_per_net(2:end,:);
maskname = '/Users/dianaperez/Desktop/HCP_operculum_Righthem_variant_cluster_mask.
dtseries.nii';
ROI_mask = ft_read_cifti_mod(maskname);
clear networks_IDs
clear variants_of_interest
clear verts_of_interest
verts_of_interest = find(ROI_mask.data==1);
variants_of_interest = netvars382(verts_of_interest, :);
network_IDs = unique(variants_of_interest);
num_verts_per_net = [];
for c = 1:length(network_IDs)
num_verts_per_net(c,1) = network_IDs(c);
num_verts = find(variants_of_interest==network_IDs(c));
```

```
num_verts_per_net(c,2) = length(num_verts);
end
num_verts_per_net = num_verts_per_net(2:end,:);
clear networks_IDs
clear variants_of_interest
clear verts_of_interest
maskname = '/Users/dianaperez/Desktop/HCP_operculum_Righthem_variant_cluster_mask_RHvsLH.
dtseries.nii';
ROI_mask = ft_read_cifti_mod(maskname);
load('/Users/dianaperez/Box/Research/Lateralization_Variants/subjectData/goodSubs384.
mat')
load('/Users/dianaperez/Box/Research/Lateralization_Variants/subjectData/goodSubs752.
mat')
goodSubs384 = sort(goodSubs384,'ascend');
goodSubs752 = sort(goodSubs752,'ascend');
help sort
load('/Users/dianaperez/Box/Research/Lateralization_Variants/subjectData/goodSubs384.
mat')
load('/Users/dianaperez/Box/Research/Lateralization_Variants/subjectData/goodSubs752.
mat')
sortrows(goodSubs384,1)
goodSubs384 = ans;
goodSubs752 = sort(goodSubs752,'ascend');
netvars384 = [];
for s = 1:length(goodSubs752)
if ismember(goodSubs752(s),goodSubs384)
netvars384 = [netvars384 netvars.data(:,s)];
end
end
netvarsLH = [];
netvarsMH = [];
netvarsRH = [];
for s = 1:length(goodSubs384)
if goodSubs384(s,2) < -28
netvarsLH = [netvarsLH; netvars384(:,s)];
elseif goodSubs384(s,2) < 48
netvarsMH = [netvarsMH; netvars384(:,s)];
elseif goodSubs384(s,2) <= 100
netvarsRH = [netvarsRH; netvars384(:,s)];
end
end
verts_of_interest = find(ROI_mask.data==1);
netvarsLH = [];
netvarsMH = [];
netvarsRH = [];
for s = 1:length(goodSubs384)
if goodSubs384(s,2) < -28
netvarsLH = [netvarsLH netvars384(:,s)];
elseif goodSubs384(s,2) < 48
netvarsMH = [netvarsMH netvars384(:,s)];
elseif goodSubs384(s,2) <= 100
netvarsRH = [netvarsRH netvars384(:,s)];
end
end
verts_of_interest = find(ROI_mask.data==1);
variants_of_interest = netvarsLH(verts_of_interest, :);
network_IDs = unique(variants_of_interest);
for c = 1:length(network_IDs)
num_verts_per_net(c,1) = network_IDs(c);
num_verts = find(variants_of_interest==network_IDs(c));
```

```
num_verts_per_net(c,2) = length(num_verts);
end
num_verts_per_net = num_verts_per_net(2:end,:);
clear verts_of_interest
clear variants_of_interest
clear networks_IDs
verts_of_interest = find(ROI_mask.data==1);
variants_of_interest = netvarsRH(verts_of_interest, :);
network_IDs = unique(variants_of_interest);
num_verts_per_net = [];
for c = 1:length(network_IDs)
num_verts_per_net(c,1) = network_IDs(c);
num_verts = find(variants_of_interest==network_IDs(c));
num_verts_per_net(c,2) = length(num_verts);
end
num_verts_per_net = num_verts_per_net(2:end,:);
maskname = '/Users/dianaperez/Desktop/HCP_operculum_Lefthem_variant_cluster_mask_RHvsLH.
dtseries.nii';
ROI_mask = ft_read_cifti_mod(maskname);
clear verts_of_interest
clear variants_of_interest
clear network_IDs
verts_of_interest = find(ROI_mask.data==1);
variants_of_interest = netvarsRH(verts_of_interest, :);
network_IDs = unique(variants_of_interest);
num_verts_per_net = [];
for c = 1:length(network_IDs)
num_verts_per_net(c,1) = network_IDs(c);
num_verts = find(variants_of_interest==network_IDs(c));
num_verts_per_net(c,2) = length(num_verts);
end
num_verts_per_net = num_verts_per_net(2:end,:);
clear network_IDs
clear verts_of_interest
clear variants_of_interest
verts_of_interest = find(ROI_mask.data==1);
variants_of_interest = netvarsLH(verts_of_interest, :);
network_IDs = unique(variants_of_interest);
num_verts_per_net = [];
for c = 1:length(network_IDs)
num_verts_per_net(c,1) = network_IDs(c);
num_verts = find(variants_of_interest==network_IDs(c));
num_verts_per_net(c,2) = length(num_verts);
end
num_verts_per_net = num_verts_per_net(2:end,:);
maskname = '/Users/dianaperez/Desktop/HCP_operculum_Righthem_variant_cluster_mask.
dtseries.nii';
ROI_mask = ft_read_cifti_mod(maskname);
verts_of_interest = find(ROI_mask.data==1);
%-- 3/28/21, 2:12 PM --%
corr_to_corr
%-- 4/8/21, 9:20 PM --%
overlapmap_gender
females(1)
overlapmap_gender
permute_flippedVars
permute_flippedVars
%-- 4/28/21, 10:05 AM --%
cd /Users/dianaperez/Box/DATA
dir(BOLD_DATA)
```

```
dir('BOLD_DATA')
ff = dir('BOLD_DATA');
folder_1 = ff.name(3:2500);
folder_1 = ff.name{3:2500};
folder_1 = ff.name(3:2500,1);
folder_names = ff.name;
folder_names = ff.name(3:end);
folder_names = ff.name{3:end};
ff.name
ff.name{1}
ff.name(1)
ff.name(:,11)
ff.name(1,1)
ff.name{1,1}
name = {ff.name}.';
folder_names = {ff.name(3:end)}
folder_names = {ff.name{3:end}};
folder_names = {ff.name{3:end},1}};
folder_names = {ff.name};
folder_names = name;
folder_names = folder_names(3:end);
folder_1 = folder_names(1:2500);
writeTable(folder_1, 'BOLD_DATA_1.txt')
writetable(folder_1, 'BOLD_DATA_1.txt')
writecell(folder_1, 'BOLD_DATA_1.txt')
folder_1 = folder_names(1:3000);
folder_1 = folder_names(1:2000);
writecell(folder_1, 'BOLD_DATA_1.txt')
writecell(folder_1, 'BOLD_DATA_2.txt')
folder_1 = folder_names(1:2500);
folder_1 = folder_names(1:3000);
writecell(folder_1, 'BOLD_DATA_1.txt')
permute_flippedVars
unique(new_cifti)
unique(right_hem)
unique(left_hem)
find(left_hem ==1)
find(cifti.data ==1)
ff = find(left_hem ==1);
ee = find(cifti.data ==1)
ee = find(cifti.data>0)
dd = find(right_hem ==1);
size(dd) + size(ff)
load('/Users/dianaperez/Desktop/Lateralization_thresh10/AvgVariantSizesByHem_10thresh.
mat')
clear all
load('/Users/dianaperez/Desktop/Lateralization_thresh10/AvgVariantSizesByHem_10thresh.
mat')
allSubs_avg = average{1,4};
histogram(allSubs_avg(:,1))
hold on
histogram(allSubs_avg(:,2))
ax = gca;
ax.FontSize = 15;
xlabel('Number of Subjects')
ylabel(
ylabel('Average Variant Size')
legend('Left Hem', 'Right Hem')
title('Average Variant Size x Hemisphere')
LH_avg = average{1,1};
```



```

histogram(LH_avg(:,1))
hold on
histogram(LH_avg(:,2))
ax = gca;
ax.FontSize = 15;
xlabel('Number of Subjects')
ylabel('Average Variant Size')
legend('Left Hem', 'Right Hem')
title('Average Variant Size x Hemisphere - Left Handers')
RH_avg = average{1,2};
histogram(RH_avg(:,1))
hold on
histogram(RH_avg(:,2))
ax = gca;
ax.FontSize = 15;
xlabel('Number of Subjects')
ylabel('Average Variant Size')
legend('Left Hem', 'Right Hem')
title('Average Variant Size x Hemisphere - Right Handers')
load('/Users/dianaperez/Desktop/Lateralization_thresh10/VariantSizesByHem_10thresh.mat')
allSubs_left = varsizehem{1,4}{1,1};
allSubs_left = varsizehem{1,4}{1,1};
allSubs_right = varsizehem{1,4}{1,2};
histogram(allSubs_left)
hold on
histogram(allSubs_right)
legend('Left Hem', 'Right Hem')
help dist
h = histogram(allSubs_left)
hold on
histogram(allSubs_right)
%-- 5/6/21, 11:34 AM --%
load('/Users/dianaperez/Desktop/Lateralization_thresh10/AvgVariantSizesByHem_10thresh.
mat')
size = average{1,1};
left = size(:,1);
left = size(:,1);
right = size(:,2);
histogram(left, 20);
histogram(left, 30);
hold on
histogram(right,30);
ax = gca;
permute_flippedVars
clear all
permute_flippedVars
%-- 5/6/21, 3:38 PM --%
corr_to_corr
network_seg
addpath(genpath('/Volumes/Research_HD/Dependencies'))
network_seg
mean(between)
mean(between, 'omitnan')
network_seg
%-- 5/7/21, 10:21 PM --%
run('/Users/dianaperez/Desktop/Week6_diana.m')
%-- 5/19/21, 2:52 PM --%
permute_flippedVars
permutations_cluster_correction
clear all

```

```
uiopen↵
('/Volumes/GRATTONLAB/HCP/Variants/networkAssignments/Templatematch_spCorr_bysubject_IDs_↵
reassigned.dtseries.nii',1)
net_assignment = ft_read_cifti_mod↵
('/Volumes/GRATTONLAB/HCP/Variants/networkAssignments/Templatematch_spCorr_bysubject_IDs_↵
reassigned.dtseries.nii');
%-- 5/22/21, 1:44 PM --%
net_assignment = ft_read_cifti_mod↵
('/Volumes/GRATTONLAB/HCP/Variants/networkAssignments/Templatematch_spCorr_bysubject_IDs_↵
reassigned.dtseries.nii');
startup
net_assignment = ft_read_cifti_mod↵
('/Volumes/GRATTONLAB/HCP/Variants/networkAssignments/Templatematch_spCorr_bysubject_IDs_↵
reassigned.dtseries.nii');
left_hem = net_assignment.data(net_assignment.brainstructure==1);
brain = net_assignment.brainstructure(net_assignment.brainstructure>0);
left_hem = net_assignment.data(brain==1);
left_hem = net_assignment.data(brain==1);
right_hem = net_assignment.data(brain==2);
left_hem_nets = unique(left_hem);
right_hem_nets = unique(right_hem);
num_verts_per_net = [];
for n = 1:length(left_hem_nets)
num_verts_per_net(n,1) = left_hem_nets(n);
num_verts = find(left_hem==left_hem_nets(n));
num_verts_per_net(n,2) = length(num_verts);
end
num_verts_per_net_left = num_verts_per_net;
for n = 1:length(right_hem_nets)
num_verts_per_net(n,1) = right_hem_nets(n);
num_verts = find(right_hem==right_hem_nets(n));
num_verts_per_net(n,2) = length(num_verts);
end
left_hem = net_assignment.data(net_assignment.brainstructure==1,:);
left_hem = net_assignment.data(net_assignment.brainstructure==1,:);
left_hem = net_assignment.data(net_assignment.brainstructure==1,:);
right_hem = net_assignment.data(net_assignment.brainstructure==2,:);
left_hem = net_assignment.data(brain==1,:);
right_hem = net_assignment.data(brain==2,:);
unique(net_assignment.data);
unique(net_assignment.data);
num_verts_per_net = [];
networks = [1:16];
clear num_verts right_hem_nets left_hem_nets
clear ans
clear n
clear num_verts_per_net_left
networks = [1:16];
num_verts_per_net = [];
for n = 1:length(networks)
num_verts_per_net(n,1) = networks(n);
num_verts_per_net(n,2) = length(find(left_hem==networks(n)));
num_verts_per_net(n,3) = length(find(right_hem==networks(n)));
end
pie(num_verts_per_net(:,2))
for n = 1:length(networks)
num_verts_per_net(n,1) = networks(n);
num_verts_per_net(n,2) = length(find(left_hem==networks(n)));
num_verts_per_net(n,3) = length(find(right_hem==networks(n)));
for s = 1:size(left_hem,2)
```

```
per_subject(s,1) = s;
per_subject(s,2) = length(find(left_hem(:,s)==networks(n)));
per_subject(s,3) = length(find(right_hem(:,s)==networks(n)));
end
end
for n = 1:length(networks)
num_verts_per_net(n,1) = networks(n);
num_verts_per_net(n,2) = length(find(left_hem==networks(n)));
num_verts_per_net(n,3) = length(find(right_hem==networks(n)));
for s = 1:size(left_hem,2)
per_subject(s,1,n) = s;
per_subject(s,2,n) = length(find(left_hem(:,s)==networks(n)));
per_subject(s,3,n) = length(find(right_hem(:,s)==networks(n)));
end
end
num_verts_per_net = [];
count = 1;
for n = 1:length(networks)
num_verts_per_net(n,1) = networks(n);
num_verts_per_net(n,2) = length(find(left_hem==networks(n)));
num_verts_per_net(n,3) = length(find(right_hem==networks(n)));
for s = 1:size(left_hem,2)
per_subject(s,count) = s;
count = count + 1;
per_subject(s,count) = length(find(left_hem(:,s)==networks(n)));
count = count + 1;
per_subject(s,count) = length(find(right_hem(:,s)==networks(n)));
count = count + 1;
end
end
clear per_subject
num_verts_per_net = [];
count = 1;
for n = 1:length(networks)
num_verts_per_net(n,1) = networks(n);
num_verts_per_net(n,2) = length(find(left_hem==networks(n)));
num_verts_per_net(n,3) = length(find(right_hem==networks(n)));
for s = 1:size(left_hem,2)
per_subject(s,count) = s;
count = count + 1;
per_subject(s,count) = length(find(left_hem(:,s)==networks(n)));
count = count + 1;
per_subject(s,count) = length(find(right_hem(:,s)==networks(n)));
count = count + 1;
end
end
count = 1;
for n = 1:length(networks)
num_verts_per_net(n,1) = networks(n);
num_verts_per_net(n,2) = length(find(left_hem==networks(n)));
num_verts_per_net(n,3) = length(find(right_hem==networks(n)));
for s = 1:size(left_hem,2)
per_subject(s,count) = length(find(left_hem(:,s)==networks(n)));
count = count + 1;
per_subject(s,count) = length(find(right_hem(:,s)==networks(n)));
count = count + 1;
end
end
clear per_subject
num_verts_per_net = [];
```

```
for n = 1:length(networks)
num_verts_per_net(n,1) = networks(n);
num_verts_per_net(n,2) = length(find(left_hem==networks(n)));
num_verts_per_net(n,3) = length(find(right_hem==networks(n)));
count = 1;
for s = 1:size(left_hem,2)
per_subject(s,count) = s;
count = count + 1;
per_subject(s,count) = length(find(left_hem(:,s)==networks(n)));
count = count + 1;
per_subject(s,count) = length(find(right_hem(:,s)==networks(n)));
count = count + 1;
end
end
clear per_subject
for s = 1:size(left_hem,2)
count = 1;
for n = 1:length(networks)
per_subject(s,count) = s;
count = count + 1;
per_subject(s,count) = length(find(left_hem(:,s)==networks(n)));
count = count + 1;
per_subject(s,count) = length(find(right_hem(:,s)==networks(n)));
count = count + 1;
end
end
networks_by_hem
%-- 5/22/21, 3:15 PM --%
networks_by_hem
startup
networks_by_hem
left_hem(:,s)
find(left_hem(:,s)==networks(n))
clear per
clear per_subject
for s = 1:size(left_hem,2)
count = 1;
per_subject(s,count) = s;
count = count + 1;
for n = 1:length(networks)
per_subject(s,count) = s;
count = count + 1;
per_subject(s,count) = length(find(left_hem(:,s)==networks(n)));
count = count + 1;
per_subject(s,count) = length(find(right_hem(:,s)==networks(n)));
count = count + 1;
end
end
clear per_subject
for s = 1:size(left_hem,2)
count = 1;
per_subject(s,count) = s;
count = count + 1;
for n = 1:length(networks)
per_subject(s,count) = length(find(left_hem(:,s)==networks(n)));
count = count + 1;
per_subject(s,count) = length(find(right_hem(:,s)==networks(n)));
count = count + 1;
end
end
```

```

histogram(per_subject(:,6))
hold on
histogram(per_subject(:,7))
ax = gca
ax.FontSize = 15;
title('Frontoparietal Network Number of Vertices')
legend('Left Hem', 'Right Hem')
xlabel('Number of Vertices')
ylabel('Number of Subjects')
histogram(per_subject(:,14))
hold on
histogram(per_subject(:,15))
ax = gca
ax.FontSize = 15;
title('Ventral Attention Network')
legend('Left Hem', 'Right Hem')
xlabel('Number of Vertices')
ylabel('Number of Subjects')
histogram(per_subject(:,18))
hold on
histogram(per_subject(:,19))
ax = gca
ax.FontSize = 15;
title('Cingulo-opercular Network')
xlabel('Number of Vertices')
ylabel('Number of Subjects')
legend('Left Hem', 'Right Hem')
network_seg
atlas_dir = '/Users/dianaperez/Box/Dependencies/';
atlas = 'Seitzman300';
atlas_params = atlas_parameters_GrattonLab(atlas,atlas_dir);
corr_to_corr
datadir = '/Users/dianaperez/Box/corrmats_Seitzman300/';
corr_to_corr
mean_corrmat(1,:,:,:)
mean_corrmat(4,:,:,:)
mean_corrmat(3,:,:,:)
maskmat = ones(atlas_params.num_rois);
maskmat = logical(triu(maskmat,1));
count = 1;
for s = 1:6
    tmp = mean_corrmat(s,:,:,:);
    corrlin(count,:) = tmp(maskmat);
    count = count+1;
end
for s = 1:6
    figure_corrmat_GrattonLab(squeeze(mean_corrmat(s,:,:,:)),atlas_params, -0.4,1);
    title(['AllSessAvg, Subject ' subs{s}]);
    colormap('jet');
    saveas(gcf, [outdir 'Corrmat_' subs{s} 'sessmean.tiff'], 'tiff');
end
for s = 1:6
    figure_corrmat_GrattonLab(squeeze(mean_corrmat(s,:,:,:)),atlas_params, -1,1);
    title(['AllSessAvg, Subject ' subs{s}]);
    saveas(gcf, [outdir 'Corrmat_' subs{s} 'sessmean.tiff'], 'tiff');
end
mean_LS02 = mean_corrmat(1,:,:,:);
clear all
datadir = '/Users/dianaperez/Box/corrmats_Seitzman300/';
data = load(sprintf('%s/sub-%s/sub-%s_sess-%d_task-rest_corrmat_Seitzman300.mat',

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datadir, subs{sub}, subs{sub}, ses));
data = load(sprintf('%s/sub-%s/sub-%s_sess-%d_task-rest_corrmat_Seitzman300.mat',
datadir, 'LS02', 'LS02', 1));
mean_LS02_mat = data;
sess1_roi_timeseries = data.sess_roi_timeseries;
sess_roi_timeseries = [];
data.sess_roi_timeseries = [];
mean_LS02_mat.sess_roi_timeseries{1} = sess1_roi_timeseries;
mean_LS02_mat.sess_roi_timeseries = [];
mean_LS02_mat.sess_roi_timeseries{1} = sess1_roi_timeseries;
data = load(sprintf('%s/sub-%s/sub-%s_sess-%d_task-rest_corrmat_Seitzman300.mat',
datadir, 'LS02', 'LS02', 2));
sess2_roi_timeseries = data.sess_roi_timeseries;
mean_LS02_mat.sess_roi_timeseries{2} = sess2_roi_timeseries;
data_1 = load(sprintf('%s/sub-%s/sub-%s_sess-%d_task-rest_corrmat_Seitzman300.mat',
datadir, 'LS02', 'LS02', 1));
ses1_corrmat = data_1.corrmat;
ses1_corrmat = data_1.corrmat;
ses2_corrmat = data.corrmat;
data_3 = load(sprintf('%s/sub-%s/sub-%s_sess-%d_task-rest_corrmat_Seitzman300.mat',
datadir, 'LS02', 'LS02', 3));
ses3_corrmat = data_3.corrmat;
mean_corrmat(1,:,:,:) = data_1.corrmat;
mean_corrmat(1,:,:,:) = data_1.corrmat;
mean_corrmat(2,:,:,:) = data.corrmat;
mean_corrmat(3,:,:,:) = data_3.corrmat;
mean_corrmat = squeeze(mean(mean_corrmat));
sub-LS02_mean_corrmat = mean_corrmat;
LS02_mean_corrmat = mean_corrmat;
clear all
datadir = '/Users/dianaperez/Box/corrmats_Seitzman300/';
data = load(sprintf('%s/sub-%s/sub-%s_sess-%d_task-rest_corrmat_Seitzman300.mat',
datadir, subs{sub}, subs{sub}, ses));
data_1 = load(sprintf('%s/sub-%s/sub-%s_sess-%d_task-rest_corrmat_Seitzman300.mat',
datadir, 'LS03', 'LS03', 1));
data_2 = load(sprintf('%s/sub-%s/sub-%s_sess-%d_task-rest_corrmat_Seitzman300.mat',
datadir, 'LS03', 'LS03', 2));
data_3 = load(sprintf('%s/sub-%s/sub-%s_sess-%d_task-rest_corrmat_Seitzman300.mat',
datadir, 'LS03', 'LS03', 3));
data_4 = load(sprintf('%s/sub-%s/sub-%s_sess-%d_task-rest_corrmat_Seitzman300.mat',
datadir, 'LS03', 'LS03', 4));
data_5 = load(sprintf('%s/sub-%s/sub-%s_sess-%d_task-rest_corrmat_Seitzman300.mat',
datadir, 'LS03', 'LS03', 5));
mean_corrmat(1,:,:,:) = data_1.corrmat;
mean_corrmat(2,:,:,:) = data_2.corrmat;
mean_corrmat(3,:,:,:) = data_3.corrmat;
mean_corrmat(4,:,:,:) = data_4.corrmat;
mean_corrmat(5,:,:,:) = data_5.corrmat;
sub-LS03_mean_corrmat = squeeze(mean(mean_corrmat));
LS03_mean_corrmat = squeeze(mean(mean_corrmat));
clear data*
datadir = '/Users/dianaperez/Box/corrmats_Seitzman300/';
clear LS03_mean_corrmat mean_corrmat
data_1 = load(sprintf('%s/sub-%s/sub-%s_sess-%d_task-rest_corrmat_Seitzman300.mat',
datadir, 'LS05', 'LS05', 1));
data_2 = load(sprintf('%s/sub-%s/sub-%s_sess-%d_task-rest_corrmat_Seitzman300.mat',
datadir, 'LS05', 'LS05', 2));
data_3 = load(sprintf('%s/sub-%s/sub-%s_sess-%d_task-rest_corrmat_Seitzman300.mat',
datadir, 'LS05', 'LS05', 3));
data_4 = load(sprintf('%s/sub-%s/sub-%s_sess-%d_task-rest_corrmat_Seitzman300.mat',
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datadir, 'LS05', 'LS05', 4));
data_5 = load(sprintf('%s/sub-%s/sub-%s_sess-%d_task-rest_corrmat_Seitzman300.mat', ↵
datadir, 'LS05', 'LS05', 5));
mean_corrmat(1,:,:,:) = data_1.corrmat;
mean_corrmat(2,:,:,:) = data_2.corrmat;
mean_corrmat(3,:,:,:) = data_3.corrmat;
mean_corrmat(4,:,:,:) = data_4.corrmat;
mean_corrmat(5,:,:,:) = data_5.corrmat;
LS05_mean_corrmat = squeeze(mean(mean_corrmat));
clear mean_corrmat data_*
clear LS05_mean_corrmat
data_1 = load(sprintf('%s/sub-%s/sub-%s_sess-%d_task-rest_corrmat_Seitzman300.mat', ↵
datadir, 'LS05', 'LS05', 1));
data_1 = load(sprintf('%s/sub-%s/sub-%s_sess-%d_task-rest_corrmat_Seitzman300.mat', ↵
datadir, 'LS07', 'LS07', 1));
data_2 = load(sprintf('%s/sub-%s/sub-%s_sess-%d_task-rest_corrmat_Seitzman300.mat', ↵
datadir, 'LS07', 'LS07', 2));
mean_corrmat(1,:,:,:) = data_1.corrmat;
mean_corrmat(2,:,:,:) = data_2.corrmat;
LS07_mean_corrmat = squeeze(mean(mean_corrmat));
atlas_dir = '/Users/dianaperez/Box/Dependencies/';
atlas_params = atlas_parameters_GrattonLab('Seitzman300',atlas_dir);
permute_flippedVars
fcimage_corrmat_volume
load('/Volumes/GRATTONLAB/HCP/Diana/HCP_variants/border0rInsert.mat')
clear all
load('/Volumes/GRATTONLAB/HCP/Diana/HCP_variants/border0rInsert.mat')
permute_flippedVars
load('net_clusters_left.mat')
load('net_clusters_right.mat')
load('cluster_info_left.mat')
load('cluster_info_right.mat')
varmap_1 = ft_read_cifti_mod↵
('/Volumes/GRATTONLAB/HCP/Variants/split/100206_uniqueIDs_afterReassign.dtseries.nii')
varmap_2 = ft_read_cifti_mod↵
('/Users/dianaperez/Desktop/Lateralization_thresh10/variantMaps_04282021/100206_Threshold↵
edVariantMap_SNRExclude_10.dtseries.nii');
corr(varmap_1.data, varmap_2.data)
find(varmap_1.data>0);
vars = find(varmap_1.data>0);
vars_2 = find(varmap_2.data>0);
varmap_1 = ft_read_cifti_mod↵
('/Volumes/GRATTONLAB/HCP/Variants/split/100206_uniqueIDs_afterReassign.dtseries.nii')
unique(varmap_1.data)
varmap_1 = ft_read_cifti_mod('/Volumes/GRATTONLAB/HCP/Variants/split/100206_reassigned.↵
dtseries.nii')
unique(varmap_1.data)
load('VariantsvsFlippedVariants_allSubs_true_spCorr_1000_thresh_10_permutations_2.mat')
load('VariantsvsFlippedVariants_allSubs_trueDiffMap_1000_thresh_10_permutations_2.mat')
permutations_cluster_correction
addpath(genpath('/Users/dianaperez/Box/Dependencies'))
permutations_cluster_correction
%-- 6/3/21, 12:23 PM --%
load↵
('/Users/dianaperez/Desktop/VariantsvsFlippedVariants_MSC_true_spCorr_1000_thresh_10_perm↵
utations_2.mat')
load↵
('/Users/dianaperez/Desktop/VariantsvsFlippedVariants_MSC_spCorr_1000_thresh_10_permutati↵
ons.mat')
figure
```

```

plot(1:1000, spCorrs, 'bo')
ylim([.1 1])
plot(500, true_spCorr, 'ro', 'MarkerFaceColor', 'r')
hold on
plot(1:1000, spCorrs, 'bo')
ylim([.1 1])
ylabel('r value');
xlabel('Permutations');
title(['Left Hem vs Pseudo Left Hem Variant Maps - MSC']);
m = findobj(gca, 'Type', 'line');
hleg1 = legend(m(1:2), 'randomized maps', 'true map', 'Location', 'SouthWest');
hleg1.FontSize = 14;
ax = gca;
ax.FontSize = 14;
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.5, 0.7]);
print(gcf,
['/Users/dianaperez/Desktop/VariantsvsFlippedVariants_MSC_spCorr_10_thresh_1000_permutati
ons_results.jpg'], '-dpng', '-r300');
close gcf
title(['Left Hem vs Pseudo Left Hem Variant Maps - MSC']);
ylabel('r value');
xlabel('Permutations');
title(['Left Hem vs Pseudo Left Hem Variant Maps - MSC']);
m = findobj(gca, 'Type', 'line');
hleg1 = legend(m(1:2), 'randomized maps', 'true map', 'Location', 'SouthWest');
hleg1.FontSize = 14;
ax = gca;
ax.FontSize = 14;
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.5, 0.7]);
print(gcf,
['/Users/dianaperez/Desktop/VariantsvsFlippedVariants_MSC_spCorr_10_thresh_1000_permutati
ons_results.jpg'], '-dpng', '-r300');
close gcf
load('/Users/dianaperez/Desktop/varcount384split.mat')
load('/Users/dianaperez/Desktop/vertcount384split.mat')
left_cluster = var_count(3,:);
left_cluster = var_count(:,3);
right_cluster = var_count(:,4);
left_verts = vert_count(:,3);
right_verts = vert_count(:,4);
histogram(left_cluster)
hold on
histogram(right_cluster)
legend('Left Hem', 'Right Hem')
title('Variant Clusters by Hemisphere, 384 HCP subs split variants')
histogram(left_verts)
hold on
histogram(right_verts)
legend('Left Hem', 'Right Hem')
title('Variant Vertices by Hemisphere, 384 HCP subs split variants')
load('/Users/dianaperez/Desktop/varcountMSC.mat')
load('/Users/dianaperez/Desktop/vertcountMSC.mat')
left_cluster = var_count(1,:);
right_cluster = var_count(2,:);
left_vert = vert_count(1,:);
right_vert = vert_count(2,:);
histogram(left_cluster)
left_cluster = var_count(:,1);
right_cluster = var_count(:,2);
left_vert = vert_count(:,1);

```



```

right_vert = vert_count(:,2);
histogram(left_cluster)
hold on
histogram(right_cluster)
HCP_assignNetworksToVariants
load('goodSubs384.mat')
HCP_assignNetworksToVariants
addpath '/Users/dianaperez/Documents/GitHub/Lifespan-Analysis'
HCP_assignNetworksToVariants
%-- 6/14/21, 12:19 AM --%
load
('Users/dianaperez/Desktop/384_original_variants/VariantsvsFlippedVariants_allSubs_true_
spCorr_1000_thresh_10_permutations_2.mat')
load
('Users/dianaperez/Desktop/384_original_variants/VariantsvsFlippedVariants_allSubs_spCor
r_1000_thresh_10_permutations.mat')
plot(1:1000, spCorrs, 'bo')
y([.2 1])
ylim([.2 1])
hold on
plot(500, true_spCorr, 'ro', 'MarkerFaceColor', 'r')
HCP_assignNetworksToVariants
addpath '/Users/dianaperez/Documents/GitHub/Lifespan-Analysis'
HCP_assignNetworksToVariants
load('goodSubs384.mat')
find(subs==955465)
find(subs==955465)
%-- 6/19/21, 11:05 AM --%
assignVars_HCP_dice
addpath '/Users/dianaperez/Documents/GitHub/Lifespan-Analysis'
assignVars_HCP_dice
addpath(genpath('/Users/dianaperez/Box/Dependencies/cifti-matlab-master/'))
assignVars_HCP_dice
read_nifti2_hdr
assignVars_HCP_dice
%-- 6/20/21, 10:43 PM --%
assignVars_HCP_dice
addpath '/Users/dianaperez/Box/Dependencies/cifti-matlab-master'
assignVars_HCP_dice
addpath '/Users/dianaperez/Documents/GitHub/Lifespan-Analysis'
assignVars_HCP_dice
%-- 6/21/21, 9:49 AM --%
assignVars_HCP_dice
%-- 6/21/21, 11:17 AM --%
assignVars_HCP_dice
addpath '/Users/dianaperez/Box/Dependencies/cifti-matlab-master'
assignVars_HCP_dice
addpath '/Users/dianaperez/Documents/GitHub/Lifespan-Analysis'
assignVars_HCP_dice
%-- 6/21/21, 3:44 PM --%
assignVars_HCP_dice
addpath '/Users/dianaperez/Documents/GitHub/Lifespan-Analysis'
assignVars_HCP_dice
load('/Users/dianaperez/Desktop/HCP384_split_WashUTemp_variants_info.mat')
load('/Users/dianaperez/Desktop/HCP384_split_WashUTemp_networksxHem.mat')
addpath(genpath('/Users/dianaperez/Box/Dependencies/cifti-matlab-master'))
addpath(genpath('/Users/dianaperez/Box/Research/Lateralization_Variants'))
dataLoc =
'/Users/dianaperez/Box/Research/Lateralization_thresh10/variantMaps_042821/reassigned/';
%location of variant maps

```

```
load('goodSubs384.mat')
subs = goodSubs384;
outfile = '/Users/dianaperez/Desktop/HCP_VANvariants_OverlapMap_384GoodSubs_split.
dtseries.nii';
for s = 1:subs
% load subject data
file = [dataLoc num2str(subs(s)) '_reassigned.dtseries.nii'];
cifti = ft_read_cifti_mod(file);
for q = 1:length(cifti.data)
if cifti.data(q) == 7
allsubs(q,s) = 1;
else
allsubs(q,s) = 0;
end
end
end
template = cifti;
groupsum = sum(allsubs,2);
disp(['The maximum number of subjects that overlap in the ' group ' group is ' num2str
(max(groupsum))])
groupmap = sum(allsubs,2)/nfiles;
disp(['The maximum proportion of subjects that overlap in the ' group ' group is '
num2str(max(groupmap))])
overlap_map = template;
overlap_map.data = groupmap;
ft_write_cifti_mod(outfile, overlap_map);
Overlap_VANvars
clear all
Overlap_VANvars
find(cifti.data==7)
sum(allsubs(:,6))
sum(find(cifti.data==7))
count(find(cifti.data==7))
find(cifti.data==7)
sum(allsubs(:,5))
template = cifti;
groupsum = sum(allsubs,2);
disp(['The maximum number of subjects that overlap in the ' group ' group is ' num2str
(max(groupsum))])
groupmap = sum(allsubs,2)/nfiles;
disp(['The maximum proportion of subjects that overlap in the ' group ' group is '
num2str(max(groupmap))])
overlap_map = template;
overlap_map.data = groupmap;
template = cifti;
groupsum = sum(allsubs,2);
disp(['The maximum number of subjects that overlap in the group is ' num2str(max
(groupsum))])
groupmap = sum(allsubs,2)/nfiles;
disp(['The maximum proportion of subjects that overlap in the group is ' num2str(max
(groupmap))])
overlap_map = template;
overlap_map.data = groupmap;
template = cifti;
groupsum = sum(allsubs,2);
disp(['The maximum number of subjects that overlap in the group is ' num2str(max
(groupsum))])
groupmap = sum(allsubs,2)/384;
disp(['The maximum proportion of subjects that overlap in the group is ' num2str(max
(groupmap))])
```

```
overlap_map = template;
overlap_map.data = groupmap;
ft_write_cifti_mod(outfile, overlap_map);
Overlap_VANvars
load('/Users/dianaperez/Desktop/HCP384_split_WashUTemp_variants_info.mat')
count(variants_info.left_hem.indiv_vars==3)
count(variants_info.left_hem.indiv_vars==3)
find(variants_info.left_hem.indiv_vars==3)
variants_info.left_hem.indiv_vars = sortrows(variants_info.left_hem.↵
indiv_vars, 'network', 'ascend');
network = variants_info.left_hem.indiv_vars{929:1309,3};
%-- 6/23/21, 3:07 PM --%
load('HCP384_split_WashUTemp_variants_info.mat')
numVars = variants_info.left_hem.group_avg{:,1};
numVars_LH = numVars;
numVars1 = variants_info.right_hem.group_avg{:,1};
numVars_RH = numVars1;
clear numVars numVars1
hist(numVars_LH)
hold on
hist(numVars_RH)
close hist
close all
histogram(numVars_LH)
hold on
histogram(numVars_RH)
title('Number of Variant Clusters in Each Hemisphere')
ylabel(
ylabel('Number of Variants')
xlabel('Number of Subjects')
m = findobj(gca, 'Type', 'line')
hleg1 = legend(m(1:2), 'Left Hem', 'Right Hem', 'Location', 'SouthWest');
legend('Left Hem', 'Right Hem', 'Location', 'SouthWest');
legend('Left Hem', 'Right Hem', 'Location', 'NorthEast');
ax = gca;
ax.FontSize = 14;
print(gcf, '/Users/dianaperez/Desktop/NumVarsxHem.jpg', '-dpng', '-r300');
clear ax m
clear numVars*
avgVarSize = variants_info.right_hem.group_avg{:,3};
avgVarSize_RH = avgVarSize;
clear avgVarSize
avgVarSize = variants_info.left_hem.group_avg{:,3};
avgVarSize_LH = avgVarSize;
clear avgVarSize
histogram(avgVarSize_LH)
close all
histogram(avgVarSize_LH)
hold on
histogram(avgVarSize_RH)
title('Average Variant Size in Each Hemisphere')
ylabel('Number of Vertices')
ylabel('Average Number of Vertices')
xlabel('Number of Subjects')
legend('Left Hem', 'Right Hem', 'Location', 'NorthEast');
ax = gca;
ax.FontSize = 14;
print(gcf, '/Users/dianaperez/Desktop/AvgVarSizexHem.jpg', '-dpng', '-r300');
[p_fdr, p_masked]= FDR([0.439504962 0.128675384 0.000274487 0.253014009 0.015014079], .↵
05);
```

```
clear ax avgVarSize* p*
numBorderVars = variants_info.left_hem.group_avg{:,4};
numBorderVars_LH = variants_info.left_hem.group_avg{:,4};
clear numBorderVars
numBorderVars_RH = variants_info.right_hem.group_avg{:,4};
histogram(numBorderVars_LH)
close all
histogram(numBorderVars_LH)
histogram(numBorderVars_RH)
histogram(numBorderVars_LH)
hold on
histogram(numBorderVars_RH)
title('Number of Border Variants in Each Hemisphere')
ylabel('Number of Variants')
xlabel('Number of Subjects')
legend('Left Hem', 'Right Hem', 'Location', 'NorthEast');
ax = gca;
ax.FontSize = 14;
print(gcf, '/Users/dianaperez/Desktop/BorderVarsxHem.jpg', '-dpng', '-r300');
clear numBorder* ax
numEctopicVars = variants_info.right_hem.group_avg{:,5};
numEctopicVars_RH = variants_info.right_hem.group_avg{:,5};
clear numEctopicVars
numEctopicVars_LH = variants_info.left_hem.group_avg{:,5};
histogram(numEctopicVars_LH)
hold on
histogram(numEctopicVars_RH)
title('Number of Ectopic Variants in Each Hemisphere')
ylabel('Number of Variants')
xlabel('Number of Subjects')
legend('Left Hem', 'Right Hem', 'Location', 'NorthEast');
ax = gca;
ax.FontSize = 14;
close all
histogram(numBorderVars_LH)
histogram(numEctopicVars_LH)
hold on
histogram(numEctopicVars_RH)
title('Number of Ectopic Variants in Each Hemisphere')
ylabel('Number of Variants')
xlabel('Number of Subjects')
legend('Left Hem', 'Right Hem', 'Location', 'NorthEast');
ax = gca;
ax.FontSize = 14;
print(gcf, '/Users/dianaperez/Desktop/EctopicVarsxHem.jpg', '-dpng', '-r300');
pvals = [0.000000011 0.121789080 0.000306024 0.086926523 0.000058774 0.468638463
0.029082589 0.234696159 0.094306337 0.041385242 1.000000000 0.622875291 0.879335992];
[p_fdr, p_masked]= FDR(pvals, .05);
load('HCP384_original_variants_info.mat')
clear ax num* p* pvals
numVars = variants_info.left_hem.group_avg{1,1};
LH_numVars = variants_info.left_hem.group_avg{1,1};
RH_numVars = variants_info.right_hem.group_avg{1,1};
clear numVars
LH_numVars = variants_info.left_hem.group_avg{1,:};
LH_numVars = variants_info.left_hem.group_avg{:,1};
RH_numVars = variants_info.right_hem.group_avg{:,1};
histogram(LH_numVars)
close all
histogram(LH_numVars)
```

```
hold on
histogram(RH_numVars)
title('Number of Variants in Each Hemisphere')
ylabel('Number of Variant Clusters')
xlabel('Number of Subjects')
legend( 'Left Hem', 'Right Hem', 'Location', 'NorthEast');
ax = gca;
ax.FontSize = 14;
print(gcf, '/Users/dianaperez/Desktop/NumVarsxHem_384HCPog.jpg', '-dpng', '-r300');
avgVarSize = variants_info.left_hem.group_avg{:,3};
LH_avgVarSize = variants_info.left_hem.group_avg{:,3};
RH_avgVarSize = variants_info.right_hem.group_avg{:,3};
close all
histogram(LH_avgVarSize)
hold on
histogram(RH_avgVarSize)
title('Average Variant Size in Each Hemisphere')
ylabel('Number of Vertices')
xlabel('Number of Subjects')
legend( 'Left Hem', 'Right Hem', 'Location', 'NorthEast');
ax = gca;
ax.FontSize = 14;
xlabel('Number of Vertices')
ylabel('Number of Subjects')
print(gcf, '/Users/dianaperez/Desktop/AvgSizexHem_384HCPog.jpg', '-dpng', '-r300');
LH_numVars = variants_info.left_hem.group_avg{:,1};
RH_numVars = variants_info.right_hem.group_avg{:,1};
histogram(LH_numVars)
hold on
histogram(RH_numVars)
title('Number of Variants in Each Hemisphere')
xlabel('Number of Vertices')
ylabel('Number of Subjects')
legend( 'Left Hem', 'Right Hem', 'Location', 'NorthEast');
ax = gca;
ax.FontSize = 14;
close all
histogram(LH_numVars)
hold on
histogram(RH_numVars)
title('Number of Variants in Each Hemisphere')
xlabel('Number of Vertices')
ylabel('Number of Subjects')
legend( 'Left Hem', 'Right Hem', 'Location', 'NorthEast');
ax.FontSize = 14;
ax = gca;
ax.FontSize = 14;
print(gcf, '/Users/dianaperez/Desktop/NumVarsxHem_384HCPog.jpg', '-dpng', '-r300');
xlabel('Number of Variant Clusters')
print(gcf, '/Users/dianaperez/Desktop/NumVarsxHem_384HCPog.jpg', '-dpng', '-r300');
Overlap_VANvars
clear all
uiopen('/Users/dianaperez/Desktop/120_LR_minsize400_recolored_manualconsensus_LR.dlabel.
nii',1)
average = ft_read_cifti_mod
('/Users/dianaperez/Desktop/120_LR_minsize400_recolored_manualconsensus_LR.dlabel.nii');
find_former_networks
for net = 1:16
num_verts = length(find(allsubs==net));
disp(['Number of ' network_names{NOI} ' variant vertices that were formerly assigned to '

```

```

network_names{net} ' is: ' num2str(num_verts));
end
network_names{NOI}
assignVars_HCP_dice
sum = length(find(allsubs>0));
sum = length(find(allsubs>0));
network_names = {'DMN' 'Vis' 'FP' 'DAN' 'VAN' 'Sal' 'CO' 'SMd' 'SML'
'Aud' 'Tpole' 'MTL' 'PMN' 'PON'};
old_nets = [];
for net = 1:16
num_verts = length(find(allsubs==net));
old_nets(1,net) = num_verts;
old_nets(1,net) = num_verts/sum;
disp(['Number of ' network_names{NOI} ' variant vertices that were formerly assigned to '
network_names{net} ' is: ' num2str(num_verts)]);
end
network_names = {'DMN' 'Vis' 'FP' 'Reward' 'Unassign' 'DAN' 'VAN' 'Sal'
'CO' 'SMd' 'SML' 'Aud' 'Tpole' 'MTL' 'PMN' 'PON'};
old_nets = [];
for net = 1:16
num_verts = length(find(allsubs==net));
old_nets(1,net) = num_verts;
old_nets(1,net) = num_verts/sum;
disp(['Number of ' network_names{NOI} ' variant vertices that were formerly assigned to '
network_names{net} ' is: ' num2str(num_verts)]);
end
old_nets = [];
for net = 1:16
num_verts = length(find(allsubs==net));
old_nets(1,net) = num_verts;
old_nets(2,net) = num_verts/sum;
disp(['Number of ' network_names{NOI} ' variant vertices that were formerly assigned to '
network_names{net} ' is: ' num2str(num_verts)]);
end
VANvars_oldNets = old_nets;
find_former_networks
DMNvars_oldNets = old_nets;
find_former_networks
FPNvars_oldNets = old_nets;
pie(DMNvars_oldNets)
close all
pie(DMNvars_oldNets)
network_names = {'DMN' 'Vis' 'FP' 'Unassigned' 'DAN' 'Unassigned2' 'VAN'
'Sal' 'CO' 'SMd' 'SML' 'Aud' 'Tpole' 'MTL' 'PMN' 'PON' 'Consensus'};
%% The colors that correspond to the Power functional networks in connectome workbench
rgb_colors = [1 0 0; %% The colors that correspond to the Power functional networks in
MATLAB
0 0 .6;
1 1 0;
.67 .67 .67;
0 .8 0;
.67 .67 .67;
0 .6 .6;
0 0 0;
.3 0 .6;
.2 1 1;
1 .5 0;
.6 .2 1;
.2 1 .2;
0 .2 .4;

```

```

0 0 1;
.8 .8 .6;
.4 0 0];
networksexist = find(allsubs > 0);
network_legend = network_names(networksexist);
find_former_networks
networksexist = unique(find(allsubs > 0));
networksexist = unique(allsubs > 0);
networksexist = unique(allsubs);
networksexist = unique(allsubs);
networksexist = networksexits(2:end);
networksexist = networksexist(2:end);
network_legend = network_names(networksexist);
color_legend = rgb_colors(networksexist,:);
filename = [network_names{NOI} '_Variant_PieChart_Old_Networks.jpg'];
h = pie(old_nets(1,:));
h = pie(old_nets(1,:));
delete(findobj(h,'Type','text'))
for k = 1:length(h)/2
set(h(k*2-1), 'FaceColor', color_legend(k,:));
end
network_names = {'DMN' 'Vis' 'FP' 'Unassigned' 'DAN' 'Unassigned2' 'VAN'↵
'Sal' 'CO' 'SMd' 'SML' 'Aud' 'Tpole' 'MTL' 'PMN' 'PON'}; %% The colors↵
that correspond to the Power functional networks in connectome workbench
rgb_colors = [1 0 0; %% The colors that correspond to the Power functional networks in↵
MATLAB
0 0 .6;
1 1 0;
.67 .67 .67;
0 .8 0;
.67 .67 .67;
0 .6 .6;
0 0 0;
.3 0 .6;
.2 1 1;
1 .5 0;
.6 .2 1;
.2 1 .2;
0 .2 .4;
0 0 1;
.8 .8 .6];
for k = 1:length(h)/2
set(h(k*2-1), 'FaceColor', color_legend(k,:));
end
find_former_networks
h = pie(old_nets(1,networksexist));
print(gcf,['/Users/dianaperez/Desktop/' filename],'-dpng','-r300');
find_former_networks
permute_flippedVars
LH_files{1}
permute_flippedVars
permute_left_right_handlers
min(spCorrs)
permutations_cluster_correction
%-- 6/29/21, 11:40 AM --%
permutations_cluster_correction
permutations_create_zmap
outfile = '/Users/dianaperez/Desktop/RightvsNonRightHanders_zmap.dtseries.nii';
ft_write_cifti_mod(outfile, template_cifti);
permute_left_right_handlers

```

```

find(spCorrs<=true_spCorr)
4/1000
load
('/Users/dianaperez/Box/Research/Lateralization_thresh10/Permutation_Tests/PseudoRightvsP
pseudoLHAnders_spCorr_1000perms_thresh_10.mat')
load
('/Users/dianaperez/Box/Research/Lateralization_thresh10/Permutation_Tests/RightvsLeftHan
ders_true_spCorr_1000_thresh_10.mat')
find(spCorrs<=true_spCorr)
4/1000
load
('/Users/dianaperez/Box/Research/Lateralization_thresh10/Permutation_Tests/PseudoRightvsP
pseudoNonRHanders_spCorr_1000perms_thresh_10.mat')
load
('/Users/dianaperez/Box/Research/Lateralization_thresh10/Permutation_Tests/RightvsnonRigh
tHanders_true_spCorr_1000_thresh_10.mat')
find(spCorrs<=true_spCorr)
1/1000
permutations_cluster_correction
permutations_create_zmap
load
('/Users/dianaperez/Box/Research/Lateralization_thresh10/Permutation_Tests/RightvsLeftHan
ders_true_RH-LH_diff_1000_thresh_10.mat')
clear all
load
('/Users/dianaperez/Box/Research/Lateralization_thresh10/Permutation_Tests/RightvsLeftHan
ders_true_RH-LH_diff_1000_thresh_10.mat')
template =
'/Volumes/GRATTONLAB/Lifespan/BIDS/Nifti/derivatives/postFCproc_CIFTI/dconn_cifti_normalw
all/sub-LS05_vs_120_allsubs_sptlCorr_cortex_smooth_2.55.dtseries.nii';
template.data = trueDiff;
template_cifti = ft_read_cifti_mod(template);
template_cifti.data = trueDiff;
ft_write_cifti_mod('/Users/dianaperez/Desktop/RH-LH_diffmap_HCP.dtseries.nii',
template_cifti);
load
('/Users/dianaperez/Box/Research/Lateralization_thresh10/Permutation_Tests/RightvsnonRigh
tHanders_true_RH-nonRH_diff_1000_thresh_10.mat')
template_cifti.data = trueDiff;
ft_write_cifti_mod('/Users/dianaperez/Desktop/RH-nonRH_diffmap_HCP.dtseries.nii',
template_cifti);
clear all
load('/Users/dianaperez/Desktop/HCP384_LH_split_networksxHem.mat')
load('/Users/dianaperez/Desktop/HCP384_LH_split_variants_info.mat')
networks_LH = networksxHem;
variants_LH = variants_info;
load('/Users/dianaperez/Desktop/HCP384_RH_split_networksxHem.mat')
load('/Users/dianaperez/Desktop/HCP384_RH_split_variants_info.mat')
networks_RH = networksxHem;
variants_RH = variants_info;
clear networksxHem variants_info
load('/Users/dianaperez/Desktop/HCP384_nonRH_split_networksxHem.mat')
load('/Users/dianaperez/Desktop/HCP384_nonRH_split_variants_info.mat')
test_mat = [];
for net = 1:16
end
count = 1;
for net = 1:16
test_mat(:,count) = networksxHem.clustersLH(net);
count = count + 1;

```



```
test_mat(:,count) = networksxHem.clustersRH(net);
count = count + 1;
end
for net = 1:16
test_mat(:,count) = networksxHem.clustersLH(:,net);
count = count + 1;
test_mat(:,count) = networksxHem.clustersRH(:,net);
count = count + 1;
end
test_mat(:,1) = networksxHem.clustersLH(:,1);
test_mat = [];
test_mat(:,1) = networksxHem.clustersLH(:,1);
test_mat = [];
for net = 1:16
test_mat(:,count) = networksxHem.clustersLH(:,net);
count = count + 1;
test_mat(:,count) = networksxHem.clustersRH(:,net);
count = count + 1;
end
for net = 1:16
test_mat(:,count) = networks_LH.clustersLH(:,net);
count = count + 1;
test_mat(:,count) = networks_LH.clustersRH(:,net);
count = count + 1;
end
test_mat = [];
for net = 1:16
test_mat(:,count) = networks_LH.clustersLH(:,net);
count = count + 1;
test_mat(:,count) = networks_LH.clustersRH(:,net);
count = count + 1;
end
test_mat = [];
for net = 1:16
test_mat(:,count) = networks_RH.clustersLH(:,net);
count = count + 1;
test_mat(:,count) = networks_RH.clustersRH(:,net);
count = count + 1;
end
test_mat = [];
for net = 1:16
test_mat(:,count) = networks_RH.clustersLH(:,net);
count = count + 1;
test_mat(:,count) = networks_RH.clustersRH(:,net);
count = count + 1;
end
test_mat = [];
count = 1;
for net = 1:16
test_mat(:,count) = networks_LH.clustersLH(:,net);
count = count + 1;
test_mat(:,count) = networks_LH.clustersRH(:,net);
```

```
count = count + 1;
end
test_mat = [];
count = 1;
for net = 1:16
test_mat(:,count) = networksxHem.clustersLH(:,net);
count = count + 1;
test_mat(:,count) = networksxHem.clustersRH(:,net);
count = count + 1;
end
test_mat = [];
count = 1;
for net = 1:16
test_mat(:,count) = networks_RH.clustersLH(:,net);
count = count + 1;
test_mat(:,count) = networks_RH.clustersRH(:,net);
count = count + 1;
end
[p_fdr, p_masked] = FDR([0.030170961
0.170336911
0.376937036
0.148030671
0.533538493
0.574763999
0.044798574
0.664306034
0.713950162
0.713950162
0.327715806
0.382384794], 0.05)
[p_fdr, p_masked] = FDR([0.00000008
0.44393710
0.00035698
0.12366324
0.00000741
0.57929040
0.01332763
0.19043900
0.10892350
0.06271794
1.00000000], 0.05)
[p_fdr, p_masked] = FDR([0.00000008
0.44393710
0.00035698
0.12366324
0.00000741
0.57929040
0.01332763
0.19043900
0.10892350
0.06271794
1.00000000], 0.05)
load('/Users/dianaperez/HCP752_LH_og_networksxHem.mat')
load('/Users/dianaperez/HCP752_LH_og_variants_info.mat')
variants_LHand = variants_info;
networks_LHand = networksxHem;
clear all
networks_Lhand = load('/Users/dianaperez/HCP752_LH_og_networksxHem.mat')
load('/Users/dianaperez/HCP752_LH_og_networksxHem.mat')
clear all
```

```
load('/Users/dianaperez/HCP752_LH_og_networksxHem.mat')
load('/Users/dianaperez/HCP752_LH_og_variants_info.mat')
variants_Lhand = variants_info;
load('/Users/dianaperez/HCP752LH_og_variants_info.mat')
variants_Rhand = variants_info;
clear variants_info networksxHem
[p_fdr, p_masked] = FDR([0.00000001
0.12178908
0.00030602
0.08692652
0.00005877
0.46863846
0.02908259
0.23469616
0.09430634
0.04138524
1.00000000
0.62287529
0.87933599], 0.05)
[p_fdr, p_masked] = FDR([0.00000001
0.12178908
0.00030602
0.08692652
0.00005877
0.46863846
0.02908259
0.23469616
0.09430634
0.04138524
1.00000000
0.62287529
0.87933599], 0.01)
[p_fdr, p_masked] = FDR([0.022453292
0.966626019
0.03172435
0.035036299], 0.05)
clear all
ft_read_cifti
ft_read_cifti_mod✓
('/Users/dianaperez/Desktop/HCP_FPNvariants_OverlapMap_384GoodSubs_split.dtseries.nii');
FPN_overlap = ans.data;
ft_read_cifti_mod✓
('/Users/dianaperez/Desktop/HCP_VANvariants_OverlapMap_384GoodSubs_split.dtseries.nii');
VAN_overlap = ans.data;
ft_read_cifti_mod✓
('/Users/dianaperez/Desktop/HCP_DMNvariants_OverlapMap_384GoodSubs_split.dtseries.nii');
DMN_overlap = ans.data;
clear ans
cifti = ft_read_cifti_mod✓
('/Users/dianaperez/Desktop/HCP_FPNvariants_OverlapMap_384GoodSubs_split.dtseries.nii');
for i = 1:length(cifti.brainstructure)
if cifti.brainstructure(i) > 0
end
end
count = 1;
for i = 1:length(cifti.brainstructure)
if cifti.brainstructure(i) > 0
if VAN_overlap(count) > 0
new_cifti(i,1) = VAN_overlap(count);
else new_cifti(i,1) = 0;
```

```
end
count = count + 1;
else new_cifti(i,1) = 0;
end
end
64984/2
VAN_new = new_cifti;
count = 1;
or i = 1:length(cifti.brainstructure)
if cifti.brainstructure(i) > 0
new_cifti(i,1) = VAN_overlap(count);
count = count + 1;
else new_cifti(i,1) = 0;
end
end
for i = 1:length(cifti.brainstructure)
if cifti.brainstructure(i) > 0
new_cifti(i,1) = VAN_overlap(count);
count = count + 1;
else new_cifti(i,1) = 0;
end
end
corr(new_cifti, VAN_new)
clear new_cifti
count = 1;
for i = 1:length(cifti.brainstructure)
if cifti.brainstructure(i) > 0
DMN_new(i,1) = DMN_overlap(count);
count = count + 1;
else DMN_new(i,1) = 0;
end
end
count = 1;
for i = 1:length(cifti.brainstructure)
if cifti.brainstructure(i) > 0
FPN_new(i,1) = FPN_overlap(count);
count = count + 1;
else FPN_new(i,1) = 0;
end
end
left_DMN = DMN_new(1:32492);
left_FPN = FPN_new(1:32492);
left_VAN = VAN_new(1:32492);
right_DMN = DMN_new(32493:end);
right_FPN = FPN_new(32493:end);
right_VAN = VAN_new(32493:end);
flip_DMN = [right_DMN; left_DMN];
flip_FPN = [right_FPN; left_FPN];
flip_VAN = [right_VAN; left_VAN];
flip_DMN = flip_DMN(cifti.brainstructure>0);
flip_FPN = flip_FPN(cifti.brainstructure>0);
flip_VAN = flip_VAN(cifti.brainstructure>0);
diff_DMN = DMN_overlap - flip_DMN;
diff_FPN = FPN_overlap - flip_FPN;
diff_VAN = VAN_overlap - flip_VAN;
template = cifti;
template.data = diff_FPN;
ft_write_cifti_mod('/Users/dianaperez/Desktop/HCP_FPNvariants_diffMap_384GoodSubs_split.
dtseries.nii', template);
template.data = diff_DMN;
```

```
ft_write_cifti_mod('/Users/dianaperez/Desktop/HCP_DMNvariants_diffMap_384GoodSubs_split.
dtseries.nii', template);
template.data = diff_VAN;
ft_write_cifti_mod('/Users/dianaperez/Desktop/HCP_VANvariants_diffMap_384GoodSubs_split.
dtseries.nii', template);
clear all
CreateVariantFiles_LS03specific
load('/Users/dianaperez/Desktop/Thesis/HCP384_LH_split_variants_info.mat')
load('/Users/dianaperez/Desktop/Thesis/HCP384_original_variants_info.mat')
load('/Users/dianaperez/Desktop/Thesis/HCP752_LH_og_variants_info.mat')
load('/Users/dianaperez/Desktop/Thesis/HCP752_RH_og_variants_info.mat')
clear all
load('/Users/dianaperez/Desktop/Thesis/HCP752_LH_og_variants_info.mat')
load('/Users/dianaperez/Desktop/Thesis/HCP752_RH_og_variants_info.mat')
LH_info = load('/Users/dianaperez/Desktop/Thesis/HCP752_LH_og_variants_info.mat');
RH_info = load('/Users/dianaperez/Desktop/Thesis/HCP752_RH_og_variants_info.mat');
addpath(genpath('/Users/dianaperez/Box/Dependencies/cifti-matlab-master'));
rootDir = '/Users/dianaperez/Box/Research/Lateralization_Variants/';
%rootDir = '/projects/p31161/lateralizationVariants/';
addpath(genpath(rootDir))
outputDir = [rootDir 'Permutation_Tests/'];
%% VARIABLES
numperms = 1000;% number of permutations
spCorrs=zeros(numperms,1);% initialize mat of values i'm permuting
%% LOAD files
LH_info = load('/Users/dianaperez/Desktop/Thesis/HCP752_LH_og_variants_info.mat');
RH_info = load('/Users/dianaperez/Desktop/Thesis/HCP752_RH_og_variants_info.mat');
LH_variantSize = [LH_info.variant_info.left_hem.group_avg(:,3); LH_info.variant_info.
right_hem.group_avg(:,3)];
RH_variantSize = [RH_info.variant_info.left_hem.group_avg(:,3); RH_info.variant_info.
right_hem.group_avg(:,3)];
LH_numVars = [LH_info.variant_info.left_hem.group_avg(:,1); LH_info.variant_info.
right_hem.group_avg(:,1)];
RH_numVars = [RH_info.variant_info.left_hem.group_avg(:,1); RH_info.variant_info.
right_hem.group_avg(:,1)];
LH_variantSize = [LH_info.variants_info.left_hem.group_avg(:,3); LH_info.variants_info.
right_hem.group_avg(:,3)];
RH_variantSize = [RH_info.variants_info.left_hem.group_avg(:,3); RH_info.variants_info.
right_hem.group_avg(:,3)];
LH_numVars = [LH_info.variants_info.left_hem.group_avg(:,1); LH_info.variants_info.
right_hem.group_avg(:,1)];
RH_numVars = [RH_info.variants_info.left_hem.group_avg(:,1); RH_info.variants_info.
right_hem.group_avg(:,1)];
numSubs = length(LH_variantSize) + length(RH_variantSize);
numSubs = height(LH_variantSize) + height(RH_variantSize);
clear all
%% PATHS
%addpath(genpath('/projects/b1081/Scripts'));
addpath(genpath('/Users/dianaperez/Box/Dependencies/cifti-matlab-master'));
rootDir = '/Users/dianaperez/Box/Research/Lateralization_Variants/';
%rootDir = '/projects/p31161/lateralizationVariants/';
addpath(genpath(rootDir))
outputDir = [rootDir 'Permutation_Tests/'];
%% VARIABLES
numperms = 1000;% number of permutations
spCorrs=zeros(numperms,1);% initialize mat of values i'm permuting
%% LOAD files
LH_info = load('/Users/dianaperez/Desktop/Thesis/HCP752_LH_og_variants_info.mat');
RH_info = load('/Users/dianaperez/Desktop/Thesis/HCP752_RH_og_variants_info.mat');
LH_variantSize = [LH_info.variants_info.left_hem.group_avg(:,3); LH_info.variants_info.
right_hem.group_avg(:,3)];
```

```
right_hem.group_avg(:,3)];
RH_variantSize = [RH_info.variants_info.left_hem.group_avg(:,3); RH_info.variants_info.
right_hem.group_avg(:,3)];
LH_numVars = [LH_info.variants_info.left_hem.group_avg(:,1); LH_info.variants_info.
right_hem.group_avg(:,1)];
RH_numVars = [RH_info.variants_info.left_hem.group_avg(:,1); RH_info.variants_info.
right_hem.group_avg(:,1)];
numSubs = length(LH_variantSize) + length(RH_variantSize);
help dir
cd /Users/dianaperez/Box
dir('105216_rfMRI_REST2_RL')
L = dir('105216_rfMRI_REST2_RL')
clear all
cd /Users/dianaperez/Box/DATA/
bold_dir = dir('BOLD_DATA');
for dir = 1:length(bold_dir)
    dir_files = dir(bold_dir.name(dir));
end
end
dir('105216_rfMRI_REST2_RL')
clear dir
dir('105216_rfMRI_REST2_RL')
d = dir('105216_rfMRI_REST2_RL')
bytes = [d.bytes].';
sum(bytes)
size_in_gig = ans/1000000000
clear all
cd /Users/dianaperez/Box/DATA/
bold_dir = dir('BOLD_DATA');
data_info = [];
count = 1;
d = 1'
dir_files = dir(bold_dir.name(d));
%% folder
data_info(count,1) = bold_dir.name(d);
%% number of files
data_info(count,2) = length(dir_files);
%% size
bytes = [dir_files.bytes].';
size_in_bytes = sum(bytes);
size_in_gigs = size_in_bytes/1000000000;
data_info(count,3) = size_in_gigs;
count = count + 1;
find_missing_HCPdata
bold_dir = bold_dir(3:end);
for i=1:4072
end
data_info = cell(4072,2);
for i=1:4072
    thisDir = dir(['/Users/dianaperez/Box/DATA/BOLD_DATA/' bold_dir(i).name]);
    data_info(i,1) = bold_dir(i).name;
    sizes = ([thisDir.bytes].')./1000000000;
    folder_sizes(i,2)=sum(sizes);
end
data_info = cell(4072,2);
for i=1:4072
    thisDir = dir(['/Users/dianaperez/Box/DATA/BOLD_DATA/' bold_dir(i).name]);
    data_info(i,1) = bold_dir(i).name;
    sizes = ([thisDir.bytes].')./1000000000;
    data_info(i,2)=sum(sizes);
```

```

end
for i=1:4072
thisDir = dir(['/Users/dianaperez/Box/DATA/BOLD_DATA/' bold_dir(i).name]);
data_info{i,1} = bold_dir(i).name;
sizes = ([thisDir.bytes].')./1000000000;
data_info{i,2}=sum(sizes);
end
clear bold_dir count d i sizes thisDir
load('/Users/dianaperez/Downloads/BOLD_DATA_folder_sizes.mat')
folder_match = cell(4072,1);
for i = 1:4072
if data_info(i,2) == folder_sizes(i,2)
folder_match(i,1) == 0
else
folder_match(i,1) == 1
end
end
for i = 1:4072
if data_info{i,2} == folder_sizes{i,2}
folder_match{i,1} == 0
else
folder_match{i,1} == 1
end
end
folder_match = [];
for i = 1:4072
if data_info{i,2} == folder_sizes{i,2}
folder_match(i,1) == 0
else
folder_match(i,1) == 1
end
end
for i = 1:4072
if data_info{i,2} == folder_sizes{i,2}
folder_match(i) == 0
else
folder_match(i) == 1
end
end
if data_info{1,2} == folder_sizes{1,2}
disp('they match')
end
find_missing_HCPdata
folder_match = folder_match';
find(folder_match==1)
data_info(1679)
data_info(3043)
data_info(3435)
data_info(1679,2 - folder_sizes{1672,2}
data_info{1679,2} - folder_sizes{1672,2}
data_info{3435,2} - folder_sizes{3435,2}
data_info{3043,2} - folder_sizes{3043,2}
clear all
load('/Users/dianaperez/Desktop/Thesis/HCP752_LH_og_variants_info.mat')
load('/Users/dianaperez/Desktop/Thesis/HCP752_RH_og_variants_info.mat')
permute_varSize
avgVarSize = LH_varSize{1,1};
permute_varSize
help interp1
permute_varSize

```

```

plot(1:1000, perm_corrs(:,1), 'bo', 'MarkerFaceColor', 'b')
hold on
plot(500, LHand_varSize_trueCorr, 'ro', 'MarkerFaceColor', 'r')
h = length(find(perm_corrs(:,1)<LHand_varSize_trueCorr))
h = length(find(perm_corrs(:,2)<RHand_varSize_trueCorr))
h = length(find(perm_corrs(:,2)<Rhand_varSize_trueCorr))
close gcf
plot(1:1000, perm_corrs(:,2), 'bo', 'MarkerFaceColor', 'b')
plot(500, Rhand_varSize_trueCorr, 'ro', 'MarkerFaceColor', 'r')
hold on
plot(1:1000, perm_corrs(:,2), 'bo', 'MarkerFaceColor', 'b')
close gcf
h = length(find(perm_corrs(:,3)<LHand_numVars_trueCorr))
h = length(find(perm_corrs(:,4)<RHand_numVars_trueCorr))
h = length(find(perm_corrs(:,4)<Rhand_numVars_trueCorr))
%-- 7/22/21, 12:32 PM --%
permute_varSize
length(find(perm_corrs(:,1)<LHand_varSize_trueCorr))/1000
disp(['P value for LHand varSize comparison is ' length(find(perm_corrs(:,1)<
<LHand_varSize_trueCorr))/1000])
disp(['P value for RHand varSize comparison is ' length(find(perm_corrs(:,2)<
<RHand_varSize_trueCorr))/1000])
disp(['P value for LHand numVars comparison is ' length(find(perm_corrs(:,3)<
<LHand_numVars_trueCorr))/1000])
disp(['P value for RHand numVars comparison is ' length(find(perm_corrs(:,4)<
<Rhand_numVars_trueCorr))/1000])
disp(['P value for LHand varSize diff comparison is ' length(find(perm_corrs(:,5)<
<LHand_varSize_diff))/1000])
disp(['P value for RHand varSize diff comparison is ' length(find(perm_corrs(:,6)<
<RHand_varSize_diff))/1000])
disp(['P value for LHand numVars diff comparison is ' length(find(perm_corrs(:,7)<
<LHand_numVars_diff))/1000])
disp(['P value for RHand numVars diff comparison is ' length(find(perm_corrs(:,8)<
<Rhand_numVars_diff))/1000])
disp(['P value for LHand varSize comparison is ' length(find(perm_corrs(:,1)<
<LHand_varSize_trueCorr))/1000])
disp(['P value for RHand varSize comparison is ' length(find(perm_corrs(:,2)<
<Rhand_varSize_trueCorr))/1000])
disp(['P value for LHand numVars comparison is ' length(find(perm_corrs(:,3)<
<LHand_numVars_trueCorr))/1000])
disp(['P value for RHand numVars comparison is ' length(find(perm_corrs(:,4)<
<Rhand_numVars_trueCorr))/1000])
disp(['P value for LHand varSize diff comparison is ' length(find(perm_corrs(:,5)<
<LHand_varSize_diff))/1000])
disp(['P value for RHand varSize diff comparison is ' length(find(perm_corrs(:,6)<
<RHand_varSize_diff))/1000])
disp(['P value for LHand numVars diff comparison is ' length(find(perm_corrs(:,7)<
<LHand_numVars_diff))/1000])
disp(['P value for RHand numVars diff comparison is ' length(find(perm_corrs(:,8)<
<Rhand_numVars_diff))/1000])
disp(['P value for LHand varSize comparison is ' num2str(length(find(perm_corrs(:,1)<
<LHand_varSize_trueCorr))/1000)])
disp(['P value for RHand varSize comparison is ' num2str(length(find(perm_corrs(:,2)<
<Rhand_varSize_trueCorr))/1000)])
disp(['P value for LHand numVars comparison is ' num2str(length(find(perm_corrs(:,3)<
<LHand_numVars_trueCorr))/1000)])
disp(['P value for RHand numVars comparison is ' num2str(length(find(perm_corrs(:,4)<
<Rhand_numVars_trueCorr))/1000)])
disp(['P value for LHand varSize diff comparison is ' num2str(length(find(perm_corrs(:,5)<
<LHand_varSize_diff))/1000)])

```



```

disp(['P value for RHand varSize diff comparison is ' num2str(length(find(perm_corrs(:,6)
<RHand_varSize_diff))/1000)])
disp(['P value for LHand varSize comparison is ' num2str(length(find(perm_corrs(:,1)
<LHand_varSize_trueCorr))/1000)])
disp(['P value for RHand varSize comparison is ' num2str(length(find(perm_corrs(:,2)
<RHand_varSize_trueCorr))/1000)])
disp(['P value for LHand numVars comparison is ' num2str(length(find(perm_corrs(:,3)
<LHand_numVars_trueCorr))/1000)])
disp(['P value for RHand numVars comparison is ' num2str(length(find(perm_corrs(:,4)
<RHand_numVars_trueCorr))/1000)])
disp(['average corr value for LHand varSize diff comparison is ' num2str(mean(perm_corrs
(:,5)))]
disp(['average corr value for RHand varSize diff comparison is ' num2str(mean(perm_corrs
(:,6)))]
disp(['average corr value for LHand numVars diff comparison is ' num2str(mean(perm_corrs
(:,7)))]
disp(['average corr value for RHand numVars diff comparison is ' num2str(mean(perm_corrs
(:,8)))]
permute_varSize
numVars = LH_info.variants_info.both_hems(:,1);
permute_varSize
help datasample
permute_varSize
disp(['P value for varSize comparison is ' num2str(length(find(perm_corrs(:,1)
<avg_varSize_trueCorr))/1000)])
disp(['P value for varSizeDiff comparison is ' num2str(length(find(perm_corrs(:,4)
<avg_varSizeDiff_trueCorr))/1000)])
disp(['P value for numVarsDiff comparison is ' num2str(length(find(perm_corrs(:,3)
<avg_numVarsDiff_trueCorr))/1000)])
length(find(perm_corrs(:,3)<avg_numVarsDiff_trueCorr))
permute_varSize
%-- 7/28/21, 1:45 PM --%
load('/Users/dianaperez/Desktop/HCP384_original_variants_info.mat')
clear all
allSubs_info = load('/Users/dianaperez/Desktop/HCP384_original_variants_info.mat');
numVars = allSubs_info.variants_info.left_hem.group_avg(:,1);
clear numVars
left_hem = allSubs_info.variants_info.left_hem.group_avg(:,1);
right_hem = allSubs_info.variants_info.right_hem.group_avg(:,1);
diff = left_hem-right_hem;
mean_diff = mean(left_hem - right_hem);
help mean
permute_varSize
disp(['Below: ' num2str(length(find(perm_diff<true_diff))/1000)])
disp(['Above: ' num2str(length(find(perm_diff>true_diff))/1000)])
permute_varSize
left_hand_diff = mean(left_hand_left_hem(:, :) - left_hand_right_hem(:, :));
permute_varSize
left_hand_left_hem = LHand_info.variants_info.left_hem.group_avg(:,1);
left_hand_right_hem = LHand_info.variants_info.right_hem.group_avg(:,1);
left_hand_diffs = left_hand_left_hem - left_hand_right_hem;
true_diff_left_hand = mean(left_hand_diffs);
right_hand_left_hem = RHand_info.variants_info.left_hem.group_avg(:,1);
right_hand_right_hem = RHand_info.variants_info.right_hem.group_avg(:,1);
right_hand_diffs = right_hand_left_hem - right_hand_right_hem;
true_diff_right_hand = mean(right_hand_diffs);
true_diff = true_diff_left_hand - true_diff_right_hand;
num_vars_diff = [left_hand_diffs; right_hand_diffs];
numSubs = 699;
flip_switch = zeros(numSubs,1);

```

```

flip_switch(1:651) = 1;
for p = 1:numperms
    rng('shuffle');
    ind = randperm(length(flip_switch));
    for s = 1:length(flip_switch)
        rand_flip_switch(s,1) = flip_switch(ind(s));
    end
    pseudo_left_handers = num_vars_diff(rand_flip_switch==0);
    pseudo_right_handers = num_vars_diff(rand_flip_switch==1);
    perm_diffs(p,1) = mean(pseudo_left_handers) - mean(pseudo_right_handers);
end
disp(['Hemisphere X Handedness, Number of Variants (permute handedness) - Below: '
num2str(length(find(perm_diffs<true_diff))/1000)])
disp(['Hemisphere X Handedness, Number of Variants (permute handedness) - Above: '
num2str(length(find(perm_diffs>true_diff))/1000)])
clear perm_diffs
left_hand_left_hem = LHand_info.variants_info.left_hem.group_avg(:,1);
left_hand_right_hem = LHand_info.variants_info.right_hem.group_avg(:,1);
left_hand_diffs = left_hand_left_hem - left_hand_right_hem;
true_diff_left_hand = mean(left_hand_diffs);
right_hand_left_hem = RHand_info.variants_info.left_hem.group_avg(:,1);
right_hand_right_hem = RHand_info.variants_info.right_hem.group_avg(:,1);
right_hand_diffs = right_hand_left_hem - right_hand_right_hem;
true_diff_right_hand = mean(right_hand_diffs);
true_diff = true_diff_left_hand - true_diff_right_hand;
num_vars_diff = [left_hand_diffs; right_hand_diffs];
numSubs = 699;
flip_switch = zeros(numSubs,1);
flip_switch(1:651) = 1;
for p = 1:numperms
    rng('shuffle');
    ind = randperm(length(flip_switch));
    for s = 1:length(flip_switch)
        rand_flip_switch(s,1) = flip_switch(ind(s));
    end
    pseudo_left_handers = num_vars_diff(rand_flip_switch==0);
    pseudo_right_handers = num_vars_diff(rand_flip_switch==1);
    perm_diffs(p,1) = mean(pseudo_left_handers) - mean(pseudo_right_handers);
end
disp(['Hemisphere X Handedness, Number of Variants (permute handedness) - Below: '
num2str(length(find(perm_diffs<true_diff))/1000)])
disp(['Hemisphere X Handedness, Number of Variants (permute handedness) - Above: '
num2str(length(find(perm_diffs>true_diff))/1000)])
permute_varSize
rng('shuffle');
ind = randperm(length(flip_switch));
rand_flip_switch = flip_switch(ind)
%-- 7/29/21, 2:28 PM --%
permute_varSize
%-- 8/6/21, 3:50 PM --%
crossRefHCP
goodSubs_col = cell2mat(goodSubs(1:752,2));
dem_info_752 = [goodSubs_all(:,1) goodSubs_all(:,2) goodSubs_all(:,12)];
crossRefHCP
%-- 8/14/21, 3:44 PM --%
crossRefHCP
goodSubs_col = cell2mat(goodSubs(1:384,1));
crossRefHCP
clear all
load('/Users/dianaperez/Desktop/384_demog_info.mat')

```

```

goodSubs384 = dem_info;
load('/Users/dianaperez/Desktop/752_demog_info.mat')
goodSubs752 = dem_info;
length(find(goodSubs384=='M'))
length(find(goodSubs384=='M'))
length(find(goodSubs384=='M'))
length(find(goodSubs384(:,3)=='M'))
length(find(goodSubs384(:,3)=='M'))
384-210
mean(goodSubs384(:,2))
crossRefHCP
%-- 8/15/21, 3:41 PM --%
assignVars_HCP_dice
%-- 8/16/21, 9:10 PM --%
load('/Users/dianaperez/Desktop/HCP384_new_split_variants_info.mat')
y = variants_info.left_hem.group_avg(:,1);
x = variants_info.right_hem.group_avg(:,1);
[h,p,stats] = ttest(x,y);
numVars = variants_info.left_hem.group_avg(:,1);
numVars_left = variants_info.left_hem.group_avg(:,1);
numVars_right = variants_info.right_hem.group_avg(:,1);
[h,p,stats] = ttest(numVars_left,numVars_right);
[h,p,stats] = ttest(numVars_right,numVars_left);
varSize_left = variants_info.left_hem.group_avg(:,3);
varSize_right = variants_info.right_hem.group_avg(:,3);
[h,p,ci,stats] = ttest(varSize_left,varSize_right);
mean(varSize_left)
mean(varSize_right)
load('/Users/dianaperez/Desktop/Thesis/HCP384_original_variants_info.mat')
numVars_left = variants_info.left_hem.group_avg(:,1);
numVars_right = variants_info.right_hem.group_avg(:,1);
[h,p,ci,stats] = ttest(numVars_left,numVars_right);
mean(numVars_left)
mean(numVars_right)
varSize_left = variants_info.left_hem.group_avg(:,3);
varSize_right = variants_info.right_hem.group_avg(:,3);
[h,p,ci,stats] = ttest(varSize_left,varSize_right);
mean(varSize_left)
mean(varSize_right)
load('/Users/dianaperez/Desktop/HCP384_new_split_variants_info.mat')
varSize_left = variants_info.left_hem.group_avg(:,3);
varSize_right = variants_info.right_hem.group_avg(:,3);
[h,p,ci,stats] = ttest(varSize_left,varSize_right);
mean(varSize_left)
mean(varSize_right)
numVars_left = variants_info.left_hem.group_avg(:,1);
numVars_right = variants_info.right_hem.group_avg(:,1);
[h,p,ci,stats] = ttest(numVars_left,numVars_right);
mean(numVars_left)
mean(numVars_right)
histogram(numVars_left)
hold on
histogram(numVars_right)
title('Number of variant clusters')
xlabel('Number of clusters')
ylabel('Frequency')
ylabel('Frequency (number of subjects)')
legend('Left Hemisphere', 'Right Hemisphere')
ax = gca;
ax.FontSize = 14;

```

```

set(gcf, 'Units', 'Normalized', 'OuterPosition', [0, 30, 0, 100]);
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0, 30, 0, 100]);
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.5, 0.7]);
print(gcf, '/Users/dianaperez/Desktop/HCP_new_split_vars_numVars.jpg', '-dpng', '-r300')
close gcf
histogram(varSize_left)
hold on
histogram(varSize_right)
title('Average variant size')
xlabel('Variant size in vertices')
ylabel('Frequency (number of subjects)')
legend('Left Hemisphere', 'Right Hemisphere')
ax = gca;
ax.FontSize = 14;
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.5, 0.7]);
print(gcf, '/Users/dianaperez/Desktop/HCP_new_split_vars_varSize.jpg', '-dpng', '-r300')
close gcf
load
load('/Users/dianaperez/Box/Research/Lateralization_thresh10/subjectData/goodSubs_addtlinfo.
mat')
load('/Users/dianaperez/Desktop/Thesis/HCP384_split_WashUTemp_networksxHem.mat')
HCP_goodSubs1 = HCP_goodSubs1(2:end);
HCP_goodSubs1 = HCP_goodSubs1{1:end};
lang = HCP_goodSubs{2:end,5};
lang = HCP_goodSubs{2:385,5};
lang_measures = HCP_goodSubs
clear all
load
load('/Users/dianaperez/Box/Research/Lateralization_thresh10/subjectData/goodSubs_addtlinfo.
mat')
load('/Users/dianaperez/Desktop/Thesis/HCP384_split_WashUTemp_networksxHem.mat')
lang_overall = cell2mat(HCP_goodSubs(2:end,5));
subs = cell2mat(HCP_goodSubs(2:end,1));
lang_scores = [subs; lang_overall];
lang_scores = [subs lang_overall];
clear lang_overall subs
load('/Users/dianaperez/Box/Research/Lateralization_thresh10/subjectData/goodSubs384.
mat')
VAN_left = clustersLH;
VAN_right = clustersRH;
diff = VAN_left-VAN_right;
VAN_diff = [goodSubs384(:,1) diff];
VAN_diff = [VAN_diff lang_scores(:,2)];
corr(VAN_diff(:,2), VAN_diff(:,3))
corrcoef(VAN_diff(:,2), VAN_diff(:,3))
R = corrcoef(VAN_diff(:,2), VAN_diff(:,3))
rsq = R(2,1)^2
plot(VAN_diff(:,2), VAN_diff(:,3))
R = corrcoef(VAN_diff1, VAN_diff2)
rsq = R(2,1)^2
plot(VAN_diff(:,2), VAN_diff(:,3))
plot(VAN_diff1, VAN_diff2)
R = corrcoef(VAN_diff3, VAN_diff2)
rsq = R(2,1)^2
plot(VAN_diff3, VAN_diff2)
scatter(VAN_diff3, VAN_diff2)
find(VAN_diff3==NaN)
find(VAN_diff2==NaN)
R = corrcoef(VAN_diff3, VAN_diff2)
rsq = R(2,1)^2

```

```

lang_VANdiff = VAN_diff;
title('Linear Regression: Language scores and # of VAN Variants (left - right)')
xlabel('language overall accuracy')
ylabel('# of VAN variants difference (left hem - right hem)')
VANdiff_hand = [goodSubs384 VAN_diff1];
R = corrcoef(VANdiff_hand(:,2), VANdiff_hand(:,3))
rsq = R(2,1)^2
R = corrcoef(lang_VANdiff(:,2), lang_VANdiff(:,4));
rsq = R(2,1)^2
load('/Users/dianaperez/Desktop/HCP384_new_split_variants_info.mat')
load('/Users/dianaperez/Box/Research/Lateralization_thresh10/subjectData/goodSubs384.
mat')
clear goodSubs384
load('/Users/dianaperez/Box/Research/Lateralization_thresh10/subjectData/goodSubs384.
mat')
numVars_diff = variants_info.left_hem.group_avg(:,1)-variants_info.right_hem.group_avg(:,
1);
numVars = variants_info.left_hem.group_avg{1,1};
numVars1 = variants_info.left_hem.group_avg{: ,1};
clear numVars
numVars = variants_info.right_hem.group_avg{: ,1};
numVars_diff = numVars1 - numVars;
R = corrcoef(goodSubs384(:,2), goodSubs384(:,3))
rsq = R(2,1)^2
scatter(goodSubs384(:,2), goodSubs384(:,3))
clear all
load('/Users/dianaperez/Desktop/HCP384_new_split_networksxHem.mat')
load('/Users/dianaperez/Desktop/HCP384_new_split_variants_info.mat')
numVars = variants_info.left_hem.group_avg{: ,1};
left_numVars = numVars;
right_numVars = variants_info.right_hem.group_avg{: ,1};
clear numVars
right_varSize = variants_info.right_hem.group_avg{: ,3};
left_varSize = variants_info.left_hem.group_avg{: ,3};
histogram(left_numVars)
hold on
histogram(right_numVars)
[p t ci stats] = ttest(left_numVars,right_numVars);
title('Number of Variants Across Hemispheres')
xlabel('
xlabel('Number of Clusters')
ylabel('Frequency (Number of Subjects)')
legend('Left Hemisphere', 'Right Hemisphere')
ax = gca;
ax.FontSize = 14;
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.5, 0.7]);
print(gcf,[outputDir outputName '.jpg'],'-dpng','-r300');
print(gcf,'/Users/dianaperez/Desktop/numVars_384subs_newsplitsvars.jpg','-dpng','-r300');
print(gcf,'/Users/dianaperez/Desktop/numVars_384subs_newsplitsvars.jpg','-dpng','-r300');
close gcf
histogram(left_varSize)
hold on
histogram(right_varSize)
[sign p ci stats] = ttest(left_numVars,right_numVars);
[sign p ci stats] = ttest(left_varSize,right_varSize);
xlabel('Variant Size (Number of Vertices)')
ylabel('Frequency (Number of Subjects)')
ax = gca;
ax.FontSize = 14;
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.5, 0.7]);

```

```

legend('Left Hemisphere', 'Right Hemisphere')
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.5, 0.7]);
ax.FontSize = 14;
title('Average Variant Size Across Hemispheres')
ax = gca;
ax.FontSize = 14;
print(gcf, '/Users/dianaperez/Desktop/varSize_384subs_newspllitvars.jpg', '-dpng', '-r300');
close gcf
left_hemi = networksxHem.clustersLH(:,2);
left_hemi = networksxHem.clustersLH(:,2);
network_names = {'DMN' 'Vis' 'FP' 'Unassigned' 'DAN' 'Unassigned2' 'VAN'
'Sal' 'CO' 'Smd' 'SML' 'Aud' 'Tpole' 'MTL' 'PMN' 'PON'};
wb_colors = [1 2 3 5 7 8 9 10 11 12 13 14 15 16];
results = [];
statistics = [];
for net = 1:numel(network_names)
left_hemi = networksxHem.clustersLH(:,net);
right_hemi = networksxHem.clustersRH(:,net);
results(net,1) = network_names{net};
results(net,2) = mean(left_hemi);
results(net,3) = mean(right_hemi);
[sig p ci stats] = ttest(left_hemi, right_hemi);
results(net,4) = p;
statistics{net} = stats;
end
network_names = {'DMN' 'Vis' 'FP' 'Unassigned' 'DAN' 'Unassigned2' 'VAN'
'Sal' 'CO' 'Smd' 'SML' 'Aud' 'Tpole' 'MTL' 'PMN' 'PON'};
wb_colors = [1 2 3 5 7 8 9 10 11 12 13 14 15 16];
results = [];
statistics = [];
for net = 1:numel(network_names)
left_hemi = networksxHem.clustersLH(:,net);
right_hemi = networksxHem.clustersRH(:,net);
results(net,1) = network_names{net};
results(net,2) = mean(left_hemi);
results(net,3) = mean(right_hemi);
[sig p ci stats] = ttest(left_hemi, right_hemi);
results(net,4) = p;
statistics{net} = stats;
end
results(1,1) = network_names{net}
results(1,1) = network_names{1}
results(1,1) = network_names(1)
results(1,1) = network_names(1,1)
results(1,1) = network_names{1,1}
results{1,1} = network_names{1,1}
for net = 1:numel(network_names)
left_hemi = networksxHem.clustersLH(:,net);
right_hemi = networksxHem.clustersRH(:,net);
results(net,1) = mean(left_hemi);
results(net,2) = mean(right_hemi);
[sig p ci stats] = ttest(left_hemi, right_hemi);
results(net,3) = p;
statistics{net} = stats;
end
for net = 1:numel(network_names)
left_hemi = networksxHem.clustersLH(:,net);
right_hemi = networksxHem.clustersRH(:,net);
results(net,1) = mean(left_hemi);
results(net,2) = mean(right_hemi);

```

```

[sig p ci stats] = ttest(left_hemi, right_hemi);
results(net,3) = p;
statistics(net) = stats;
end
results = [];
statistics = [];
for net = 1:numel(network_names)
left_hemi = networksxHem.clustersLH(:,net);
right_hemi = networksxHem.clustersRH(:,net);
results(net,1) = mean(left_hemi);
results(net,2) = mean(right_hemi);
[sig p ci stats] = ttest(left_hemi, right_hemi);
results(net,3) = p;
statistics{net} = stats;
end
results = [];
statistics = [];
for net = 1:numel(network_names)
left_hemi = networksxHem.clustersLH(:,net);
right_hemi = networksxHem.clustersRH(:,net);
results(net,1) = mean(left_hemi);
results(net,2) = mean(right_hemi);
[sig p ci stats] = ttest(left_hemi, right_hemi);
results(net,3) = sig;
results(net,4) = p;
statistics{net} = stats;
end
[p_fdr, p_masked] = FDR(results(:,4), .05)
find(p_masked==1);
[p_fdr, results(:,5)] = FDR(results(:,4), .05)
bar(1:16, results(:,1:2))
legend('Left Hem', 'Right Hem')
ax = gca;
XTick(network_names)
xticks(network_names)
xticklabels(network_names)
xticklabels(network_names)
network_names = {'DMN' 'Vis' 'FP' ' ' 'DAN' ' ' 'VAN' 'Sal' 'CO' 'SMD'
'SML' 'Aud' 'Tpole' 'MTL' 'PMN' 'PON'};
xticklabels(network_names)
/Volumes/fsmresfiles/PBS/Gratton_Lab/Lifespan/Post-
COVID/BIDS/DICOMS/LS08_1/FMRI_STUDIES_GRATTON_20210820_092658_908000
cd /Volumes/fsmresfiles/PBS/Gratton_Lab/Lifespan/Post-
COVID/BIDS/DICOMS/LS08_1/FMRI_STUDIES_GRATTON_20210820_092658_908000
info = dicominfo('LS08_1.MR.FMRI_STUDIES_GRATTON.
0006.0002.2021.08.20.11.04.25.695290.6470031.IMA')
cd /Users/dianaperez/Documents/GitHub/Lateralization_Variants/
permute_handedness
run('/Users/dianaperez/Documents/GitHub/General_Scripts/untitled4.m')
clear all
load('/Users/dianaperez/Desktop/HCP384_new_split_networksxHem_wBE.mat')
load('/Users/dianaperez/Desktop/HCP384_new_split_variants_info_wBE.mat')
numVars = variants_info.left_hem.group_avg{1,1};
LHem_numVars = variants_info.left_hem.group_avg{1,1};
RHem_numVars = variants_info.right_hem.group_avg{1,1};
clear numVars
avgVarSize = variants_info.left_hem.group_avg{: ,3};
LHem_avgVarSize = variants_info.left_hem.group_avg{: ,3};
RHem_avgVarSize = variants_info.right_hem.group_avg{: ,3};
clear avgVarSize

```



```

RHem_numBorder = variants_info.right_hem.group_avg{:,4};
RHem_numEctopic = variants_info.right_hem.group_avg{:,5};
LHem_numBorder = variants_info.left_hem.group_avg{:,4};
LHem_numEctopic = variants_info.left_hem.group_avg{:,5};
[sig p ci stats] = ttest(LHem_avgVarSize, RHem_avgVarSize);
p_vals = [];
p_vals = [p_vals; p];
[sig p ci stats] = ttest(LHem_numVars, RHem_numVars);
LHem_numVars = variants_info.left_hem.group_avg{1,1};
LHem_numVars = variants_info.left_hem.group_avg{:,1};
RHem_numVars = variants_info.right_hem.group_avg{:,1};
[sig p ci stats] = ttest(LHem_numVars, RHem_numVars);
[sig p ci stats] = ttest(LHem_numBorder, RHem_numVars);
[sig p ci stats] = ttest(LHem_numBorder, RHem_numBorder);
[sig p ci stats] = ttest(LHem_numEctopic, RHem_numEctopic);
mean(LH_avgVarSize)
mean(LHem_avgVarSize)
mean(RHem_avgVarSize)
mean(LHem_numVars)
mean(RHem_numVars)
mean(RHem_numEctopic)
mean(RHem_numBorder)
mean(LHem_numBorder)
mean(LHem_numEctopic)
[sig p ci stats] = ttest(LHem_numVars, RHem_numVars);
p_vals = [p_vals; p];
[sig p ci stats] = ttest(LHem_numBorder, RHem_numBorder);
p_vals = [p_vals; p];
[sig p ci stats] = ttest(LHem_numEctopic, RHem_numEctopic);
p_vals = [p_vals; p];
[p_fdr, p_sig] = FDR(p_vals, .05)
addpath '/Users/dianaperez/Desktop'
[p_fdr, p_sig] = FDR(p_vals, .05)
clear LHem_numBorder LHem_numEctopic RHem_numBorder RHem_numEctopic
clear p*
clear sig stats ci ans
clear variants_info
histograms(LHem_avgVarSize)
histogram(LHem_avgVarSize)
hold on
network_names = {'DMN' 'Vis' 'FP' 'DAN' 'VAN' 'Sal' 'CO' 'SMd' 'SMl'
'Aud' 'Tpole' 'MTL' 'PMN' 'PON'};
wb_colors = [1 2 3 5 7 8 9 10 11 12 13 14 15 16];
results = [];
statistics = [];
for net = 1:numel(network_names)
left_hemi = networksxHem.clustersLH(:,net);
right_hemi = networksxHem.clustersRH(:,net);
results(net,1) = mean(left_hemi);
results(net,2) = mean(right_hemi);
[sig p ci stats] = ttest(left_hemi, right_hemi);
results(net,3) = sig;
results(net,4) = p;
statistics{net} = stats;
end
[p_fdr, results(:,5)] = FDR(results(:,4), .05);
network_names = {'DMN' 'Vis' 'FP' 'Reward' 'DAN' 'VAN' ' ' 'Sal' 'CO' 'SMd'
'SMl' 'Aud' 'Tpole' 'MTL' 'PMN' 'PON'};
%network_names = {'DMN' 'Vis' 'FP' 'DAN' 'VAN' 'Sal' 'CO' 'SMd' 'SMl'
'Aud' 'Tpole' 'MTL' 'PMN' 'PON'};

```



```

results = [];
statistics = [];
for net = 1:numel(network_names)
left_hemi = networksxHem.clustersLH(:,net);
right_hemi = networksxHem.clustersRH(:,net);
results(net,1) = mean(left_hemi);
results(net,2) = mean(right_hemi);
[sig p ci stats] = ttest(left_hemi, right_hemi);
results(net,3) = sig;
results(net,4) = p;
statistics{net} = stats;
end
[p_fdr, results(:,5)] = FDR(results(:,4), .05);
bar(1:16, results(:,1:2))
legend('Left Hem', 'Right Hem')
bar(1:16, results(:,1:2))
legend('Left Hem', 'Right Hem')
bar(network_names, results(:,1:2))
bar(1:16, results(:,1:2))
legend('Left Hem', 'Right Hem')
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.5, 0.7]);
xticks(1:16)
xticklabels(network_names)
network_names = {'DMN' 'Vis' 'FP' 'Reward' 'DAN' ' ' 'VAN' 'Sal' 'CO' 'SMd'
'SML' 'Aud' 'Tpole' 'MTL' 'PMN' 'PON'};
xticklabels(network_names)
ax = gca;
ax.FontSize = 14;
network_names = {'DMN' 'Vis' 'FP' ' ' 'DAN' ' ' 'VAN' 'Sal' 'CO' 'SMd'
'SML' 'Aud' 'Tpole' 'MTL' 'PMN' 'PON'};
xticklabels(network_names)
ylabel('Average Number of Variants Assigned to Network');
xlabel('Network');
title('Variants Assigned to Each Network Across Hemispheres');
print(gcf, '/Users/dianaperez/Desktop/Networks_by_Hem_new_split_vars.jpg', '-dpng', '-r300');
print(gcf, '/Users/dianaperez/Desktop/Networks_by_Hem_new_split_vars.jpg', '-dpng', '-r300');
Overlap_VANvars
DMN_overlap = ft_read_cifti_mod(
('/Users/dianaperez/Desktop/HCP_newsplit_DMNvariants_OverlapMap_384GoodSubs.dtseries.
nii')
flip_DMN_overlap = insert_nonbrain(DMN_overlap.data, 'both', DMN_overlap);
64984/2
> left = flip_DMN_overlap(1:32492);
left = flip_DMN_overlap(1:32492);
right = flip_DMN_overlap(32493:end);
DMN_cifti_1 = [left; right];
DMN_cifti_1 = DMN_cifti_1(DMN_overlap.brainstructure>0);
DMN_cifti_2 = [right; left];
DMN_cifti_2 = DMN_cifti_2(DMN_overlap.brainstructure>0);
DMN_diff = DMN_cifti_1 - DMN_cifti_2;
DMN_diff = DMN_cifti_1 - DMN_cifti_2;
template = DMN_overlap;
template.data = DMN_diff;
ft_write_cifti_mod('/Users/dianaperez/Desktop/DMN_diff_map.dtseries.nii', template);
VIS_overlap = ft_read_cifti_mod(
('/Users/dianaperez/Desktop/HCP_newsplit_VISvariants_OverlapMap_384GoodSubs.dtseries.
nii')
flip_VIS_overlap = insert_nonbrain(VIS_overlap.data, 'both', VIS_overlap);

```

```
> left = flip_VIS_overlap(1:32492);
left = flip_VIS_overlap(1:32492);
right = flip_VIS_overlap(32493:end);
VIS_cifti_1 = [left; right];
VIS_cifti_2 = [right; left];
VIS_cifti_1 = VIS_cifti_1(VIS_overlap.brainstructure>0);
VIS_cifti_2 = VIS_cifti_2(VIS_overlap.brainstructure>0);
VIS_diff = VIS_cifti_1 - VIS_cifti_2;
template.data = VIS_diff;
ft_write_cifti_mod('/Users/dianaperez/Desktop/VIS_diff_map.dtseries.nii', template);
VAN_overlap = ft_read_cifti_mod(
('/Users/dianaperez/Desktop/HCP_newsplit_VANvariants_OverlapMap_384GoodSubs.dtseries.
nii'))
all_verts = insert_nonbrain(VAN_overlap.data, 'both', VAN_overlap);
left = all_verts(1:32492);
right = all_verts(32493:end);
cifti_1 = [left; right];
cifti_2 = [right; left];
cifti_1 = cifti_1(VIS_overlap.brainstructure>0);
cifti_2 = cifti_2(VIS_overlap.brainstructure>0);
diff = cifti_1 - cifti_2;
template.data = diff;
ft_write_cifti_mod('/Users/dianaperez/Desktop/VAN_diff_map.dtseries.nii', template);
overlap = ft_read_cifti_mod(
('/Users/dianaperez/Desktop/HCP_newsplit_CONvariants_OverlapMap_384GoodSubs.dtseries.
nii'))
all_verts = insert_nonbrain(overlap.data, 'both', overlap);
left = all_verts(1:32492);
right = all_verts(32493:end);
cifti_1 = [left; right];
cifti_2 = [right; left];
cifti_1 = cifti_1(VIS_overlap.brainstructure>0);
cifti_2 = cifti_2(VIS_overlap.brainstructure>0);
diff = cifti_1 - cifti_2;
template.data = diff;
ft_write_cifti_mod('/Users/dianaperez/Desktop/CON_diff_map.dtseries.nii', template);
overlap = ft_read_cifti_mod(
('/Users/dianaperez/Desktop/HCP_newsplit_SMLvariants_OverlapMap_384GoodSubs.dtseries.
nii'))
all_verts = insert_nonbrain(overlap.data, 'both', overlap);
left = all_verts(1:32492);
right = all_verts(32493:end);
cifti_1 = [left; right];
cifti_2 = [right; left];
cifti_1 = cifti_1(VIS_overlap.brainstructure>0);
cifti_2 = cifti_2(VIS_overlap.brainstructure>0);
diff = cifti_1 - cifti_2;
template.data = diff;
ft_write_cifti_mod('/Users/dianaperez/Desktop/SML_diff_map.dtseries.nii', template);
Overlap_VANvars
unique(overlap_map(26000:end))
load('/Users/dianaperez/Desktop/HCP384_new_split_networksxHem.mat')
load('/Users/dianaperez/Desktop/HCP384_new_split_variants_info.mat')
clear al
clear all
load('/Users/dianaperez/Desktop/HCP384_new_split_networksxHem.mat')
load('/Users/dianaperez/Desktop/HCP384_new_split_variants_info.mat')
LHem = variants_info.left_hem.group_avg{1,1};
RHem = variants_info.right_hem.group_avg{1,1};
[sig p ci stats] = ttest(LHem, RHem);
```

```
LHem = variants_info.left_hem.group_avg{:,1};
RHem = variants_info.right_hem.group_avg{:,1};
[sig p ci stats] = ttest(LHem, RHem);
LHem_mean = mean(LHem);
RHem_mean = mean(RHem);
[sig p ci stats] = ttest(LHem, RHem);
disp(p)
if sig == 1
disp(['Comparison is significant with p = ' num2str(p)])
end
LHem_mean = mean(LHem)
RHem_mean = mean(RHem)
histogram(LHem)
hold on
histogram(RHem)
title(title_str)
xlabel(x_label)
ylabel(y_label)
legend('Left Hemisphere', 'Right Hemisphere')
ax = gca;
ax.FontSize = 14;
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.5, 0.7]);
LHem = variants_info.left_hem.group_avg{:,1};
RHem = variants_info.right_hem.group_avg{:,1};
title_str = 'Number of Variants Across Hemispheres';
x_label = 'Number of Clusters';
y_label = 'Frequency (Number of Subjects)';
outputName = 'HCP384_new_split_vars_numVars.jpg';
histogram(LHem)
hold on
histogram(RHem)
title(title_str)
xlabel(x_label)
ylabel(y_label)
legend('Left Hemisphere', 'Right Hemisphere')
ax = gca;
ax.FontSize = 14;
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.5, 0.7]);
LHem = variants_info.left_hem.group_avg{:,3};
RHem = variants_info.right_hem.group_avg{:,3};
title_str = 'Average Variant Size Across Hemispheres';
x_label = 'Average Variant Size (Number of Vertices)';
y_label = 'Frequency (Number of Subjects)';
outputName = 'HCP384_new_split_vars_avgVarSize.jpg';
[sig p ci stats] = ttest(LHem, RHem);
disp(p)
if sig == 1
disp(['Comparison is significant with p = ' num2str(p)])
end
LHem_mean = mean(LHem)
RHem_mean = mean(RHem)
histogram(LHem)
hold on
histogram(RHem)
title(title_str)
xlabel(x_label)
ylabel(y_label)
legend('Left Hemisphere', 'Right Hemisphere')
ax = gca;
ax.FontSize = 14;
```

```

set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.5, 0.7]);
network_names = {'DMN' 'Vis' 'FP' ' ' 'DAN' ' ' 'VAN' 'Sal' 'CO' 'SMd'↵
'SMl' 'Aud' 'Tpole' 'MTL' 'PMN' 'PON'};
%network_names = {'DMN' 'Vis' 'FP' 'DAN' 'VAN' 'Sal' 'CO' 'SMd' 'SMl'↵
'Aud' 'Tpole' 'MTL' 'PMN' 'PON'};
results = [];
statistics = [];
for net = 1:numel(network_names)
left_hemi = networksxHem.clustersLH(:,net);
right_hemi = networksxHem.clustersRH(:,net);
results(net,1) = mean(left_hemi);
results(net,2) = mean(right_hemi);
[sig p ci stats] = ttest(left_hemi, right_hemi);
results(net,3) = sig;
results(net,4) = p;
statistics{net} = stats;
end
[p_fdr, results(:,5)] = FDR(results(:,4), .05);
bar(1:16, results(:,1:2))
legend('Left Hem', 'Right Hem')
xticks(1:16)
xticklabels(network_names)
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.5, 0.7]);
ylabel('Average Number of Variants Assigned to Network');
xlabel('Network');
title_str('Variants Assigned to Each Network Across Hemispheres');
Overlap_VANvars
%-- 9/2/21, 9:53 AM --%
permute_varSize
left_hem = allSubs_info.variants_info.left_hem.group_avg{:,4};
right_hem = allSubs_info.variants_info.right_hem.group_avg{:,4};
true_diff = mean(left_hem - right_hem);
flip_switch = zeros(length(left_hem),1);
flip_switch(1:(length(left_hem))/2) = 1;
for p = 1:numperms
rng('shuffle');
ind = randperm(length(flip_switch));
rand_flip_switch = flip_switch(ind);
for s = 1:length(left_hem)
if rand_flip_switch(s) == 1
pseudo_left_hem(s,1) = left_hem(s);
pseudo_right_hem(s,1) = right_hem(s);
elseif rand_flip_switch(s) == 0
pseudo_right_hem(s,1) = left_hem(s);
pseudo_left_hem(s,1) = right_hem(s);
end
end
perm_diffs(p) = mean(pseudo_left_hem - pseudo_right_hem);
end
disp(['Average Number of Border Variants (permute hemispheres) - Below: ' num2str(length↵
(find(perm_diffs<true_diff))/1000)])
disp(['Average Number of Border Variants (permute hemispheres) - Above: ' num2str(length↵
(find(perm_diffs>true_diff))/1000)])
clear left_hem right_hem true_diff flip_switch rand_flip_switch pseudo_left_hem↵
pseudo_right_hem
%% Test diff in number of ectopic variants across hemispheres in all subjects
left_hem = allSubs_info.variants_info.left_hem.group_avg{:,5};
right_hem = allSubs_info.variants_info.right_hem.group_avg{:,5};
true_diff = mean(left_hem - right_hem);
flip_switch = zeros(length(left_hem),1);

```

```

flip_switch(1:(length(left_hem))/2) = 1;
for p = 1:numperms
    rng('shuffle');
    ind = randperm(length(flip_switch));
    rand_flip_switch = flip_switch(ind);
    for s = 1:length(left_hem)
        if rand_flip_switch(s) == 1
            pseudo_left_hem(s,1) = left_hem(s);
            pseudo_right_hem(s,1) = right_hem(s);
        elseif rand_flip_switch(s) == 0
            pseudo_right_hem(s,1) = left_hem(s);
            pseudo_left_hem(s,1) = right_hem(s);
        end
    end
perm_diffs(p) = mean(pseudo_left_hem - pseudo_right_hem);
end
disp(['Average Number of Ectopic Variants (permute hemispheres) - Below: ' num2str(length(
(find(perm_diffs<true_diff))/1000))]
disp(['Average Number of Ectopic Variants (permute hemispheres) - Above: ' num2str(length(
(find(perm_diffs>true_diff))/1000))]
clear left_hem right_hem true_diff flip_switch rand_flip_switch pseudo_left_hem
pseudo_right_hem
permute_variantproperties
net_info.networksxHem.clustersLH{:,1}
net_info.networksxHem.clustersLH{1}
net_info.networksxHem.clustersLH(:,1)
permute_variantproperties
LHand_info.variants_info.both_hems{:,3}
%-- 9/10/21, 12:43 PM --%
load
('/Users/dianaperez/Desktop/Thesis/Spatial_Location/HCP/VariantsvsFlippedVariants_allSubs
_spCorr_1000_thresh_10_permutations.mat')
load
('/Users/dianaperez/Desktop/Thesis/Spatial_Location/HCP/VariantsvsFlippedVariants_allSubs
_true_spCorr_1000_thresh_10_permutations.mat')
scatter(spCorrs, 'jitter', 'on')
help scatter
scatter(spCorrs, 1:1000, 'jitter', 'on')
scatter(1:1000, spCorrs, 'jitter', 'on', 'jitterAmount', 0.5)
help plotspread
plotSpread
cd /Users/dianaperez/Documents/MATLAB/plotSpread/
plotSpread(spCorrs)
hold on
plotSpread(true_spCorr)
plotSpread(spCorrs, spCorr)
plotSpread(spCorrs, spCorrs)
data = {spCorrs, spCorrs, spCorrs};
plotSpread(data,[],[],{'HCP', 'HCP split', 'MSC'});
plotSpread(data,[],[],{'HCP', 'HCP split', 'MSC'});
figure
plotSpread(data,[],[],{'HCP', 'HCP split', 'MSC'});
plotSpread(data,'categoryLabels',{'HCP', 'HCP split', 'MSC'});
true = {true_spCorr, true_spCorr, true_spCorr};
hold on
plotSpread(true)
close(gcf)
figure
plotSpread(data,'categoryLabels',{'HCP', 'HCP split', 'MSC'}, 'categoryMarkers',
{'o','o','o'}, 'categoryColors',{'r','b','g'});

```

```

plotSpread(data,'categoryLabels',{'HCP', 'HCP split', 'MSC'}, 'categoryMarkers',↵
{'o','o','o'}, 'categoryColors',{'r','r','r'});
data = [randn(50,1);randn(50,1)+3.5]*[1 1];
catIdx = [ones(50,1);zeros(50,1);randi([0,1],[100,1])];
figure
plotSpread(data,'categoryIdx',catIdx,...
'categoryMarkers',{'o','+'}, 'categoryColors',{'r','b'})
close gcf
data = [spCorrs; spCorrs];
catIdx = [ones(1000,1);zeros(1000,1)];
catIdx = [ones(1000,1);zeros(1000,1)];
figure
plotSpread(data,'categoryIdx',catIdx,...
'categoryMarkers',{'o','+'}, 'categoryColors',{'r','b'})
data = [true_spCorr; spCorrs];
catIdx = [1;zeros(1000,1)];
plotSpread(data,'categoryIdx',catIdx,...
'categoryMarkers',{'o','+'}, 'categoryColors',{'r','b'})
'categoryMarkers',{'o','+'}, 'categoryColors',{'r','r'})
plotSpread(data,'categoryIdx',catIdx,...
'categoryMarkers',{'o','+'}, 'categoryColors',{'r','r'})
load('/Users/dianaperez/Desktop/Thesis/Variant_Properties/HCP384_new_split_variants_info.↵
mat')
numVars = variants_info.left_hem.group_avg{:},1];
variants_info.left_hem.group_avg.numVars(1) = 1;
LH = numVars;
RH = variants_info.right_hem.group_avg{:},1];
x = [LH RH];
al_goodplot(x)
diff = LH-RH;
al_goodplot(diff)
al_goodplot(x,[],[],[], 'man')
close gcf
al_goodplot(LH)
al_goodplot(RH,[],[],[], 'man')
close gcf
al_goodplot(LH,[],[],[], 'left')
al_goodplot(RH,[],[],[], 'right')
al_goodplot([],[],[],[], 'man')
close gcf
al_goodplot(LH,0,.2,[], 'left')
al_goodplot(RH,0,.2,[], 'right')
legend('Left Hemisphere', 'Right Hemisphere')
legend('Right Hemisphere')
legend('Left Hemisphere',[], 'Right Hemisphere')
help findobj
h = findobj
m = findobj(gca, 'Type', 'Scatter');
legend(m(1:2), 'left hem', 'right hem')
legend(m(1:2), 'right hem', 'left hem')
close gcf
help distributionPlot
plot(spCorrs, 'MarkerEdgeColor', 'black')
close gcf
scatter(spCorrs, 'MarkerEdgeColor', 'black')
help scatter
scatter(spCorrs,1:1000, 'MarkerEdgeColor', 'black')
scatter(1:1000,spCorrs, 'MarkerEdgeColor', 'black')
hold on
scatter(500,true_spCorr, 'MarkerEdgeColor', 'red', 'MarkerFaceColor', 'red', 'SizeData',↵

```

```

50)
scatter(500,true_spCorr, 'MarkerEdgeColor', 'red', 'MarkerFaceColor', 'red', 'SizeData', ↵
100)
scatter(1:1000,spCorrs, 'MarkerEdgeColor', 'black', 'XJitter', 'density')
scatter(1:1000,spCorrs, 'MarkerEdgeColor', 'black', 'XJitter', 'density')
close(gcf)
scatter(1:1000,spCorrs, 'MarkerEdgeColor', 'black', 'XJitter', 'density')
hold on
scatter(500,true_spCorr, 'MarkerEdgeColor', 'red', 'MarkerFaceColor', 'red', 'SizeData', ↵
100)
axis([0,1000])
axis([0, 1000, .6, 1])
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.5, 0.7]);
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.7, 0.7, 0.5, 0.7]);
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.7, 0.7, 0.7, 0.7]);
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.7, 0.7, 0.3, 0.7]);
close(gcf)
scatter(1:1000,spCorrs, 'MarkerEdgeColor', 'black', 'SizeData', 50)
scatter(500,true_spCorr, 'MarkerEdgeColor', 'red', 'MarkerFaceColor', 'red', 'SizeData', ↵
100)
hold on
scatter(1:1000,spCorrs, 'MarkerEdgeColor', 'black', 'SizeData', 50)
axis([0, 1000, .6, 1])
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.7, 0.7, 0.3, 0.7]);
scatter(1:1000,spCorrs, 'MarkerEdgeColor', 'black', 'SizeData', 100)
close(gcf)
scatter(1:1000,spCorrs, 'MarkerEdgeColor', 'black', 'SizeData', 150)
hold on
scatter(500,true_spCorr, 'MarkerEdgeColor', 'red', 'MarkerFaceColor', 'red', 'SizeData', ↵
300)
close(gcf)
scatter(1:1000,spCorrs, 'MarkerEdgeColor', 'black', 'SizeData', 100)
scatter(500,true_spCorr, 'MarkerEdgeColor', 'red', 'MarkerFaceColor', 'red', 'SizeData', ↵
200)
hold on
scatter(1:1000,spCorrs, 'MarkerEdgeColor', 'black', 'SizeData', 100)
axis([0, 1000, .6, 1])
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.7, 0.5, 0.3, 0.7]);
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.7, 0.7, 0.3, 0.7]);
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.7, 0.7, 0.3, 0.3]);
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.7, 0.7, 0.3, 0.9]);
axis([0, 1000, .65, 1])
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.7, 0.7, 0.3, 0.8]);
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.7, 0.7, 0.3, 0.7]);
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.7, 0.5, 0.3, 0.9]);
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.7, 0.3, 0.3, 0.9]);
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.7, 0.5, 0.3, 0.9]);
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.3, 0.9]);
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.3, 0.7]);
ylabel('Correlation')
ax = gca;
ax.FontSize = 14;
ax.FontSize = 24;
XTick()
xticks()
xticks(0)
xticks( )
xticks([])
yticks([1, .75,.65])
yticks([.65,.75,1])

```



```

print(gcf, '
/Users/dianaperez/Desktop/Thesis/Figures/Figure1_permutations_scatterplot_HCP.jpg', '-
dpng', '-r300')
scatter(1:3,spCorrs(1:3), 'MarkerEdgeColor', 'black', 'SizeData', 100)
close gcf
scatter(1:3,spCorrs(1:3), 'MarkerEdgeColor', 'black', 'SizeData', 100)
scatter(1:1000,spCorrs, 'MarkerEdgeColor', 'black', 'SizeData', 100)
close gcf
scatter(500,true_spCorr, 'MarkerEdgeColor', 'red', 'MarkerFaceColor', 'red', 'SizeData',
200)
hold on
scatter(1:1000,spCorrs, 'MarkerEdgeColor', 'black', 'SizeData', 100)
axis([0, 1000, .65, 1])
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.3, 0.7]); %first and second
control position on screen, third controls width, and fourth controls height
ylabel('Correlation')
xticks([])
yticks([.65,.75,1])
ax = gca;
ax.FontSize = 24;
m = findobj(gca, 'Type', 'Scatter')
m = findobj(gca, 'Type', 'Scatter')
hleg1 = legend(m(1:2), 'TrueCorrelation', 'Permuted Correlatoin Values', 'Location',
'SouthWest');
hleg1 = legend(m(1:2), 'Permuted Correlatoin Values', 'TrueCorrelation', 'Location',
'Right');
hleg1 = legend(m(1:2), 'Permuted Correlatoin Values', 'TrueCorrelation', 'Location',
'westoutside');
hleg1 = legend(m(1:2), 'Permuted Correlatoin Values', 'TrueCorrelation', 'Location',
'eastoutside');
hleg1 = legend(m(1:2), 'Permuted Correlation Values', 'True Correlation', 'Location',
'eastoutside');
hleg1 = legend(m(1:2), 'Permuted Correlation Values', 'True Correlation', 'Location',
'southoutside');
leg_marks = findobj(hleg1, 'Type', 'Marker');
leg_marks = findobj(hleg1, 'Marker');
leg_marks = findobj(hleg1, 'Marker', 'none', '-xor');
leg_marks = findobj(hleg1, 'Type', 'Scatter');
help findobj
leg_marks = findobj(hleg1, 'Type', 'legend');
marks = findobj(leg_marks, 'Marker', 'none', '-xor')
set(marks, 'MarkerSize', 20)
m = findobj(hleg1, 'Type', 'patch');
set(m, 'MarkerSize', 50)
set(m, 'MarkerSize', 150)
m = findobj(gcf, 'Type', 'patch');
set(m, 'MarkerSize', 150)
m = findobj(gcf, 'type', 'patch');
m = findobj(gcf);
m = findobj(gcf)
m_l = findobj(m.Legend)
m
m.Legend
m = findobj(gcf, 'type', 'legend')
l = findobj(m, 'type', 'patch')
print(gcf, '
/Users/dianaperez/Desktop/Thesis/Figures/Figure1_permutations_scatterplot_HCP.jpg', '-
dpng', '-r300')
overlapmap
overlap = ft_read_cifti_mod

```



```
('/Users/dianaperez/Desktop/HCP_allSubs_OverlapMap_384GoodSubs_threshold_10.dtseries.
nii');
diffmat_file = overlap.data;
permutations_cluster_correction
trueDiff_cc = zeros(size(trueDiff));
size(trueDiff)
permutations_cluster_correction
size(diffmats)
clear size
size(diffmats)
permutations_cluster_correction
trueDiff_cc = zeros(size(trueDiff));
trueDiff_cc(truediff_bin==1) = trueDiff(truediff_bin==1);
template.data = trueDiff_cc;
ft_write_cifti_mod(outfile_diffmap, template);
%-- 9/13/21, 5:50 PM --%
permute_variantproperties
net_perms = [];
net_perms(1,1) = true_diff;
permute_variantproperties
net_perms(:,net)= [true_diff; perm_diffs'];
net_perms = [];
net_perms(:,net)= [true_diff; perm_diffs'];
permute_variantproperties
help save
permute_variantproperties
length(find(net_perms(2:end,1)<net_perms(1,1)))/1000
length(find(net_perms(2:end,1)<net_perms(1,1)))
find(net_perms(2:end,1)<net_perms(1,1))
1/1000
save([outputDir 'network_perms_numVars.mat'], 'net_perms');
3/10
permute_variantproperties
clear all
load('/Users/dianaperez/Desktop/Thesis/Variant_Properties/HCP384_new_split_variants_info.
mat')
avgVarSize = variants_info.left_hem.group_avg(:,3);
leftHem = variants_info.left_hem.group_avg(:,3);
rightHem = variants_info.right_hem.group_avg(:,3);
[h,mu,sigma,q,notch] = al_goodplot(leftHem,0,'left');
addpath '/Users/dianaperez/Documents/MATLAB/upload'
[h,mu,sigma,q,notch] = al_goodplot(leftHem,0,'left');
[h,mu,sigma,q,notch] = al_goodplot(leftHem,0,[],'left');
[h,mu,sigma,q,notch] = al_goodplot(leftHem,0,[],[],'left');
[h,mu,sigma,q,notch] = al_goodplot(rightHem,0,[],[],'right');
[h,mu,sigma,q,notch] = al_goodplot(leftHem,0,.2,[],'left');
[h,mu,sigma,q,notch] = al_goodplot(rightHem,0,.2,[],'right');
axis([-0.5, 0.5, 0, 400])
axis([-0.4, 0.4, 0, 400])
xticks([])
axis([-0.4, 0.4, 50, 400])
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.3, 0.7]);
axis([-0.4, 0.4, 50, 350])
ylabel('Average Variant Size in Vertices')
ax = gca;
ax.FontSize = 24;
hleg1 = legend(m(1:2), 'Right Hemisphere', 'Left Hemisphere', 'Location',
'SouthOutside');
hleg1.FontSize = 14;
m = findobj(gca);
```

```

findobj(gca)
m = findobj(gca, 'Type', 'Scatter');
hleg1 = legend(m(1:2), 'Right Hemisphere', 'Left Hemisphere', 'Location', 'SouthOutside');
hleg1 = legend(m(1:2), 'Permuted Correlation Values', 'True Correlation', 'Location', 'SouthOutside');
hleg1 = legend(m(1:2), 'Right Hemisphere', 'Left Hemisphere', 'Location', 'SouthOutside');
[h,mu,sigma,q,notch] = al_goodplot(rightHem,0,.2,[],'man');
[h,mu,sigma,q,notch] = al_goodplot(leftHem,0,.2,[],'left');
[h,mu,sigma,q,notch] = al_goodplot(rightHem,0,.2,[],'right');
xticks([])
axis([-0.4, 0.4, 50, 400])
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.3, 0.7]); %first and second control position on screen, third controls width, and fourth controls height
m = findobj(gca, 'Type', 'Scatter');
hleg1 = legend(m(1:2), 'Right Hemisphere', 'Left Hemisphere', 'Location', 'SouthOutside');
ax = gca;
ax.FontSize = 24;
ylabel('Average Variant Size in Vertices')
print(gcf, '/Users/dianaperez/Desktop/Thesis/Figures/Figure2_violin_avgVarSize_HCP.jpg', '-dpng', '-r300')
numVarsLH = variants_info.left_hem.group_avg(:,1);
numVarsRH = variants_info.right_hem.group_avg(:,1);
numBorderLH = variants_info.left_hem.group_avg(:,4);
numBorderRH = variants_info.right_hem.group_avg(:,4);
numEctopicLH = variants_info.left_hem.group_avg(:,5);
numEctopicRH = variants_info.right_hem.group_avg(:,5);
leftHem = [numVarsLH numBorderLH numEctopicLH];
rightHem = [numVarsRH numBorderRH numEctopicRH];
[h,mu,sigma,q,notch] = al_goodplot(leftHem,0,.2,[],'left');
[h,mu,sigma,q,notch] = al_goodplot(rightHem,0,.2,[],'right');
xticks([])
axis([-0.4, 0.4, 50, 400])
ax = gca;
ax.FontSize = 24;
ylabel('Average Variant Size in Vertices')
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.3, 0.7]); %first and second control position on screen, third controls width, and fourth controls height
m = findobj(gca, 'Type', 'Scatter');
hleg1 = legend(m(1:2), 'Right Hemisphere', 'Left Hemisphere', 'Location', 'SouthOutside');
numVarsLH = variants_info.left_hem.group_avg(:,1);
numVarsRH = variants_info.right_hem.group_avg(:,1);
numBorderLH = variants_info.left_hem.group_avg(:,4);
numBorderRH = variants_info.right_hem.group_avg(:,4);
numEctopicLH = variants_info.left_hem.group_avg(:,5);
numEctopicRH = variants_info.right_hem.group_avg(:,5);
leftHem = [numVarsLH numBorderLH numEctopicLH];
rightHem = [numVarsRH numBorderRH numEctopicRH];
[h,mu,sigma,q,notch] = al_goodplot(leftHem,[],.2,[],'left');
[h,mu,sigma,q,notch] = al_goodplot(rightHem,[],.2,[],'right');
xticks([])
%axis([-0.4, 0.4, 50, 400])
ax = gca;
ax.FontSize = 24;
[h,mu,sigma,q,notch] = al_goodplot(leftHem,[],.2,[],'left');
[h,mu,sigma,q,notch] = al_goodplot(leftHem,[-1,0,1],.2,[],'left');
[h,mu,sigma,q,notch] = al_goodplot(rightHem,[-1,0,1],.2,[],'right');

```

```

colorOrder = get(gca, 'ColorOrder');
colorOrder = get(gca, 'ColorOrder');
[h,mu,sigma,q,notch] = al_goodplot(rightHem,[-1,0,1],.2,[ 0    0.4470 0.7410], 'right');
[h,mu,sigma,q,notch] = al_goodplot(rightHem,[-1,0,1],.2,[ 0.8500    0.3250 0.0980], 'right');
[h,mu,sigma,q,notch] = al_goodplot(rightHem,[-1,0,1],.2,[ 0.9290    0.6940 0.1250], 'right');
[h,mu,sigma,q,notch] = al_goodplot(rightHem,[-1,0,1],.2,[0.4940    0.1840 0.5560], 'right');
[h,mu,sigma,q,notch] = al_goodplot(rightHem,[-1,0,1],.2,[0.4660    0.6740 0.1880], 'right');
[h,mu,sigma,q,notch] = al_goodplot(rightHem,[-1,0,1],.2,[    0.3010    0.7450 0.9330], 'right');
figure
[h,mu,sigma,q,notch] = al_goodplot(rightHem,[-1,0,1],.2,[ 0    0.4470 0.7410], 'right');
[h,mu,sigma,q,notch] = al_goodplot(leftHem,[-1,0,1],.2,[ 0    0.4470    0.7410], 'left');
[h,mu,sigma,q,notch] = al_goodplot(rightHem,[-1,0,1],.2,[0.4660    0.6740 0.1880], 'right');
xticks([])
ax = gca;
ax.FontSize = 24;
ylabel('Number of Variants')
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.7, 0.7]);
yticks([-1,0,1])
yticklabels({'All Variants', 'Border Shifts', 'Ectopic Intrusions'})
xticklabels({'All Variants', 'Border Shifts', 'Ectopic Intrusions'})
xticks([-1,0,1])
[h,mu,sigma,q,notch] = al_goodplot(leftHem,[-1,0,1],.2,[ 0    0.4470    0.7410], 'left');
[h,mu,sigma,q,notch] = al_goodplot(rightHem,[-1,0,1],.2,[0.4660    0.6740 0.1880], 'right');
%axis([-0.4, 0.4, 50, 400])
ax = gca;
ax.FontSize = 24;
ylabel('Number of Variants')
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.7, 0.7]); %first and second
control position on screen, third controls width, and fourth controls height
xticklabels({'All Variants', 'Border Shifts', 'Ectopic Intrusions'})
xticks([-1,0,1])
m = findobj(gca, 'Type', 'Scatter')
hleg1 = legend(m(1:2), 'Right Hemisphere', 'Left Hemisphere', 'Location', 'SouthOutside');
print(gcf,
'/Users/dianaperez/Desktop/Thesis/Figures/Figure2_violin_numVarsBorderEctopic_HCP.jpg',
'-dpng', '-r300')
[h,mu,sigma,q,notch] = al_goodplot(leftHem,[-1,0,1],.2,[ 0    0.4470    0.7410], 'left');
[h,mu,sigma,q,notch] = al_goodplot(rightHem,[-1,0,1],.2,[0.4660    0.6740 0.1880], 'right');
%axis([-0.4, 0.4, 50, 400])
ax = gca;
ax.FontSize = 24;
ylabel('Number of Variants')
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.7, 0.7]); %first and second
control position on screen, third controls width, and fourth controls height
xticklabels({'All Variants', 'Border Shifts', 'Ectopic Intrusions'})
xticks([-1,0,1])
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.5, 0.7]); %first and second
control position on screen, third controls width, and fourth controls height
print(gcf,

```

```

'/Users/dianaperez/Desktop/Thesis/Figures/Figure2_violin_numVarsBorderEctopic_HCP.jpg',↵
'-dpng', '-r300')
leftHem = variants_info.left_hem.group_avg{:,3};
rightHem = variants_info.right_hem.group_avg{:,3};
[h,mu,sigma,q,notch] = al_goodplot(leftHem,0,.2,[],'left');
[h,mu,sigma,q,notch] = al_goodplot(rightHem,0,.2,[],'right');
xticks([])
axis([-0.4, 0.4, 50, 400])
ax = gca;
ax.FontSize = 24;
ylabel('Average Variant Size in Vertices')
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.3, 0.7]); %first and second↵
control position on screen, third controls width, and fourth controls height
% m = findobj(gca, 'Type', 'Scatter');
% hleg1 = legend(m(1:2), 'Right Hemisphere', 'Left Hemisphere', 'Location',↵
'SouthOutside');
print(gcf, '/Users/dianaperez/Desktop/Thesis/Figures/Figure2_violin_avgVarSize_HCP.jpg',↵
'-dpng', '-r300')
leftHem = variants_info.left_hem.group_avg{:,3};
rightHem = variants_info.right_hem.group_avg{:,3};
[h,mu,sigma,q,notch] = al_goodplot(leftHem,0,.2,[ 0      0.4470      0.7410],'left');
[h,mu,sigma,q,notch] = al_goodplot(rightHem,0,.2,[0.4660      0.6740      0.1880],'right');
xticks([0])
xticklabels('Average Variant Size')
axis([-0.4, 0.4, 50, 400])
ax = gca;
ax.FontSize = 24;
ylabel('Number of Vertices')
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.3, 0.7]); %first and second↵
control position on screen, third controls width, and fourth controls height
% m = findobj(gca, 'Type', 'Scatter');
% hleg1 = legend(m(1:2), 'Right Hemisphere', 'Left Hemisphere', 'Location',↵
'SouthOutside');
print(gcf, '/Users/dianaperez/Desktop/Thesis/Figures/Figure2_violin_avgVarSize_HCP.jpg',↵
'-dpng', '-r300')
load↵
('/Users/dianaperez/Desktop/Thesis/Variant_Properties/Permutation_Tests/HCP384_LhemvsRhem↵
_num_ectopic_ints.mat')
load↵
('/Users/dianaperez/Desktop/Thesis/Variant_Properties/Permutation_Tests/HCP384_LhemvsRhem↵
_num_border_shifts.mat')
load↵
('/Users/dianaperez/Desktop/Thesis/Variant_Properties/Permutation_Tests/HCP384_LhemvsRhem↵
_var_size.mat')
load↵
('/Users/dianaperez/Desktop/Thesis/Variant_Properties/Permutation_Tests/HCP384_LhemvsRhem↵
_num_vars.mat')
clear all
load↵
('/Users/dianaperez/Desktop/Thesis/Variant_Properties/Permutation_Tests/HCP384_LhemvsRhem↵
_num_ectopic_ints.mat')
perms_ectopic = output_mat;
perms_border = load↵
('/Users/dianaperez/Desktop/Thesis/Variant_Properties/Permutation_Tests/HCP384_LhemvsRhem↵
_num_border_shifts.mat');
load↵
('/Users/dianaperez/Desktop/Thesis/Variant_Properties/Permutation_Tests/HCP384_LhemvsRhem↵
_num_border_shifts.mat')
perms_border = output_mat;
load↵

```

```
( '/Users/dianaperez/Desktop/Thesis/Variant_Properties/Permutation_Tests/HCP384_LhemvsRhem
_var_size.mat')
perms_varSize = output_mat;
load
( '/Users/dianaperez/Desktop/Thesis/Variant_Properties/Permutation_Tests/HCP384_LhemvsRhem
_num_vars.mat')
perms_numVars = output_mat;
clear output_mat
all_perms = [perms_varSize perms_numVars perms_border perms_ectopic];
ind = zeros(1:1001);
ind = zeros(1001,1);
ind(1,1) = 1;
handles = plotSpread(all_perms, 'categoryIdx', [ind ind ind ind])
addpath '/Users/dianaperez/Documents/MATLAB/plotSpread/plotSpread'
handles = plotSpread(all_perms, 'categoryIdx', [ind ind ind ind])
handles = plotSpread(all_perms, 'categoryIdx', [ind ind ind ind], 'distributionMarkers',
{'Variant Size', 'Number of Variants', 'Number of Border Shifts', 'Number of Ectopic
Intrusions'})
handles = plotSpread(all_perms, 'distributionMarkers', {'Variant Size', 'Number of
Variants', 'Number of Border Shifts', 'Number of Ectopic Intrusions'}, 'categoryIdx',
[ind ind ind ind])
dist_labels = {'Variant Size', 'Number of Variants', 'Number of Border Shifts', 'Number
of Ectopic Intrusions'};
handles = plotSpread(all_perms, 'distributionMarkers', dist_labels, 'categoryIdx', [ind
ind ind ind])
handles = plotSpread(all_perms, 'categoryMarkers', dist_labels, 'categoryIdx', [ind ind
ind ind])
handles = plotSpread(all_perms, 'categoryMarkers', {'Variant Size', 'Number of Variants',
'Number of Border Shifts', 'Number of Ectopic Intrusions'}, 'categoryIdx', [ind ind ind
ind])
handles = plotSpread(all_perms, 'categoryMarkers', {'True Correlation', 'Permuted
Correlations'}, 'categoryIdx', [ind ind ind ind])
handles = plotSpread(all_perms, 'categoryLabels', {'True Correlation', 'Permuted
Correlations'}, 'categoryIdx', [ind ind ind ind])
all_perms = all_perms(:,2:4);
handles = plotSpread(all_perms, 'categoryLabels', {'Permuted Correlations', 'True
Correlation'}, 'categoryIdx', [ind ind ind])
handles = plotSpread(all_perms, 'categoryLabels', {'Permuted Correlations', 'True
Correlation'}, 'categoryIdx', [ind ind ind], 'xNames', {'Number of Variants', 'Number of
Border Variants', 'Number of Ectopic Variants'})
handles = plotSpread(all_perms, 'categoryLabels', {'Permuted Correlations', 'True
Correlation'}, 'categoryIdx', [ind ind ind], 'xNames', {'Number of Variants', 'Number of
Border Variants', 'Number of Ectopic Variants'}, 'xValues', [1,3,5])
plotColors(:,1) = [0,0,0];
plotColors(:,2) = [255,0,0];
plotColors(:,1) = mat2cell([0,0,0]);
for x = 1:nData
plotColors(nData,1) = [0,0,0];
plotColors(nData,2) = [255,0,0];
end
help applycform
help repmat
black_markers = [0,0,0];
red_markers = [255,0,0];
plotColors = repmat([black_markers red_markers], nData,1);
plotColors = repmat([black_markers red_markers]', nData,1);
black_markers = '[0,0,0]';
red_markers = '[255,0,0]';
plotColors = repmat([black_markers red_markers], nData,1);
plotSpread
```

```
handles = plotSpread(all_perms, 'categoryLabels', {'Permuted Correlations','True  
Correlation'}, 'categoryIdx', [ind ind ind], 'xNames', {'Number of Variants', 'Number of  
Border Variants', 'Number of Ectopic Variants'}, 'xValues', [1,3,5])  
plotColors{1,1} = [0,0,0];  
plotColors{2,1} = [0,0,0];  
plotColors{3,1} = [0,0,0];  
plotColors{1,2} = [255,0,0];  
plotColors{2,2} = [255,0,0];  
plotColors{3,2} = [255,0,0];  
ph(iData,iCategory) = plot(ah,data(currentIdx,1),...  
data(currentIdx,2),...  
'marker',plotMarkers{iData,iCategory},...  
'markeredgecolor',plotColors{iData,iCategory},...  
%'lineStyle','none',...  
'sizedata', 100,...  
'DisplayName',plotLabels{iData,iCategory});  
ph(iData,iCategory) = plot(ah,data(currentIdx,1),...  
data(currentIdx,2),...  
'marker',plotMarkers{iData,iCategory},...  
'markeredgecolor',plotColors{iData,iCategory},...  
'sizedata', 100,...  
'DisplayName',plotLabels{iData,iCategory});  
plotSpread  
handles = plotSpread(all_perms, 'categoryLabels', {'Permuted Correlations','True  
Correlation'}, 'categoryIdx', [ind ind ind], 'xNames', {'Number of Variants', 'Number of  
Border Variants', 'Number of Ectopic Variants'}, 'xValues', [1,3,5])  
plotSpread  
handles = plotSpread(all_perms, 'categoryLabels', {'Permuted Correlations','True  
Correlation'}, 'categoryIdx', [ind ind ind], 'xNames', {'Number of Variants', 'Number of  
Border Variants', 'Number of Ectopic Variants'}, 'xValues', [1,3,5])  
plotSpread  
handles = plotSpread(all_perms, 'categoryLabels', {'Permuted Correlations','True  
Correlation'}, 'categoryIdx', [ind ind ind], 'xNames', {'Number of Variants', 'Number of  
Border Variants', 'Number of Ectopic Variants'}, 'xValues', [1,3,5])  
handles = plotSpread(all_perms, 'categoryMarkers', {'bo', 'r'}, 'categoryLabels',  
{'Permuted Correlations','True Correlation'}, 'categoryIdx', [ind ind ind], 'xNames',  
{'Number of Variants', 'Number of Border Variants', 'Number of Ectopic Variants'},  
'xValues', [1,3,5])  
handles = plotSpread(all_perms, 'categoryMarkers', {'o', 'o'}, 'categoryLabels',  
{'Permuted Correlations','True Correlation'}, 'categoryIdx', [ind ind ind], 'xNames',  
{'Number of Variants', 'Number of Border Variants', 'Number of Ectopic Variants'},  
'xValues', [1,3,5])  
handles = plotSpread(all_perms, 'categoryMarkers', {'o', ''}, 'categoryLabels',  
{'Permuted Correlations','True Correlation'}, 'categoryIdx', [ind ind ind], 'xNames',  
{'Number of Variants', 'Number of Border Variants', 'Number of Ectopic Variants'},  
'xValues', [1,3,5])  
handles = plotSpread(all_perms, 'categoryMarkers', {'o', '.'}, 'categoryLabels',  
{'Permuted Correlations','True Correlation'}, 'categoryIdx', [ind ind ind], 'xNames',  
{'Number of Variants', 'Number of Border Variants', 'Number of Ectopic Variants'},  
'xValues', [1,3,5])  
plotSpread  
handles = plotSpread(all_perms, 'categoryMarkers', {'o', '.'}, 'categoryLabels',  
{'Permuted Correlations','True Correlation'}, 'categoryIdx', [ind ind ind], 'xNames',  
{'Number of Variants', 'Number of Border Variants', 'Number of Ectopic Variants'},  
'xValues', [1,3,5])  
plot(all_perms(1,:), 'MarkerEdgeColor', 'red', 'MarkerFaceColor', 'red', 'MarkerSize',  
50)  
plot(all_perms(1,:), 'MarkerEdgeColor', 'red', 'MarkerFaceColor', 'red', 'MarkerSize',  
100, 'linestyle', 'none')  
handles = plotSpread(all_perms, 'categoryMarkers', {'o', '.'}, 'categoryLabels',
```



```
{'Permuted Correlations','True Correlation'}, [ind ind ind], 'xNames',↵
{'Number of Variants', 'Number of Border Variants', 'Number of Ectopic Variants'},↵
'xValues', [1,3,5])
plot(all_perms(1,:), 'MarkerEdgeColor', 'red', 'MarkerFaceColor', 'red', 'MarkerSize',↵
100, 'linestyle', 'none')
plot([1,3,5],all_perms(1,:), 'MarkerEdgeColor', 'red', 'MarkerFaceColor', 'red',↵
'MarkerSize', 100, 'linestyle', 'none')
handles = plotSpread(all_perms(2:end,:), 'categoryMarkers', {'o'}, 'xNames', {'Number of↵
Variants', 'Number of Border Variants', 'Number of Ectopic Variants'}, 'xValues',↵
[1,3,5])
hold on
plot([1,3,5],all_perms(1,:), 'MarkerEdgeColor', 'red', 'MarkerFaceColor', 'red',↵
'MarkerSize', 100, 'linestyle', 'none')
plot([1,3,5],all_perms(1,:), 'MarkerEdgeColor', 'red', 'MarkerFaceColor', 'r',↵
'MarkerSize', 12, 'linestyle', 'none')
plot([1,3,5],all_perms(1,:), 'MarkerFaceColor', 'r', 'MarkerSize', 12, 'linestyle',↵
'none')
scatter([1,3,5],all_perms(1,:), 'MarkerFaceColor', 'r', 'SizeData', 50)
handles = plotSpread(all_perms, 'categoryMarkers', {'o', '.'}, 'categoryLabels',↵
{'Permuted Correlations','True Correlation'}, 'categoryIdx', [ind ind ind], 'xNames',↵
{'Number of Variants', 'Number of Border Variants', 'Number of Ectopic Variants'},↵
'xValues', [1,3,5])
scatter([1,3,5],all_perms(1,:), 'MarkerEdgeColor', 'red', 'MarkerFaceColor', 'red',↵
'SizeData', 50)
ax = gca;
ax.FontSize = 24;
ylabel('Number of Variants')
xticklabels({'All Variants', 'Border Shifts', 'Ectopic Intrusions'})
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.5, 0.7]); %first and second↵
control position on screen, third controls width, and fourth controls height
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.7, 0.7]); %first and second↵
control position on screen, third controls width, and fourth controls height
print(gcf,↵
'/Users/dianaperez/Desktop/Thesis/Figures/Figure3_spreadPlot_numVarsBorderEctopic_HCP.↵
jpg', '-dpng', '-r300')
handles = plotSpread(perms_varSize, 'categoryMarkers', {'o', '.'}, 'categoryLabels',↵
{'Permuted Correlations','True Correlation'}, 'categoryIdx', ind, 'xNames', {'Average↵
Variant Size'}, 'xValues', [1])
scatter(1,perms_varSize(1,:), 'MarkerEdgeColor', 'red', 'MarkerFaceColor', 'red',↵
'SizeData', 50)
ax = gca;
ax.FontSize = 24;
axis([-0.4, 0.4, 50, 400])
axis([-0.4, 0.4, 0, 1])
handles = plotSpread(perms_varSize, 'categoryMarkers', {'o', '.'}, 'categoryLabels',↵
{'Permuted Correlations','True Correlation'}, 'categoryIdx', ind, 'xNames', {'Average↵
Variant Size'}, 'xValues', [1])
scatter(1,perms_varSize(1,:), 'MarkerEdgeColor', 'red', 'MarkerFaceColor', 'red',↵
'SizeData', 50)
ax = gca;
ax.FontSize = 24;
axis([-0.4, 0.4, -25, 10])
axis([0, 2, -25, 10])
axis([0.5, 1.5, -25, 10])
axis([0, 2, -25, 10])
axis([0.25, 1.25, -25, 10])
axis([0.75, 1.25, -25, 10])
axis([0.5, 1.5, -25, 10])
ylabel('Number of Vertices')
ylabel('Difference in Number of Vertices (Left Hem - Right Hem')
```

```

ylabel('Difference in Number of Vertices')
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.7, 0.7]); %first and second
control position on screen, third controls width, and fourth controls height
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.3, 0.7]); %first and second
control position on screen, third controls width, and fourth controls height
print(gcf, '/Users/dianaperez/Desktop/Thesis/Figures/Figure3_spreadPlot_varSize_HCP.jpg',
'-dpng', '-r300')
m = findobj(gca, 'Type', 'Scatter')
handles = plotSpread(all_perms, 'categoryMarkers', {'o', '.'}, 'categoryLabels',
{'Permuted Correlations', 'True Correlation'}, 'categoryIdx', [ind ind ind], 'xNames',
{'Number of Variants', 'Number of Border Variants', 'Number of Ectopic Variants'},
'xValues', [1,3,5])
scatter([1,3,5],all_perms(1,:), 'MarkerEdgeColor', 'red', 'MarkerFaceColor', 'red',
'SizeData', 50)
ax = gca;
ax.FontSize = 24;
ylabel('Difference in Number of Variants')
xticklabels({'All Variants', 'Border Shifts', 'Ectopic Intrusions'})
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.7, 0.7]); %first and second
control position on screen, third controls width, and fourth controls height
print(gcf,
'/Users/dianaperez/Desktop/Thesis/Figures/Figure3_spreadPlot_numVarsBorderEctopic_HCP.
jpg', '-dpng', '-r300')
handles = plotSpread(all_perms, 'categoryMarkers', {'o', '.'}, 'categoryLabels',
{'Permuted Correlations', 'True Correlation'}, 'categoryIdx', [ind ind ind], 'xNames',
{'Number of Variants', 'Number of Border Variants', 'Number of Ectopic Variants'},
'xValues', [1,3,5])
scatter([1,3,5],all_perms(1,:), 'MarkerEdgeColor', 'red', 'MarkerFaceColor', 'red',
'SizeData', 50)
ax = gca;
ax.FontSize = 24;
ylabel('Difference in Number of Variants')
xticklabels({'All Variants', 'Border Shifts', 'Ectopic Intrusions'})
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.7, 0.7]); %first and second
control position on screen, third controls width, and fourth controls height
axis([0.5, 5.5, -.6, .6])
print(gcf,
'/Users/dianaperez/Desktop/Thesis/Figures/Figure3_spreadPlot_numVarsBorderEctopic_HCP.
jpg', '-dpng', '-r300')
axis([0.25, 5.75, -.6, .6])
axis([0, 6, -.6, .6])
print(gcf,
'/Users/dianaperez/Desktop/Thesis/Figures/Figure3_spreadPlot_numVarsBorderEctopic_HCP.
jpg', '-dpng', '-r300')
handles = plotSpread(all_perms, 'categoryMarkers', {'o', '.'}, 'categoryLabels',
{'Permuted Correlations', 'True Correlation'}, 'categoryColors', {'k','r'}, 'categoryIdx',
[ind ind ind], 'xNames', {'Number of Variants', 'Number of Border Variants', 'Number of
Ectopic Variants'}, 'xValues', [1,3,5])
load
('/Users/dianaperez/Desktop/Thesis/Variant_Properties/Network_Comparisons/network_perms_n
umVars.mat')
good_nets = [net_perms(:,1:3) net_perms(:,5) net_perms(:,7:12) net_perms(:,14:end)];
good_nets = [net_perms(:,1:3) net_perms(:,5) net_perms(:,7:end)];
rgb_colors = [1 0 0;
0 0 .6;
1 1 0;
.67 .67 .67;
0 .8 0;
.67 .67 .67;
0 .6 .6;

```



```

0 0 0;
.3 0 .6;
.2 1 1;
1 .5 0;
.6 .2 1;
.2 1 .2;
0 .2 .4;
0 0 1;
.8 .8 .6;
.4 0 0];
rgb_colors = [1 0 0;
0 0 .6;
1 1 0;
0 .8 0;
0 .6 .6;
0 0 0;
.3 0 .6;
.2 1 1;
1 .5 0;
.6 .2 1;
.2 1 .2;
0 .2 .4;
0 0 1;
.8 .8 .6];
handles = plotSpread(all_perms, 'categoryMarkers', {'o', '.'}, 'categoryLabels', ↵
{'Permuted Correlations','True Correlation'}, 'categoryColors', {'k','r'}, 'categoryIdx', ↵
[ind ind ind], 'xNames', {'Number of Variants', 'Number of Border Variants', 'Number of ↵
Ectopic Variants'}, 'xValues', [1,3,5])
network_names = {'DMN' 'Vis' 'FP' 'DAN' 'VAN' 'Sal' 'CO' 'SMd' 'SML' 'Aud' ↵
'Tpole' 'MTL' 'PMN' 'PON'};
handles = plotSpread(good_nets, 'categoryMarkers', {'x', '.'}, 'categoryLabels', ↵
{'Permuted Differences','True Difference'}, 'distributionColors', rgb_colors, ↵
'categoryIdx', [ind ind ind ind ind ind ind ind ind ind ind ind ind ind ind], 'xNames', ↵
network_names)
for n = 1:numel(network_names)
    rgb_for_plot{n} = rgb_color(n,:);
end
for n = 1:numel(network_names)
    rgb_for_plot{n} = rgb_colors(n,:);
end
handles = plotSpread(good_nets, 'categoryMarkers', {'x', '.'}, 'categoryLabels', ↵
{'Permuted Differences','True Difference'}, 'distributionColors', rgb_for_plot, ↵
'categoryIdx', [ind ind ind ind ind ind ind ind ind ind ind ind ind ind ind], 'xNames', )
handles = plotSpread(good_nets, 'categoryMarkers', {'x', '.'}, 'categoryLabels', ↵
{'Permuted Differences','True Difference'}, 'distributionColors', rgb_for_plot, ↵
'categoryIdx', [ind ind ind ind ind ind ind ind ind ind ind ind ind ind ind], 'xNames')
handles = plotSpread(good_nets, 'categoryMarkers', {'x', '.'}, 'categoryLabels', ↵
{'Permuted Differences','True Difference'}, 'distributionColors', rgb_for_plot, ↵
'categoryIdx', [ind ind ind ind ind ind ind ind ind ind ind ind ind ind ind])
handles = plotSpread(good_nets, 'categoryMarkers', {'x', '.'}, 'categoryLabels', ↵
{'Permuted Differences','True Difference'}, 'distributionColors', rgb_for_plot, ↵
'categoryIdx', [ind ind ind ind ind ind ind ind ind ind ind ind ind ind ind])
rgb_colors = [1 0 0;
0 0 .6;
1 1 0;
0 .8 0;
0 .6 .6;
0 0 0;
.3 0 .6;
.2 1 1;

```

```
1 .5 0;
.6 .2 1;
.2 1 .2;
0 .2 .4;
0 0 1;
.4 0 0];
rgb_for_plot = {};
for n = 1:numel(network_names)
    rgb_for_plot{n} = rgb_colors(n,:);
end
handles = plotSpread(good_nets, 'categoryMarkers', {'x', '.'}, 'categoryLabels', ↵
{'Permuted Differences', 'True Difference'}, 'distributionColors', rgb_for_plot, ↵
'categoryIdx', [ind ind ind ind ind ind ind ind ind ind ind ind ind ind], 'xNames', )
handles = plotSpread(good_nets, 'categoryMarkers', {'x', '.'}, 'categoryLabels', ↵
{'Permuted Differences', 'True Difference'}, 'distributionColors', rgb_for_plot', ↵
'categoryIdx', [ind ind ind ind ind ind ind ind ind ind ind ind ind ind])
rgb_colors = [1 0 0;
0 0 .6;
1 1 0;
0 .8 0;
0 .6 .6;
0 0 0;
.3 0 .6;
.2 1 1;
1 .5 0;
.6 .2 1;
.2 1 .2;
0 .2 .4;
0 0 1;
.8 .8 .6];
net_inds = [ind ind ind ind ind ind ind ind ind ind ind ind ind ind];
handles = plotSpread(good_nets, 'categoryMarkers', {'x', '.'}, 'categoryLabels', ↵
{'Permuted Differences', 'True Difference'}, 'distributionColors', rgb_for_plot', ↵
'categoryIdx', net_inds, 'xNames', network_names)
rgb_colors = [1 0 0;
0 0 .6;
1 1 0;
0 .8 0;
0 .6 .6;
0 0 0;
.3 0 .6;
.2 1 1;
1 .5 0;
.6 .2 1;
.2 1 .2;
0 .2 .4;
0 0 1;
.8 .8 .6];
handles = plotSpread(good_nets, 'categoryMarkers', {'x', '.'}, 'categoryLabels', ↵
{'Permuted Differences', 'True Difference'}, 'distributionColors', rgb_for_plot', ↵
'categoryIdx', net_inds, 'xNames', network_names)
rgb_for_plot = {};
for n = 1:numel(network_names)
    rgb_for_plot{n} = rgb_colors(n,:);
end
handles = plotSpread(good_nets, 'categoryMarkers', {'x', '.'}, 'categoryLabels', ↵
{'Permuted Differences', 'True Difference'}, 'distributionColors', rgb_for_plot', ↵
'categoryIdx', net_inds, 'xNames', network_names)
scatter(1,perms_varSize(1,:), 'MarkerEdgeColor', 'red', 'MarkerFaceColor', 'red', ↵
'SizeData', 50)
```

```

handles = plotSpread(good_nets, 'categoryMarkers', {'x', '.'}, 'categoryLabels',
{'Permuted Differences', 'True Difference'}, 'distributionColors', rgb_for_plot',
'categoryIdx', net_inds, 'xNames', network_names)
for n = 1:numel(network_names)
scatter(n,good_nets(1,n), 'MarkerEdgeColor', rgb_for_plot{1,n}, 'MarkerFaceColor',
rgb_for_plot{1,n}, 'SizeData', 50)
end
scatter([1:14],good_nets(1,:), 'MarkerEdgeColor', 'k', 'MarkerFaceColor', 'k',
'SizeData', 50)
scatter(6,good_nets(1,6), 'MarkerEdgeColor', 'w', 'MarkerFaceColor', 'w', 'SizeData', 50)
ylabel('Difference in Number of Variants')
xlabel('Assigned Network')
ax = gca;
ax.FontSize = 24;
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.7, 0.7]); %first and second
control position on screen, third controls width, and fourth controls height
axis([0, 15, -0.4, .7])
print(gcf, '/Users/dianaperez/Desktop/Thesis/Figures/Figure5_spreadPlot_networks_HCP.
jpg', '-dpng', '-r300')
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.9, 0.7]); %first and second
control position on screen, third controls width, and fourth controls height
print(gcf, '/Users/dianaperez/Desktop/Thesis/Figures/Figure5_spreadPlot_networks_HCP.
jpg', '-dpng', '-r300')
load('/Users/dianaperez/Desktop/Thesis/Variant_Properties/HCP384_new_split_networksxHem.
mat')
left_hem = networksxHem.clustersLH;
left_hem = [networksxHem.clustersLH(:,1:3) networksxHem.clustersLH(:,5) networksxHem.
clustersLH(:,7:end)];
right_hem = [networksxHem.clustersRH(:,1:3) networksxHem.clustersRH(:,5) networksxHem.
clustersRH(:,7:end)];
[h,mu,sigma,q,notch] = al_goodplot(leftHem,1:14,.2,[ 0 0.4470 0.7410],'left');
[h,mu,sigma,q,notch] = al_goodplot(rightHem,1:14,.2,[0.4660 0.6740
0.1880],'right');
[h,mu,sigma,q,notch] = al_goodplot(left_hem,1:14,.2,[ 0 0.4470 0.7410],'left');
rightHem = [networksxHem.clustersRH(:,1:3) networksxHem.clustersRH(:,5) networksxHem.
clustersRH(:,7:end)];
[h,mu,sigma,q,notch] = al_goodplot(rightHem,1:14,.2,[0.4660 0.6740
0.1880],'right');
axis([0, 15, -0.4, .7])
axis([0, 15, -0.1, 1])
8
axis([0, 15, -0.1, 8])
axis([0, 15, -0.1, 10])
5
axis([0, 15, -0.1, 15])
axis([0, 15, -0.1, 10])
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.9, 0.7]); %first and second
control position on screen, third controls width, and fourth controls height
load
('/Users/dianaperez/Desktop/Thesis/Spatial_Location/HCP/Handedness/LeftvsRightHanders_tru
e_spCorr_1000_permutations.mat')
load
('/Users/dianaperez/Desktop/Thesis/Spatial_Location/HCP/Handedness/PseudoRightvsPseudoLef
tHanders_spCorr_1000_permutations.mat')
clear all
load
('/Users/dianaperez/Desktop/Thesis/Spatial_Location/HCP/Handedness/LeftvsRightHanders_tru
e_spCorr_1000_permutations.mat')
load
('/Users/dianaperez/Desktop/Thesis/Spatial_Location/HCP/Handedness/PseudoRightvsPseudoLef

```

```
tHanders_spCorr_1000_permutations.mat')
scatter(500,true_spCorr, 'MarkerEdgeColor', 'red', 'MarkerFaceColor', 'red', 'SizeData', 200)
hold on
scatter(1:1000,spCorrs, 'MarkerEdgeColor', 'black', 'SizeData', 100)
axis([0, 1000, .65, 1])
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.3, 0.7]); %first and second
control position on screen, third controls width, and fourth controls height
ylabel('Correlation')
xticks([])
yticks([.65,.75,1])
ax = gca;
ax.FontSize = 24;
m = findobj(gca, 'Type', 'Scatter');
hleg1 = legend(m(1:2), 'Permuted Correlation Values', 'True Correlation', 'Location', 'SouthOutside');
hleg1.FontSize = 14;
hleg1.FontSize = 24;
scatter(500,true_spCorr, 'MarkerEdgeColor', 'red', 'MarkerFaceColor', 'red', 'SizeData', 200)
close gcf
scatter(1:1000,spCorrs, 'MarkerEdgeColor', 'black', 'SizeData', 100)
hold on
scatter(500,true_spCorr, 'MarkerEdgeColor', 'red', 'MarkerFaceColor', 'red', 'SizeData', 200)
axis([0, 1000, .65, 1])
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.3, 0.7]); %first and second
control position on screen, third controls width, and fourth controls height
ylabel('Correlation')
xticks([])
yticks([.65,.75,1])
ax = gca;
ax.FontSize = 24;
m = findobj(gca, 'Type', 'Scatter');
hleg1 = legend(m(1:2), 'Permuted Correlation Values', 'True Correlation', 'Location', 'SouthOutside');
hleg1.FontSize = 24;
print(gcf, '/Users/dianaperez/Desktop/Thesis/Figures/Figure5_spreadPlot_networks_HCP.jpg', '-dpng', '-r300')
print(gcf, '/Users/dianaperez/Desktop/Thesis/Figures/Figure6_perms_handedness_sptlLocation_HCP.jpg', '-dpng', '-r300')
load
('/Users/dianaperez/Desktop/Thesis/Spatial_Location/HCP/VariantsvsFlippedVariants_allSubs_true_spCorr_1000_thresh_10_permutations.mat')
load
('/Users/dianaperez/Desktop/Thesis/Spatial_Location/HCP/VariantsvsFlippedVariants_allSubs_spCorr_1000_thresh_10_permutations.mat')
mean(spCorrs)
load('/Users/dianaperez/Desktop/Thesis/Variant_Properties/HCP384_new_split_variants_info.mat')
clear all
load('/Users/dianaperez/Desktop/Thesis/Variant_Properties/HCP384_new_split_variants_info.mat')
avgVarSize = variants_info.left_hem.group_avg(:,3);
mean(avgVarSize)
avgVarSize = variants_info.right_hem.group_avg(:,3);
mean(avgVarSize)
135.9740 - 115.7615
load
```

```
( '/Users/dianaperez/Desktop/Thesis/Variant_Properties/Permutation_Tests/HCP384_LhemvsRhem
_var_size.mat')
true = output_mat(1,1);
perms = output_mat(2:end,1)
length(find(perms<true))
length(find(perms>true))
numVars = variants_info.left_hem.group_avg{:,1};
mean(numVars)
numVars = variants_info.right_hem.group_avg{:,1};
mean(numVars)
load
( '/Users/dianaperez/Desktop/Thesis/Variant_Properties/Permutation_Tests/HCP384_LhemvsRhem
_num_vars.mat')
true = output_mat(1,1);
perms = output_mat(2:end,1);
length(find(perms<true))
length(find(perms>true))
2/1000
load
( '/Users/dianaperez/Desktop/Thesis/Variant_Properties/Permutation_Tests/HCP384_LhemvsRhem
_num_border_shifts.mat')
LH = variants_info.left_hem.group_avg{:,4};
RH = variants_info.right_hem.group_avg{:,4};
mean(LH)
mean(RH)
true = output_mat(1,1);
perms = output_mat(2:end,1);
length(find(perms<true))
length(find(perms>true))
19/1000
LH = variants_info.left_hem.group_avg{:,5};
RH = variants_info.right_hem.group_avg{:,5};
mean(LH)
mean(RH)
load
( '/Users/dianaperez/Desktop/Thesis/Variant_Properties/Permutation_Tests/HCP384_LhemvsRhem
_num_ectopic_ints.mat')
true = output_mat(1,1);
perms = output_mat(2:end,1);
length(find(perms<true))
length(find(perms>true))
44/1000
setpref('Internet', 'SMTP_Server', 'smtp.gmail.com')
setpref('Internet', 'E-Mail', 'sdianaperezrivera@gmail.com')
setpref('Internet', 'E-Mail', 'dianaperezrivera@gmail.com')
sendmail('dianaperezrivera@gmail.com', 'Hey Loser')
mail = 'dianaperezrivera@gmail.com';
password = 'cambrea1';
password = 'Cambrea1!';
server = 'smtp.gmail.com';
props = java.lang.System.getProperties;
props.setProperty('mail.smtp.port', '587');
props.setProperty('mail.smtp.auth','true');
props.setProperty('mail.smtp.starttls.enable','true');
setpref('Internet','E-mail', mail);
setpref('Internet','SMTP_Server', server);
setpref('Internet','SMTP_Username', mail);
setpref('Internet','SMTP_Password', password);
% Send the email
sendmail('recipient_email_address@example.com', 'Test', 'Msg from MATLAB');
```

```

mail = 'dianaperezrivera2024@u.northwestern.edu';
props = java.lang.System.getProperties;
props.setProperty('mail.smtp.port', '587');
props.setProperty('mail.smtp.auth', 'true');
props.setProperty('mail.smtp.starttls.enable', 'true');
setpref('Internet', 'E_mail', mail);
setpref('Internet', 'SMTP_Server', server);
setpref('Internet', 'SMTP_Username', mail);
setpref('Internet', 'SMTP_Password', password);
% Send the email
sendmail('dianaperezrivera@gmail.com', 'Test', 'Msg from MATLAB');
help system
BIDSify_deface
cd(nifti_path)
BIDSify_deface
cd ~
BIDSify_deface
[result, output] = system('dcm2bids')
addpath('/Users/dianaperez/opt/anaconda3/bin/dcm2bids')
[result, output] = system('dcm2bids')
setenv('PATH', [getenv('PATH') '/Users/dianaperez/opt/anaconda3/bin/dcm2bids']);
[result, output] = system('dcm2bids')
[result, output] = system('echo $PATH')
cd Documents/GitHub/Lateralization_Variants/
N = 5;
x = [randn(N,1); 100+randn(N,1)]; % 5 points toward the left of the plot, and 5 toward
the right
y = sort(randn(2*N,1));
% Scatter plot, and then connect each marker on the left
% to corresponding one on the right
figure
scatter(x,y)
line([x(1:N) x(N+1:end)]', [y(1:N) y(N+1:end)]')
help line
close(gcf)
load('/Users/dianaperez/Desktop/Thesis/Variant_Properties/HCP384_new_split_variants_info.
mat')
leftHem = variants_info.left_hem.group_avg{:,3};
rightHem = variants_info.right_hem.group_avg{:,3};
[h,mu,sigma,q,notch] = al_goodplot(leftHem,0.2,[ 0 0.4470 0.7410], 'left');
[h,mu,sigma,q,notch] = al_goodplot(rightHem,0.2,[0.4660 0.6740 0.1880], 'right');
xticks([0])
xticklabels('Average Variant Size')
axis([-0.4, 0.4, 50, 400])
ax = gca;
ax.FontSize = 24;
ylabel('Number of Vertices')
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.3, 0.7]); %first and second
control position on screen, third controls width, and fourth controls height
close(gcf)
[h,mu,sigma,q,notch] = al_goodplot(leftHem,0.2,[ 0 0.4470 0.7410], 'left');
[h,mu,sigma,q,notch] = al_goodplot(rightHem,0.2,[0.4660 0.6740 0.1880], 'right');
close(gcf)
[h,mu,sigma,q,notch] = al_goodplot(leftHem,0.1,[ 0 0.4470 0.7410], 'left');
[h,mu,sigma,q,notch] = al_goodplot(rightHem,0.2,[0.4660 0.6740 0.1880], 'right');
line(leftHem', rightHem')
line(leftHem, rightHem)
N = 184;
line([leftHem(1:N) leftHem(N+1:end)]', [rightHem(1:N) rightHem(N+1:end)]')
N = 5;

```

```

x = [randn(N,1); 100+randn(N,1)]; % 5 points toward the left of the plot, and 5 toward
the right
y = sort(randn(2*N,1));
% Scatter plot, and then connect each marker on the left
% to corresponding one on the right
figure
scatter(x,y)
line([x(1:N) x(N+1:end)]',[y(1:N) y(N+1:end)]')
randn(N,1)
help randn
100+randn(N,1)
y = sort(randn(2*N,1));
y = sort(randn(2*N,1))
[x(1:N) x(N+1:end)]'
[y(1:N) y(N+1:end)]'
close(gcf)
[h,mu,sigma,q,notch] = al_goodplot(leftHem,0.1,.2,[ 0 0.4470 0.7410],'left');
[h,mu,sigma,q,notch] = al_goodplot(rightHem,0.2,.2,[0.4660 0.6740 0.1880],'right');
y = zeros(384,2);
y(:,1) = 0.1;
y(:,2) = 0.2;
x(:,1) = leftHem;
x = [leftHem rightHem];
line([x(1:384) x(385:end)]',[y(1:384) y(385:end)]')
close(gcf)
[h,mu,sigma,q,notch] = al_goodplot(leftHem,0.1,.2,[ 0 0.4470 0.7410],'left');
[h,mu,sigma,q,notch] = al_goodplot(rightHem,0.2,.2,[0.4660 0.6740 0.1880],'right');
line(x',y')
x = x';
y = y';
z = y;
y = x;
x = z;
close(gcf)
[h,mu,sigma,q,notch] = al_goodplot(leftHem,0.1,.2,[ 0 0.4470 0.7410],'left');
[h,mu,sigma,q,notch] = al_goodplot(rightHem,0.2,.2,[0.4660 0.6740 0.1880],'right');
line(x,y)
clear all
load('/Users/dianaperez/Desktop/Thesis/Variant_Properties/HCP384_new_split_variants_info.
mat')
varID = variants_info.left_hem.indiv_vars{1,1};
LHem_numVarVerts = sum(variants_info.left_hem.indiv_vars{1,1});
varSize = variants_info.left_hem.indiv_vars{: ,2};
LHem_numVarVerts = sum(variants_info.left_hem.indiv_vars{: ,2});
length(variants_info.left_hem.indiv_vars)
height(variants_info.left_hem.indiv_vars)
variant_verts = [];
for var = 1:height(variants_info.left_hem.indiv_vars)
for sub = 1:384
sub_varVerts = [];
if variants_info.left_hem.indiv_vars{var,1} > variants_info.left_hem.indiv_vars{var+1,1}
sub_varVerts = [sub_varVerts + variants_info.left_hem.indiv_vars{var,2}];
else
variant_verts(sub) = sub_varVerts;
end
end
end
count_variant_vertices
variants_info.left_hem.indiv_vars{var,1}
count_variant_vertices

```

```
variants_info.left_hem.indiv_vars{var,2}
0 + variants_info.left_hem.indiv_vars{var,2}
count_variant_vertices
variants_info.left_hem.indiv_vars{var,1}
count_variant_vertices
varSize1 = variants_info.left_hem.indiv_vars{1:17,2};
sum(varSize1)
count_variant_vertices
variant_verts = variant_verts';
variant_verts(384,1) = sub_varVerts;
LH_variant_verts = variant_verts;
count_variant_vertices
RH_variant_verts = variant_verts;
variant_verts = [LH_variant_verts RH_variant_verts];
permute_variantproperties
mean(allSubs_info.variants_info.left_hem.group_avg{:,3})
mean(allSubs_info.variants_info.right_hem.group_avg{:,3})
mean(allSubs_info.variants_info.left_hem.group_avg{:,2})
mean(allSubs_info.variants_info.left_hem.group_avg{:,3})
mean(allSubs_info.variants_info.left_hem.group_avg{:,2})
mean(allSubs_info.variants_info.right_hem.group_avg{:,2})
%-- 9/18/21, 5:05 PM --%
operc_cluster
overlap_bin(vertex)
neigh_verts = neigh(vertex);
neigh_verts = neigh(:,vertex);
neigh_verts = neigh(vertex,:);
operc_cluster
neigh_verts = neigh(vertices(ver),:);
operc_cluster
vertices = unique(vertices);
for x=1:20
for vert = 1:length(vertices)
neigh_verts = neigh(vertices(vert),:);
for n = 1:length(neigh_verts)
if overlap_bin(neigh_verts(n)) == 1
vertices = [vertices; neigh_verts(n)];
end
end
end
vertices = unique(vertices);
end
vertices = unique(vertices);
neigh_verts(7)
if neigh_verts(7)==NaN
disp('hello')
end
maybe_mask = vertices;
operc_cluster
help isnan
operc_cluster
vertices = unique(vertices);
operc_cluster
maybe_maybe_mask = vertices;
operc_mask(vertices) = 1;
template = overlap;
template.data = operc_mask;
ft_write_cifti_mod('Users/dianaperez/Desktop/operc_mask_v1.dtseries.nii', template)
ft_write_cifti_mod('/Users/dianaperez/Desktop/operc_mask_v1.dtseries.nii', template)
operc_mask(maybe_maybe_mask) = 1;
```



```
template.data = operc_mask;
ft_write_cifti_mod('/Users/dianaperez/Desktop/operc_mask_v2_20_iterations.dtseries.nii', ↵
template)
operc_mask(vertices) = 1;
template.data = operc_mask;
ft_write_cifti_mod('/Users/dianaperez/Desktop/operc_mask_v2_30_iterations.dtseries.nii', ↵
template)
operc_mask(maybe_mask) = 1;
template.data = operc_mask;
ft_write_cifti_mod('/Users/dianaperez/Desktop/operc_mask_v1.dtseries.nii', template)
operc_mask(maybe_mask) = 1;
template.data = operc_mask;
ft_write_cifti_mod('/Users/dianaperez/Desktop/operc_mask_v1_958_verts.dtseries.nii', ↵
template)
operc_mask = zeros(numVerts,1);
operc_mask(maybe_mask) = 1;
template.data = operc_mask;
ft_write_cifti_mod('/Users/dianaperez/Desktop/operc_mask_v1_958_verts.dtseries.nii', ↵
template)
operc_mask = zeros(numVerts,1);
operc_mask(maybe_mask) = 1;
template.data = operc_mask;
ft_write_cifti_mod('/Users/dianaperez/Desktop/operc_mask_v2_1113_verts.dtseries.nii', ↵
template)
operc_mask = zeros(numVerts,1);
operc_mask(vertices) = 1;
template.data = operc_mask;
ft_write_cifti_mod('/Users/dianaperez/Desktop/operc_mask_v3_1889_verts.dtseries.nii', ↵
template)
operc_cluster
operc_mask = zeros(numVerts,1);
operc_mask(vertices) = 1;
template.data = operc_mask;
ft_write_cifti_mod('/Users/dianaperez/Desktop/operc_mask_v4.dtseries.nii', template)
operc_cluster
operc_mask = zeros(numVerts,1);
operc_mask(vertices) = 1;
template.data = operc_mask;
ft_write_cifti_mod('/Users/dianaperez/Desktop/operc_mask_v5.dtseries.nii', template)
operc_cluster
operc_mask = zeros(numVerts,1);
operc_mask(vertices) = 1;
template.data = operc_mask;
ft_write_cifti_mod('/Users/dianaperez/Desktop/operc_mask_v6.dtseries.nii', template)
operc_cluster
ft_write_cifti_mod('/Users/dianaperez/Desktop/operc_mask_moreRHvars.dtseries.nii', ↵
template)
operc_cluster
if unique_IDs(1) == 0
unique_IDs(1) = [];
end
operc_cluster
find(var_map.data==sub_data(var,1));
operc_cluster
LH_cluster = [];
LH_cluster.number = LH_cluster_number;
LH_cluster.proportion = LH_cluster_proportion;
save([output_dir 'operc_nets_lefthemcluster.mat'], 'LH_cluster')
output_dir = '/Users/dianaperez/Desktop';
save([output_dir 'operc_nets_lefthemcluster.mat'], 'LH_cluster')
```

```
RH_cluster = [];  
RH_cluster.number = RH_cluster_number;  
RH_cluster.proportion = RH_cluster_proportion;  
save([output_dir 'operc_nets_righthemcluster.mat'], 'RH_cluster')  
operc_cluster  
RH_mask_flipped = [RH_mask_gifti(32493:end); RH_mask_gifti(1:32492)];  
LH_mask_flipped = [LH_mask_gifti(32493:end); LH_mask_gifti(1:32492)];  
RH_mask_flipped = [RH_mask_gifti(32493:end); RH_mask_gifti(1:32492)];  
LH_mask_flipped = [LH_mask_gifti(32493:end); LH_mask_gifti(1:32492)];  
RH_mask_flipped = RH_mask_flipped(diffmap.brainstructure>0);  
LH_mask_flipped = LH_mask_flipped(diffmap.brainstructure>0);  
operc_cluster  
load([output_dir 'operc_nets_lefthemcluster_righthem.mat']);  
load([output_dir 'operc_nets_lefthemcluster_righthem.mat']);  
clear all  
load([output_dir 'operc_nets_lefthemcluster_righthem.mat']);  
output_dir = '/Users/dianaperez/Desktop/Thesis/Spatial_Location/';  
load([output_dir 'operc_nets_lefthemcluster_righthem.mat']);  
leftclust_righthem = LH_cluster;  
load([output_dir 'operc_nets_lefthemcluster_lefthem.mat']);  
leftclust_lefthem = LH_cluster;  
clear LH_cluster  
t = tiledlayout(1,2,'TileSpacing', 'compact');  
ax1 = nexttile;  
pie(ax1,leftclust_lefthem)  
title('Left Hem')  
pie(ax1,leftclust_lefthem.number)  
title('Left Hem')  
pie(ax1,leftclust_lefthem.number{:},3)  
number = leftclust_lefthem.number{1,3}  
number = leftclust_lefthem.number(:,3);  
close(gcf)  
t = tiledlayout(1,2,'TileSpacing', 'compact');  
ax1 = nexttile;  
pie(ax1,leftclust_lefthem.number(:,3))  
title('Left Hem')  
close(gcf)  
for net = 1:16  
nets(net) = length(find(leftclust_lefthem.number(:,3)==net));  
end  
t = tiledlayout(1,2,'TileSpacing', 'compact');  
ax1 = nexttile;  
pie(ax1,nets)  
title('Left Hem')  
for net = 1:16  
nets_right(net) = length(find(leftclust_righthem.number(:,3)==net));  
end  
ax2 = nexttile;  
pie(ax2,nets_right)  
title('Right Hem')  
ax = gca;  
clear all  
rgb_colors = [1 0 0;  
0 0 .6;  
1 1 0;  
0 .8 0;  
0 .6 .6;  
0 0 0;  
.3 0 .6;  
.2 1 1;
```

```
1 .5 0;
.6 .2 1;
.2 1 .2;
0 .2 .4;
0 0 1;
.8 .8 .6];
load([output_dir 'operc_nets_lefthemcluster_righthem.mat']);
leftclust_righthem = LH_cluster;
load([output_dir 'operc_nets_lefthemcluster_lefthem.mat']);
leftclust_lefthem = LH_cluster;
clear LH_cluster
for net = 1:16
nets_left(net) = length(find(leftclust_lefthem.number(:,3)==net));
end
nets_left(4) = [];
nets_left(6) = [];
t = tiledlayout(1,2,'TileSpacing', 'compact');
ax1 = nexttile;
h = pie(ax1,nets_left)
title('Left Hem')
for net = 1:16
nets_right(net) = length(find(leftclust_righthem.number(:,3)==net));
end
nets_right(4) = [];
nets_right(6) = [];
ax2 = nexttile;
h = pie(ax2,nets_right)
title('Right Hem')
output_dir = '/Users/dianaperez/Desktop/Thesis/Spatial_Location/';
load([output_dir 'operc_nets_lefthemcluster_righthem.mat']);
leftclust_righthem = LH_cluster;
load([output_dir 'operc_nets_lefthemcluster_lefthem.mat']);
leftclust_lefthem = LH_cluster;
clear LH_cluster
for net = 1:16
nets_left(net) = length(find(leftclust_lefthem.number(:,3)==net));
end
nets_left(4) = [];
nets_left(6) = [];
t = tiledlayout(1,2,'TileSpacing', 'compact');
ax1 = nexttile;
h = pie(ax1,nets_left)
title('Left Hem')
for net = 1:16
nets_right(net) = length(find(leftclust_righthem.number(:,3)==net));
end
nets_right(4) = [];
nets_right(6) = [];
ax2 = nexttile;
h = pie(ax2,nets_right)
title('Right Hem')
patchHande = findobj(h, 'Type', 'Patch');
set(patchHand, {'FaceColor'}, mat2cell(rgb_colors, ones(size(rgb_colors,1),1),3)
set(patchHande, {'FaceColor'}, mat2cell(rgb_colors, ones(size(rgb_colors,1),1),3))
close(gcf)
output_dir = '/Users/dianaperez/Desktop/Thesis/Spatial_Location/';
load([output_dir 'operc_nets_lefthemcluster_righthem.mat']);
leftclust_righthem = LH_cluster;
load([output_dir 'operc_nets_lefthemcluster_lefthem.mat']);
leftclust_lefthem = LH_cluster;
```

```

clear LH_cluster
for net = 1:16
nets_left(net) = length(find(leftclust_lefthem.number(:,3)==net));
end
nets_left(4) = [];
nets_left(6) = [];
t = tiledlayout(1,2,'TileSpacing', 'compact');
ax1 = nexttile;
h = pie(ax1,nets_left);
title('Left Hem')
patchHandle = findobj(h, 'Type', 'Patch');
set(patchHandle, {'FaceColor'}, mat2cell(rgb_colors, ones(size(rgb_colors,1),1),3))
for net = 1:16
nets_right(net) = length(find(leftclust_righthem.number(:,3)==net));
end
nets_right(4) = [];
nets_right(6) = [];
ax2 = nexttile;
h = pie(ax2,nets_right);
title('Right Hem')
patchHandle = findobj(h, 'Type', 'Patch');
set(patchHandle, {'FaceColor'}, mat2cell(rgb_colors, ones(size(rgb_colors,1),1),3))
for net = 1:16
nets_left(net) = length(find(leftclust_lefthem.number(:,3)==net));
end
nets_left(6) = [];
nets_left(4) = [];
t = tiledlayout(1,2,'TileSpacing', 'compact');
ax1 = nexttile;
h = pie(ax1,nets_left);
title('Left Hem')
patchHandle = findobj(h, 'Type', 'Patch');
set(patchHandle, {'FaceColor'}, mat2cell(rgb_colors, ones(size(rgb_colors,1),1),3))
%network_names = {'DMN' 'Vis' 'FP' 'DAN' 'Lang' 'Sal' 'CO' 'SMd' 'SML' 'Aud'};
for net = 1:16
nets_right(net) = length(find(leftclust_righthem.number(:,3)==net));
end
nets_right(6) = [];
nets_right(4) = [];
ax2 = nexttile;
h = pie(ax2,nets_right);
title('Right Hem')
patchHandle = findobj(h, 'Type', 'Patch');
set(patchHandle, {'FaceColor'}, mat2cell(rgb_colors, ones(size(rgb_colors,1),1),3))
close(gcf)
rgb_colors = [1 0 0;
1 1 0;
0 .8 0;
0 .6 .6;
0 0 0;
.3 0 .6;
1 .5 0;];
output_dir = '/Users/dianaperez/Desktop/Thesis/Spatial_Location/';
load([output_dir 'operc_nets_lefthemcluster_righthem.mat']);
leftclust_righthem = LH_cluster;
load([output_dir 'operc_nets_lefthemcluster_lefthem.mat']);
leftclust_lefthem = LH_cluster;
clear LH_cluster
for net = 1:16
nets_left(net) = length(find(leftclust_lefthem.number(:,3)==net));

```

```

end
nets_left(6) = [];
nets_left(4) = [];
t = tiledlayout(1,2,'TileSpacing', 'compact');
ax1 = nexttile;
h = pie(ax1,nets_left, network_names);
title('Left Hem')
patchHandle = findobj(h, 'Type', 'Patch');
set(patchHandle, {'FaceColor'}, mat2cell(rgb_colors, ones(size(rgb_colors,1),1),3))
network_names = {'DMN' 'FP' 'DAN' 'Lang' 'Sal' 'CO' 'SML'};
for net = 1:16
nets_right(net) = length(find(leftclust_righthem.number(:,3)==net));
end
nets_right(6) = [];
nets_right(4) = [];
ax2 = nexttile;
h = pie(ax2,nets_right, network_names);
title('Right Hem')
patchHandle = findobj(h, 'Type', 'Patch');
set(patchHandle, {'FaceColor'}, mat2cell(rgb_colors, ones(size(rgb_colors,1),1),3))
labels_legend = {'Default Mode' 'Frontoparietal', 'Dorsal Attention', 'Language', 'Salience', 'Cingulo-opercular', 'Somatomotor lateral'};
lgd = legend(labels_legend);
lgd.Layout.Tile = 'east';
rgb_colors = [1 0 0;
1 1 0;
0 .8 0;
0 .6 .6;
0 0 0;
.3 0 .6;
1 .5 0;];
output_dir = '/Users/dianaperez/Desktop/Thesis/Spatial_Location/';
load([output_dir 'operc_nets_lefthemcluster_righthem.mat']);
leftclust_righthem = LH_cluster;
load([output_dir 'operc_nets_lefthemcluster_lefthem.mat']);
leftclust_lefthem = LH_cluster;
clear LH_cluster
network_names = {'DMN' 'FP' 'DAN' 'Lang' 'Sal' 'CO' 'SML'};
for net = 1:16
nets_left(net) = length(find(leftclust_lefthem.number(:,3)==net));
end
nets_left(6) = [];
nets_left(4) = [];
t = tiledlayout(1,2,'TileSpacing', 'compact');
ax1 = nexttile;
h = pie(ax1,nets_left, network_names);
title('Left Hem')
patchHandle = findobj(h, 'Type', 'Patch');
set(patchHandle, {'FaceColor'}, mat2cell(rgb_colors, ones(size(rgb_colors,1),1),3))
for net = 1:16
nets_right(net) = length(find(leftclust_righthem.number(:,3)==net));
end
nets_right(6) = [];
nets_right(4) = [];
ax2 = nexttile;
h = pie(ax2,nets_right, network_names);
title('Right Hem')
patchHandle = findobj(h, 'Type', 'Patch');
set(patchHandle, {'FaceColor'}, mat2cell(rgb_colors, ones(size(rgb_colors,1),1),3))
labels_legend = {'Default Mode' 'Frontoparietal', 'Dorsal Attention', 'Language', 'Salience', 'Cingulo-opercular', 'Somatomotor lateral'};

```

```

'Salience', 'Cingulo-opercular', 'Somatomotor lateral'};
lgd = legend(labels_legend);
lgd.Layout.Tile = 'east';
rgb_colors = [1 0 0;
1 1 0;
0 .8 0;
0 .6 .6;
0 0 0;
.3 0 .6;
1 .5 0];
output_dir = '/Users/dianaperez/Desktop/Thesis/Spatial_Location/';
load([output_dir 'operc_nets_lefthemcluster_righthem.mat']);
leftclust_righthem = LH_cluster;
load([output_dir 'operc_nets_lefthemcluster_lefthem.mat']);
leftclust_lefthem = LH_cluster;
clear LH_cluster
network_names = {'DMN' 'FP' 'DAN' 'Lang' 'Sal' 'CO' 'SML'};
for net = 1:16
nets_left(net) = length(find(leftclust_lefthem.number(:,3)==net));
end
nets_left(10:end) = [];
nets_left(8) = [];
nets_left(6) = [];
nets_left(4) = [];
nets_left(2) = [];
t = tiledlayout(1,2,'TileSpacing', 'compact');
ax1 = nexttile;
h = pie(ax1,nets_left, network_names);
title('Left Hem')
patchHandle = findobj(h, 'Type', 'Patch');
set(patchHandle, {'FaceColor'}, mat2cell(rgb_colors, ones(size(rgb_colors,1),1),3))
for net = 1:16
nets_right(net) = length(find(leftclust_righthem.number(:,3)==net));
end
nets_right(10:end) = [];
nets_right(8) = [];
nets_right(6) = [];
nets_right(4) = [];
nets_right(2) = [];
ax2 = nexttile;
h = pie(ax2,nets_right, network_names);
title('Right Hem')
patchHandle = findobj(h, 'Type', 'Patch');
set(patchHandle, {'FaceColor'}, mat2cell(rgb_colors, ones(size(rgb_colors,1),1),3))
labels_legend = {'Default Mode' 'Frontoparietal', 'Dorsal Attention', 'Language',
'Salience', 'Cingulo-opercular', 'Somatomotor lateral'};
lgd = legend(labels_legend);
lgd.Layout.Tile = 'east';
for net = 1:16
nets_left(net) = length(find(leftclust_lefthem.number(:,3)==net));
end
for net = 1:16
nets_right(net) = length(find(leftclust_righthem.number(:,3)==net));
end
nets_left(6) = [];
nets_left(4) = [];
nets_right(6) = [];
nets_right(4) = [];
nets_left(10:end) = [];
nets_right(10:end) = [];

```

```

nets_left(8) = [];
nets_right(8) = [];
nets_left(2) = [];
nets_right(2) = [];
t = tiledlayout(1,2,'TileSpacing', 'compact');
ax1 = nexttile;
h = pie(ax1,nets_left, network_names);
title('Left Hem')
patchHandle = findobj(h, 'Type', 'Patch');
set(patchHandle, {'FaceColor'}, mat2cell(rgb_colors, ones(size(rgb_colors,1),1),3))
h = pie(ax2,nets_right, network_names);
title('Right Hem')
patchHandle = findobj(h, 'Type', 'Patch');
set(patchHandle, {'FaceColor'}, mat2cell(rgb_colors, ones(size(rgb_colors,1),1),3))
labels_legend = {'Default Mode' 'Frontoparietal', 'Dorsal Attention', 'Language', 'Salience', 'Cingulo-opercular', 'Somatomotor lateral'};
lgd = legend(labels_legend);
lgd.Layout.Tile = 'east';
close(gcf)
rgb_colors = [1 0 0;
1 1 0;
0 .8 0;
0 .6 .6;
0 0 0;
.3 0 .6;
1 .5 0;];
output_dir = '/Users/dianaperez/Desktop/Thesis/Spatial_Location/';
load([output_dir 'operc_nets_lefthemcluster_righthem.mat']);
leftclust_righthem = LH_cluster;
load([output_dir 'operc_nets_lefthemcluster_lefthem.mat']);
leftclust_lefthem = LH_cluster;
clear LH_cluster
network_names = {'DMN' 'FP' 'DAN' 'Lang' 'Sal' 'CO' 'SML'};
for net = 1:16
nets_left(net) = length(find(leftclust_lefthem.number(:,3)==net));
end
nets_left(6) = [];
nets_left(4) = [];
nets_left(10:end) = [];
nets_left(8) = [];
nets_left(2) = [];
t = tiledlayout(1,2,'TileSpacing', 'compact');
ax1 = nexttile;
h = pie(ax1,nets_left, network_names);
title('Left Hem')
patchHandle = findobj(h, 'Type', 'Patch');
set(patchHandle, {'FaceColor'}, mat2cell(rgb_colors, ones(size(rgb_colors,1),1),3))
for net = 1:16
nets_right(net) = length(find(leftclust_righthem.number(:,3)==net));
end
nets_right(6) = [];
nets_right(4) = [];
nets_right(10:end) = [];
nets_right(8) = [];
nets_right(2) = [];
ax2 = nexttile;
h = pie(ax2,nets_right, network_names);
title('Right Hem')
patchHandle = findobj(h, 'Type', 'Patch');
set(patchHandle, {'FaceColor'}, mat2cell(rgb_colors, ones(size(rgb_colors,1),1),3))

```

```
labels_legend = {'Default Mode' 'Frontoparietal', 'Dorsal Attention', 'Language', 'Salience', 'Cingulo-opercular', 'Somatomotor lateral'};
lgd = legend(labels_legend);
lgd.Layout.Tile = 'east';
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.9, 0.7]); %first and second control position on screen, third controls width, and fourth controls height
ax1.FontSize = 24;
ax2.FontSize = 24;
h.FontSize = 21
lgd.FontSize = 21
lgd.FontSize = 24
print(gcf, 'Users/dianaperez/Desktop/Thesis/Figures/Figure7_pieChart_operc_cluster_networks_lefthem_HCP.jpg', '-dpng', '-r300')
output_dir = '/Users/dianaperez/Desktop/Thesis/Spatial_Location/';
load([output_dir 'operc_nets_righthemcluster_righthem.mat']);
leftclust_righthem = LH_cluster;
load([output_dir 'operc_nets_righthemcluster_lefthem.mat']);
leftclust_lefthem = LH_cluster;
clear LH_cluster
leftclust_righthem = RH_cluster;
output_dir = '/Users/dianaperez/Desktop/Thesis/Spatial_Location/';
load([output_dir 'operc_nets_righthemcluster_righthem.mat']);
leftclust_righthem = RH_cluster;
load([output_dir 'operc_nets_righthemcluster_lefthem.mat']);
leftclust_lefthem = RH_cluster;
clear RH_cluster
clear all
rgb_colors = [1 0 0;
1 1 0;
0 .8 0;
0 .6 .6;
0 0 0;
.3 0 .6;
1 .5 0;];
output_dir = '/Users/dianaperez/Desktop/Thesis/Spatial_Location/';
load([output_dir 'operc_nets_righthemcluster_righthem.mat']);
rightclust_righthem = RH_cluster;
load([output_dir 'operc_nets_righthemcluster_lefthem.mat']);
rightclust_lefthem = RH_cluster;
clear RH_cluster
for net = 1:16
nets_left(net) = length(find(rightclust_lefthem.number(:,3)==net));
end
nets_left(6) = [];
nets_left(4) = [];
for net = 1:16
nets_right(net) = length(find(leftclust_righthem.number(:,3)==net));
end
for net = 1:16
nets_right(net) = length(find(rightclust_righthem.number(:,3)==net));
end
nets_right(6) = [];
nets_right(4) = [];
clear all
rgb_colors = [1 0 0;
1 1 0;
0 .8 0;
0 .6 .6;
0 0 0;
```



```

.3 0 .6;
1 .5 0;
.8 .8 .6;];
output_dir = '/Users/dianaperez/Desktop/Thesis/Spatial_Location/';
load([output_dir 'operc_nets_righthemcluster_righthem.mat']);
rightclust_righthem = RH_cluster;
load([output_dir 'operc_nets_righthemcluster_lefthem.mat']);
rightclust_lefthem = RH_cluster;
clear RH_cluster
network_names = {'DMN' 'FP' 'DAN' 'Lang' 'Sal' 'CO' 'SMl' 'PON'};
for net = 1:16
nets_left(net) = length(find(rightclust_lefthem.number(:,3)==net));
end
nets_left(6) = [];
nets_left(4) = [];
nets_left(10:13) = [];
nets_left(8) = [];
nets_left(2) = [];
t = tiledlayout(1,2,'TileSpacing', 'compact');
ax1 = nexttile;
h = pie(ax1,nets_left, network_names);
title('Left Hem')
patchHandle = findobj(h, 'Type', 'Patch');
set(patchHandle, {'FaceColor'}, mat2cell(rgb_colors, ones(size(rgb_colors,1),1),3))
for net = 1:16
nets_right(net) = length(find(rightclust_righthem.number(:,3)==net));
end
nets_right(6) = [];
nets_right(4) = [];
nets_right(10:13) = [];
nets_right(8) = [];
nets_right(2) = [];
ax2 = nexttile;
h = pie(ax2,nets_right, network_names);
title('Right Hem')
patchHandle = findobj(h, 'Type', 'Patch');
set(patchHandle, {'FaceColor'}, mat2cell(rgb_colors, ones(size(rgb_colors,1),1),3))
labels_legend = {'Default Mode' 'Frontoparietal', 'Dorsal Attention', 'Language',
'Salience', 'Cingulo-Opercular', 'Somatomotor Ventral' 'Parieto-Occipital'};
lgd = legend(labels_legend);
lgd.Layout.Tile = 'east';
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.9, 0.7]); %first and second
control position on screen, third controls width, and fourth controls height
ax1.FontSize = 24;
ax2.FontSize = 24;
lgd.FontSize = 24;
print(gcf,
'/Users/dianaperez/Desktop/Thesis/Figures/Figure7_pieChart_operc_cluster_networks_righthe
m_HCP.jpg', '-dpng', '-r300')
%-- 9/20/21, 7:08 PM --%
load('/Users/dianaperez/Desktop/Thesis/Variant_Properties/HCP384_new_split_networksxHem.
mat')
network_names = {'DMN' 'Vis' 'FP' ' ' 'DAN' ' ' 'VAN' 'Sal' 'CO' 'SMd'
'SML' 'Aud' 'Tpole' 'MTL' 'PMN' 'PON'};
%network_names = {'DMN' 'Vis' 'FP' 'DAN' 'VAN' 'Sal' 'CO' 'SMd' 'SMl'
'Aud' 'Tpole' 'MTL' 'PMN' 'PON'};
results = [];
statistics = [];
for net = 1:numel(network_names)
left_hemi = networksxHem.clustersLH(:,net);

```

```

right_hemi = networksxHem.clustersRH(:,net);
results(net,1) = mean(left_hemi);
results(net,2) = mean(right_hemi);
[sig p ci stats] = ttest(left_hemi, right_hemi);
results(net,3) = sig;
results(net,4) = p;
statistics{net} = stats;
end
get(gca, 'ColorOrder')
load('/Users/dianaperez/Desktop/Thesis/Variant_Properties/HCP384_new_split_variants_info.
mat')
numVarsLH = variants_info.left_hem.group_avg{:,1};
numVarsRH = variants_info.right_hem.group_avg{:,1};
numBorderLH = variants_info.left_hem.group_avg{:,4};
numBorderRH = variants_info.right_hem.group_avg{:,4};
numEctopicLH = variants_info.left_hem.group_avg{:,5};
numEctopicRH = variants_info.right_hem.group_avg{:,5};
leftHem = [numVarsLH numBorderLH numEctopicLH];
rightHem = [numVarsRH numBorderRH numEctopicRH];
[h,mu,sigma,q,notch] = al_goodplot(leftHem,[-1,0,1],.2,[ 0    0.4470    0.7410],'left');
[h,mu,sigma,q,notch] = al_goodplot(rightHem,[-1,0,1],.2,[0.8500    0.3250
0.0980],'right');
ax = gca;
ax.FontSize = 24;
ylabel('Number of Variants')
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.5, 0.7]); %first and second
control position on screen, third controls width, and fourth controls height
xticklabels({'All Variants', 'Border Shifts', 'Ectopic Intrusions'})
xticks([-1,0,1])
m = findobj(gca, 'Type', 'Scatter')
hleg1 = legend(m(1:2), 'Right Hemisphere', 'Left Hemisphere', 'Location',
'SouthOutside');
print(gcf,
'/Users/dianaperez/Desktop/Thesis/Figures/Figure2_violin_numVarsBorderEctopic_HCP.jpg',
'-dpng', '-r300')
load('/Users/dianaperez/Desktop/Thesis/Variant_Properties/HCP384_new_split_variants_info.
mat')
leftHem = variants_info.left_hem.group_avg{:,3};
rightHem = variants_info.right_hem.group_avg{:,3};
[h,mu,sigma,q,notch] = al_goodplot(leftHem,0.1,.2,[ 0    0.4470    0.7410],'left');
[h,mu,sigma,q,notch] = al_goodplot(rightHem,0.2,.2,[0.8500    0.3250    0.0980],'right');
xticks([0])
xticklabels('Average Variant Size')
axis([-0.4, 0.4, 50, 400])
ax = gca;
ax.FontSize = 24;
ylabel('Number of Vertices')
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.3, 0.7]); %first and second
control position on screen, third controls width, and fourth controls height
% m = findobj(gca, 'Type', 'Scatter');
% hleg1 = legend(m(1:2), 'Right Hemisphere', 'Left Hemisphere', 'Location',
'SouthOutside');
print(gcf, '/Users/dianaperez/Desktop/Thesis/Figures/Figure2_violin_avgVarSize_HCP.jpg',
'-dpng', '-r300')
load('/Users/dianaperez/Desktop/Thesis/Variant_Properties/HCP384_new_split_variants_info.
mat')
leftHem = variants_info.left_hem.group_avg{:,3};
rightHem = variants_info.right_hem.group_avg{:,3};
[h,mu,sigma,q,notch] = al_goodplot(leftHem,0.1,.2,[ 0    0.4470    0.7410],'left');
[h,mu,sigma,q,notch] = al_goodplot(rightHem,0.2,.2,[0.8500    0.3250    0.0980],'right');

```

```

xticks([0])
xlabel('Average Variant Size')
axis([-0.2, 0.5, 50, 400])
ax = gca;
ax.FontSize = 24;
ylabel('Number of Vertices')
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.3, 0.7]); %first and second
control position on screen, third controls width, and fourth controls height
% m = findobj(gca, 'Type', 'Scatter');
% hleg1 = legend(m(1:2), 'Right Hemisphere', 'Left Hemisphere', 'Location',
'SouthOutside');
print(gcf, '/Users/dianaperez/Desktop/Thesis/Figures/Figure2_violin_avgVarSize_HCP.jpg',
'-dpng', '-r300')
load('/Users/dianaperez/Desktop/Thesis/Variant_Properties/HCP384_new_split_variants_info.
mat')
leftHem = variants_info.left_hem.group_avg(:,3);
rightHem = variants_info.right_hem.group_avg(:,3);
[h,mu,sigma,q,notch] = al_goodplot(leftHem,0.1,.2,[ 0 0.4470 0.7410],'left');
[h,mu,sigma,q,notch] = al_goodplot(rightHem,0.2,.2,[0.8500 0.3250 0.0980],'right');
xticks([0])
xlabel('Average Variant Size')
axis([-0.2, 0.5, 50, 400])
ax = gca;
ax.FontSize = 24;
ylabel('Number of Vertices')
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.3, 0.7]); %first and second
control position on screen, third controls width, and fourth controls height
m = findobj(gca, 'Type', 'Scatter');
hleg1 = legend(m(1:2), 'Right Hemisphere', 'Left Hemisphere', 'Location',
'SouthOutside');
print(gcf, '/Users/dianaperez/Desktop/Thesis/Figures/Figure2_violin_avgVarSize_HCP.jpg',
'-dpng', '-r300')
%-- 10/2/21, 7:40 PM --%
permute_variantproperties
avgSurfArea = allSubs_info.variants_info.left_hem.group_avg(:,4);
mean(avgSurfArea)
avgSurfArea = allSubs_info.variants_info.right_hem.group_avg(:,4);
mean(avgSurfArea)
[true_diff perm_diffs] = permute_values(LHand_info.variants_info.both_hems(:,4),
RHand_info.variants_info.both_hems(:,4), flip_switch, numperms, 1, 0, 1);
[true_diff perm_diffs] = permute_values(allSubs_info.variants_info.left_hem.group_avg(:,4),
allSubs_info.variants_info.right_hem.group_avg(:,4), flip_switch, numperms, 0, 1, 0);
disp(['Average variant surface area (permute hemispheres) - Below: ' num2str(length(find
(perm_diffs<true_diff))/1000)])
disp(['Average variant surface area (permute hemispheres) - Above: ' num2str(length(find
(perm_diffs>true_diff))/1000)])
output_mat = [true_diff; perm_diffs'];
save([outputDir str 'avgSurfArea.mat'], 'output_mat'); clear output_mat
5:3
1:2
3:5
3/5
10/7
left_hem_ratio = allSubs_info.variants_info.left_hem.group_avg(:,5)./allSubs_info.
variants_info.left_hem.group_avg(:,7);
right_hem_ratio = allSubs_info.variants_info.right_hem.group_avg(:,5)./allSubs_info.
variants_info.right_hem.group_avg(:,7);
[true_diff perm_diffs] = permute_values(left_hem_ratio, right_hem_ratio, flip_switch,
numperms, 0, 1, 0);
disp(['Border:Ectopic Ratio (permute hemispheres) - Below: ' num2str(length(find

```

```

(perm_diffs<true_diff))/1000)]
disp(['Border:Ectopic Ratio (permute hemispheres) - Above: ' num2str(length(find(
(perm_diffs>true_diff))/1000)]
output_mat = [true_diff; perm_diffs'];
save([outputDir str 'border_ectopic_ratio.mat'], 'output_mat'); clear output_mat
mean(left_hem_ratio)
mean(right_hem_ratio)
network_names = {'DMN' 'Vis' 'FP' ' ' 'DAN' ' ' 'VAN' 'Sal' 'CO' 'SMd'
'SMl' 'Aud' 'Tpole' 'MTL' 'PMN' 'PON'};
%network_names = {'DMN' 'Vis' 'FP' 'DAN' 'VAN' 'Sal' 'CO' 'SMd' 'SMl'
'Aud' 'Tpole' 'MTL' 'PMN' 'PON'};
results = [];
statistics = [];
for net = 1:numel(network_names)
left_hemi = networksxHem.clustersLH(:,net);
right_hemi = networksxHem.clustersRH(:,net);
results(net,1) = mean(left_hemi);
results(net,2) = mean(right_hemi);
[sig p ci stats] = ttest(left_hemi, right_hemi);
results(net,3) = sig;
results(net,4) = p;
statistics{net} = stats;
end
load([rootDir str '_networksxHem.mat']);
str = 'HCP384_new_split';
load([rootDir str '_networksxHem.mat']);
network_names = {'DMN' 'Vis' 'FP' ' ' 'DAN' ' ' 'VAN' 'Sal' 'CO' 'SMd'
'SMl' 'Aud' 'Tpole' 'MTL' 'PMN' 'PON'};
%network_names = {'DMN' 'Vis' 'FP' 'DAN' 'VAN' 'Sal' 'CO' 'SMd' 'SMl'
'Aud' 'Tpole' 'MTL' 'PMN' 'PON'};
results = [];
statistics = [];
for net = 1:numel(network_names)
left_hemi = networksxHem.clustersLH(:,net);
right_hemi = networksxHem.clustersRH(:,net);
results(net,1) = mean(left_hemi);
results(net,2) = mean(right_hemi);
[sig p ci stats] = ttest(left_hemi, right_hemi);
results(net,3) = sig;
results(net,4) = p;
statistics{net} = stats;
end
p_vals = ~isnan(results(:,4));
p_vals = results(~isnan(results(:,4)),4);
p_vals = [p_vals; 0.001; 0.002; 0.001; 0.14; 0.02; 0.05; 0.001; 0.001];
[p_fdr, p_masked] = FDR(p_vals, .05);
p_masked(1,2) = 'DMN';
%-- 10/4/21, 11:29 AM --%
check_MRI_data
%-- 10/6/21, 10:57 AM --%
load
('/Users/dianaperez/Desktop/Thesis/Variant_Properties/HCP384_new_split_variants_info_wSur
fArea.mat')
proportionBorders = variants_info.left_hem.group_avg(:,6);
proportionBorders1 = variants_info.left_hem.group_avg(:,6);
proportionEctopics = variants_info.left_hem.group_avg(:,8);
mean(proportionBorders)
mean(proportionEctopics)
proportionBorders = variants_info.right_hem.group_avg(:,6);
mean(proportionBorders)

```

```

permute_handedness
load
('/Users/dianaperez/Desktop/Thesis/Variant_Properties/HCP384_new_split_variants_info_wSurfArea.mat')
border_LHem = variants_info.left_hem.indiv_vars(variants_info.left_hem.indiv_vars{:,5}==1);
bord_ectop = variants_info.left_hem.indiv_vars{:,5};
border_LHem = variants_info.left_hem.indiv_vars(:,variants_info.left_hem.indiv_vars{:,5}==1);
border_LHem = variants_info.left_hem.indiv_vars(:,bord_ectop==1);
border_LHem = variants_info.left_hem.indiv_vars(variants_info.left_hem.indiv_vars{:,5}==1,:);
ectopic_LHem = variants_info.left_hem.indiv_vars(variants_info.left_hem.indiv_vars{:,5}==2,:);
ectopic_RHem = variants_info.right_hem.indiv_vars(variants_info.right_hem.indiv_vars{:,5}==2,:);
border_RHem = variants_info.right_hem.indiv_vars(variants_info.right_hem.indiv_vars{:,5}==1,:);
size_bord_LH = border_LHem(:,3);
size_bord_RH = border_RHem(:,3);
size_ectop_RH = ectopic_RHem(:,3);
size_ectop_LH = ectopic_LHem(:,3);
mean(size_bord_LH)
varSize_mm2 = border_LHem(:,3);
mean(border_LHem)
mean(varSize_mm2)
varSize_mm2 = border_RHem(:,3);
mean(varSize_mm2)
varSize_mm2 = ectopic_LHem(:,3);
mean(varSize_mm2)
varSize_mm2 = ectopic_RHem(:,3);
mean(varSize_mm2)
permute_variantproperties
load
('/Users/dianaperez/Desktop/Thesis/Spatial_Location/operc_nets_lefthemcluster_lefthem.mat')
clear all
load
('/Users/dianaperez/Desktop/Thesis/Spatial_Location/operc_nets_lefthemcluster_lefthem.mat')
load
('/Users/dianaperez/Desktop/Thesis/Spatial_Location/operc_nets_lefthemcluster_righthem.mat')
LH_cluster_RH = LH_cluster;
load
('/Users/dianaperez/Desktop/Thesis/Spatial_Location/operc_nets_lefthemcluster_lefthem.mat')
LH_cluster_LH = LH_cluster;
load
('/Users/dianaperez/Desktop/Thesis/Spatial_Location/operc_nets_righthemcluster_lefthem.mat')
RH_cluster_LH = RH_cluster;
load
('/Users/dianaperez/Desktop/Thesis/Spatial_Location/operc_nets_righthemcluster_righthem.mat')
RH_cluster_RH = RH_cluster;
clear LH_cluster RH_cluster
length(find(LH_cluster_LH.number==7))
length(find(LH_cluster_RH.number==7))
length(find(RH_cluster_RH.number==7))

```

```

length(find(RH_cluster_LH.number==7))
length(find(LH_cluster_LH.number==7))/length(LH_cluster_LH.number)
length(find(LH_cluster_RH.number==7))/length(LH_cluster_RH.number)
length(find(RH_cluster_RH.number==7))/length(RH_cluster_RH.number)
length(find(RH_cluster_LH.number==7))/length(RH_cluster_LH.number)
rgb_colors = [1 0 0;
1 1 0;
0 .8 0;
0 .6 .6;
0 0 0;
.3 0 .6;
1 .5 0;
.8 .8 .6];
output_dir = '/Users/dianaperez/Desktop/Thesis/Spatial_Location/';
load([output_dir 'operc_nets_righthemcluster_righthem.mat']);
rightclust_righthem = RH_cluster;
load([output_dir 'operc_nets_righthemcluster_lefthem.mat']);
rightclust_lefthem = RH_cluster;
clear RH_cluster
network_names = {'DMN' 'FP' 'DAN' 'Lang' 'Sal' 'CO' 'SML' 'PON'};
for net = 1:16
nets_left(net) = length(find(rightclust_lefthem.number(:,3)==net));
end
nets_left(6) = [];
nets_left(4) = [];
nets_left(10:13) = [];
nets_left(8) = [];
nets_left(2) = [];
sum(nets_left)
nets_left(4)/sum(nets_left)
for net = 1:16
nets_right(net) = length(find(rightclust_righthem.number(:,3)==net));
end
nets_right(6) = [];
nets_right(4) = [];
nets_right(10:13) = [];
nets_right(8) = [];
nets_right(2) = [];
nets_right(4)/sum(nets_right)
for net = 1:16
nets_left(net) = length(find(rightclust_lefthem.number(:,3)==net));
end
nets_left(7)/sum(nets_left)
length(find(RH_cluster_LH.number==7))
for net = 1:16
nets_right(net) = length(find(rightclust_righthem.number(:,3)==net));
end
nets_right(7)/sum(nets_right)
for net = 1:16
nets_left(net) = length(find(LH_cluster_LH.number(:,3)==net));
end
nets_left(7)/sum(nets_left)
for net = 1:16
nets_right(net) = length(find(LH_cluster_RH.number(:,3)==net));
end
nets_right(7)/sum(nets_right)
permute_variantproperties
[p_fdr, p_masked] = FDR([0.083, .906, .003, .996, .776, .224, .833,.167, .001, .999, .
161, .807,.769,.175,.675, .088,.051,.243,.702,.406,.518,.072,.899,.005,.995,.746,.2,.
148,.837,.667,.303,.224,.739,.315,.634,.106,.875],.05)

```

```

p_vals = [0.083, .906, .003, .996, .776, .224, .833,.167, .001, .999, .161, .807,.769,.
175,.675, .088,.051,.243,.702,.406,.518,.072,.899,.005,.995,.746,.2,.148,.837,.667,.303,.
224,.739,.315,.634,.106,.875];
permute_variantproperties
p_vals = [];
p_vals = [p_vals; 1 1];
clear p_vals
permute_variantproperties
for length(p_vals)
for y = 1:length(p_vals)
if p_vals(y,1)==0 && p_vals(y,2)==0
continue;
elseif p_vals(y,1) < p_vals(y,2) || p_vals(y,1) == p_vals(y,2)
p_vals(y,3) == p_vals(y,1);
elseif p_vals(y,1) > p_vals(y,2)
p_vals(y,3) == p_vals(y,2);
end
[p_fdr p_masked] = FDR(p_vals(:,3), .05);
end
y = 1;
p_vals(y,1)==0 && p_vals(y,2)==0
p_vals(y,1) < p_vals(y,2) || p_vals(y,1) == p_vals(y,2)
p_vals(y,1) > p_vals(y,2)
for y = 1:length(p_vals)
if p_vals(y,1)==0 && p_vals(y,2)==0
continue;
elseif p_vals(y,1) < p_vals(y,2) || p_vals(y,1) == p_vals(y,2)
p_vals(y,3) = p_vals(y,1);
elseif p_vals(y,1) > p_vals(y,2)
p_vals(y,3) = p_vals(y,2);
end
[p_fdr p_masked] = FDR(p_vals(:,3), .05);
end
final_pvals = p_vals(find(p_vals(:,3)>0),3);
[p_fdr p_masked] = FDR(final_pvals, .05);
permute_variantproperties
[p_fdr p_masked] = FDR(final_pvals, .05);
permute_variantproperties
p_vals = [p_vals(:,1); p_vals(:,2)];
p_fdr = FDR(p_vals,0.05)
permute_variantproperties
load
('/Users/dianaperez/Desktop/Thesis/Spatial_Location/HCP/Handedness/LeftvsRightHanders_tru
e_spCorr_1000_permutations.mat')
load
('/Users/dianaperez/Desktop/Thesis/Spatial_Location/HCP/Handedness/PseudoRightvsPseudoLef
tHanders_spCorr_1000_permutations.mat')
clear all
load
('/Users/dianaperez/Desktop/Thesis/Spatial_Location/HCP/Handedness/LeftvsRightHanders_tru
e_spCorr_1000_permutations.mat')
load
('/Users/dianaperez/Desktop/Thesis/Spatial_Location/HCP/Handedness/PseudoRightvsPseudoLef
tHanders_spCorr_1000_permutations.mat')
length(find(spCorrs<true_spCorr))/length(spCorrs)
length(find(spCorrs>true_spCorr))/length(spCorrs)
%-- 10/8/21, 1:00 PM --%
load('/Users/dianaperez/Desktop/Thesis/Variant_Properties/HCP384_new_split_networksxHem.
mat')
rgb_colors = [1 0 0; %DMN

```



```

0 0 .6; %Vis
1 1 0; %FP
0 .8 0; %DAN
0 .6 .6; %VAN
0 0 0; %Sal
.3 0 .6; %CON
.2 1 1; %SMd
1 .5 0; %SML
.6 .2 1; %Aud
.2 1 .2; %Tpole
0 .2 .4; %MTL
0 0 1; %PMN
.8 .8 .6]; %PON
leftHem = [networksxHem.clustersLH(:,1:3) networksxHem.clustersLH(:,5) networksxHem.
clustersLH(:,7:end)];
mean(leftHem)
rightHem = [networksxHem.clustersRH(:,1:3) networksxHem.clustersRH(:,5) networksxHem.
clustersRH(:,7:end)];
mean_RH = mean(rightHem);
data = [leftHem' rightHem'];
data = [mean_LH' mean_RH'];
mean_LH = mean(leftHem);
data = [mean_LH' mean_RH'];
data(11,:) = [];
bar(data)
close(gcf)
bar(data, .7)
bar(data, 1)
bar(data, 1.5)
bar(data, 1)
bar(data,1)
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.3, 0.9]);
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.9, 0.7]);
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.7, 0.7]);
xlabel('Assigned Network')
ylabel('Number of Variants')
ax = gca;
ax.FontSize = 24;
xticklabels({'DMN' 'Vis' 'FP' 'DAN' 'VAN' 'Sal' 'CO' 'SMd' 'SML' 'Aud'
'MTL' 'PMN' 'PON'});
findobj(gca)
m = findobj(gca, 'Type', 'Bar')
hleg1 = legend(m(1:2), 'Right Hemisphere', 'Left Hemisphere', 'Location',
'SouthOutside');
print(gcf, '/Users/dianaperez/Desktop/Thesis/Figures/Figure5_barGraph_networks_HCP.jpg',
'-dpng', '-r300')
load
('/Users/dianaperez/Desktop/Thesis/Variant_Properties/Network_Comparisons/network_perms_n
umVars.mat')
good_nets = [net_perms(:,1:3) net_perms(:,5) net_perms(:,7:end)];
network_names = {'DMN' 'Vis' 'FP' 'DAN' 'Lang' 'Sal' 'CO' 'SMd' 'SML' 'Aud'
'Tpole' 'MTL' 'PMN' 'PON'};
rgb_for_plot = {};
for n = 1:numel(network_names)
    rgb_for_plot{n} = rgb_colors(n,:);
end
net_inds = [ind ind ind ind ind ind ind ind ind ind ind ind ind];
handles = plotSpread(good_nets, 'categoryMarkers', {'x', '.'}, 'categoryLabels',
{'Permuted Differences', 'True Difference'}, 'distributionColors', rgb_for_plot,
'categoryIdx', net_inds, 'xNames', network_names)

```



```
% for n = 1:numel(network_names)
%     scatter(n,good_nets(1,n), 'MarkerEdgeColor', rgb_for_plot{1,n}, 'MarkerFaceColor',↵
rgb_for_plot{1,n}, 'SizeData', 50)
% end
scatter([1:14],good_nets(1,:), 'MarkerEdgeColor', 'k', 'MarkerFaceColor', 'k',↵
'SizeData', 50)
scatter(6,good_nets(1,6), 'MarkerEdgeColor', 'w', 'MarkerFaceColor', 'w', 'SizeData', 50)
ylabel('Difference in Number of Variants')
xlabel('Assigned Network')
axis([0, 15, -0.4, .7])
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.7, 0.7]);
print(gcf, '/Users/dianaperez/Desktop/Thesis/Figures/Figure5_spreadPlot_networks_HCP.↵
jpg', '-dpng', '-r300')
load↵
('/Users/dianaperez/Desktop/Thesis/Variant_Properties/Network_Comparisons/network_perms_n↵
umVars.mat')
good_nets = [net_perms(:,1:3) net_perms(:,5) net_perms(:,7:end)];
network_names = {'DMN' 'Vis' 'FP' 'DAN' 'Lang' 'Sal' 'CO' 'SMd' 'SML' 'Aud'↵
'Tpole' 'MTL' 'PMN' 'PON'};
rgb_for_plot = {};
for n = 1:numel(network_names)
rgb_for_plot{n} = rgb_colors(n,:);
end
ind = zeros(1001,1);
ind(1,1) = 1;
net_inds = [ind ind ind ind ind ind ind ind ind ind ind ind ind ind];
handles = plotSpread(good_nets, 'categoryMarkers', {'x', '.'}, 'categoryLabels',↵
{'Permuted Differences','True Difference'}, 'distributionColors', rgb_for_plot',↵
'categoryIdx', net_inds, 'xNames', network_names)
% for n = 1:numel(network_names)
%     scatter(n,good_nets(1,n), 'MarkerEdgeColor', rgb_for_plot{1,n}, 'MarkerFaceColor',↵
rgb_for_plot{1,n}, 'SizeData', 50)
% end
scatter([1:14],good_nets(1,:), 'MarkerEdgeColor', 'k', 'MarkerFaceColor', 'k',↵
'SizeData', 50)
scatter(6,good_nets(1,6), 'MarkerEdgeColor', 'w', 'MarkerFaceColor', 'w', 'SizeData', 50)
ylabel('Difference in Number of Variants')
xlabel('Assigned Network')
axis([0, 15, -0.4, .7])
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.7, 0.7]);
addpath '/Users/dianaperez/Documents/MATLAB/plotSpread/plotSpread'
load↵
('/Users/dianaperez/Desktop/Thesis/Variant_Properties/Network_Comparisons/network_perms_n↵
umVars.mat')
good_nets = [net_perms(:,1:3) net_perms(:,5) net_perms(:,7:end)];
network_names = {'DMN' 'Vis' 'FP' 'DAN' 'Lang' 'Sal' 'CO' 'SMd' 'SML' 'Aud'↵
'Tpole' 'MTL' 'PMN' 'PON'};
rgb_for_plot = {};
for n = 1:numel(network_names)
rgb_for_plot{n} = rgb_colors(n,:);
end
ind = zeros(1001,1);
ind(1,1) = 1;
net_inds = [ind ind ind ind ind ind ind ind ind ind ind ind ind ind];
handles = plotSpread(good_nets, 'categoryMarkers', {'x', '.'}, 'categoryLabels',↵
{'Permuted Differences','True Difference'}, 'distributionColors', rgb_for_plot',↵
'categoryIdx', net_inds, 'xNames', network_names)
% for n = 1:numel(network_names)
%     scatter(n,good_nets(1,n), 'MarkerEdgeColor', rgb_for_plot{1,n}, 'MarkerFaceColor',↵
rgb_for_plot{1,n}, 'SizeData', 50)
```

```
% end
scatter([1:14],good_nets(1,:), 'MarkerEdgeColor', 'k', 'MarkerFaceColor', 'k',
'SizeData', 50)
scatter(6,good_nets(1,6), 'MarkerEdgeColor', 'w', 'MarkerFaceColor', 'w', 'SizeData', 50)
ylabel('Difference in Number of Variants')
xlabel('Assigned Network')
axis([0, 15, -0.4, .7])
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.7, 0.7]);
clear gca
load
('/Users/dianaperez/Desktop/Thesis/Variant_Properties/Network_Comparisons/network_perms_n
umVars.mat')
good_nets = [net_perms(:,1:3) net_perms(:,5) net_perms(:,7:end)];
network_names = {'DMN' 'Vis' 'FP' 'DAN' 'Lang' 'Sal' 'CO' 'SMd' 'SML' 'Aud'
'Tpole' 'MTL' 'PMN' 'PON'};
rgb_for_plot = {};
for n = 1:numel(network_names)
    rgb_for_plot{n} = rgb_colors(n,:);
end
ind = zeros(1001,1);
ind(1,1) = 1;
net_inds = [ind ind ind ind ind ind ind ind ind ind ind ind ind ind];
handles = plotSpread(good_nets, 'categoryMarkers', {'x', '.'}, 'categoryLabels',
{'Permuted Differences', 'True Difference'}, 'distributionColors', rgb_for_plot,
'categoryIdx', net_inds, 'xNames', network_names)
% for n = 1:numel(network_names)
%     scatter(n,good_nets(1,n), 'MarkerEdgeColor', rgb_for_plot{1,n}, 'MarkerFaceColor',
rgb_for_plot{1,n}, 'SizeData', 50)
% end
scatter([1:14],good_nets(1,:), 'MarkerEdgeColor', 'k', 'MarkerFaceColor', 'k',
'SizeData', 50)
scatter(6,good_nets(1,6), 'MarkerEdgeColor', 'w', 'MarkerFaceColor', 'w', 'SizeData', 50)
ylabel('Difference in Number of Variants')
xlabel('Assigned Network')
axis([0, 15, -0.4, .7])
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.7, 0.7]);
ax = gca;
ax.FontSize = 24;
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.9, 0.7]);
print(gcf, '/Users/dianaperez/Desktop/Thesis/Figures/Figure3_spreadPlot_networks_HCP.
jpg', '-dpng', '-r300')
load('/Users/dianaperez/Desktop/Thesis/Variant_Properties/HCP384_new_split_networksxHem.
mat')
leftHem = [networksxHem.clustersLH(:,1:3) networksxHem.clustersLH(:,5) networksxHem.
clustersLH(:,7:end)];
mean_LH = mean(leftHem);
rightHem = [networksxHem.clustersRH(:,1:3) networksxHem.clustersRH(:,5) networksxHem.
clustersRH(:,7:end)];
mean_RH = mean(rightHem);
data = [mean_LH' mean_RH'];
data(11,:) = []; %delete Tpole data bc zero
%[h,mu,sigma,q,notch] = al_goodplot(leftHem,1:14,.2,[ 0 0.4470 0.7410],'left');
%[h,mu,sigma,q,notch] = al_goodplot(rightHem,1:14,.2,[0.4660 0.6740
0.1880],'right');
bar(data,1)
axis([0, 15, -0.1, 10])
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.9, 0.7]);
xlabel('Assigned Network')
ylabel('Number of Variants')
ax = gca;
```

```

ax.FontSize = 24;
xticklabels({'DMN' 'Vis' 'FP' 'DAN' 'Lang' 'Sal' 'CO' 'SMd' 'SML' 'Aud'
'MTL' 'PMN' 'PON'});
m = findobj(gca, 'Type', 'Bar')
hleg1 = legend(m(1:2), 'Right Hemisphere', 'Left Hemisphere', 'Location',
'SouthOutside');
print(gcf, '/Users/dianaperez/Desktop/Thesis/Figures/Figure5_barGraph_networks_HCP.jpg',
'-dpng', '-r300')
load('/Users/dianaperez/Desktop/Thesis/Variant_Properties/HCP384_new_split_networksxHem.
mat')
leftHem = [networksxHem.clustersLH(:,1:3) networksxHem.clustersLH(:,5) networksxHem.
clustersLH(:,7:end)];
mean_LH = mean(leftHem);
rightHem = [networksxHem.clustersRH(:,1:3) networksxHem.clustersRH(:,5) networksxHem.
clustersRH(:,7:end)];
mean_RH = mean(rightHem);
data = [mean_LH' mean_RH'];
data(11,:) = []; %delete Tpole data bc zero
%[h,mu,sigma,q,notch] = al_goodplot(leftHem,1:14,.2,[ 0 0.4470 0.7410],'left');
%[h,mu,sigma,q,notch] = al_goodplot(rightHem,1:14,.2,[0.4660 0.6740
0.1880],'right');
bar(data,1)
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.9, 0.7]);
xlabel('Assigned Network')
ylabel('Number of Variants')
ax = gca;
ax.FontSize = 24;
xticklabels({'DMN' 'Vis' 'FP' 'DAN' 'Lang' 'Sal' 'CO' 'SMd' 'SML' 'Aud'
'MTL' 'PMN' 'PON'});
m = findobj(gca, 'Type', 'Bar')
hleg1 = legend(m(1:2), 'Right Hemisphere', 'Left Hemisphere', 'Location',
'SouthOutside');
print(gcf, '/Users/dianaperez/Desktop/Thesis/Figures/Figure3_barGraph_networks_HCP.jpg',
'-dpng', '-r300')
clear all
RH_clust = ft_read_cifti_mod('/Users/dianaperez/Desktop/Thesis/operc_mask_moreRHvars.
dtseries.nii');
LH_clust = ft_read_cifti_mod('/Users/dianaperez/Desktop/Thesis/operc_mask_moreLHvars.
dtseries.nii');
LH_clust_gifti = insert_nonbrain(LH_clust.data, 'both', LH_clust);
left_hem = LH_clust_gifti(1:(64984/2));
right_hem = LH_clust_gifti(32493:end);
new_cifti = [right_hem;left_hem];
new_cifti = new_cifti(LH_clust.brainstructure>0);
LH_clust.data = new_cifti;
ft_write_cifti_mod('/Users/dianaperez/Desktop/Thesis/operc_mask_moreLHvars_onRH.dtseries.
nii', LH_clust)
RH_clust_gifti = insert_nonbrain(RH_clust.data, 'both', RH_clust);
left_hem = RH_clust_gifti(1:(64984/2));
right_hem = RH_clust_gifti(32493:end);
new_cifti = [right_hem;left_hem];
new_cifti = new_cifti(RH_clust.brainstructure>0);
RH_clust.data = new_cifti;
ft_write_cifti_mod('/Users/dianaperez/Desktop/Thesis/operc_mask_moreRHvars_onLH.dtseries.
nii', RH_clust)
clear all
load('/Users/dianaperez/Desktop/Thesis/Variant_Properties/HCP384_new_split_networksxHem.
mat')
output_dir = '/Users/dianaperez/Desktop/Thesis/Spatial_Location/';
load([output_dir 'operc_nets_lefthemcluster_righthem.mat']);

```

```
leftclust_righthem = LH_cluster;
load([output_dir 'operc_nets_lefthemcluster_lefthem.mat']);
leftclust_lefthem = LH_cluster;
clear LH_cluster
network_names = {'DMN' 'FP' 'DAN' 'Lang' 'Sal' 'CO' 'SMI'};
for net = 1:16
nets_left(net) = length(find(leftclust_lefthem.number(:,3)==net));
end
nets_left(6) = [];
nets_left(4) = [];
nets_left(10:end) = [];
nets_left(8) = [];
nets_left(2) = [];
operc_cluster
addpath '/Users/dianaperez/Documents/GitHub/Lateralization_Variants/thesis_scripts'
RH_mask_gifti = insert_nonbrain(operc_mask_RH, 'both', diffmap);
LH_mask_gifti = insert_nonbrain(operc_mask_LH, 'both', diffmap);
RH_mask_flipped = [RH_mask_gifti(32493:end); RH_mask_gifti(1:32492)];
LH_mask_flipped = [LH_mask_gifti(32493:end); LH_mask_gifti(1:32492)];
RH_mask_flipped = RH_mask_flipped(diffmap.brainstructure>0);
LH_mask_flipped = LH_mask_flipped(diffmap.brainstructure>0);
RH_vertices = find(RH_mask_flipped == 1);
RH_cluster_vars = [];
for sub = 1:length(subs)
sub_data = [];
var_map = ft_read_cifti_mod([data_dir num2str(subs(sub)) '_uniqueIDs_afterReassign.↵
dtseries.nii']);
net_assign = ft_read_cifti_mod([data_dir num2str(subs(sub)) '_reassigned.dtseries.nii']);
unique_IDs = unique(var_map.data(RH_vertices));
if unique_IDs(1) == 0
unique_IDs(1) = [];
end
sub_data(:,1) = unique_IDs;
for var = 1:length(unique_IDs)
verts = find(var_map.data==sub_data(var,1));
sub_data(var,2) = verts(1);
sub_data(var,3) = net_assign.data(verts(1));
end
RH_cluster_vars = [RH_cluster_vars; sub_data];
for net = 1:16
if isempty(sub_data)
RH_cluster_proportion(sub,net) = 0;
RH_cluster_number(sub,net) = 0;
else
RH_cluster_proportion(sub,net) = length(find(sub_data(:,3)==net))/size(sub_data,1);
RH_cluster_number(sub,net) = length(find(sub_data(:,3)==net));
end
end
end
RH_cluster = [];
RH_cluster.vars = RH_cluster_vars;
RH_cluster.number = RH_cluster_number;
RH_cluster.proportion = RH_cluster_proportion;
save([output_dir 'operc_nets_righthemcluster_righthem.mat'], 'RH_cluster')
LH_vertices = find(LH_mask_flipped == 1);
LH_cluster_vars = [];
for sub = 1:length(subs)
sub_data = [];
var_map = ft_read_cifti_mod([data_dir num2str(subs(sub)) '_uniqueIDs_afterReassign.↵
dtseries.nii']);
```

```

net_assign = ft_read_cifti_mod([data_dir num2str(subs(sub)) '_reassigned.dtseries.nii']);
unique_IDs = unique(var_map.data(LH_vertices));
if unique_IDs(1) == 0
unique_IDs(1) = [];
end
sub_data(:,1) = unique_IDs;
for var = 1:length(unique_IDs)
verts = find(var_map.data==sub_data(var,1));
sub_data(var,2) = verts(1);
sub_data(var,3) = net_assign.data(verts(1));
end
LH_cluster_vars = [LH_cluster_vars; sub_data];
for net = 1:16
if isempty(sub_data)
LH_cluster_proportion(sub,net) = 0;
LH_cluster_number(sub,net) = 0;
else
LH_cluster_proportion(sub,net) = length(find(sub_data(:,3)==net))/size(sub_data,1);
LH_cluster_number(sub,net) = length(find(sub_data(:,3)==net));
end
end
end
LH_cluster = [];
LH_cluster.vars = LH_cluster_vars;
LH_cluster.number = LH_cluster_number;
LH_cluster.proportion = LH_cluster_proportion;
save([output_dir 'operc_nets_lefthemcluster_righthem.mat'], 'LH_cluster')
clear all
load↵
('/Users/dianaperez/Desktop/Thesis/Spatial_Location/operc_nets_lefthemcluster_lefthem.↵
mat')
load↵
('/Users/dianaperez/Desktop/Thesis/Spatial_Location/operc_nets_righthemcluster_lefthem.↵
mat')
LH_cluster_LHem
LH_cluster_LHem = LH_cluster;
RH_cluster_LHem = RH_cluster;
load↵
('/Users/dianaperez/Desktop/Thesis/Spatial_Location/operc_nets_lefthemcluster_righthem.↵
mat')
load↵
('/Users/dianaperez/Desktop/Thesis/Spatial_Location/operc_nets_righthemcluster_righthem.↵
mat')
LH_cluster_RHem = LH_cluster;
RH_cluster_RHem = RH_cluster;
clear LH_cluster RH_cluster
LH_cluster = LH_cluster_LHem.number;
input_for_anova = [];
for net = 1:16
numvars = LH_cluster(:,net);
numvars = numvars(numvars>0);
hemi = ones(length(numvars),1);
network = zeros(length(numvars),1);
network(:, :) = net;
input_for_anova = [hemi network numvars];
end
net = 1
numvars = LH_cluster(:,net);
numvars = numvars(numvars>0);
hemi = ones(length(numvars),1);

```

```
network = zeros(length(numvars),1);
unique(network)
network(:, :) = net;
unique(network)
input_for_anova = [input_for_anova; hemi network numvars];
input_for_anova = [];
for net = 1:16
    numvars = LH_cluster(:,net);
    numvars = numvars(numvars>0);
    hemi = ones(length(numvars),1);
    network = zeros(length(numvars),1);
    network(:, :) = net;
    input_for_anova = [input_for_anova; hemi network numvars];
end
RH_cluster = LH_cluster_RHem.number;
for net = 1:16
    numvars = RH_cluster(:,net);
    numvars = numvars(numvars>0);
    hemi = ones(length(numvars),1);
    hemi(:, :) = 2;
    network = zeros(length(numvars),1);
    network(:, :) = net;
    input_for_anova = [input_for_anova; hemi network numvars];
end
input_for_anova = [];
LH_cluster = RH_cluster_LHem.number;
for net = 1:16
    numvars = LH_cluster(:,net);
    numvars = numvars(numvars>0);
    hemi = ones(length(numvars),1);
    network = zeros(length(numvars),1);
    network(:, :) = net;
    input_for_anova = [input_for_anova; hemi network numvars];
end
RH_cluster = RH_cluster_RHem.number;
for net = 1:16
    numvars = RH_cluster(:,net);
    numvars = numvars(numvars>0);
    hemi = ones(length(numvars),1);
    hemi(:, :) = 2;
    network = zeros(length(numvars),1);
    network(:, :) = net;
    input_for_anova = [input_for_anova; hemi network numvars];
end
input_for_anova = [];
LH_cluster = RH_cluster_LHem.number;
for net = 1:16
    numvars = LH_cluster(:,net);
    %numvars = numvars(numvars>0);
    hemi = ones(length(numvars),1);
    network = zeros(length(numvars),1);
    network(:, :) = net;
    input_for_anova = [input_for_anova; hemi network numvars];
end
RH_cluster = RH_cluster_RHem.number;
for net = 1:16
    numvars = RH_cluster(:,net);
    %numvars = numvars(numvars>0);
    hemi = ones(length(numvars),1);
    hemi(:, :) = 2;
```

```
network = zeros(length(numvars),1);
network(:, :) = net;
input_for_anova = [input_for_anova; hemi network numvars];
end
input_for_anova = [];
LH_cluster = RH_cluster_LHem.number;
for net = 1:16
    numvars = LH_cluster(:,net);
    %numvars = numvars(numvars>0);
    if unique(numvars)==0
        continue
    end
    hemi = ones(length(numvars),1);
    network = zeros(length(numvars),1);
    network(:, :) = net;
    input_for_anova = [input_for_anova; hemi network numvars];
end
RH_cluster = RH_cluster_RHem.number;
for net = 1:16
    numvars = RH_cluster(:,net);
    %numvars = numvars(numvars>0);
    if unique(numvars)==0
        continue
    end
    hemi = ones(length(numvars),1);
    hemi(:, :) = 2;
    network = zeros(length(numvars),1);
    network(:, :) = net;
    input_for_anova = [input_for_anova; hemi network numvars];
end
input_for_anova = [];
LH_cluster = RH_cluster_LHem.number;
for net = 1:16
    numvars = LH_cluster(:,net);
    %numvars = numvars(numvars>0);
    %     if unique(numvars)==0
    %         continue
    %     end
    %     hemi = ones(length(numvars),1);
    hemi = 1;
    %     network = zeros(length(numvars),1);
    %     network(:, :) = net;
    input_for_anova = [input_for_anova; hemi net length(numvars)];
end
input_for_anova = [];
LH_cluster = RH_cluster_LHem.number;
for net = 1:16
    numvars = LH_cluster(:,net);
    %numvars = numvars(numvars>0);
    %     if unique(numvars)==0
    %         continue
    %     end
    %     hemi = ones(length(numvars),1);
    hemi = 1;
    %     network = zeros(length(numvars),1);
    %     network(:, :) = net;
    input_for_anova = [input_for_anova; hemi net sum(numvars)];
end
H_cluster = RH_cluster_RHem.number;
for net = 1:16
```

```
numvars = RH_cluster(:,net);
%numvars = numvars(numvars>0);
if unique(numvars)==0
continue
end
%hemi = ones(length(numvars),1);
hemi = 2;
%network = zeros(length(numvars),1);
%network(:, :) = net;
input_for_anova = [input_for_anova; hemi net sum(numvars)];
end
input_for_anova = [];
LH_cluster = RH_cluster_LHem.number;
for net = 1:16
numvars = LH_cluster(:,net);
%numvars = numvars(numvars>0);
%    if unique(numvars)==0
%        continue
%    end
%    hemi = ones(length(numvars),1);
hemi = 1;
%    network = zeros(length(numvars),1);
%    network(:, :) = net;
input_for_anova = [input_for_anova; hemi net sum(numvars)];
end
RH_cluster = RH_cluster_RHem.number;
for net = 1:16
numvars = RH_cluster(:,net);
%numvars = numvars(numvars>0);
%    if unique(numvars)==0
%        continue
%    end
%hemi = ones(length(numvars),1);
hemi = 2;
%network = zeros(length(numvars),1);
%network(:, :) = net;
input_for_anova = [input_for_anova; hemi net sum(numvars)];
end
input_for_anova = [];
LH_cluster = LH_cluster_LHem.number;
for net = 1:16
numvars = LH_cluster(:,net);
%numvars = numvars(numvars>0);
%    if unique(numvars)==0
%        continue
%    end
%    hemi = ones(length(numvars),1);
hemi = 1;
%    network = zeros(length(numvars),1);
%    network(:, :) = net;
input_for_anova = [input_for_anova; hemi net sum(numvars)];
end
RH_cluster = LH_cluster_RHem.number;
for net = 1:16
numvars = RH_cluster(:,net);
%numvars = numvars(numvars>0);
%    if unique(numvars)==0
%        continue
%    end
%hemi = ones(length(numvars),1);
```



```
hemi = 2;
%network = zeros(length(numvars),1);
%network(:, :) = net;
input_for_anova = [input_for_anova; hemi net sum(numvars)];
end
sum(LH_cluster_LHem.number, 1)
sum(LH_cluster_LHem.number, 2);
mean(ans)
sum(LH_cluster_LHem.number, 2);
sum(LH_cluster_RHem.number, 2);
mean(ans)
sum(RH_cluster_LHem.number, 2);
mean(ans)
sum(RH_cluster_RHem.number, 2);
mean(ans)
input_for_anova = [];
LH_cluster = LH_cluster_LHem.number;
count = 1;
for net = 1:16
    numvars = LH_cluster(:,net);
    %numvars = numvars(numvars>0);
    if unique(numvars)==0
        continue
    end
    hemi = ones(length(numvars),1);
    %hemi = 1;
    network = zeros(length(numvars),1);
    network(:, :) = net;
    input_for_anova = [input_for_anova; count hemi network sum(numvars)];
    count = count + 1;
end
input_for_anova = [];
LH_cluster = LH_cluster_LHem.number;
count = 1;
for net = 1:16
    numvars = LH_cluster(:,net);
    %numvars = numvars(numvars>0);
    if unique(numvars)==0
        continue
    end
    hemi = ones(length(numvars),1);
    %hemi = 1;
    network = zeros(length(numvars),1);
    network(:, :) = net;
    sub = zeros(length(numvars),1);
    sub(:, :) = count;
    input_for_anova = [input_for_anova; sub hemi network sum(numvars)];
    count = count + 1;
end
LH_cluster = LH_cluster_LHem.number;
count = 1;
net = 1;
numvars = LH_cluster(:,net);
if unique(numvars)==0
    continue
end
hemi = ones(length(numvars),1);
hemi(:, :) = 1;
sub = [1:384];
network = zeros(length(numvars),1);
```

```
network(:, :) = net;
sub = [1:384]';
input_for_anova = [input_for_anova; sub hemi network numvars];
input_for_anova = [];
LH_cluster = LH_cluster_LHem.number;
for net = 1:16
    numvars = LH_cluster(:, net);
    %numvars = numvars(numvars>0);
    if unique(numvars)==0
        continue
    end
    hemi = ones(length(numvars), 1);
    hemi(:, :) = 1;
    network = zeros(length(numvars), 1);
    network(:, :) = net;
    sub = [1:384]';
    input_for_anova = [input_for_anova; sub hemi network numvars];
end
RH_cluster = LH_cluster_RHem.number;
for net = 1:16
    numvars = RH_cluster(:, net);
    %numvars = numvars(numvars>0);
    if unique(numvars)==0
        continue
    end
    sub = [1:384]';
    hemi = ones(length(numvars), 1);
    hemi = 2;
    network = zeros(length(numvars), 1);
    network(:, :) = net;
    input_for_anova = [input_for_anova; sub hemi net numvars];
end
RH_cluster = LH_cluster_RHem.number;
for net = 1:16
    numvars = RH_cluster(:, net);
    %numvars = numvars(numvars>0);
    if unique(numvars)==0
        continue
    end
    sub = [1:384]';
    hemi = ones(length(numvars), 1);
    hemi(:, :) = 2;
    network = zeros(length(numvars), 1);
    network(:, :) = net;
    input_for_anova = [input_for_anova; sub hemi net numvars];
end
RH_cluster = LH_cluster_RHem.number;
for net = 1:16
    numvars = RH_cluster(:, net);
    %numvars = numvars(numvars>0);
    if unique(numvars)==0
        continue
    end
    sub = [1:384]';
    hemi = ones(length(numvars), 1);
    hemi(:, :) = 2;
    network = zeros(length(numvars), 1);
    network(:, :) = net;
    input_for_anova = [input_for_anova; sub hemi network numvars];
end
```

```

input_for_anova = [];
LH_cluster = RH_cluster_LHem.number;
for net = 1:16
numvars = LH_cluster(:,net);
%numvars = numvars(numvars>0);
if unique(numvars)==0
continue
end
hemi = ones(length(numvars),1);
hemi(:, :) = 1;
network = zeros(length(numvars),1);
network(:, :) = net;
sub = [1:384]';
input_for_anova = [input_for_anova; sub hemi network numvars];
end
RH_cluster = RH_cluster_RHem.number;
for net = 1:16
numvars = RH_cluster(:,net);
%numvars = numvars(numvars>0);
if unique(numvars)==0
continue
end
sub = [1:384]';
hemi = ones(length(numvars),1);
hemi(:, :) = 2;
network = zeros(length(numvars),1);
network(:, :) = net;
input_for_anova = [input_for_anova; sub hemi network numvars];
end
help anova1
load
('/Users/dianaperez/Desktop/Thesis/Variant_Properties/Permutation_Tests/HCP384_LhemvsRhem
_avgSurfArea.mat')
clear all
load
('/Users/dianaperez/Desktop/Thesis/Variant_Properties/Permutation_Tests/HCP384_LhemvsRhem
_avgSurfArea.mat')
length(find(output_mat(2:end,:)<output_mat(1,1)))
length(find(output_mat(2:end,:)>output_mat(1,1)))
load
('/Users/dianaperez/Desktop/Thesis/Variant_Properties/HCP384_new_split_variants_info_wSur
fArea.mat')
leftHem = variants_info.left_hem.group_avg(:,4);
rightHem = variants_info.right_hem.group_avg(:,4);
[h,mu,sigma,q,notch] = al_goodplot(leftHem,0.1,.2,[ 0      0.4470      0.7410],'left');
[h,mu,sigma,q,notch] = al_goodplot(rightHem,0.2,.2,[0.8500      0.3250      0.0980],'right');
xticks([0])
xticklabels('Average Variant Size')
axis([-0.2, 0.5, 50, 400])
ax = gca;
ax.FontSize = 24;
addpath '/Users/dianaperez/Documents/MATLAB/upload'
leftHem = variants_info.left_hem.group_avg(:,4);
rightHem = variants_info.right_hem.group_avg(:,4);
[h,mu,sigma,q,notch] = al_goodplot(leftHem,0.1,.2,[ 0      0.4470      0.7410],'left');
[h,mu,sigma,q,notch] = al_goodplot(rightHem,0.2,.2,[0.8500      0.3250      0.0980],'right');
xticks([0])
xticklabels('Average Variant Size')
axis([-0.2, 0.5, 50, 400])
ax = gca;

```

```

ax.FontSize = 24;
close(gcf)
load('/Users/dianaperez/Desktop/Thesis/Variant_Properties/HCP384_new_split_variants_info.mat')
leftHem = variants_info.left_hem.group_avg{:,4};
rightHem = variants_info.right_hem.group_avg{:,4};
[h,mu,sigma,q,notch] = al_goodplot(leftHem,0.1,.2,[ 0    0.4470    0.7410],'left');
[h,mu,sigma,q,notch] = al_goodplot(rightHem,0.15,.2,[0.8500    0.3250
0.0980],'right');
xticks([0])
xlabel('Average Variant Size')
axis([-0.2, 0.5, 50, 400])
ax = gca;
ax.FontSize = 24;
close(gcf)
load(
('/Users/dianaperez/Desktop/Thesis/Variant_Properties/HCP384_new_split_variants_info_wSurfArea.mat')
leftHem = variants_info.left_hem.group_avg{:,4};
rightHem = variants_info.right_hem.group_avg{:,4};
[h,mu,sigma,q,notch] = al_goodplot(leftHem,0.1,.2,[ 0    0.4470    0.7410],'left');
[h,mu,sigma,q,notch] = al_goodplot(rightHem,0.15,.2,[0.8500    0.3250
0.0980],'right');
xticks([0])
xlabel('Average Variant Size')
axis([-0.2, 0.5, 50, 400])
ax = gca;
ax.FontSize = 24;
close(gcf)
leftHem = variants_info.left_hem.group_avg{:,4};
rightHem = variants_info.right_hem.group_avg{:,4};
[h,mu,sigma,q,notch] = al_goodplot(leftHem,0.1,.2,[ 0    0.4470    0.7410],'left');
[h,mu,sigma,q,notch] = al_goodplot(rightHem,0.11,.2,[0.8500    0.3250
0.0980],'right');
xticks([0])
xlabel('Average Variant Size')
axis([-0.2, 0.5, 50, 400])
ax = gca;
ax.FontSize = 24;
close(gcf)
leftHem = variants_info.left_hem.group_avg{:,4};
rightHem = variants_info.right_hem.group_avg{:,4};
[h,mu,sigma,q,notch] = al_goodplot(leftHem,0.1,.2,[ 0    0.4470    0.7410],'left');
[h,mu,sigma,q,notch] = al_goodplot(rightHem,0.11,.2,[0.8500    0.3250
0.0980],'right');
xticks([0])
xlabel('Average Variant Size')
axis([-0.2, 0.5, 100, 700])
ax = gca;
ax.FontSize = 24;
max(variants_info.left_hem.group_avg{:,4});
max(variants_info.left_hem.group_avg{:,4})
max(variants_info.right_hem.group_avg{:,4})
close(gcf)
leftHem = variants_info.left_hem.group_avg{:,4};
rightHem = variants_info.right_hem.group_avg{:,4};
[h,mu,sigma,q,notch] = al_goodplot(leftHem,0.1,.2,[ 0    0.4470    0.7410],'left');
[h,mu,sigma,q,notch] = al_goodplot(rightHem,0.11,.2,[0.8500    0.3250
0.0980],'right');
xticks([0])

```

```
xticklabels('Average Variant Size')
axis([-0.2, 0.5, 100, 800])
ax = gca;
ax.FontSize = 24;
ylabel('Surface Area (mm^2)')
xticks([1.105])
xticklabels('Average Variant Size')
close gcf
leftHem = variants_info.left_hem.group_avg{:,4};
rightHem = variants_info.right_hem.group_avg{:,4};
[h,mu,sigma,q,notch] = al_goodplot(leftHem,0.1,.2,[ 0    0.4470    0.7410],'left');
[h,mu,sigma,q,notch] = al_goodplot(rightHem,0.11,.2,[0.8500    0.3250
0.0980],'right');
xticks([1.105])
xticklabels('Average Variant Size')
axis([-0.2, 0.5, 100, 800])
ax = gca;
ax.FontSize = 24;
ylabel('Surface Area (mm^2)')
xticks([0.105])
ax = gca;
ax.FontSize = 24;
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.3, 0.7]); %first and second
control position on screen, third controls width, and fourth controls height
print(gcf, '/Users/dianaperez/Desktop/Thesis/Figures/Figure2_violin_avgVarSize_HCP.jpg',
'-dpng', '-r300')
perms_varSize = output_mat;
ind = zeros(1001,1);
ind(1,1) = 1;
handles = plotSpread(all_perms, 'categoryMarkers', {'o', '.'}, 'categoryLabels',
{'Permuted Correlations','True Correlation'}, 'categoryColors', {'k','r'}, 'categoryIdx',
[ind], 'xNames', {'Average Variant Size'}, 'xValues', [1])
scatter(1,perms_varSize(1,:), 'MarkerEdgeColor', 'red', 'MarkerFaceColor', 'red',
'SizeData', 50)
ax = gca;
ax.FontSize = 24;
axis([0.5, 1.5, -25, 10])
handles = plotSpread(perms_varSize, 'categoryMarkers', {'o', '.'}, 'categoryLabels',
{'Permuted Correlations','True Correlation'}, 'categoryColors', {'k','r'}, 'categoryIdx',
[ind], 'xNames', {'Average Variant Size'}, 'xValues', [1])
scatter(1,perms_varSize(1,:), 'MarkerEdgeColor', 'red', 'MarkerFaceColor', 'red',
'SizeData', 50)
ax = gca;
ax.FontSize = 24;
axis([0.5, 1.5, -25, 10])
close gcf
handles = plotSpread(perms_varSize, 'categoryMarkers', {'o', '.'}, 'categoryLabels',
{'Permuted Correlations','True Correlation'}, 'categoryColors', {'k','r'}, 'categoryIdx',
[ind], 'xNames', {'Average Variant Size'}, 'xValues', [1])
scatter(1,perms_varSize(1,:), 'MarkerEdgeColor', 'red', 'MarkerFaceColor', 'red',
'SizeData', 50)
ax = gca;
ax.FontSize = 24;
ylabel('Difference in Surface Area (mm^2)')
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.3, 0.7]); %first and second
control position on screen, third controls width, and fourth controls height
print(gcf, '/Users/dianaperez/Desktop/Thesis/Figures/Figure2_spreadPlot_varSize_HCP.jpg',
'-dpng', '-r300')
axis([0.5, 1.5, -60, 20])
print(gcf, '/Users/dianaperez/Desktop/Thesis/Figures/Figure2_spreadPlot_varSize_HCP.jpg',
```

```
'-dpng', '-r300')
load
('/Users/dianaperez/Desktop/Thesis/Spatial_Location/operc_nets_lefthemcluster_lefthem.
mat')
clear all
load
('/Users/dianaperez/Desktop/Thesis/Spatial_Location/operc_nets_lefthemcluster_lefthem.
mat')
LH_cluster_onLH = LH_cluster;
load
('/Users/dianaperez/Desktop/Thesis/Spatial_Location/operc_nets_lefthemcluster_righthem.
mat')
LH_cluster_onRH = LH_cluster;
clear LH_cluster
load
('/Users/dianaperez/Desktop/Thesis/Spatial_Location/operc_nets_righthemcluster_lefthem.
mat')
RH_cluster_onLH = RH_cluster;
load
('/Users/dianaperez/Desktop/Thesis/Spatial_Location/operc_nets_righthemcluster_righthem.
mat')
RH_cluster_onRH = RH_cluster;
clear RH_cluster
sum(LH_cluster_onLH(:,1))
sum(LH_cluster_onLH.number(:,1))
sum(LH_cluster_onLH.number(:,3))
sum(LH_cluster_onLH.number(:,5))
sum(LH_cluster_onLH.number(:,7\))
sum(LH_cluster_onLH.number(:,7))
sum(LH_cluster_onLH.number(:,8))
sum(LH_cluster_onLH.number(:,9))
sum(LH_cluster_onLH.number(:,11))
sum(LH_cluster_onRH.number(:,1))
sum(LH_cluster_onRH.number(:,3))
sum(LH_cluster_onRH.number(:,5))
sum(LH_cluster_onRH.number(:,7))
sum(LH_cluster_onRH.number(:,8))
sum(LH_cluster_onRH.number(:,9))
sum(LH_cluster_onRH.number(:,11))
for net = 1:16
nets_left(net) = length(find(LH_cluster_onLH.number(:,3)==net));
end
nets_left(6) = [];
nets_left(4) = [];
nets_left(10:end) = [];
nets_left(8) = [];
nets_left(2) = [];
for net = 1:16
nets_left(net) = length(find(LH_cluster_onLH.vars(:,3)==net));
end
nets_left(6) = [];
nets_left(4) = [];
nets_left(10:end) = [];
nets_left(8) = [];
nets_left(2) = [];
sum(nets_left
sum(nets_left)
total_vars = 306;
for net = 1:length(nets_left)
perc = nets_left(net)/total_vars
```

```
end
for net = 1:16
nets_right(net) = length(find(LH_cluster_onRH.vars(:,3)==net));
end
nets_right(6) = [];
nets_right(4) = [];
nets_right(10:end) = [];
nets_right(8) = [];
nets_right(2) = [];
sum(nets_right)
total_vars = 254;
for net = 1:length(nets_left)
perc = nets_right(net)/total_vars
end
for net = 1:16
nets_right(net) = length(find(RH_cluster_onRH.vars(:,3)==net));
end
nets_right(6) = [];
nets_right(4) = [];
nets_right(10:end) = [];
nets_right(8) = [];
nets_right(2) = [];
sum(nets_right)
total_vars = 541;
for net = 1:length(nets_left)
perc = nets_right(net)/total_vars
end
for net = 1:16
nets_right(net) = length(find(RH_cluster_onRH.vars(:,3)==net));
end
nets_right(6) = [];
nets_right(4) = [];
nets_right(10:15) = [];
nets_right(8) = [];
nets_right(2) = [];
for net = 1:16
nets_right(net) = length(find(RH_cluster_onRH.vars(:,3)==net));
end
nets_right(6) = [];
nets_right(4) = [];
nets_right(8) = [];
nets_right(2) = [];
nets_right(9:11) = [];
for net = 1:length(nets_left)
perc = nets_right(net)/total_vars
end
total_vars = sum(nets_right);
for net = 1:length(nets_right)
perc = nets_right(net)/total_vars
end
for net = 1:16
nets_right(net) = length(find(RH_cluster_onLH.vars(:,3)==net));
end
nets_right(6) = [];
nets_right(4) = [];
nets_right(8) = [];
nets_right(2) = [];
total_vars = sum(nets_right);
for net = 1:length(nets_right)
perc = nets_right(net)/total_vars
```

```
end
clear all
load
(' /Users/dianaperez/Desktop/Thesis/Variant_Properties/HCP384_new_split_variants_info_wSurfArea.mat')
bordervars_LH = variants_info.left_hem.indiv_vars(variants_info.left_hem.indiv_vars==1,:);
bordervars_LH = variants_info.left_hem.indiv_vars(variants_info.left_hem.indiv_vars==1,:);
bordervars_LH = variants_info.left_hem.indiv_vars(variants_info.left_hem.indiv_vars(:,5)==1,:);
bord_ectop = variants_info.left_hem.indiv_vars(:,5);
bordervars_LH = variants_info.left_hem.indiv_vars(bord_ectop==1,:);
ectopicvars_LH = variants_info.left_hem.indiv_vars(bord_ectop==2,:);
bord_ectop1 = variants_info.right_hem.indiv_vars(:,5);
bordervars_RH = variants_info.right_hem.indiv_vars(bord_ectop==1,:);
bordervars_RH = variants_info.right_hem.indiv_vars(bord_ectop1==1,:);
ectopicvars_RH = variants_info.right_hem.indiv_vars(bord_ectop1==2,:);
clear bord_ectop
clear bord_ectop1
numperms = 1000;
left = bordervars_LH;
right = bordervars_RH;
flip_switch = zeros(length(left)+length(right),1);
flip_switch = zeros(height(left)+height(right),1);
flip_switch(1:length(flip_switch)/2) = 1;
true_mean_LH = mean(left);
true_mean_LH = mean(left(:,5));
true_mean_RH = mean(right(:,5));
true_diff = true_mean_LH - true_mean_RH;
surfArea_LH = left(:,5);
surfArea_RH = right(:,5);
for p = 1:numperms
    rng('shuffle');
    ind = randperm(length(flip_switch));
    rand_flip_switch = flip_switch(ind);
    for s = 1:length(flip_switch)
    end
end
variable = [surfArea_LH; surfArea_RH];
flip_switch = zeros(height(left)+height(right),1);
flip_switch(1:length(surfArea_LH)/2) = 1;
clear all
if rand_flip_switch(s) == 1
    pseudo_left(s,1) = left(s);
    pseudo_right(s,1) = right(s);
elseif rand_flip_switch(s) == 0
    pseudo_right(s,1) = left(s);
    pseudo_left(s,1) = right(s);
end
end
perm_diffs(p) = mean(pseudo_left - pseudo_right);
end
end
load
(' /Users/dianaperez/Desktop/Thesis/Variant_Properties/HCP384_new_split_variants_info_wSurfArea.mat')
bord_ectop = variants_info.left_hem.indiv_vars(:,5);
bordervars_LH = variants_info.left_hem.indiv_vars(bord_ectop==1,:);
ectopicvars_LH = variants_info.left_hem.indiv_vars(bord_ectop==2,:);
bord_ectop1 = variants_info.right_hem.indiv_vars(:,5);
```



```
bordervars_RH = variants_info.right_hem.indiv_vars(bord_ectop1==1,:);
ectopicvars_RH = variants_info.right_hem.indiv_vars(bord_ectop1==2,:);
numperms = 1000;
left = bordervars_LH{:,5};
right = bordervars_RH{:,5};
true_diff = mean(left)- mean(right);
flip_switch = zeros(length(left)+length(right),1);
flip_switch(1:length(left)/2) = 1;
all = [left; right];
for p = 1:numperms
    rng('shuffle');
    ind = randperm(length(flip_switch));
    rand_flip_switch = flip_switch(ind);
    pseudo_left = [];
    pseudo_right = [];
    for s = 1:length(all)
        if rand_flip_switch(s) == 1
            pseudo_left = [pseudo_left; all(s)];
        elseif rand_flip_switch(s) == 0
            pseudo_right(s,1) = [pseudo_right; all(s)];
        end
    end
    perm_diffs(p) = mean(pseudo_left - pseudo_right);
end
permute_bord_ectop_size_byHem
p = length(find(perm_diffs<true_diff))
p = length(find(perm_diffs>true_diff))
left = ectopicvars_LH{:,3};
right = ectopicvars_RH{:,3};
true_diff = mean(left)- mean(right);
flip_switch = zeros(length(left)+length(right),1);
flip_switch(1:length(left)/2) = 1;
all = [left; right];
for p = 1:numperms
    rng('shuffle');
    ind = randperm(length(flip_switch));
    rand_flip_switch = flip_switch(ind);
    pseudo_left = [];
    pseudo_right = [];
    for s = 1:length(all)
        if rand_flip_switch(s) == 1
            pseudo_left = [pseudo_left; all(s)];
        elseif rand_flip_switch(s) == 0
            pseudo_right = [pseudo_right; all(s)];
        end
    end
    perm_diffs(p) = mean(pseudo_left) - mean(pseudo_right);
end
p = length(find(perm_diffs<true_diff))
clear all
load
('Users/dianaperez/Desktop/Thesis/Variant_Properties/Permutation_Tests/HCP384_LhemvsRhem
_num_vars.mat')
clear all
load
('Users/dianaperez/Desktop/Thesis/Variant_Properties/Permutation_Tests/HCP384_LhemvsRhem
_num_vars.mat')
clear output_mat
ind = zeros(1001,1);
ind(1,1) = 1;
```

```

handles = plotSpread(perms_varSize, 'categoryMarkers', {'o', '.'}, 'categoryLabels', ↵
{'Permuted Correlations','True Correlation'}, 'categoryColors', {'k','r'}, 'categoryIdx', ↵
[ind], 'xNames', {'Average Variant Size'}, 'xValues', [1])
scatter(1,perms_varSize(1,:), 'MarkerEdgeColor', 'red', 'MarkerFaceColor', 'red', ↵
'SizeData', 50)
ax = gca;
ax.FontSize = 24;
axis([0.5, 1.5, -60, 20])
handles = plotSpread(perms_varSize, 'categoryMarkers', {'o', '.'}, 'categoryLabels', ↵
{'Permuted Correlations','True Correlation'}, 'categoryColors', {'k','r'}, 'categoryIdx', ↵
[ind], 'xNames', {'Number of Variants'}, 'xValues', [1])
load ↵
('/Users/dianaperez/Desktop/Thesis/Variant_Properties/Permutation_Tests/HCP384_LhemvsRhem ↵
_num_vars.mat')
perms_numVars = output_mat;
clear output_mat
handles = plotSpread(perms_numVars, 'categoryMarkers', {'o', '.'}, 'categoryLabels', ↵
{'Permuted Correlations','True Correlation'}, 'categoryColors', {'k','r'}, 'categoryIdx', ↵
[ind], 'xNames', {'Number of Variants'}, 'xValues', [1])
scatter(1,perms_numVars(1,:), 'MarkerEdgeColor', 'red', 'MarkerFaceColor', 'red', ↵
'SizeData', 50)
ax = gca;
ax.FontSize = 24;
axis([0.5, 1.5, -60, 20])
ylabel('Difference in Number of Variants')
handles = plotSpread(perms_numVars, 'categoryMarkers', {'o', '.'}, 'categoryLabels', ↵
{'Permuted Correlations','True Correlation'}, 'categoryColors', {'k','r'}, 'categoryIdx', ↵
[ind], 'xNames', {'Number of Variants'}, 'xValues', [1])
scatter(1,perms_numVars(1,:), 'MarkerEdgeColor', 'red', 'MarkerFaceColor', 'red', ↵
'SizeData', 50)
ax = gca;
ax.FontSize = 24;
axis([0.5, 1.5, -60, 20])
ylabel('Difference in Number of Variants')
close gcf
clear all
load ↵
('/Users/dianaperez/Desktop/Thesis/Variant_Properties/HCP384_new_split_variants_info_wSur ↵
fArea.mat')
numVarsLH = variants_info.left_hem.group_avg{:,1};
numVarsRH = variants_info.right_hem.group_avg{:,1};
leftHem = numVarsLH;
rightHem = numVarsRH;
[h,mu,sigma,q,notch] = al_goodplot(leftHem,[0],.2,[ 0    0.4470    0.7410], 'left');
[h,mu,sigma,q,notch] = al_goodplot(rightHem,[0.1],.2,[0.8500    0.3250 ↵
0.0980], 'right'); % red-ish
close gcf
[h,mu,sigma,q,notch] = al_goodplot(leftHem,[0],.2,[ 0    0.4470    0.7410], 'left');
[h,mu,sigma,q,notch] = al_goodplot(rightHem,[0.01],.2,[0.8500    0.3250 ↵
0.0980], 'right'); % red-ish
xticks([0.105])
xticklabels('Number of Variants')
axis([-0.2, 0.5, 100, 800])
ax = gca;
ax.FontSize = 24;
close gcf
[h,mu,sigma,q,notch] = al_goodplot(leftHem,[0],.2,[ 0    0.4470    0.7410], 'left');
[h,mu,sigma,q,notch] = al_goodplot(rightHem,[0.01],.2,[0.8500    0.3250 ↵
0.0980], 'right'); % red-ish
xticks([0.105])

```

```
xticklabels('Number of Variants')
xticks([0.005])
ax = gca;
ax.FontSize = 24;
axis([-0.2, 0.5, -5, 25])
axis([-0.2, 0.2, -5, 25])
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.3, 0.7]); %first and second
control position on screen, third controls width, and fourth controls height
print(gcf, '/Users/dianaperez/Desktop/FigureforPres_numVars_HCP.jpg', '-dpng', '-r300')
numBorderLH = variants_info.left_hem.group_avg{:,4};
numBorderRH = variants_info.right_hem.group_avg{:,4};
numEctopicLH = variants_info.left_hem.group_avg{:,5};
numEctopicRH = variants_info.right_hem.group_avg{:,5};
leftHem = [numBorderLH numEctopicLH];
rightHem = [numBorderRH numEctopicRH];
[h,mu,sigma,q,notch] = al_goodplot(leftHem,[-1,0],.2,[ 0    0.4470    0.7410],'left');
[h,mu,sigma,q,notch] = al_goodplot(leftHem,[-1,1],.2,[ 0    0.4470    0.7410],'left');
numBorderLH = variants_info.left_hem.group_avg{:,5};
numBorderRH = variants_info.right_hem.group_avg{:,5};
numEctopicLH = variants_info.left_hem.group_avg{:,7};
numEctopicRH = variants_info.right_hem.group_avg{:,7};
leftHem = [numBorderLH numEctopicLH];
rightHem = [numBorderRH numEctopicRH];
[h,mu,sigma,q,notch] = al_goodplot(leftHem,[-1,1],.2,[ 0    0.4470    0.7410],'left');
[h,mu,sigma,q,notch] = al_goodplot(rightHem,[-1,1],.2,[0.8500    0.3250
0.0980],'right'); % red-ish
close(gcf)
[h,mu,sigma,q,notch] = al_goodplot(leftHem,[-1,0],.2,[ 0    0.4470    0.7410],'left');
[h,mu,sigma,q,notch] = al_goodplot(rightHem,[-1,0],.2,[0.8500    0.3250
0.0980],'right'); % red-ish
close(gcf)
[h,mu,sigma,q,notch] = al_goodplot(rightHem,[-1,-.6],.2,[0.8500    0.3250
0.0980],'right'); % red-ish
[h,mu,sigma,q,notch] = al_goodplot(leftHem,[-1,-.6],.2,[ 0    0.4470    0.7410],'left');
ax = gca;
ax.FontSize = 24;
ylabel('Number of Variants')
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.5, 0.7]); %first and second
control position on screen, third controls width, and fourth controls height
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.3, 0.7]); %first and second
control position on screen, third controls width, and fourth controls height
xticklabels({'Border Shifts', 'Ectopic Intrusions'})
xticks([-1,-.6])
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.4, 0.7]); %first and second
control position on screen, third controls width, and fourth controls height
print(gcf, '/Users/dianaperez/Desktop/FigureforPres_bordEctopic_HCP.jpg', '-dpng', '-r
300')
ft_read_cifti_mod
('/Users/dianaperez/Desktop/Thesis/old_Stuff/newsplit_vs_oldsplit_comparison/571144_Thres
holdedVariantMap_SNRExclude_networksAssigned_10.dtseries.nii')
cifti = ans;
bin_cifti = logical(cifti.data);
unique(bin_cifti)
template = cifti;
template.data = bin_cifti;
ft_write_cifti_mod('/Users/dianaperez/Desktop/binarized_varmap_HCPsub571144.dtseries.
nii', template)
ft_read_cifti_mod
('/Users/dianaperez/Desktop/Thesis/old_Stuff/newsplit_vs_oldsplit_comparison/571144_reass
igned.dtseries.nii')
```

```
cifti = ans;
find(cifti.data==11)
unique(cifti.data)
ft_read_cifti_mod
('/Volumes/RESEARCH_HD/HCP_Variants/new_split_vars/571144_revised_uniqueIDs_minsize30.
dtseries.nii')
cifti = ans;
find(cifti.data==11)
verts = ans;
verts_var_11 = ans;
verts_var_10 = find(cifti.data==10);
bin_cifti = zeroes(size(cifti.data));
bin_cifti = zeros(size(cifti.data));
bin_cifti(verts_var_10) = 1;
bin_cifti(verts_var_11) = 1;
template.data = bin_cifti;
ft_write_cifti_mod('/Users/dianaperez/Desktop/binarized_varmap_HCPsub571144_splitvars.
dtseries.nii', template)
bin_cifti(verts_var_10) = 7;
template.data = bin_cifti;
ft_write_cifti_mod
('/Users/dianaperez/Desktop/binarized_varmap_HCPsub571144_splitvars_var10_lang.dtseries.
nii', template)
bin_cifti(verts_var_10) = 1;
bin_cifti(verts_var_11) = 7;
template.data = bin_cifti;
ft_write_cifti_mod
('/Users/dianaperez/Desktop/binarized_varmap_HCPsub571144_splitvars_var11_lang.dtseries.
nii', template)
cifti = ft_read_cifti_mod
('/Users/dianaperez/Desktop/Thesis/old_Stuff/newsplit_vs_oldsplit_comparison/571144_Thres
holdedVariantMap_SNRExclude_networksAssigned_10.dtseries.nii')
bin_cifti = zeroes(size(cifti.data));
bin_cifti = zeros(size(cifti.data));
verts = find(cifti.data==24);
bin_cifti(verts) = 1;
template.data = bin_cifti;
ft_write_cifti_mod('/Users/dianaperez/Desktop/binarized_varmap_HCPsub571144_bigVarOnly.
dtseries.nii', template)
unique(cifti.data)
%-- 10/19/21, 11:58 AM --%
load
('/Users/dianaperez/Box/Research/Lateralization_thresh10/subjectData/goodSubs_addtlinfo.
mat')
lang_scores = HCP_goodSubs{2:end,5};
lang_scores = HCP_goodSubs{:,5};
HCP_goodSubs{:,5}
HCP_goodSubs{5}
HCP_goodSubs{1:384,5}
lang_scores = cell2mat(HCP_goodSubs{2:end,5});
lang_scores = cell2mat(HCP_goodSubs{:,5});
lang_scores = HCP_goodSubs1{2:end};
lang_scores = cell2mat(HCP_goodSubs1{2:end});
load
('/Users/dianaperez/Desktop/Thesis/Variant_Properties/HCP384_new_split_variants_info_wSur
fArea.mat')
sub = 2;
lang_scores = [];
for s = 1:384
lang_scores(s,1) = cell2mat(HCP_goodSubs{sub,5});
```

```
sub = sub+1;
end
sub = 2;
lang_scores = [];
for s = 1:384
    lang_scores(s,1) = cell2mat(HCP_goodSubs{sub,5});
    sub = sub+1;
end
lang_scores = cell2mat(HCP_goodSubs1(2:end));
clear HCP_goodSubs1
clear s sub ans
numVars = variants_info.left_hem.group_avg{1,1};
diff_score = variants_info.left_hem.group_avg{1,1} - variants_info.right_hem.group_avg{1,1}
diff_scores = variants_info.left_hem.group_avg{:,1} - variants_info.right_hem.group_avg{:,1};
corr(lang_scores, diff_scores)
corrcoeff(lang_scores, diff_scores)
corrcoef(lang_scores, diff_scores)
find(isnan(diff_scores))
find(isnan(diff_scores))
find(isnan(lang_scores))
backup = [lang_scores diff_scores];
backup(ans,:) = [];
find(isnan(backup))
lang_scores(nans,:) = [];
diff_scores(nans,:) = [];
nans = find(isnan(lang_scores));
lang_scores(nans,:) = [];
diff_scores(nans,:) = [];
corrcoef(lang_scores, diff_scores)
[R P] = corrcoef(lang_scores, diff_scores)
diff_scores = variants_info.left_hem.group_avg{:,2} - variants_info.right_hem.group_avg{:,2};
diff_scores(nans,:) = [];
[R P] = corrcoef(lang_scores, diff_scores)
diff_scores = variants_info.left_hem.group_avg{:,3} - variants_info.right_hem.group_avg{:,3};
diff_scores(nans,:) = [];
[R P] = corrcoef(lang_scores, diff_scores)
diff_scores = variants_info.left_hem.group_avg{:,4} - variants_info.right_hem.group_avg{:,4};
diff_scores(nans,:) = [];
[R P] = corrcoef(lang_scores, diff_scores)
diff_scores = variants_info.left_hem.group_avg{:,1} - variants_info.right_hem.group_avg{:,1};
diff_scores(nans,:) = [];
[R P] = corrcoef(lang_scores, diff_scores);
plot(lang_scores, diff_scores)
scatter(lang_scores,diff_scores)
clear all
ft_read_cifti_mod
('Users/dianaperez/Downloads/VariantSplittingFiles/variantsPCA_allSubs_HCPfull_topEigenv
alue.dtseries.nii')
load
('Users/dianaperez/Box/Research/Lateralization_thresh10/subjectData/goodSubs_addtlinf
mat')
lang_scores = cell2mat(HCP_goodSubs1(2:end));
load
('Users/dianaperez/Desktop/Thesis/Variant_Properties/HCP384_new_split_variants_info_wSur
```

```
fArea.mat')
diff_scores = variants_info.left_hem.group_avg(:,1) - variants_info.right_hem.group_avg{
(:,1);
nans = find(isnan(lang_scores));
lang_scores(nans,:) = [];
diff_scores(nans,:) = [];
[R P] = corrcoef(lang_scores, diff_scores);
diff_scores = variants_info.left_hem.group_avg(:,2) - variants_info.right_hem.group_avg{
(:,2);
%-- 10/25/21, 12:04 PM --%
load('/Users/dianaperez/Desktop/Thesis/Variant_Properties/HCP384_new_split_variants_info.
mat')
numVarsLH = variants_info.left_hem.group_avg(:,1);
numVarsRH = variants_info.right_hem.group_avg(:,1);
leftHem = numVarsLH;
rightHem = numVarsRH;
[h,mu,sigma,q,notch] = al_goodplot(leftHem,.1,.2,[ 0    0.4470    0.7410],'left');
addpath '/Users/dianaperez/Documents/MATLAB/upload'
[h,mu,sigma,q,notch] = al_goodplot(rightHem,0.11,.2,[0.8500    0.3250
0.0980],'right');
xticks([0.105])
[h,mu,sigma,q,notch] = al_goodplot(leftHem,0.1,.2,[ 0    0.4470    0.7410],'left');
xlabel('Number of Variants')
axis([-0.2, 0.5, 0, 20])
axis([-0.2, 0.5, 2, 20])
ax = gca;
ax.FontSize = 24;
xlabel('Variant Frequency Across Hemispheres')
ylabel('Number of Variant Regions')
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.3, 0.7]); %first and second
control position on screen, third controls width, and fourth controls height
print(gcf, '/Users/dianaperez/Desktop/posterfig_numVars_HCP.jpg', '-dpng', '-r300')
axis([-0.2, 0.5, 0, 20])
print(gcf, '/Users/dianaperez/Desktop/posterfig_numVars_HCP.jpg', '-dpng', '-r300')
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.3, 0.9]); %first and second
control position on screen, third controls width, and fourth controls height
print(gcf, '/Users/dianaperez/Desktop/posterfig_numVars_HCP.jpg', '-dpng', '-r300')
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.75]); %first and
second control position on screen, third controls width, and fourth controls height
print(gcf, '/Users/dianaperez/Desktop/posterfig_numVars_HCP.jpg', '-dpng', '-r300')
xlabel('Number of Variants')
print(gcf, '/Users/dianaperez/Desktop/posterfig_numVars_HCP.jpg', '-dpng', '-r300')
operc_cluster
addpath '/Users/dianaperez/Documents/GitHub/Lateralization_Variants/thesis_scripts'
operc_cluster
output_dir = '/Users/dianaperez/Desktop/Thesis/Spatial_Location/';
load([output_dir 'operc_nets_lefthemcluster_righthem.mat']);
leftclust_righthem = LH_cluster;
load([output_dir 'operc_nets_lefthemcluster_lefthem.mat']);
leftclust_lefthem = LH_cluster;
clear LH_cluster
clear all
output_dir = '/Users/dianaperez/Desktop/Thesis/Spatial_Location/';
load([output_dir 'operc_nets_lefthemcluster_righthem.mat']);
leftclust_righthem = LH_cluster;
load([output_dir 'operc_nets_lefthemcluster_lefthem.mat']);
leftclust_lefthem = LH_cluster;
clear LH_cluster
for net = 1:16
nets_left(net) = length(find(leftclust_lefthem.number(:,3)==net));
```

```

end
length(find(leftclust_lefthem.number(:,3)==7))
for net = 1:16
nets_left(net) = length(find(leftclust_lefthem.vars(:,3)==net));
end
for net = 1:16
nets_right(net) = length(find(leftclust_righthem.vars(:,3)==net));
end
network_names = {'DMN' 'Vis' 'FP' 'DAN' 'Lang' 'CO' 'SMd' 'Aud' 'PON'};
nets_left(13:15) = [];
nets_left(11) = [];
nets_left(8) = [];
nets_left(6) = [];
nets_left(4) = [];
t = tiledlayout(1,2,'TileSpacing', 'compact');
ax1 = nexttile;
h = pie(ax1,nets_left, network_names);
title('Left Hem')
patchHandle = findobj(h, 'Type', 'Patch');
set(patchHandle, {'FaceColor'}, mat2cell(rgb_colors, ones(size(rgb_colors,1),1),3))
rgb_colors = [1 0 0; %DMN
0 0 .6; %Vis
1 1 0; %FP
0 .8 0; %DAN
0 .6 .6;%VAN
.3 0 .6;%CON
.2 1 1;%SMd
.6 .2 1;%Aud
.8 .8 .6]; %PON
t = tiledlayout(1,2,'TileSpacing', 'compact');
ax1 = nexttile;
h = pie(ax1,nets_left, network_names);
title('Left Hem')
patchHandle = findobj(h, 'Type', 'Patch');
set(patchHandle, {'FaceColor'}, mat2cell(rgb_colors, ones(size(rgb_colors,1),1),3))
nets_right(13:15) = [];
nets_right(11) = [];
nets_right(8) = [];
nets_right(6) = [];
nets_right(4) = [];
ax2 = nexttile;
h = pie(ax2,nets_right, network_names);
title('Right Hem')
patchHandle = findobj(h, 'Type', 'Patch');
set(patchHandle, {'FaceColor'}, mat2cell(rgb_colors, ones(size(rgb_colors,1),1),3))
labels_legend = {'Default Mode', 'Visual', 'Frontoparietal', 'Dorsal Attention', '
'Language', 'Cingulo-opercular', 'Somatomotor Dorsal', 'Auditory', 'Parietal',
Occipital'};
lgd = legend(labels_legend);
lgd.Layout.Tile = 'east';
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.9, 0.7]); %first and second
control position on screen, third controls width, and fourth controls height
ax1.FontSize = 24;
lgd.FontSize = 24;
ax2.FontSize = 24;
print(gcf, '
/Users/dianaperez/Desktop/Thesis/Figures/Figure7_pieChart_operc_cluster_networks_lefthem
_HCP.jpg', '-dpng', '-r300')
output_dir = '/Users/dianaperez/Desktop/Thesis/Spatial_Location/';
load([output_dir 'operc_nets_righthemcluster_righthem.mat']);

```



```
rightclust_righthem = RH_cluster;
load([output_dir 'operc_nets_righthemcluster_lefthem.mat']);
rightclust_lefthem = RH_cluster;
clear RH_cluster
for net = 1:16
nets_left(net) = length(find(rightclust_lefthem.vars(:,3)==net));
end
for net = 1:16
nets_right(net) = length(find(rightclust_righthem.vars(:,3)==net));
end
network_names = {'DMN' 'Vis' 'FP' 'DAN' 'Lang' 'CO' 'SMd' 'Aud' 'PON'};
nets_left(13:15) = [];
nets_left(11) = [];
nets_left(8) = [];
nets_left(6) = [];
nets_left(4) = [];
for net = 1:16
nets_left(net) = length(find(rightclust_lefthem.vars(:,3)==net));
end
rgb_colors = [1 0 0;
1 1 0;
0 .8 0;
0 .6 .6;
0 0 0;
.3 0 .6;
1 .5 0;
.8 .8 .6];
output_dir = '/Users/dianaperez/Desktop/Thesis/Spatial_Location/';
load([output_dir 'operc_nets_righthemcluster_righthem.mat']);
rightclust_righthem = RH_cluster;
load([output_dir 'operc_nets_righthemcluster_lefthem.mat']);
rightclust_lefthem = RH_cluster;
clear RH_cluster
network_names = {'DMN' 'FP' 'DAN' 'Lang' 'Sal' 'CO' 'SMl' 'PON'};
for net = 1:16
nets_left(net) = length(find(rightclust_lefthem.vars(:,3)==net));
end
nets_left(12:16) = [];
nets_left(10) = [];
nets_left(6) = [];
nets_left(4) = [];
nets_left(2) = [];
t = tiledlayout(1,2,'TileSpacing', 'compact');
ax1 = nexttile;
h = pie(ax1,nets_left, network_names);
title('Left Hem')
patchHandle = findobj(h, 'Type', 'Patch');
set(patchHandle, {'FaceColor'}, mat2cell(rgb_colors, ones(size(rgb_colors,1),1),3))
for net = 1:16
nets_left(net) = length(find(rightclust_lefthem.vars(:,3)==net));
end
nets_left(12:15) = [];
nets_left(10) = [];
nets_left(6) = [];
nets_left(4) = [];
nets_left(2) = [];
t = tiledlayout(1,2,'TileSpacing', 'compact');
ax1 = nexttile;
h = pie(ax1,nets_left, network_names);
title('Left Hem')
```



```

patchHandle = findobj(h, 'Type', 'Patch');
set(patchHandle, {'FaceColor'}, mat2cell(rgb_colors, ones(size(rgb_colors,1),1),3))
for net = 1:16
nets_right(net) = length(find(rightclust_righthem.number(:,3)==net));
end
for net = 1:16
nets_right(net) = length(find(rightclust_righthem.vars(:,3)==net));
end
nets_right(12:15) = [];
nets_left(10) = [];
nets_left(6) = [];
nets_left(4) = [];
nets_left(2) = [];
ax2 = nexttile;
h = pie(ax2,nets_right, network_names);
title('Right Hem')
patchHandle = findobj(h, 'Type', 'Patch');
set(patchHandle, {'FaceColor'}, mat2cell(rgb_colors, ones(size(rgb_colors,1),1),3))
labels_legend = {'Default Mode' 'Frontoparietal', 'Dorsal Attention', 'Language', 'Salience', 'Cingulo-Opercular', 'Somatomotor Ventral' 'Parieto-Occipital'};
lgd = legend(labels_legend);
lgd.Layout.Tile = 'east';
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.9, 0.7]); %first and second control position on screen, third controls width, and fourth controls height
ax1.FontSize = 24;
ax2.FontSize = 24;
lgd.FontSize = 24;
nets_right(12:15) = [];
nets_right(10) = [];
nets_right(6) = [];
nets_right(4) = [];
nets_right(2) = [];
for net = 1:16
nets_right(net) = length(find(rightclust_righthem.vars(:,3)==net));
end
nets_right(12:15) = [];
nets_right(10) = [];
nets_right(6) = [];
nets_right(4) = [];
nets_right(2) = [];
ax2 = nexttile;
h = pie(ax2,nets_right, network_names);
title('Right Hem')
patchHandle = findobj(h, 'Type', 'Patch');
set(patchHandle, {'FaceColor'}, mat2cell(rgb_colors, ones(size(rgb_colors,1),1),3))
labels_legend = {'Default Mode' 'Frontoparietal', 'Dorsal Attention', 'Language', 'Salience', 'Cingulo-Opercular', 'Somatomotor Ventral' 'Parieto-Occipital'};
lgd = legend(labels_legend);
lgd.Layout.Tile = 'east';
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.9, 0.7]); %first and second control position on screen, third controls width, and fourth controls height
ax1.FontSize = 24;
ax2.FontSize = 24;
lgd.FontSize = 24;
print(gcf, '
/Users/dianaperez/Desktop/Thesis/Figures/Figure7_pieChart_operc_cluster_networks_righthe
m_HCP.jpg', '-dpng', '-r300')
sum(nets_left)
for n = 1:length(nets_left)
proportion(n) = nets_left(n)/sum(nets_left);

```

```

end
for net = 1:16
nets_right(net) = length(find(leftclust_righthem.vars(:,3)==net));
end
nets_right(13:15) = [];
nets_right(11) = [];
nets_right(8) = [];
nets_right(6) = [];
nets_right(4) = [];
for n = 1:length(nets_right)
proportion(n) = nets_right(n)/sum(nets_right);
end
sum(nets_right)
network_names = {'DMN' 'Vis' 'FP' 'DAN' 'Lang' 'CO' 'SMd' 'Aud' 'PON'};
for net = 1:16
nets_left(net) = length(find(leftclust_lefthem.vars(:,3)==net));
end
nets_left(13:15) = [];
nets_left(11) = [];
nets_left(8) = [];
nets_left(6) = [];
nets_left(4) = [];
sum(nets_left)
for n = 1:length(nets_left)
proportion(n) = nets_left(n)/sum(nets_left);
end
for net = 1:16
nets_right(net) = length(find(rightclust_righthem.vars(:,3)==net));
end
nets_right(12:15) = [];
nets_right(10) = [];
nets_right(6) = [];
nets_right(4) = [];
nets_right(2) = [];
sum(nets_right)
for n = 1:length(nets_right)
proportion(n) = nets_right(n)/sum(nets_right);
end
%-- 10/30/21, 6:39 PM --%
permute_variantproperties
%-- 10/31/21, 7:17 PM --%
permute_variantproperties
avgSurfArea = allSubs_info.variants_info.left_hem.group_avg(:,4);
avgSurfArea = mean(allSubs_info.variants_info.left_hem.group_avg(:,4));
avgSurfArea_RH = mean(allSubs_info.variants_info.right_hem.group_avg(:,4))
disp(['Hemisphere X Handedness, Average Variant Surface Area (permute handedness) -<
Below: ' num2str(length(find(perm_diffs<true_diff))/numperms)])
disp(['Hemisphere X Handedness, Average Variant Surface Area (permute handedness) -<
Above: ' num2str(length(find(perm_diffs>true_diff))/numperms)])
true_diff = perm_diffs(1);
perm_diffs = perm_diffs(2:end);
disp(['Hemisphere X Handedness, Average Variant Surface Area (permute handedness) -<
Below: ' num2str(length(find(perm_diffs<true_diff))/numperms)])
disp(['Hemisphere X Handedness, Average Variant Surface Area (permute handedness) -<
Above: ' num2str(length(find(perm_diffs>true_diff))/numperms)])
[true_diff perm_diffs] = permute_values(allSubs_info.variants_info.left_hem.group_avg(:,<
4), allSubs_info.variants_info.right_hem.group_avg(:,4), flip_switch, numperms, 0, 1, 0);
disp(['Average Variant Size (permute hemispheres) - Below: ' num2str(length(find<
(perm_diffs<true_diff))/1000)])
disp(['Average Variant Size (permute hemispheres) - Above: ' num2str(length(find<

```

```
(perm_diffs>true_diff))/1000)]
permute_variantproperties
save([outputDir str 'surf_area_proportion.mat'], 'output_mat');
check_MRI_data
pwd
dcm_dir = '/Volumes/fsmresfiles/PBS/Gratton_Lab/Lifespan/Post-COVID/BIDS/DICOM';
cd dcm_dir
cd ~/
clear all
load('/Volumes/fsmresfiles/PBS/Gratton_Lab/Lifespan/Post-
COVID/BIDS/derivatives/preproc_FCProc/corrmats_Seitzman300/sub-LS02/sub-LS02_sess-1_task-
rest_corrmat_Seitzman300.mat')
LS02_ses-1 = corrmats;
LS02_ses1 = corrmats;
load('/Volumes/fsmresfiles/PBS/Gratton_Lab/Lifespan/Post-
COVID/BIDS/derivatives/preproc_FCProc/corrmats_Seitzman300/sub-LS02/sub-LS02_sess-2_task-
rest_corrmat_Seitzman300.mat')
LS02_ses2 = corrmats;
load('/Volumes/fsmresfiles/PBS/Gratton_Lab/Lifespan/Post-
COVID/BIDS/derivatives/preproc_FCProc/corrmats_Seitzman300/sub-LS02/sub-LS02_sess-3_task-
rest_corrmat_Seitzman300.mat')
LS02_ses3 = corrmats;
load('/Volumes/fsmresfiles/PBS/Gratton_Lab/Lifespan/Post-
COVID/BIDS/derivatives/preproc_FCProc/corrmats_Seitzman300/sub-LS02/sub-LS02_sess-4_task-
rest_corrmat_Seitzman300.mat')
LS02_ses4 = corrmats;
load('/Volumes/fsmresfiles/PBS/Gratton_Lab/Lifespan/Post-
COVID/BIDS/derivatives/preproc_FCProc/corrmats_Seitzman300/sub-LS02/sub-LS02_sess-5_task-
rest_corrmat_Seitzman300.mat')
LS02_ses5 = corrmats;
corr(LS02_ses1,LS02_ses2)
corrcoef(LS02_ses1,LS02_ses2)
corrs_LS02 = [];
corrs_LS02(1) = corrcoef(LS02_ses1,LS02_ses2);
corrs_LS02(1) = 0.67;
corrcoef(LS02_ses1,LS02_ses3)
corrs_LS02(2) = 0.66;
corrcoef(LS02_ses1,LS02_ses4)
corrs_LS02(3) = 0.66;
corrcoef(LS02_ses1,LS02_ses5)
corrs_LS02(3) = 0.66;
corrs_LS02(4) = 0.66;
corrcoef(LS02_ses2,LS02_ses5)
corrs_LS02(5) = 0.66;
corrcoef(LS02_ses2,LS02_ses4)
corrs_LS02(6) = 0.73;
corrcoef(LS02_ses2,LS02_ses3)
corrs_LS02(7) = 0.73;
corrcoef(LS02_ses3,LS02_ses4)
corrs_LS02(8) = 0.76;
corrcoef(LS02_ses3,LS02_ses5)
corrs_LS02(9) = 0.64;
corrcoef(LS02_ses4,LS02_ses5)
corrs_LS02(10) = 0.66;
mean(corrs_LS02)
clear ans corrmats sess_roi* tmask* LS02*
load('/Volumes/fsmresfiles/PBS/Gratton_Lab/Lifespan/Pre-
COVID/BIDS/Nifti/derivatives/preproc_FCProc/corrmats_Seitzman300/sub-LS02/sub-LS02_sess-
1_task-rest_corrmat_Seitzman300.mat')
sub = 'LS02';
```

```
pre_dir = '/Volumes/fsmresfiles/PBS/Gratton_Lab/Lifespan/Pre-  
COVID/BIDS/Nifti/derivatives/preproc_FCProc/cormats_Seitzman300/';  
post_dir = '/Volumes/fsmresfiles/PBS/Gratton_Lab/Lifespan/Post-  
COVID/BIDS/derivatives/preproc_FCProc/cormats_Seitzman300/';  
pre_sessions = 3;  
post_sessions = 5;  
for pre_ses = 1:pre_sessions  
load([pre_dir 'sub-' sub '/sub-' sub '_sess-' pre_ses '_task-rest_corrmat_Seitzman300.  
mat'])  
cormats_pre(:, :, pre_ses) = cormat;  
end  
for post_ses = 1:post_sessions  
load([post_dir 'sub-' sub '/sub-' sub '_sess-' post_ses '_task-rest_corrmat_Seitzman300.  
mat'])  
cormats_post(:, :, post_ses) = cormat;  
end  
%% compare pre and post covid cormats  
sub = 'LS02';  
pre_dir = '/Volumes/fsmresfiles/PBS/Gratton_Lab/Lifespan/Pre-  
COVID/BIDS/Nifti/derivatives/preproc_FCProc/cormats_Seitzman300/';  
post_dir = '/Volumes/fsmresfiles/PBS/Gratton_Lab/Lifespan/Post-  
COVID/BIDS/derivatives/preproc_FCProc/cormats_Seitzman300/';  
pre_sessions = 3;  
post_sessions = 5;  
for pre_ses = 1:pre_sessions  
load([pre_dir 'sub-' sub '/sub-' sub '_sess-' num2str(pre_ses) '_task-  
rest_corrmat_Seitzman300.mat'])  
cormats_pre(:, :, pre_ses) = cormat;  
end  
for post_ses = 1:post_sessions  
load([post_dir 'sub-' sub '/sub-' sub '_sess-' num2str(post_ses) '_task-  
rest_corrmat_Seitzman300.mat'])  
cormats_post(:, :, post_ses) = cormat;  
end  
clear cormat sess_roi* tmask* LS02*  
corr = corrcoef(cormats_post(:, :, 1), cormats_post(:, :, 2));  
corr(2)  
combos = combntns(1:3, 2)  
%-- 11/4/21, 5:57 PM --%  
combos = combntns(1:3, 2)  
length(combos)  
combos(1, 1)  
combos(1, 2)  
combos(2, 1)  
prevspost_compare  
clear all  
prevspost_compare  
%-- 11/10/21, 6:14 PM --%  
rgb_colors = [1 0 0; %DMN  
0 0 .6; %Vis  
1 1 0; %FP  
.67 .67 .67; %Unassigned  
0 .8 0; %DAN  
.67 .67 .67; %Unassigned2  
0 .6 .6; %VAN  
0 0 0; % Sal  
.3 0 .6; %CON  
.2 1 1; %SMd  
1 .5 0; % SMI  
.6 .2 1; %Aud
```

```

.2 1 .2; %Tpole
0 .2 .4; %MTL
0 0 1; %PMN
.8 .8 .6; %PON
.8 .8 .6]; %PON
network_names = {'DMN' 'Vis' 'FP' 'DAN' 'Lang' 'Sal' 'CO' 'SMd' 'SML' 'Aud' ↵
'Tpole' 'MTL' 'PMN' 'PON'};
rgb_colors = [1 0 0; %DMN
0 0 .6; %Vis
1 1 0; %FP
.67 .67 .67; %Unassigned
0 .8 0; %DAN
.67 .67 .67; %Unassigned2
0 .6 .6; %VAN
0 0 0; % Sal
.3 0 .6; %CON
.2 1 1; %SMd
1 .5 0; % SML
.6 .2 1; %Aud
.2 1 .2; %Tpole
0 .2 .4; %MTL
0 0 1; %PMN
.8 .8 .6]; %PON
network_names = {'DMN' 'Vis' 'FP' 'Reward' 'DAN' 'Unassigned' 'Lang' 'Sal' 'CO' ↵
'SMd' 'SML' 'Aud' 'Tpole' 'MTL' 'PMN' 'PON'};
load('/Users/dianaperez/Desktop/network_sizes.mat')
LH_vert_prop = size_of_net(:,2);
LH_surf_prop = size_of_net(:,4);
RH_vert_prop = size_of_net(:,6);
RH_surf_prop = size_of_net(:,8);
vert_perc = [LH_vert_prop RH_vert_prop];
surf_perc = [LH_surf_prop RH_surf_prop];
plot(surf_perc)
scatter(surf_perc)
scatter(LH_surf_prop, RH_surf_prop)
scatter(LH_surf_prop, RH_surf_prop, 10, rgb_colors)
scatter(LH_surf_prop, RH_surf_prop, 50, rgb_colors, 'filled')
scatter(LH_surf_prop, RH_surf_prop, 150, rgb_colors, 'filled')
scatter(LH_surf_prop, RH_surf_prop, 100, rgb_colors, 'filled')
help axis
axis([0 0.20 0 0.20])
axis([0 0.18 0 0.18])
refline(1,0)
xlabel('Left Hemisphere')
ylabel('Right Hemisphere')
line = refline(1,0);
line.Color = 'black';
title('Percent surface area by hemisphere')
ax = gca;
ax.FontSize = 24;
m = findobj(gca, 'Type', 'Scatter');
hleg1 = legend(m, network_names, 'Location', 'SouthOutside');
hleg1.FontSize = 14;
scatter(LH_surf_prop, RH_surf_prop, 100, rgb_colors, 'filled')
axis([0 0.18 0 0.18])
m = findobj(gca, 'Type', 'Scatter');
hleg1 = legend(m, network_names, 'Location', 'SouthOutside');
hleg1.FontSize = 14;
m = findobj(gca, 'Type', 'Scatter')
m(1:16)

```

```

scatter(LH_surf_prop, RH_surf_prop, 100, rgb_colors, 'filled')
axis([0 0.18 0 0.18])
legend(network_names)
scatter(LH_surf_prop, RH_surf_prop, 100, rgb_colors, 'filled')
axis([0 0.18 0 0.18])
line = reffline(1,0);
xlabel('Left Hemisphere')
ylabel('Right Hemisphere')
title('Percent surface area by hemisphere')
ax = gca;
ax.FontSize = 24;
print(gcf, '✓
'/Users/dianaperez/Desktop/Thesis/Figures/Figure1_permutations_scatterplot_HCP.jpg', '-✓
dpng', '-r300')
close gcf
scatter(LH_surf_prop, RH_surf_prop, 100, rgb_colors, 'filled')
axis([0 0.18 0 0.18])
line = reffline(1,0);
xlabel('Left Hemisphere')
ylabel('Right Hemisphere')
title('Percent surface area by hemisphere')
ax = gca;
ax.FontSize = 24;
print(gcf, '/Users/dianaperez/Desktop/Thesis/Figures/network_surfarea_groupavg.jpg', '-✓
dpng', '-r300')
scatter(LH_surf_prop, RH_surf_prop, 100, rgb_colors, 'filled')
axis([0 0.18 0 0.18])
line = reffline(1,0);
line.Color = 'black';
xlabel('Left Hemisphere')
ylabel('Right Hemisphere')
title('Percent surface area by hemisphere')
ax = gca;
ax.FontSize = 24;
print(gcf, '/Users/dianaperez/Desktop/Thesis/Figures/network_surfarea_groupavg.jpg', '-✓
dpng', '-r300')
scatter(LH_vert_prop, RH_vert_prop, 100, rgb_colors, 'filled')
axis([0 0.18 0 0.18])
line = reffline(1,0);
line.Color = 'black';
xlabel('Left Hemisphere')
ylabel('Right Hemisphere')
title('Percent number of vertices by hemisphere')
ax = gca;
ax.FontSize = 24;
print(gcf, '/Users/dianaperez/Desktop/Thesis/Figures/network_numverts_groupavg.jpg', '-✓
dpng', '-r300')
scatter(LH_surf_prop, RH_surf_prop, 80, rgb_colors, 'filled')
axis([0 0.18 0 0.18])
line = reffline(1,0);
line.Color = 'black';
xlabel('Left Hemisphere')
ylabel('Right Hemisphere')
title('Percent surface area by hemisphere')
% ax = gca;
% ax.FontSize = 24;
print(gcf, '/Users/dianaperez/Desktop/Thesis/Figures/network_surfarea_groupavg.jpg', '-✓
dpng', '-r300')
scatter(LH_vert_prop, RH_vert_prop, 80, rgb_colors, 'filled')
axis([0 0.18 0 0.18])

```

```

line = reffline(1,0);
line.Color = 'black';
xlabel('Left Hemisphere')
ylabel('Right Hemisphere')
title('Percent number of vertices by hemisphere')
% ax = gca;
% ax.FontSize = 24;
print(gcf, '/Users/dianaperez/Desktop/Thesis/Figures/network_numverts_groupavg.jpg', '-r300', '-r300')
scatter(LH_surf_prop, RH_surf_prop, 80, rgb_colors, 'filled')
axis([0 0.18 0 0.18])
line = reffline(1,0);
line.Color = 'black';
xlabel('Left Hemisphere')
ylabel('Right Hemisphere')
title('Percent surface area by hemisphere')
ax = gca;
ax.FontSize = 24;
print(gcf, '/Users/dianaperez/Desktop/Thesis/Figures/network_surfarea_groupavg.jpg', '-r300', '-r300')
clear all
network_names = {'DMN' 'Vis' 'FP' 'Reward' 'DAN' 'Unassigned' 'Lang' 'Sal' 'CO' 'SMd' 'SML' 'Aud' 'Tpole' 'MTL' 'PMN' 'PON'};
rgb_colors = [1 0 0; %DMN
0 0 .6; %Vis
1 1 0; %FP
.67 .67 .67; %Unassigned
0 .8 0; %DAN
.67 .67 .67; %Unassigned2
0 .6 .6; %VAN
0 0 0; % Sal
.3 0 .6; %CON
.2 1 1; %SMd
1 .5 0; % SML
.6 .2 1; %Aud
.2 1 .2; %Tpole
0 .2 .4; %MTL
0 0 1; %PMN
.8 .8 .6]; %PON
% load the matrix with sizes
load('/Users/dianaperez/Desktop/MSc_network_sizes.mat')
LH_vert_prop = size_of_net(:,2);
LH_surf_prop = size_of_net(:,4);
RH_vert_prop = size_of_net(:,6);
RH_surf_prop = size_of_net(:,8);
scatter(LH_surf_prop, RH_surf_prop, 80, rgb_colors, 'filled')
axis([0 0.18 0 0.18])
line = reffline(1,0);
line.Color = 'black';
xlabel('Left Hemisphere')
ylabel('Right Hemisphere')
title('Percent surface area by hemisphere')
ax = gca;
ax.FontSize = 24;
close(gcf)
scatter(LH_surf_prop, RH_surf_prop, 80, rgb_colors, 'filled')
%axis([0 0.18 0 0.18])
line = reffline(1,0);
line.Color = 'black';
xlabel('Left Hemisphere')

```

```
ylabel('Right Hemisphere')
title('Percent surface area by hemisphere')
ax = gca;
ax.FontSize = 24;
print(gcf, '/Users/dianaperez/Desktop/Thesis/Figures/MSC_network_surfarea_groupavg.jpg',
'-dpng', '-r300')
load('/Users/dianaperez/Desktop/HCP_network_sizes.mat')
LH_vert_prop = size_of_net(:,2);
LH_surf_prop = size_of_net(:,4);
RH_vert_prop = size_of_net(:,6);
RH_surf_prop = size_of_net(:,8);
scatter(LH_surf_prop, RH_surf_prop, 80, rgb_colors, 'filled')
%axis([0 0.18 0 0.18])
line = reffline(1,0);
line.Color = 'black';
xlabel('Left Hemisphere')
ylabel('Right Hemisphere')
title('Percent surface area by hemisphere')
ax = gca;
ax.FontSize = 24;
print(gcf, '/Users/dianaperez/Desktop/Thesis/Figures/HCP_network_surfarea_groupavg.jpg',
'-dpng', '-r300')
load('/Users/dianaperez/Desktop/Thesis/Variant_Properties/HCP384_new_split_networksxHem.
mat')
avg_numVars = mean(networksxHem.clustersLH(:,,:));
avg_numVars_RH = mean(networksxHem.clustersRH(:,,:));
scatter(avg_numVars, avg_numVars_RH, 80, rgb_colors, 'filled')
axis([0 2.5 0 2.5])
line = reffline(1,0);
line.Color = 'black';
xlabel('Left Hemisphere')
ylabel('Right Hemisphere')
title('Average Number of Variants')
ax = gca;
ax.FontSize = 24;
print(gcf, '/Users/dianaperez/Desktop/Thesis/Figures/HCP_numVarsbyHem.jpg', '-dpng', '-
r300')
help pie
h = pie(avg_numVars, network_names);
patchHandle = findobj(h, 'Type', 'Patch');
set(patchHandle, {'FaceColor'}, mat2cell(rgb_colors, ones(size(rgb_colors,1),1),3))
lgd = legend(network_names);
h = pie(avg_numVars, network_names)
patchHandle = findobj(h, 'Type', 'Patch')
set(patchHandle, {'FaceColor'}, mat2cell(rgb_colors, ones(size(rgb_colors,1),1),3))
lgd = legend(network_names);
lgd.Layout.Tile = 'east';
lgd.LayoutTile = 'east';
lgd.Layout.Tile = 'east';
lgd.FontSize = 24;
print(gcf, '/Users/dianaperez/Desktop/Thesis/Figures/Legend.jpg', '-dpng', '-r300')
load('/Users/dianaperez/Desktop/network_sizes.mat')
LH_vert_prop = size_of_net(:,2);
LH_surf_prop = size_of_net(:,4);
RH_vert_prop = size_of_net(:,6);
RH_surf_prop = size_of_net(:,8);
scatter(LH_surf_prop, RH_surf_prop, 80, rgb_colors, 'filled')
axis([0 0.18 0 0.18])
line = reffline(1,0);
line.Color = 'black';
```



```
xlabel('Left Hemisphere')
ylabel('Right Hemisphere')
ax = gca;
ax.FontSize = 24;
print(gcf, '/Users/dianaperez/Desktop/Thesis/Figures/network_surfarea_groupavg.jpg', '-d
png', '-r300')
title('Percent surface area by hemisphere')
print(gcf, '/Users/dianaperez/Desktop/Thesis/Figures/network_surfarea_groupavg.jpg', '-d
png', '-r300')
this.metadata = [];
this.label = [];
this.data = [];
this_LH = this;
this_RH = this;
this_LH = gifti_read
('/Users/dianaperez/Box/Dependencies/32k_ConteAtlas_v2_distribute/RSN-networks.L.
32k_fs_LR.label.gii', this);
this_RH = gifti_read
('/Users/dianaperez/Box/Dependencies/32k_ConteAtlas_v2_distribute/RSN-networks.R.
32k_fs_LR.label.gii', this);
unique(this_LH.data{1,1}.data)
unique(this_LH.data{1,2}.data)
unique(this_LH.data{1,3}.data)
unique(this_LH.data{1,4}.data)
unique(this_RH.data{1,1}.data)
cifti_w_nonbrain = [this_LH.data{1,1}.data; this_RH.data{1,1}.data];
template = ft_read_cifti_mod
('/Users/dianaperez/Box/Dependencies/Important_Files/120_templates.dtseries.nii');
cifti = cifti_w_nonbrain(template.brainstructure>0);
template.data = cifti;
ft_write_cifti_mod('/Users/dianaperez/Desktop/Yeo_nets.dtseries.nii', template)
for v = 1:length(this_RH.data{1,1}.data)
end
medial = find(this_RH.data{1,1}.data==35);
this_RH.data{1,1}.data(medial) = 0;
unique(this_RH.data{1,1}.data)
net1 = find(this_RH.data{1,1}.data==36);
this_RH.data{1,1}.data(net1) = 42;
this_RH.data{1,1}.data(net1) = 39;
unique(this_RH.data{1,1}.data)
net2 = find(this_RH.data{1,1}.data==37);
this_RH.data{1,1}.data(net2) = 40;
unique(this_RH.data{1,1}.data)
net3 = find(this_RH.data{1,1}.data==38);
this_RH.data{1,1}.data(net3) = 41;
unique(this_RH.data{1,1}.data)
net4 = find(this_RH.data{1,1}.data==39);
this_RH.data{1,1}.data(net4) = 42;
net5 = find(this_RH.data{1,1}.data==40);
this_RH.data{1,1}.data(net5) = 43;
unique(this_RH.data{1,1}.data)
net6 = find(this_RH.data{1,1}.data==41);
this_RH.data{1,1}.data(net6) = 44;
unique(this_RH.data{1,1}.data)
this_RH = gifti_read
('/Users/dianaperez/Box/Dependencies/32k_ConteAtlas_v2_distribute/RSN-networks.R.
32k_fs_LR.label.gii', this);
network_seg
atlas_dir = '/Users/dianaperez/Box/Dependencies/';
atlas = 'Seitzman300';
```

```
% load atlas that contains roi info (including which rois belong to each network)
atlas_params = atlas_parameters_GrattonLab(atlas,atlas_dir);
for n = 1:size(atlas_params.networks,1) % go through each network
    rois = atlas_params.mods{1,n}; %extract the rois belonging to system n
    num_rois = length(rois); % number of rois in system n
    net_size(n) = num_rois;
end
sum(net_size)
this.metadata = [];
this_RH = gifti_read✓
('/Users/dianaperez/Box/Dependencies/32k_ConteAtlas_v2_distribute/RSN-networks.R.✓
32k_fs_LR.label.gii', this);
addpath '/Users/dianaperez/Documents/GitHub/General_Scripts/gifti-master/@gifti/private'
this_RH = gifti_read✓
('/Users/dianaperez/Box/Dependencies/32k_ConteAtlas_v2_distribute/RSN-networks.R.✓
32k_fs_LR.label.gii', this);
addpath '/Users/dianaperez/Documents/GitHub/General_Scripts/gifti-master/@gifti/private'
this_RH = gifti_read✓
('/Users/dianaperez/Box/Dependencies/32k_ConteAtlas_v2_distribute/RSN-networks.R.✓
32k_fs_LR.label.gii', this);
cd '/Users/dianaperez/Documents/GitHub/General_Scripts/gifti-master/@gifti/private'
this_RH = gifti_read✓
('/Users/dianaperez/Box/Dependencies/32k_ConteAtlas_v2_distribute/RSN-networks.R.✓
32k_fs_LR.label.gii', this);
this.data = [];
this_RH = gifti_read✓
('/Users/dianaperez/Box/Dependencies/32k_ConteAtlas_v2_distribute/RSN-networks.R.✓
32k_fs_LR.label.gii', this);
this_LH = gifti_read✓
('/Users/dianaperez/Box/Dependencies/32k_ConteAtlas_v2_distribute/RSN-networks.R.✓
32k_fs_LR.label.gii', this);
cifti_w_nonbrain = [this_LH.data{1,1}.data; this_RH.data{1,1}.data];
template = ft_read_cifti_mod✓
('/Users/dianaperez/Box/Dependencies/Important_Files/120_templates.dtseries.nii');
cifti = cifti_w_nonbrain(template.brainstructure>0);
unique(this_LH.data{1,1}.data)
medial = find(this_LH.data{1,1}.data==35);
this_LH.data{1,1}.data(medial) = 0;
net1 = find(this_LH.data{1,1}.data==36);
this_LH.data{1,1}.data(net1) = 1;
net2 = find(this_LH.data{1,1}.data==37);
this_LH.data{1,1}.data(net2) = 2;
unique(this_LH.data{1,1}.data)
net3 = find(this_LH.data{1,1}.data==38);
this_LH.data{1,1}.data(net3) = 3;
net4 = find(this_LH.data{1,1}.data==39);
net5 = find(this_LH.data{1,1}.data==40);
net6 = find(this_LH.data{1,1}.data==41);
net7 = find(this_LH.data{1,1}.data==42);
this_LH.data{1,1}.data(net4) = 4;
this_LH.data{1,1}.data(net5) = 5;
this_LH.data{1,1}.data(net6) = 6;
this_LH.data{1,1}.data(net7) = 7;
unique(this_LH.data{1,1}.data)
unique(this_RH.data{1,1}.data)
medial = find(this_RH.data{1,1}.data==35);
net1 = find(this_RH.data{1,1}.data==36);
net2 = find(this_RH.data{1,1}.data==37);
net3 = find(this_RH.data{1,1}.data==38);
net4 = find(this_RH.data{1,1}.data==39);
```

```
net5 = find(this_RH.data{1,1}.data==40);
net6 = find(this_RH.data{1,1}.data==41);
net7 = find(this_RH.data{1,1}.data==42);
this_RH.data{1,1}.data(net7) = 7;
this_RH.data{1,1}.data(net6) = 6;
this_RH.data{1,1}.data(net5) = 5;
this_RH.data{1,1}.data(net4) = 4;
this_RH.data{1,1}.data(net3) = 3;
this_RH.data{1,1}.data(net2) = 2;
this_RH.data{1,1}.data(net1) = 1;
this_RH.data{1,1}.data(medial) = 0;
cifti_w_nonbrain = [this_LH.data{1,1}.data; this_RH.data{1,1}.data];
cifti = cifti_w_nonbrain(template.brainstructure>0);
template.data = cifti;
ft_write_cifti_mod('/Users/dianaperez/Desktop/Yeo_nets.dtseries.nii', template);
cifti_w_nonbrain = [this_LH.data{1,1}.data; this_RH.data{1,1}.data];
template.data = cifti_w_nonbrain;
ft_write_cifti_mod('/Users/dianaperez/Desktop/Yeo_nets_64k.dtseries.nii', template);
this_LH = gifti_read(
    ('/Users/dianaperez/Box/Dependencies/32k_ConteAtlas_v2_distribute/RSN-networks.L.
32k_fs_LR.label.gii', this);
unique(this_LH.data{1,1}.data)
medial = find(this_LH.data{1,1}.data==37);
this_LH.data{1,1}.data(medial) = 0;
net1 = find(this_LH.data{1,1}.data==38);
this_LH.data{1,1}.data(net1) = 1;
net2 = find(this_LH.data{1,1}.data==39);
this_LH.data{1,1}.data(net2) = 2;
net3 = find(this_LH.data{1,1}.data==40);
this_LH.data{1,1}.data(net3) = 3;
net4 = find(this_LH.data{1,1}.data==41);
net5 = find(this_LH.data{1,1}.data==42);
net6 = find(this_LH.data{1,1}.data==43);
net7 = find(this_LH.data{1,1}.data==44);
this_LH.data{1,1}.data(net4) = 4;
this_LH.data{1,1}.data(net5) = 5;
this_LH.data{1,1}.data(net6) = 6;
this_LH.data{1,1}.data(net7) = 7;
unique(this_LH.data{1,1}.data)
cifti_w_nonbrain = [this_LH.data{1,1}.data; this_RH.data{1,1}.data];
cifti = cifti_w_nonbrain(template.brainstructure>0);
template.data = cifti;
ft_write_cifti_mod('/Users/dianaperez/Desktop/Yeo_nets.dtseries.nii', template);
unique(this_LH.data{1,2}.data)
og_ids = unique(this_LH.data{1,2}.data);
new_ids = 1:1:length(og_ids);
new_ids = 0:1:length(og_ids);
new_ids = 0:1:length(og_ids)-1;
this_LH = gifti_read(
    ('/Users/dianaperez/Box/Dependencies/32k_ConteAtlas_v2_distribute/RSN-networks.R.
32k_fs_LR.label.gii', this);
gifti_LH = this_LH.data{1,2}.data;
og_ids = unique(gifti_LH);
new_ids = 0:1:length(og_ids)-1;
for i = 1:length(og_ids)
    verts = find(gifti_LH==og_ids(i));
    gifti_LH(verts) = new_ids(i);
end
unique(gifti_LH)
this_RH = gifti_read
```

```
( '/Users/dianaperez/Box/Dependencies/32k_ConteAtlas_v2_distribute/RSN-networks.R. ↵
32k_fs_LR.label.gii', this);
gifti_RH = this_RH.data{1,2}.data;
og_ids = unique(gifti_RH);
new_ids = 0:1:length(og_ids)-1;
for i = 1:length(og_ids)
verts = find(gifti_RH==og_ids(i));
gifti_RH(verts) = new_ids(i);
end
unique(gifti_RH)
template = ft_read_cifti_mod↵
('/Users/dianaperez/Box/Dependencies/Important_Files/120_templates.dtseries.nii');
cifti_w_nonbrain = [gifti_LH; gifti_RH];
cifti = cifti_w_nonbrain(template.brainstructure>0);
template.data = cifti;
ft_write_cifti_mod('/Users/dianaperez/Desktop/Yeo_17nets.dtseries.nii', template);
this_LH = gifti_read↵
('/Users/dianaperez/Box/Dependencies/32k_ConteAtlas_v2_distribute/RSN-networks.L. ↵
32k_fs_LR.label.gii', this);
gifti_LH = this_LH.data{1,2}.data;
og_ids = unique(gifti_LH);
new_ids = 0:1:length(og_ids)-1;
for i = 1:length(og_ids)
verts = find(gifti_LH==og_ids(i));
gifti_LH(verts) = new_ids(i);
end
this_RH = gifti_read↵
('/Users/dianaperez/Box/Dependencies/32k_ConteAtlas_v2_distribute/RSN-networks.R. ↵
32k_fs_LR.label.gii', this);
gifti_RH = this_RH.data{1,2}.data;
og_ids = unique(gifti_RH);
new_ids = 0:1:length(og_ids)-1;
for i = 1:length(og_ids)
verts = find(gifti_RH==og_ids(i));
gifti_RH(verts) = new_ids(i);
end
template = ft_read_cifti_mod↵
('/Users/dianaperez/Box/Dependencies/Important_Files/120_templates.dtseries.nii');
cifti_w_nonbrain = [gifti_LH; gifti_RH];
cifti = cifti_w_nonbrain(template.brainstructure>0);
template.data = cifti;
ft_write_cifti_mod('/Users/dianaperez/Desktop/Yeo_17nets.dtseries.nii', template);
this_LH = gifti_read↵
('/Users/dianaperez/Box/Dependencies/32k_ConteAtlas_v2_distribute/RSN-networks.L. ↵
32k_fs_LR.label.gii', this);
gifti_LH = this_LH.data{1,2}.data;
og_ids = unique(gifti_LH);
new_ids = 0:1:length(og_ids)-1;
for i = 1:length(og_ids)
verts = find(gifti_LH==og_ids(i));
numverts(i,1) = length(verts);
gifti_LH(verts) = new_ids(i);
end
this_RH = gifti_read↵
('/Users/dianaperez/Box/Dependencies/32k_ConteAtlas_v2_distribute/RSN-networks.R. ↵
32k_fs_LR.label.gii', this);
gifti_RH = this_RH.data{1,2}.data;
og_ids = unique(gifti_RH);
new_ids = 0:1:length(og_ids)-1;
for i = 1:length(og_ids)
```

```

verts = find(gifti_RH==og_ids(i));
numverts(i,2) = length(verts);
gifti_RH(verts) = new_ids(i);
end
clear
load('/Users/dianaperez/Desktop/Thesis/Variant_Properties/HCP384_new_split_networksxHem.
mat')
for net = 1:16
se_lh(net) = std(networksxHem.clustersLH(:,net))/sqrt(384);
se_rh(net) = std(networksxHem.clustersRH(:,net))/sqrt(384);
end
network_names = {'DMN' 'Vis' 'FP' 'Reward' 'DAN' 'Unassigned' 'Lang' 'Sal' 'CO'
'SMd' 'SMl' 'Aud' 'Tpole' 'MTL' 'PMN' 'PON'};
rgb_colors = [1 0 0; %DMN
0 0 .6; %Vis
1 1 0; %FP
.67 .67 .67; %Unassigned
0 .8 0; %DAN
.67 .67 .67; %Unassigned2
0 .6 .6; %VAN
0 0 0; % Sal
.3 0 .6; %CON
.2 1 1; %SMd
1 .5 0; % SMl
.6 .2 1; %Aud
.2 1 .2; %Tpole
0 .2 .4; %MTL
0 0 1; %PMN
.8 .8 .6]; %PON
lh_numvars = mean(networksxHem.clustersLH);
rh_numvars = mean(networksxHem.clustersRH);
errorbar(lh_numvars, rh_numvars, se_rh, se_rh, se_lh, se_lh, 'o')
hold on
scatter(lh_numvars, rh_numvars, 80, rgb_colors, 'filled')
axis([0 2.5 0 2.5])
axis([0 2.5 0 2.5])
line = refline(1,0);
line.Color = 'black';
xlabel('Left Hemisphere')
ylabel('Right Hemisphere')
title('Average number of variants')
ax = gca;
ax.FontSize = 24;
print(gcf, '/Users/dianaperez/Desktop/Thesis/Figures/HCP_numvars_scatter.jpg', '-dpng',
'-r300')
%-- 11/19/21, 3:36 PM --%
load(goodSubs384.mat)
load('goodSubs384.mat')
cd '/Users/dianaperez/Documents/GitHub/Lateralization_Variants'
load('goodSubs384.mat')
load
('/Users/dianaperez/Desktop/Thesis/Variant_Properties/HCP384_new_split_variants_info_wSur
fArea.mat')
numVars = variants_info.both_hems{1,1};
numVars1 = variants_info.both_hems{: ,1};
handedness = goodSubs3841;
numVars = numVars1;
clear numVars1
clear goodSubs3841
corr(goodSubs384, handedness)

```

```

corrcoef(handedness, numVars)
[r, p] = corrcoef(handedness, numVars)
avgVarSize = variants_info.both_hems(:,2);
[r, p] = corrcoef(handedness, avgVarSize)
avgVarSize = variants_info.both_hems(:,3);
[r, p] = corrcoef(handedness, avgVarSize)
numborder = variants_info.both_hems(:,4);
[r, p] = corrcoef(handedness, numborder)
numectopic = variants_info.both_hems(:,5);
[r, p] = corrcoef(handedness, numectopic)
numVars = variants_info.left_hem.group_avg(:,1);
[r, p] = corrcoef(handedness, numVars)
numVerts = variants_info.left_hem.group_avg(:,2);
[r, p] = corrcoef(handedness, numVerts)
x = variants_info.left_hem.group_avg(:,3);
[r, p] = corrcoef(handedness, x)
x = variants_info.left_hem.group_avg(:,4);
[r, p] = corrcoef(handedness, x)
x = variants_info.left_hem.group_avg(:,5);
[r, p] = corrcoef(handedness, x)
x = variants_info.left_hem.group_avg(:,7);
[r, p] = corrcoef(handedness, x)
x = variants_info.right_hem.group_avg(:,7);
[r, p] = corrcoef(handedness, x)
x = variants_info.right_hem.group_avg(:,5);
[r, p] = corrcoef(handedness, x)
x = variants_info.right_hem.group_avg(:,4);
[r, p] = corrcoef(handedness, x)
x = variants_info.right_hem.group_avg(:,3);
[r, p] = corrcoef(handedness, x)
x = variants_info.right_hem.group_avg(:,2);
[r, p] = corrcoef(handedness, x)
x = variants_info.right_hem.group_avg(:,1);
[r, p] = corrcoef(handedness, x)
allSubs = table2cell(readtable('/Users/dianaperez/Desktop/Thesis/HCP_lang_measures.
xlsx'));
for x = 1:size(allSubs,1)
sub = cell2mat(allSubs(x,1));
if ismember(sub, goodSubs_col)
goodSubs_all(count,:) = allSubs(x,:);
count = count + 1;
end
end
for x = 1:size(allSubs,1)
sub = cell2mat(allSubs(x,1));
if ismember(sub, goodSubs384)
goodSubs_all(count,:) = allSubs(x,:);
count = count + 1;
end
end
count = 1;
for x = 1:size(allSubs,1)
sub = cell2mat(allSubs(x,1));
if ismember(sub, goodSubs384)
goodSubs_all(count,:) = allSubs(x,:);
count = count + 1;
end
end
HCP_384_goodsubs = goodSubs384;
find(goodSubs_all(:,1)==HCP_384_goodsubs(1,1))

```

```
find(goodSubs_all(:,1)=HCP_384_goodsups(1,1))
find(goodSubs_all(:,1)==HCP_384_goodsups(1,1))
HCP_384_goodsups(:,3) = variants_info.left_hem.group_avg{:,1}
HCP_384_goodsups(:,4) = variants_info.left_hem.group_avg{:,2};
HCP_384_goodsups(:,5) = variants_info.left_hem.group_avg{:,3};
HCP_384_goodsups(:,6) = variants_info.left_hem.group_avg{:,4};
HCP_384_goodsups(:,7) = variants_info.left_hem.group_avg{:,5};
HCP_384_goodsups(:,8) = variants_info.left_hem.group_avg{:,6};
HCP_384_goodsups(:,9) = variants_info.left_hem.group_avg{:,7};
HCP_384_goodsups(:,10) = variants_info.left_hem.group_avg{:,9};
HCP_384_goodsups(:,10) = variants_info.left_hem.group_avg{:,8};
HCP_384_goodsups(:,11) = variants_info.right_hem.group_avg{:,1}
HCP_384_goodsups(:,12) = variants_info.right_hem.group_avg{:,2};
HCP_384_goodsups(:,13) = variants_info.right_hem.group_avg{:,3};
HCP_384_goodsups(:,14) = variants_info.right_hem.group_avg{:,4};
HCP_384_goodsups(:,15) = variants_info.right_hem.group_avg{:,5};
HCP_384_goodsups(:,16) = variants_info.right_hem.group_avg{:,6};
HCP_384_goodsups(:,17) = variants_info.right_hem.group_avg{:,7};
HCP_384_goodsups(:,18) = variants_info.right_hem.group_avg{:,8};
HCP_384_goodsups(:,19) = variants_info.both_hems{:,1};
HCP_384_goodsups(:,19) = variants_info.both_hems{:,2};
HCP_384_goodsups(:,19) = variants_info.both_hems{:,1};
HCP_384_goodsups(:,20) = variants_info.both_hems{:,2};
HCP_384_goodsups(:,21) = variants_info.both_hems{:,3};
HCP_384_goodsups(:,22) = variants_info.both_hems{:,4};
HCP_384_goodsups(:,23) = variants_info.both_hems{:,5};
test = HCP_384_goodsups;
[B, I] = sort(test(:,1), 'ascending')
[B, I] = sort(test(:,1), 'ascend')
[B, I] = sort(test(:,1), 'ascend');
HCP_sorted = test(I);
HCP_sorted = test(I,:);
HCP_sorted{:,24} = goodSubs_all{:,2};
HCP_sorted{:,24} = goodSubs_all1;
HCP_sorted{:,24} = goodSubs_all1{:,};
HCP_sorted{:,24} = cell2mat(goodSubs_all{:,2});
testing = cell2mat(goodSubs_all)
HCP_sorted{:,24} = testing(:,2);
HCP_sorted(:,24) = testing(:,2);
HCP_sorted(:,25) = testing(:,3);
HCP_sorted(:,26) = testing(:,4);
HCP_sorted(:,27) = testing(:,5);
HCP_sorted(:,28) = testing(:,6);
HCP_sorted(:,29) = testing(:,7);
[r, p] = corrcoef(HCP_sorted(:,3), HCP_sorted(:,29))
find(isnan(HCP_sorted))
[r, p] = corrcoef(HCP_sorted(:,3), HCP_sorted(:,29), 'rows', 'complete')
[r, p] = corrcoef(HCP_sorted(:,3), HCP_sorted(:,28), 'rows', 'complete')
[r, p] = corrcoef(HCP_sorted(:,3), HCP_sorted(:,27), 'rows', 'complete')
[r, p] = corrcoef(HCP_sorted(:,3), HCP_sorted(:,26), 'rows', 'complete')
[r, p] = corrcoef(HCP_sorted(:,3), HCP_sorted(:,25), 'rows', 'complete')
[r, p] = corrcoef(HCP_sorted(:,3), HCP_sorted(:,24), 'rows', 'complete')
[r, p] = corrcoef(HCP_sorted(:,4), HCP_sorted(:,24), 'rows', 'complete')
[r, p] = corrcoef(HCP_sorted(:,4), HCP_sorted(:,25), 'rows', 'complete')
[r, p] = corrcoef(HCP_sorted(:,4), HCP_sorted(:,26), 'rows', 'complete')
[r, p] = corrcoef(HCP_sorted(:,4), HCP_sorted(:,27), 'rows', 'complete')
[r, p] = corrcoef(HCP_sorted(:,4), HCP_sorted(:,28), 'rows', 'complete')
[r, p] = corrcoef(HCP_sorted(:,4), HCP_sorted(:,29), 'rows', 'complete')
[r, p] = corrcoef(HCP_sorted(:,5), HCP_sorted(:,29), 'rows', 'complete')
[r, p] = corrcoef(HCP_sorted(:,5), HCP_sorted(:,28), 'rows', 'complete')
```


[illegible]


```
[r, p] = corrcoef(HCP_sorted(:,14), HCP_sorted(:,28), 'rows', 'complete')
[r, p] = corrcoef(HCP_sorted(:,15), HCP_sorted(:,28), 'rows', 'complete')
[r, p] = corrcoef(HCP_sorted(:,16), HCP_sorted(:,28), 'rows', 'complete')
[r, p] = corrcoef(HCP_sorted(:,17), HCP_sorted(:,28), 'rows', 'complete')
[r, p] = corrcoef(HCP_sorted(:,18), HCP_sorted(:,28), 'rows', 'complete')
[r, p] = corrcoef(HCP_sorted(:,19), HCP_sorted(:,28), 'rows', 'complete')
[r, p] = corrcoef(HCP_sorted(:,20), HCP_sorted(:,28), 'rows', 'complete')
[r, p] = corrcoef(HCP_sorted(:,21), HCP_sorted(:,28), 'rows', 'complete')
[r, p] = corrcoef(HCP_sorted(:,22), HCP_sorted(:,28), 'rows', 'complete')
[r, p] = corrcoef(HCP_sorted(:,23), HCP_sorted(:,28), 'rows', 'complete')
plot(HCP_sorted(:,23), HCP_sorted(:,variant_info.both_hems{:,1}))
plot(HCP_sorted(:,23), HCP_sorted(:,variants_info.both_hems{:,1}))
plot(HCP_sorted(:,2), HCP_sorted(:,variants_info.both_hems{:,1}))
numVars1 = variants_info.both_hems{:,1};
plot(HCP_sorted(:,2), variants_info.both_hems{:,1})
plot(HCP_sorted(:,2), variants_info.both_hems{:,1})
scatter(HCP_sorted(:,2), variants_info.both_hems{:,1})
diff_numvars = HCP_sorted(:,3) - HCP_sorted(:,11);
[r, p]=corrcoef(diff_numvars, HCP_sorted(:,24));
[r, p]=corrcoef(diff_numvars, HCP_sorted(:,25));
[r, p]=corrcoef(diff_numvars, HCP_sorted(:,26))
[r, p]=corrcoef(diff_numvars, HCP_sorted(:,26), 'rows', 'complete')
[r, p]=corrcoef(diff_numvars, HCP_sorted(:,27), 'rows', 'complete')
[r, p]=corrcoef(diff_numvars, HCP_sorted(:,28), 'rows', 'complete')
[r, p]=corrcoef(diff_numvars, HCP_sorted(:,29), 'rows', 'complete')
[r, p]=corrcoef(diff_numvars, HCP_sorted(:,24), 'rows', 'complete')
[r, p]=corrcoef(diff_numvars, HCP_sorted(:,25), 'rows', 'complete')
diff_numvars = HCP_sorted(:,4) - HCP_sorted(:,12);
[r, p]=corrcoef(diff_numvars, HCP_sorted(:,24), 'rows', 'complete')
[r, p]=corrcoef(diff_numvars, HCP_sorted(:,25), 'rows', 'complete')
[r, p]=corrcoef(diff_numvars, HCP_sorted(:,26), 'rows', 'complete')
[r, p]=corrcoef(diff_numvars, HCP_sorted(:,27), 'rows', 'complete')
[r, p]=corrcoef(diff_numvars, HCP_sorted(:,28), 'rows', 'complete')
[r, p]=corrcoef(diff_numvars, HCP_sorted(:,29), 'rows', 'complete')
cd /Volumes/Research_HD
cd Lifespan_taskfMRI/
clear all
cd ~/Desktop
DownloadPsychtoolbox
DownloadPsychtoolbox.m
cd ~/Downloads
DownloadPsychtoolbox
%-- 11/22/21, 1:10 PM --%
DownloadPsychtoolbox
yes
cd /Volumes/Research_HD
%-- 11/22/21, 8:44 PM --%
cd /Volumes/Research_HD
%-- 11/24/21, 9:34 AM --%
cd /Volumes/RESEARCH_HD/
mixed_short
test
0
mixed
test
0
sca
help HideCursor
ShowCursor()
mixed
```

```
test
0
sca
mixed
test
0
mixed
test
0
sca
%-- 11/24/21, 10:41 AM --%
mixed
test
0
%-- 11/24/21, 10:42 AM --%
mixed
test
0
%-- 11/24/21, 11:03 AM --%
mixed
test
0
mixed
test
0
mixed
test
0
mixed
test
0
mixed
test
1
%-- 11/24/21, 12:13 PM --%
group_avg = ft_read_cifti_mod('/Users/dianaperez/Desktop/Yeo_7nets.dtseries.nii');
left_hem = group_avg.data(1:29696);
right_hem = group_avg.data(29697:end);
surf_areas = ft_read_cifti_mod('/Users/dianaperez/Desktop/Thesis/surf_areas_verts.↵
dtseries.nii');
lh_surf = surf_areas.data(1:29696);
rh_surf = surf_areas.data(29697:end);
num_nets = unique(group_avg.data);
net_size = [];
for net = 1:num_nets
    lh_verts = find(left_hem==num_nets(net));
    rh_verts = find(right_hem==num_nets(net));
    net_size(net,1) = length(lh_verts);
    net_size(net,2) = length(rh_verts);
    net_size(net,3) = sum(lh_surf(lh_verts));
    net_size(net,4) = sum(rh_surf(rh_verts));
end
net_size_across_hems
%-- 11/24/21, 2:20 PM --%
mixed
test
1
y
sca
mixed
```

```

test
1
y
sca
%-- 11/29/21, 3:13 PM --%
net_size_across_hems
rgb_colors = [1 0 0;    %DMN
0 0 .6; %Vis
1 1 0; %FP
0 .8 0; %DAN
0 .6 .6;%VAN
0 0 0;%SAl
.3 0 .6;%CON
.2 1 1;%SMd
1 .5 0;%SML
.6 .2 1;%Aud
.2 1 .2;%Tpole
0 .2 .4;%MTL
0 0 1; %PMN
.8 .8 .6]; %PON
scatter(net_size(:,4), net_size(:,8), 80, rgb_colors, 'filled')
rgb_colors = [1 0 0; %DMN
0 0 .6; %Vis
1 1 0; %FP
.67 .67 .67; %Unassigned
0 .8 0; %DAN
.67 .67 .67; %Unassigned2
0 .6 .6;%VAN
0 0 0; % Sal
.3 0 .6; %CON
.2 1 1; %SMd
1 .5 0; % SML
.6 .2 1; %Aud
.2 1 .2; %Tpole
0 .2 .4; %MTL
0 0 1; %PMN
.8 .8 .6]; %PON
this.data = [];
this.label = [];
this.metadata = [];
uiopen('/Users/dianaperez/Box/Dependencies/32k_ConteAtlas_v2_distribute/RSN-networks.L.32k_fs_LR.label.gii',1)
Yeo_l = gifti_read('/Users/dianaperez/Box/Dependencies/32k_ConteAtlas_v2_distribute/RSN-networks.L.32k_fs_LR.label.gii', this);
rgb_medialwall = Yeo_l.label.rgba(38,1:3);
rgb_yeo_7nets = [rgb_medialwall; Yeo_l.label.rgba(39:45,1:3)];
rgb_yeo_17nets = [rgb_medialwall; Yeo_l.label.rgba(46:62,1:3)];
rgb_yeo_17nets
scatter(net_size(:,4), net_size(:,8), 80, rgb_yeo_17nets, 'filled')
axis([0 .14 0 .14])
line = reffline(1,0);
line.Color = 'black';
xlabel('Left Hemisphere')
ylabel('Right Hemisphere')
ax = gca;
ax.FontSize = 24;
print(gcf, '/Users/dianaperez/Desktop/Thesis/Figures/Yeo17nets_groupavg_surfarea_scatter.jpg', '-dpng', '-r300')
group_avg = ft_read_cifti_mod('/Users/dianaperez/Desktop/Yeo_7nets.dtseries.nii');
left_hem = group_avg.data(1:29696);

```

```

right_hem = group_avg.data(29697:end);
%load surf area file
surf_areas = ft_read_cifti_mod('/Users/dianaperez/Desktop/Thesis/surf_areas_verts.
dtseries.nii');
lh_surf = surf_areas.data(1:29696);
rh_surf = surf_areas.data(29697:end);
num_nets = unique(group_avg.data);
net_size = [];
for net = 1:length(num_nets)
lh_verts = find(left_hem==num_nets(net));
rh_verts = find(right_hem==num_nets(net));
net_size(net,1) = length(lh_verts);
net_size(net,2) = length(lh_verts)/length(left_hem);
net_size(net,3) = sum(lh_surf(lh_verts));
net_size(net,4) = sum(lh_surf(lh_verts))/sum(lh_surf);
net_size(net,5) = length(rh_verts);
net_size(net,6) = length(rh_verts)/length(right_hem);
net_size(net,7) = sum(rh_surf(rh_verts));
net_size(net,8) = sum(rh_surf(rh_verts))/sum(rh_surf);
end
scatter(net_size(:,4), net_size(:,8), 80, rgb_yeo_7nets, 'filled')
close(gcf)
scatter(net_size(:,4), net_size(:,8), 80, rgb_yeo_7nets, 'filled')
axis([0 .25 0 .25])
line = reffline(1,0);
line.Color = 'black';
xlabel('Left Hemisphere')
ylabel('Right Hemisphere')
ax = gca;
ax.FontSize = 24;
print(gcf, '/Users/dianaperez/Desktop/Thesis/Figures/Yeo7nets_groupavg_surfareas_scatter.
jpg', '-dpng', '-r300')
network_names_yeo17 = {'Medial Wall', 'Visual Central', 'DMN A', 'DAN B', 'Control A',
'Limbic A', 'DAN A', 'Control B', 'DMN B', 'DMN C', 'Visual Peripheral', 'Somatomotor B',
'Limbic B', 'Control C', 'Somatomotor A', 'Salience', 'VAN', 'DMN D (Aud)'};
network_names_yeo7 = {'Medial Wall', 'DAN', 'FPN', 'DMN', 'Visual', 'Limbic',
'Somatomotor', 'VAN'};
pie(net_size(:,4), network_names_yeo7);
close(gcf)
pie(net_size(:,4), network_names_yeo7);
patchHandle = findobj(h, 'Type', 'Patch');
set(patchHandle, {'FaceColor'}, mat2cell(rgb_yeo_7nets, ones(size(rgb_colors,1),1),3))
set(patchHandle, {'FaceColor'}, mat2cell(rgb_yeo_7nets, ones(size(rgb_yeo_7nets,1),1),3))
h = pie(net_size(:,4), network_names_yeo7);
patchHandle = findobj(h, 'Type', 'Patch');
set(patchHandle, {'FaceColor'}, mat2cell(rgb_yeo_7nets, ones(size(rgb_yeo_7nets,1),1),3))
lgd = legend(network_names_yeo7);
lgd.Layout.Tile = 'east';
lgd.FontSize = 24;
net_size_across_hems
h = pie(net_size(:,4), network_names_yeo17);
close(gcf)
h = pie(net_size(:,4), network_names_yeo17);
patchHandle = findobj(h, 'Type', 'Patch');
set(patchHandle, {'FaceColor'}, mat2cell(rgb_yeo_17nets, ones(size(rgb_yeo_17nets,1),1),
3))
lgd = legend(network_names_yeo17);
lgd.FontSize = 24;
network_names_yeo7
clear all

```

```
rgb_colors = [1 0 0;    %DMN
0 0 .6; %Vis
1 1 0; %FP
0 .8 0; %DAN
0 .6 .6; %VAN
.3 0 .6; %CON
.2 1 1; %SMd
.6 .2 1; %Aud
.8 .8 .6]; %PON
output_dir = '/Users/dianaperez/Desktop/Thesis/Spatial_Location/';
load([output_dir 'operc_nets_lefthemcluster_righthem.mat']);
leftclust_righthem = LH_cluster;
load([output_dir 'operc_nets_lefthemcluster_lefthem.mat']);
leftclust_lefthem = LH_cluster;
clear LH_cluster
%-- 12/2/21, 12:18 PM --%
rgb_colors = [1 0 0;
1 1 0;
0 .8 0;
0 .6 .6;
0 0 0;
.3 0 .6;
1 .5 0;
.8 .8 .6];
output_dir = '/Users/dianaperez/Desktop/Thesis/Spatial_Location/';
load([output_dir 'operc_nets_righthemcluster_righthem.mat']);
rightclust_righthem = RH_cluster;
load([output_dir 'operc_nets_righthemcluster_lefthem.mat']);
rightclust_lefthem = RH_cluster;
clear RH_cluster
network_names = {'DMN' 'FP' 'DAN' 'Lang' 'Sal' 'CO' 'SML' 'PON'};
for net = 1:16
nets_left(net) = length(find(rightclust_lefthem.vars(:,3)==net));
end
nets_left(12:15) = [];
nets_left(10) = [];
nets_left(6) = [];
nets_left(4) = [];
nets_left(2) = [];
means = [];
network_names = {'DMN' 'FP' 'DAN' 'Lang' 'Sal' 'CO' 'SML' 'PON'};
for net = 1:16
numVars = find(rightclust_lefthem.vars(:,3)==net);
nets_left(net) = mean(numVars);
end
nets_left(12:15) = [];
nets_left(10) = [];
nets_left(6) = [];
nets_left(4) = [];
nets_left(2) = [];
for net = 1:16
numVars = find(rightclust_righthem.vars(:,3)==net);
nets_right(net) = mean(numVars);
end
nets_right(12:15) = [];
nets_right(10) = [];
nets_right(6) = [];
nets_right(4) = [];
nets_right(2) = [];
numVars = find(rightclust_lefthem.vars(:,3)==net);
```

```

numVars = find(rightclust_lefthem.vars(:,3)==1);
%-- 12/17/21, 3:27 PM --%
cd /Volumes
cd RESEARCH_HD/
cd HCP_Variants/
cd new_split_vars/
var_freq_size_net
clear surf_areas
var_freq_size_net
subs{1}
var_freq_size_net
clear all
var_freq_size_net
subs{1,1}
var_freq_size_net
subs{1,1}{1}
subs{1,1}{1,1}
subs{1,1}(1)
var_freq_size_net
[data_location num2str(subs{g,1}(n)) bordEctop_str]
var_freq_size_net
numel(subs)
length(subs{1})
length(subs{1})
var_freq_size_net
clear all
load
('/Users/dianaperez/Desktop/lateralization_code/testing_output/HCP384_new_split_networksx
Hem.mat')
load
('/Users/dianaperez/Desktop/lateralization_code/testing_output/HCP384_new_split_variants_
info.mat')
var_freq_size_net
load
('/Users/dianaperez/Desktop/lateralization_code/testing_output/HCP752_new_split_RH_matche
d_variants_info.mat')
clear all
load
('/Users/dianaperez/Desktop/lateralization_code/testing_output/HCP752_new_split_RH_matche
d_variants_info.mat')
%-- 12/20/21, 5:39 PM --%
var_freq_size_net
floor(1:9/2)
%-- 12/21/21, 11:15 AM --%
in_str = 'HCP384';
out_str = append(in_str, '_LHandvsRHand_');
mixed
test
1
mixed
test
1
cd /Volumes/RESEARCH_HD/Lifespan_taskfMRI/
mixed
test
1
y
sca
mixed
test

```

```

1
y
sca
mixed
%-- 12/21/21, 11:32 AM --%
mixed
test
1
y
%-- 12/21/21, 12:58 PM --%
perms_var_freq_size
var_freq_size_net
perms_var_freq_size
var_freq_size_net
perms_var_freq_size
length(allSubs_info.variants_info.left_hem.group_avg)
height(allSubs_info.variants_info.left_hem.group_avg)
var_freq_size_net
perms_var_freq_size
var_freq_size_net
perms_var_freq_size
length(find(perm_diffs<true_diff))/numperms
length(find(perm_diffs>true_diff))/numperms
perms_var_freq_size
mixed
test
1
y
%-- 12/21/21, 2:28 PM --%
perms_var_freq_size
plotSpread
perms_var_freq_size
handles = plotSpread(perm_diffs, 'categoryMarkers', {'o', '.'}, 'categoryLabels', \
{'Permuted Difference', 'True Difference'}, 'categoryColors', {'k', 'r'}, 'xValues', [1])
handles = plotSpread(perm_diffs, 'categoryMarkers', 'o', 'categoryLabels', 'Permuted \
Difference', 'categoryColors', 'k', 'xValues', [1])
handles = plotSpread(perm_diffs, 'categoryMarkers', 'o', 'categoryLabels', 'Permuted \
Difference', 'categoryColors', 'k')
perms_var_freq_size
handles = plotSpread(perm_diffs, 'categoryMarkers', 'o', 'categoryLabels', 'Permuted \
Difference', 'categoryColors', 'k')
perms_var_freq_size
axis([0.5, 1.5, min(perm_diffs)-5, max(perm_diffs)+5])
ax.FontSize = 20;
ax.FontSize = 12;
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.3, 0.7]); %first and second \
control position on screen, third controls width, and fourth controls height
print(gcf, outfile, '-dpng', '-r300');
perms_var_freq_size
LHand_info.variants_info.both_hems{: ,4}
var_freq_size_net
subs{g,1}
subs{g,1}(n)
subs{g,1}{n}
var_freq_size_net
perms_var_freq_size
[p_fdr p_masked] = FDR(p_vals, .025);
perms_var_freq_size
permute_net_assignment
save([output_dir 'network_perms_numVars.mat'], 'output_mat');

```

```

rgb_colors = [1 0 0; %DMN
0 0 .6; %Vis
1 1 0; %FP
.67 .67 .67; %Unassigned
0 .8 0; %DAN
.67 .67 .67; %Unassigned2
0 .6 .6; %VAN
0 0 0; % Sal
.3 0 .6; %CON
.2 1 1; %SMd
1 .5 0; % SMI
.6 .2 1; %Aud
.2 1 .2; %Tpole
0 .2 .4; %MTL
0 0 1; %PMN
.8 .8 .6]; %PON
good_nets = [output_mat(:,1:3) output_mat(:,5) output_mat(:,7:end)];
rgb_for_plot = {};
for n = 1:numel(network_names)
    rgb_for_plot{n} = rgb_colors(n,:);
end
ind = zeros(1001,1);
ind(1,1) = 1;
net_inds = [ind ind ind ind ind ind ind ind ind ind ind ind ind ind];
handles = plotSpread(good_nets, 'categoryMarkers', {'x', '.'}, 'categoryLabels', ↵
{'Permuted Differences', 'True Difference'}, 'distributionColors', rgb_for_plot', ↵
'categoryIdx', net_inds, 'xNames', network_names)
ax = gca;
ax.FontSize = 24;
scatter([1:14], good_nets(1,:), 'MarkerEdgeColor', 'k', 'MarkerFaceColor', 'k', ↵
'SizeData', 50)
scatter(6, good_nets(1,6), 'MarkerEdgeColor', 'w', 'MarkerFaceColor', 'w', 'SizeData', 50)
ylabel('Difference in Number of Variants')
xlabel('Assigned Network')
axis([0, 15, -0.4, .7])
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.9, 0.7]);
%-- 12/21/21, 9:54 PM --%
permute_net_assignment
\rgb_colors = [1 0 0; %DMN
0 0 .6; %Vis
1 1 0; %FP
.67 .67 .67; %Unassigned
0 .8 0; %DAN
.67 .67 .67; %Unassigned2
0 .6 .6; %VAN
0 0 0; % Sal
.3 0 .6; %CON
.2 1 1; %SMd
1 .5 0; % SMI
.6 .2 1; %Aud
.2 1 .2; %Tpole
0 .2 .4; %MTL
0 0 1; %PMN
.8 .8 .6]; %PON
good_nets = [output_mat(:,1:3) output_mat(:,5) output_mat(:,7:end)];
rgb_good_nets = [rgb_color(:,1:3); rgb_color(:,5); rgb_color(:,7:end)];
rgb_for_plot = {};
for n = 1:14
    rgb_for_plot{n} = rgb_good_nets(n,:);
end

```



```

rgb_colors = [1 0 0; %DMN
0 0 .6; %Vis
1 1 0; %FP
.67 .67 .67; %Unassigned
0 .8 0; %DAN
.67 .67 .67; %Unassigned2
0 .6 .6; %VAN
0 0 0; % Sal
.3 0 .6; %CON
.2 1 1; %SMd
1 .5 0; % SML
.6 .2 1; %Aud
.2 1 .2; %Tpole
0 .2 .4; %MTL
0 0 1; %PMN
.8 .8 .6]; %PON
good_nets = [output_mat(:,1:3) output_mat(:,5) output_mat(:,7:end)];
rgb_good_nets = [rgb_color(:,1:3); rgb_color(:,5); rgb_color(:,7:end)];
rgb_for_plot = {};
for n = 1:14
    rgb_for_plot{n} = rgb_good_nets(n,:);
end
rgb_good_nets = [rgb_colors(:,1:3); rgb_colors(:,5); rgb_colors(:,7:end)];
rgb_good_nets = [rgb_colors(1:3,:); rgb_colors(5,:); rgb_colors(7:end,:)];
rgb_for_plot = {};
for n = 1:14
    rgb_for_plot{n} = rgb_good_nets(n,:);
end
ind = zeros(1001,1);
ind(1,1) = 1;
net_inds = [ind ind ind ind ind ind ind ind ind ind ind ind ind ind];
handles = plotSpread(good_nets, 'categoryMarkers', {'x', '.'}, 'categoryLabels', ↵
{'Permuted Differences', 'True Difference'}, 'distributionColors', rgb_for_plot', ↵
'categoryIdx', net_inds, 'xNames', network_names)
ax = gca;
ax.FontSize = 24;
scatter([1:14], good_nets(1,:), 'MarkerEdgeColor', 'k', 'MarkerFaceColor', 'k', ↵
'SizeData', 50)
scatter(6, good_nets(1,6), 'MarkerEdgeColor', 'w', 'MarkerFaceColor', 'w', 'SizeData', 50)
ylabel('Difference in Number of Variants')
xlabel('Assigned Network')
axis([0, 15, -0.4, .7])
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.9, 0.7]);
print(gcf, [output_dir out_str '_PermutationTesting_networkAssignment.jpg'], '-dpng', '-↵
r300');
allSubs_info = load([data_location in_str '_variants_info.mat']);
width(allSubs_info.variants_info.left_hem.group_avg)
help stdev
var_freq_size_net
min([leftHem; rightHem]) - std([leftHem; rightHem])
max([leftHem; rightHem]) + std([leftHem; rightHem])
axis([-0.2, 0.5, min([leftHem; rightHem]) - std([leftHem; rightHem]), max([leftHem; ↵
rightHem]) + std([leftHem; rightHem])])
var_freq_size_net
leftHem = variants_info.left_hem.group_avg{: , 1};
rightHem = variants_info.right_hem.group_avg{: , 1};
[h,mu,sigma,q,notch] = al_goodplot(leftHem, 0.1, .2, [ 0 0.4470 0.7410], 'left');
[h,mu,sigma,q,notch] = al_goodplot(rightHem, 0.11, .2, [0.8500 0.3250 ↵
0.0980], 'right');
xticks([0.105])

```

```
xticklabels('Number of Variant Regions')
axis([-0.2, 0.5, min([leftHem; rightHem])-std([leftHem; rightHem]), max([leftHem;
rightHem])+std([leftHem; rightHem])])
ax = gca;
ax.FontSize = 24;
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.3, 0.9]); %first and second
control position on screen, third controls width, and fourth controls height
m = findobj(gca, 'Type', 'Scatter');
hleg1 = legend(m(1:2), 'Right Hemisphere', 'Left Hemisphere', 'Location',
'SouthOutside');
print(gcf, [output_dir out_str '_numVars.jpg'], '-dpng', '-r300')
out_str = {'HCP384'};
print(gcf, [output_dir out_str '_numVars.jpg'], '-dpng', '-r300')
[output_dir out_str '_numVars.jpg']
[output_dir cell2mat(out_str) '_numVars.jpg']
print(gcf, [output_dir cell2mat(out_str) '_numVars.jpg'], '-dpng', '-r300')
close gcf
leftHem = variants_info.left_hem.group_avg{:,2};
rightHem = variants_info.right_hem.group_avg{:,2};
[h,mu,sigma,q,notch] = al_goodplot(leftHem,0.1,.2,[ 0    0.4470    0.7410],'left');
[h,mu,sigma,q,notch] = al_goodplot(rightHem,0.11,.2,[0.8500    0.3250
0.0980],'right');
xticks([0.105])
xticklabels('Number of Variant Vertices')
axis([-0.2, 0.5, min([leftHem; rightHem])-std([leftHem; rightHem]), max([leftHem;
rightHem])+std([leftHem; rightHem])])
ax = gca;
ax.FontSize = 24;
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.3, 0.9]); %first and second
control position on screen, third controls width, and fourth controls height
m = findobj(gca, 'Type', 'Scatter');
hleg1 = legend(m(1:2), 'Right Hemisphere', 'Left Hemisphere', 'Location',
'SouthOutside');
print(gcf, [output_dir cell2mat(out_str) '_numVarVerts.jpg'], '-dpng', '-r300')
axis([-0.2, 0.5, min([leftHem; rightHem])-std([leftHem; rightHem]), max([leftHem;
rightHem])+std([leftHem; rightHem])])
close gcf
leftHem = variants_info.left_hem.group_avg{:,3};
rightHem = variants_info.right_hem.group_avg{:,3};
[h,mu,sigma,q,notch] = al_goodplot(leftHem,0.1,.2,[ 0    0.4470    0.7410],'left');
[h,mu,sigma,q,notch] = al_goodplot(rightHem,0.11,.2,[0.8500    0.3250
0.0980],'right');
xticks([0.105])
xticklabels('Average Variant Size')
ylabel('Number of vertices')
axis([-0.2, 0.5, min([leftHem; rightHem])-std([leftHem; rightHem]), max([leftHem;
rightHem])+std([leftHem; rightHem])])
ax = gca;
ax.FontSize = 24;
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.3, 0.9]); %first and second
control position on screen, third controls width, and fourth controls height
m = findobj(gca, 'Type', 'Scatter');
hleg1 = legend(m(1:2), 'Right Hemisphere', 'Left Hemisphere', 'Location',
'SouthOutside');
print(gcf, [output_dir cell2mat(out_str) '_avgSizeVerts.jpg'], '-dpng', '-r300')
close gcf
leftHem = variants_info.left_hem.group_avg{:,4};
rightHem = variants_info.right_hem.group_avg{:,4};
[h,mu,sigma,q,notch] = al_goodplot(leftHem,0.1,.2,[ 0    0.4470    0.7410],'left');
[h,mu,sigma,q,notch] = al_goodplot(rightHem,0.11,.2,[0.8500    0.3250
```

```

0.0980], 'right');
xticks([0.105])
xticklabels('Average Variant Size')
ylabel('Surface area (mm^2)')
axis([-0.2, 0.5, min([leftHem; rightHem])-std([leftHem; rightHem]), max([leftHem;
rightHem])+std([leftHem; rightHem])])
ax = gca;
ax.FontSize = 24;
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.3, 0.9]); %first and second
control position on screen, third controls width, and fourth controls height
m = findobj(gca, 'Type', 'Scatter');
hleg1 = legend(m(1:2), 'Right Hemisphere', 'Left Hemisphere', 'Location',
'SouthOutside');
print(gcf, [output_dir cell2mat(out_str) '_avgSizeSurfArea.jpg'], '-dpng', '-r300')
close gcf
network_names = {'DMN' 'Vis' 'FP' 'DAN' 'VAN' 'Sal' 'CO' 'SMd' 'SML'
'Aud' 'Tpole' 'MTL' 'PMN' 'PON'};
good_nets_LH = [networksxHem.clustersLH(:,1:3) networksxHem.clustersLH(:,5) networksxHem.
clustersLH(:,7:end)];
good_nets_RH = [networksxHem.clustersRH(:,1:3) networksxHem.clustersRH(:,5) networksxHem.
clustersRH(:,7:end)];
good_nets_LH = mean(good_nets_LH);
good_nets_RH = mean(good_nets_RH);
means = [good_nets_LH good_nets_RH];
means = [good_nets_LH' good_nets_RH'];
bar(1:14, means)
legend('Left Hem', 'Right Hem')
xticks(1:14)
xticklabels(network_names)
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.5, 0.7]);
ylabel('Average Number of Variants Assigned to Network');
xlabel('Network');
title_str('Variants Assigned to Each Network Across Hemispheres');
print(gcf, [output_dir cell2mat(out_str) '_netAssignment.jpg'], '-dpng', '-r300');
title('Variants Assigned to Each Network Across Hemispheres');
print(gcf, [output_dir cell2mat(out_str) '_netAssignment.jpg'], '-dpng', '-r300');
print(gcf, [output_dir cell2mat(out_str) '_netAssignment.jpg'], '-dpng', '-r300']);
ax = gca;
ax.FontSize = 24;
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.9, 0.7]);
print(gcf, [output_dir cell2mat(out_str) '_netAssignment.jpg'], '-dpng', '-r300');
[output_dir cell2mat(out_str) '_netAssignment.jpg']
print(gcf, [output_dir cell2mat(out_str) '_netAssignment.jpg'], '-dpng', '-r300');
close gcf
%-- 12/28/21, 9:24 AM --%
help exists
help stdev
ft_read_cifti_mod
('/Volumes/RESEARCH_HD/HCP_Variants/new_split_vars/reassigned/100206_borderlectopic2.
dtseries.nii')
cifti = ans;
template = cifti;
clear cifti
template.data = [];
template.data = zeros(size(ans.data));
template.data = zeros(size(ans.data));
ft_write_cifti_mod('/Users/dianaperez/Desktop/lateralization_code/needed_files/template.
dtseries.nii', template)
open ft_write_cifti_mod
ft_write_cifti_mod('/Users/dianaperez/Desktop/lateralization_code/needed_files/template.

```

```
dtseries.nii', template)
filename = '/Users/dianaperez/Desktop/lateralization_code/needed_files/template.dtseries.
nii';
source = template;
ft_write_cifti_mod
ft_write_cifti_mod(filename, template)
permute_spatial_dist_handedness
help exists
permute_spatial_dist_handedness
permute_spatial_distribution
length(subs{g})
save(left_cifti_loc, 'true_left_hems')
save(right_cifti_loc, 'true_right_hems')
permute_spatial_distribution
%-- 1/6/22, 8:39 AM --%
make_spCorr_Map(1, '/Volumes/Back_Up/Lifespan/dconns/sub-LS02_allsess_tmasked.dconn.nii',
'/Volumes/Back_Up/Lifespan/dconns/sub-LS02_allsess_tmasked_sptlCorr.dtseries.
nii', '/Volumes/Back_Up/Lifespan/dconns/120_allsubs_corr.dconn.nii')
make_spCorr_Map
make_spCorr_Map(1, '/Volumes/Back_Up/Lifespan/dconns/sub-LS02_allsess_tmasked.dconn.nii',
'/Volumes/Back_Up/Lifespan/dconns/sub-LS02_allsess_tmasked_sptlCorr.dtseries.
nii', '/Volumes/Back_Up/Lifespan/dconns/120_allsubs_corr.dconn.nii')
dconn_dat
make_spCorr_Map(1, '/Volumes/Back_Up/Lifespan/dconns/sub-LS02_allsess_tmasked.dconn.nii',
'/Volumes/Back_Up/Lifespan/dconns/sub-LS02_allsess_tmasked_sptlCorr.dtseries.
nii', '/Volumes/Back_Up/Lifespan/dconns/120_allsubs_corr.dconn.nii')
make_spCorr_Map
make_spCorr_Map(1, '/Volumes/Back_Up/Lifespan/dconns/sub-LS02_allsess_tmasked.dconn.nii',
'/Volumes/Back_Up/Lifespan/dconns/sub-LS02_allsess_tmasked_sptlCorr.dtseries.
nii', '/Volumes/Back_Up/Lifespan/dconns/120_allsubs_corr.dconn.nii')
dconn_dat
make_spCorr_Map(1, '/Volumes/Back_Up/Lifespan/dconns/sub-LS02_allsess_tmasked.dconn.nii',
'/Volumes/Back_Up/Lifespan/dconns/sub-LS02_allsess_tmasked_sptlCorr.dtseries.
nii', '/Volumes/Back_Up/Lifespan/dconns/120_allsubs_corr.dconn.nii')
make_spCorr_Map(1, '/Volumes/Back_Up/Lifespan/dconns/sub-LS02_allsess_tmasked.dconn.nii',
'/Volumes/Back_Up/Lifespan/dconns/sub-LS02_allsess_tmasked_sptlCorr.dtseries.
nii', '/Volumes/Back_Up/Lifespan/dconns/120_allsubs_corr.dconn.nii')
dconn_dat
%-- 1/6/22, 1:43 PM --%
similarity_analysis
hline_new([3,8,13,15,16,21,26,28,33,34,39,44]+0.5, 'k', 2);
vline_new([3,8,13,15,16,21,26,28,33,34,39,44]+0.5, 'k', 2);
hline_new([3,8,13,18,19,24,29,31,36,37,42,47]+0.5, 'k', 2);
close(gcf)
figure('Position', [1 1 1000 800]);
imagesc(simmat, [0 1]); colormap('jet');
hline_new([3,8,13,18,19,24,29,31,36,37,42,47]+0.5, 'k', 2);
vline_new([3,8,13,18,19,24,29,31,36,37,42,47]+0.5, 'k', 2);
close(gcf)
figure('Position', [1 1 1000 800]);
imagesc(simmat, [0 1]); colormap('jet');
hline_new([8,18,19,29,31,36,37,42,47]+0.5, 'k', 2);
vline_new([8,18,19,29,31,36,37,42,47]+0.5, 'k', 2);
set(gca, 'XTick', [4,13.5,19,24.5,30.5,34,37,40,45], 'YTick',
[4,13.5,19,24.5,30.5,34,37,40,45], 'XTickLabel', ...
{'LS02', 'LS03', 'LS04', 'LS05', 'LS07', 'LS08', 'LS10', 'LS11', 'LS14'}, 'YTickLabel',
{'LS02', 'LS03', 'LS04', 'LS05', 'LS07', 'LS08', 'LS10', 'LS11', 'LS14'});
axis square;
colorbar;
close(gcf)
```

```
figure('Position',[1 1 1000 800]);
imagesc(simmat,[0 1]); colormap('jet');
hline_new([8,18,19,29,31,36,37,42]+0.5,'k',2);
vline_new([8,18,19,29,31,36,37,42]+0.5,'k',2);
set(gca,'XTick',[4,13.5,19,24.5,30.5,34,37,40,45], 'YTick',↵
[4,13.5,19,24.5,30.5,34,37,40,45], 'XTickLabel',...
{'LS02', 'LS03', 'LS04', 'LS05', 'LS07', 'LS08', 'LS10', 'LS11', 'LS14'}, 'YTickLabel',↵
{'LS02', 'LS03', 'LS04', 'LS05', 'LS07', 'LS08', 'LS10', 'LS11', 'LS14'});
axis square;
colorbar;
title('Correlation Matrix Similarity');
saveas(gcf,[outDir 'SimilarityMat_rest.tiff'],'tiff');
maskmat = ones(atlas_params.num_rois);
maskmat = logical(triu(maskmat,1));
count = 1;
for s = 1:9
    tmp = mean_corrmat(s,:,:,:);
    corrlin(count,:) = tmp(maskmat);
    count = count+1;
end
simmat = corr(corrlin');
figure('Position',[1 1 1000 800]);
imagesc(simmat,[0 1]); colormap('jet');
close gcf
maskmat = ones(atlas_params.num_rois);
maskmat = logical(triu(maskmat,1));
count = 1;
for s = 1:9
    tmp = mean_corrmat(s,:,:,:);
    corrlin(count,:) = tmp(maskmat);
    count = count+1;
end
simmat = corr(corrlin');
figure('Position',[1 1 1000 800]);
imagesc(simmat,[0 1]); colormap('jet');
maskmat = ones(atlas_params.num_rois);
maskmat = logical(triu(maskmat,1));
count = 1;
tmp = mean_corrmat(1,:,:,:);
corrlin(count,:) = tmp(maskmat);
count = count+1;
tmp = mean_corrmat(2,:,:,:);
corrlin(count,:) = tmp(maskmat);
count = count+1;
tmp = mean_corrmat(3,:,:,:);
clear corrlin
count = 1;
tmp = mean_corrmat(1,:,:,:);
corrlin(count,:) = tmp(maskmat);
count = count+1;
clear corrlin
for s = 1:9
    tmp = mean_corrmat(s,:,:,:);
    corrlin(count,:) = tmp(maskmat);
    count = count+1;
end
clear corrlin
maskmat = ones(atlas_params.num_rois);
maskmat = logical(triu(maskmat,1));
count = 1;
```

```

for s = 1:9
tmp = mean_corrmat(s,:,:,:);
corrln(count,:) = tmp(maskmat);
count = count+1;
end
simmat = corr(corrln');
figure('Position',[1 1 1000 800]);
imagesc(simmat,[0 1]); colormap('jet');
hline_new([1:8]+0.5,'k',2);
vline_new([1:8]+0.5,'k',2);
set(gca,'XTick',[1:9], 'YTick', [1:9], 'XTickLabel',...
{'LS02', 'LS03', 'LS04', 'LS05', 'LS07', 'LS08', 'LS10', 'LS11', 'LS14'}, 'YTickLabel',
{'LS02', 'LS03', 'LS04', 'LS05', 'LS07', 'LS08', 'LS10', 'LS11', 'LS14'});
axis square;
colorbar;
title('Correlation Matrix Similarity');
saveas(gcf,[outdir 'SimilarityMat_averaged.tiff'],'tiff');
clear corrmat
clear corrmat
clear corrln
count = 1;
subs = {'LS02', 'LS03', 'LS05'};
ses_pre = [3,5,5];
ses_post = [5,5,5];
count = 1;
for sub = 1:numel(subs)
for ses = 1:ses_pre(sub)
data = load(sprintf('%s/sub-%s/sub-%s_sess-%d_task-rest_corrmat_Seitzman300.mat',
datadir, subs{sub}, subs{sub}, ses));
corrmat(count,:,:,:) = data.corrmat;
sub_corrmat(ses,:,:,:) = data.corrmat;
count = count + 1;
end
if ses > 1
mean_corrmat(sub,:,:,:) = squeeze(mean(sub_corrmat));
clear sub_corrmat
else
mean_corrmat(sub,:,:,:)= squeeze(sub_corrmat);
end
end
%-- 1/8/22, 2:16 PM --%
datadir = '/Volumes/Back_Up/Lifespan/corrmats/';
outdir = '/Volumes/Back_Up/Lifespan/CNS_analyses/';
atlas_dir = '/Users/dianaperez/Box/Dependencies/';
addpath(genpath('/Users/dianaperez/Documents/GitHub/GrattonLab-General-Repo/FCPROCESS'));
addpath(genpath('/Users/dianaperez/Box/Scripts'))
addpath(genpath('/Users/dianaperez/Box/Dependencies/cifti-matlab-master'))
addpath(genpath('/Users/dianaperez/Box/Quest_Backup/Darmouth_MIND2'))
atlas_params = atlas_parameters_GrattonLab('Seitzman300',atlas_dir);
subs = {'LS02', 'LS03', 'LS05'};
ses_pre = [3,5,5];
ses_post = [5,5,5];
count = 1;
for sub = 1:numel(subs)
for ses = 1:ses_pre(sub)
data = load(sprintf('%s/sub-%s/sub-%s_sess-%d_task-rest_corrmat_Seitzman300.mat',
datadir, subs{sub}, subs{sub}, ses));
corrmat(count,:,:,:) = data.corrmat;
sub_corrmat(ses,:,:,:) = data.corrmat;
count = count + 1;

```

```
end
if ses > 1
mean_corrmat(sub, :, :, :) = squeeze(mean(sub_corrmat));
clear sub_corrmat
else
mean_corrmat(sub, :, :, :) = squeeze(sub_corrmat);
end
end
maskmat = ones(atlas_params.num_rois);
maskmat = logical(triu(maskmat,1));
count = 1;
for sub = 1:numel(subs)
for ses = 1:ses_pre(sub)
data = load(sprintf('%s/sub-%s/sub-%s_sess-%d_task-rest_corrmat_Seitzman300.mat', ↵
datadir, subs{sub}, subs{sub}, ses));
corrmat(count, :, :, :) = data.corrmat;
sub_corrmat(ses, :, :, :) = data.corrmat;
count = count + 1;
end
if ses > 1
mean_pre(sub, :, :, :) = squeeze(mean(sub_corrmat));
clear sub_corrmat
else
mean_pre(sub, :, :, :) = squeeze(sub_corrmat);
end
end
count = 1;
for sub = 1:numel(subs)
for ses = ses_pre(sub)+1:ses_pre(sub)+ses_post(sub)
data = load(sprintf('%s/sub-%s/sub-%s_sess-%d_task-rest_corrmat_Seitzman300.mat', ↵
datadir, subs{sub}, subs{sub}, ses));
corrmat(count, :, :, :) = data.corrmat;
sub_corrmat(ses, :, :, :) = data.corrmat;
count = count + 1;
end
if ses > 1
mean_post(sub, :, :, :) = squeeze(mean(sub_corrmat));
clear sub_corrmat
else
mean_post(sub, :, :, :) = squeeze(sub_corrmat);
end
end
maskmat = ones(atlas_params.num_rois);
maskmat = logical(triu(maskmat,1));
count = 1;
for sub = 1:numel(subs)
tmp = mean_pre(count, :, :, :);
corrlin_pre(count, :) = single(tmp(maskmat));
tmp = mean_post(count, :, :, :);
corrlin_post(count, :) = single(tmp(maskmat));
count = count + 1;
end
simmat = corr(corrlin_pre', corrlin_post');
figure('Position',[1 1 1000 800]);
imagesc(simmat,[0 1]); colormap('jet');
hline_new([1:2]+0.5,'k',2);
vline_new([1:2]+0.5,'k',2);
set(gca,'XTick',[1:3], 'YTick', [1:3], 'XTickLabel',...
{'LS02', 'LS03', 'LS05'}, 'YTickLabel', {'LS02', 'LS03', 'LS05'});
axis square;
```



```
colorbar;
title('Correlation Matrix Similarity - 1 year apart');
saveas(gcf,[outDir 'SimilarityMat_longitudinal.tiff'],'tiff');
reliability_analysis
addpath '/Users/dianaperez/Documents/GitHub/Lifespan-Analysis'
reliability_analysis
plot(times,corr)
plot(times,corr(1,:), 'LineWidth',3)
plot(times(1:20),corr(1,:), 'LineWidth',3)
length(times(1:20))
length(times(1:20))
length(corr(1,:))
plot(times(1:20),corr(1,1:20), 'Color',[1, 0.5, 0], 'LineWidth', 3)
hold on
plot(times(1:20),corr(2,1:20), 'Color',[0, 0, 1], 'LineWidth', 3)
plot(times(1:20),corr(3,1:20), 'Color',[0, 1, 0], 'LineWidth', 3)
close gcf
times =[5:5:100];
figure;
plot(times(1:20),corr(1,1:20), 'Color',[1, 0, 0], 'LineWidth', 3) %LS02
ylim([0 1]);
hold on
plot(times(1:20),corr(2,1:20), 'Color',[0, 1, 0], 'LineWidth', 3) %LS03
hold on
plot(times(1:20),corr(3,1:20), 'Color',[0, 0, 1], 'LineWidth', 3)%LS05
hold on
plot(times(1:20),corr(4,1:20), 'Color',[0, 1, 1], 'LineWidth', 3)%LS08
hold on
plot(times(1:20),corr(4,1:20), 'Color',[1, 0, 1], 'LineWidth', 3)%LS11
hold on
plot(times(1:20),corr(4,1:20), 'Color',[0, 0, 0], 'LineWidth', 3)%LS14
close gcf
times =[5:5:100];
figure;
plot(times(1:20),corr(1,1:20), 'Color',[1, 0, 0], 'LineWidth', 3) %LS02
hold on
plot(times(1:20),corr(2,1:20), 'Color',[0, 1, 0], 'LineWidth', 3) %LS03
hold on
plot(times(1:20),corr(3,1:20), 'Color',[0, 0, 1], 'LineWidth', 3)%LS05
hold on
plot(times(1:3),corr(4,1:3), 'Color',[0, 1, 1], 'LineWidth', 3)%LS08
hold on
plot(times(1:20),corr(4,1:20), 'Color',[1, 0, 1], 'LineWidth', 3)%LS11
hold on
close gcf
times =[5:5:100];
figure;
plot(times(1:20),corr(1,1:20), 'Color',[1, 0, 0], 'LineWidth', 3) %LS02
hold on
plot(times(1:20),corr(2,1:20), 'Color',[0, 1, 0], 'LineWidth', 3) %LS03
hold on
plot(times(1:20),corr(3,1:20), 'Color',[0, 0, 1], 'LineWidth', 3)%LS05
hold on
plot(times(1:3),corr(4,1:3), 'Color',[0, 1, 1], 'LineWidth', 3)%LS08
hold on
plot(times(1:20),corr(5,1:20), 'Color',[1, 0, 1], 'LineWidth', 3)%LS11
hold on
plot(times(1:20),corr(6,1:20), 'Color',[0, 0, 0], 'LineWidth', 3)%LS14
ylabel('Pearson Correlation (r)');
xlabel('Time (Minutes)');
```



```
m = findobj(gca,'Type','line');
hleg1 = legend(m(1:6), 'LS14', 'LS11', 'LS08', 'LS05', 'LS03', 'LS02', 'Location',
'SouthEast');
reliability_analysis
print(gcf, '/Volumes/Back_Up/Lifespan/CNS_analyses/reliability_figure.jpg', '-dpng', '-r300');
reliability_analysis
corrs_for_mean = [corr(1:3,1:20; corr(5:6,1:20));
mean = mean(corrs_for_mean);
corrs_for_mean = [corr(1:3,1:20); corr(5:6,1:20)];
mean = mean(corrs_for_mean);
times =[5:5:100];
figure;
plot(times(1:20),corr(1,1:20),'Color',[1, 0, 0],'LineWidth', 3) %LS02
hold on
plot(times(1:20),corr(2,1:20),'Color',[0, 1, 0],'LineWidth', 3) %LS03
hold on
plot(times(1:20),corr(3,1:20),'Color',[0, 0, 1],'LineWidth', 3)%LS05
hold on
plot(times(1:3),corr(4,1:3),'Color',[0, 1, 1],'LineWidth', 3)%LS08
hold on
plot(times(1:20),corr(5,1:20),'Color',[1, 0, 1],'LineWidth', 3)%LS11
hold on
plot(times(1:20),corr(6,1:20),'Color',[0, 0, 0],'LineWidth', 3)%LS14
hold on
plot(times(1:20),mean, ':', 'Color', [0.4940 0.1840 0.5560], 'LineWidth',3) %average
ylabel('Pearson Correlation (r)');
xlabel('Time (Minutes)');
m = findobj(gca,'Type','line');
hleg1 = legend(m(1:7), 'Average', 'LS14', 'LS11', 'LS08', 'LS05', 'LS03', 'LS02',
'Location', 'SouthEast');
hleg1.FontSize = 14;
ax = gca;
ax.FontSize = 17;
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.5, 0.7]);
print(gcf, '/Volumes/Back_Up/Lifespan/CNS_analyses/reliability_figure.jpg', '-dpng', '-r300');
similarity_analysis
clear all
similarity_analysis
hline_new([3,13,24]+0.5,'k',1);
vline_new([3,13,24]+0.5,'k',1);
similarity_analysis
% longitudinal analysis
subs = {'LS02', 'LS03', 'LS05'};
sessions = [8,10,10];
clear corrmats
count = 1;
for sub = 1:numel(subs)
for ses = 1:sessions(sub)
data = load(sprintf('%s/sub-%s/sub-%s_sess-%d_task-rest_corrmats_Seitzman300.mat',
datadir, subs{sub}, subs{sub}, ses));
corrmats(count,:,:,:) = data.corrmats;
sub_corrmats(ses,:,:,:) = data.corrmats;
count = count + 1;
end
end
figure('Position',[1 1 1000 800]);
imagesc(simmat,[0 1]); colormap('jet');
hline_new([8,18]+0.5,'k',2);
```

```
hline_new([3,13,23]+0.5,'k',.5);
vline_new([8,18]+0.5,'k',2);
vline_new([3,13,23]+0.5,'k',.5);
set(gca,'XTick',[4,13.5,23.5], 'YTick', [4,13.5,23.5], 'XTickLabel',...
{'LS02', 'LS03', 'LS05'}, 'YTickLabel', {'LS02', 'LS03', 'LS05'});
axis square;
colorbar;
title('Correlation Matrix Similarity');
similarity_analysis
clear corrlin
maskmat = ones(atlas_params.num_rois);
maskmat = logical(triu(maskmat,1));
count = 1;
for sub = 1:numel(subs)
for ses = 1:sessions(sub)
tmp = corrmatrix(count, :, :, :);
if fisher
corrlin(count, :) = single(FisherTransform(tmp(maskmat)));
else
corrlin(count, :) = single(tmp(maskmat));
end
count = count + 1;
end
end
simmat = corr(corrlin');
similarity_analysis
clear corrlin
similarity_analysis
%-- 1/12/22, 11:05 AM --%
ft_read_cifti_mod('/Volumes/RESEARCH_HD/Lifespan/CNS_analyses/sub-
LS02_vs_120_avg_corr_LR_cortex_corr.dtseries.nii')
spCorr = ans;
clear ans
template = ft_read_cifti_mod('/Volumes/fsmresfiles/PBS/Gratton_Lab/Lifespan/Pre-
COVID/BIDS/Nifti/derivatives/postFCproc_CIFTI/dconn_cifti_normalwall/spatialCorrMapLS03_v
s_120_allsubs_corr_cortex_smooth_2.55.dtseries.nii');
template.data = spCorr.data;
ft_write_cifti_mod('/Users/dianaperez/Desktop/sub-LS02_spCorrMap.dtseries.nii',
template);
spCorr = ft_read_cifti_mod('/Volumes/RESEARCH_HD/Lifespan/CNS_analyses/sub-
LS03_vs_120_avg_corr_LR_cortex_corr.dtseries.nii')
template.data = spCorr.data;
ft_write_cifti_mod('/Users/dianaperez/Desktop/sub-LS03_spCorrMap.dtseries.nii',
template);
spCorr = ft_read_cifti_mod('/Volumes/RESEARCH_HD/Lifespan/CNS_analyses/sub-
LS05_vs_120_avg_corr_LR_cortex_corr.dtseries.nii');
template.data = spCorr.data;
ft_write_cifti_mod('/Users/dianaperez/Desktop/sub-LS05_spCorrMap.dtseries.nii',
template);
pre = spCorr.data;
clear all
pre = ft_read_cifti_mod('/Volumes/fsmresfiles/PBS/Gratton_Lab/Lifespan/Pre-
COVID/BIDS/Nifti/derivatives/postFCproc_CIFTI/dconn_cifti_normalwall/sub-
LS02_vs_120_allsubs_corr_cortex_corr.dtseries.nii');
post = ft_read_cifti_mod('/Users/dianaperez/Desktop/sub-LS02_spCorrMap.dtseries.nii');
spCorr = corr(pre.data, post.data);
pre = ft_read_cifti_mod('/Volumes/fsmresfiles/PBS/Gratton_Lab/Lifespan/Pre-
COVID/BIDS/Nifti/derivatives/postFCproc_CIFTI/dconn_cifti_normalwall/sub-
LS03_vs_120_allsubs_sptlCorr_cortex_smooth_2.55.dtseries.nii');
post = ft_read_cifti_mod('/Users/dianaperez/Desktop/sub-LS03_spCorrMap.dtseries.nii');
```

```
spCorr = corr(pre.data, post.data);
pre = ft_read_cifti_mod('/Volumes/fsmresfiles/PBS/Gratton_Lab/Lifespan/Pre-
COVID/BIDS/Nifti/derivatives/postFCproc_CIFTI/dconn_cifti_normalwall/sub-
LS05_vs_120_allsubs_sptlCorr_cortex_smooth_2.55.dtseries.nii');
post = ft_read_cifti_mod('/Users/dianaperez/Desktop/sub-LS05_spCorrMap.dtseries.nii');
spCorr = corr(pre.data, post.data);
CreateVariantFiles_LS03specific
unique(cifti_rest.data)
CreateVariantFiles_LS03specific
clear all
EmReg_task
%-- 1/15/22, 4:26 PM --%
EmReg_task
sca
KbName('return')
KbName('enter')
[keyIsDown,secs, keyCode] = KbCheck
find(keyCode>0)
unique(keyCode)
[keyIsDown,secs, keyCode] = KbCheck
vleftKey = KbName('LeftArrow');
rightKey = KbName('RightArrow');
spaceBar = KbName('space');
%-- 1/24/22, 6:38 PM --%
help rand
permute_spatial_dist_handedness
rand_LH = all_data(:,ind(1:length(LH)));
rand_RH = all_data(:,ind(length(LH)+1:end));
corr(rand_LH, rand_RH)
ind = randperm(numSubs)
permute_spatial_dist_handedness
figure;
ind = zeros(length(perm_diffs),1);
ind(1,1) = 1;
handles = plotSpread(spCorrs, 'categoryMarkers', {'o', '.'}, 'categoryLabels', {'Permuted
Correlation', 'True Correlation'}, 'categoryColors', {'k', 'r'}, 'categoryIdx', [ind])
scatter(1,spCorrs(1,:), 'MarkerEdgeColor', 'red', 'MarkerFaceColor', 'red', 'SizeData',
50)
ax = gca;
ax.FontSize = 12;
axis([0.5, 1.5, min(spCorrs)-std(spCorrs), max(spCorrs)+std(spCorrs)])
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.3, 0.7]); %first and second
control position on screen, third controls width, and fourth controls height
print(gcf,[output_dir out_str '_permuted_spatialCorrs_' num2str(numperms)
'_permutations_figure.jpg'],'-dpng','-r300');
close gcf
help logical
network_names =
{'DMN','Vis','FP','Unassigned','DAN','Unassigned','VAN','Sal','CO','SMd','SMl','Aud','Tpo
le','MTL','PMN','PON'};
network_names{0}
overlapmap
unique(group_map)
unique(variant_map_mat)
overlapmap
num_vars = find(variant_map_mat>0);
num_vars_check = find(group_map>0);
clear num_var*
overlapmap
num_vars = find(variant_map_mat==net_id);
```

```
num_vars_check = find(group_map==1);
clear num_var*
net_size_across_hems
help exists
net_size_across_hems
axis([0 .5 0 .5])
axis([0 .2 0 .2])
axis([0 .18 0 .18])
ax.FontSize = 24;
legend(network_names)
close gcf
scatter(net_size(:,4), net_size(:,8), 80, rgb_colors, 'filled')
axis([0 .2 0 .2])
legend(network_names)
close gcf
scatter(net_size(:,4), net_size(:,8), 80, rgb_colors, 'filled')
axis([0 .2 0 .2])
line = reffline(1,0);
line.Color = 'black';
xlabel('Left Hemisphere')
ylabel('Right Hemisphere')
ax = gca;
ax.FontSize = 24;
print(gcf, [out_dir out_str '_groupavg_nets_surfareas_scatter.jpg'], '-dpng', '-r300')
close gcf
net_size_across_hems
ax_max = max([x;y])+std([x;y])
ax_max = max([x;y])+std([x;y])
x = net_size(:,4); y = net_size(:,8);
ax_max = max([x;y])+std([x;y])
axis([0 ax_max 0 ax_max])
ax_max = max([x;y])+(std([x;y])/2)
axis([0 ax_max 0 ax_max])
net_size_across_hems
EmReg_task
help inputdlg
save('/Volumes/RESEARCH_HD/LifespanProjectTasks/EmoReg/emotions_table.mat', 'myExp')
EmReg_task
sca
EmReg_task
sca
EmReg_task
sca
%-- 1/26/22, 12:58 PM --%
EmReg_task
sca
EmReg_task
sca
EmReg_task
sca
%-- 1/26/22, 1:03 PM --%
EmReg_task
sca
%-- 1/26/22, 1:06 PM --%
EmReg_task
%-- 1/26/22, 1:07 PM --%
EmReg_task
sca
%-- 1/26/22, 2:28 PM --%
EmReg_task
```

```
sca
PsychHID
clear all
PsychHID
elements=PsychHID('Elements',deviceNumber)
rc = PsychHID('KeyboardHelper', commandCode)
rc = PsychHID('KeyboardHelper', commandCode)
rc = PsychHID('Devices'))
rc = PsychHID('Devices')
[navail] = PsychHID('KbQueueFlush')
PsychHID('KbQueueCreate')
%-- 1/26/22, 2:40 PM --%
EmReg_task
sca
%-- 1/26/22, 2:57 PM --%
KbCheck
EmReg_task
sca
KbWait
KbCheck
EmReg_task
sca
EmReg_task
sca
%-- 1/26/22, 5:35 PM --%
KbWait
%-- 1/26/22, 5:36 PM --%
KbWait
%-- 1/26/22, 5:39 PM --%
KbWait
k;jsdjsla;c
EmReg_task
sca
EmReg_task
sca
KbName('space')
KbCheck
pause(.5);KbCheck
pause(.5);KbChec
pause(.5);KbCheck
[keyIsDown,secs, keyCode] = KbCheck;
clear all
PsychHID('Devices')
clear PsychHID
dev = PsychHID('Devices')
[secs, keyCode] = KbWait
WaitSecs(0.2);[secs, key] = KbWait
[a,b,c] = GetKeyboardIndices
KbName('UnifyKeyNames')
KbName('UnifyKeyNames')
WaitSecs(0.2);[secs, key] = KbWait
jj jbnjnmnkjh j
WaitSecs(0.2);[secs, key] = KbWait(3)
WaitSecs(0.2);[secs, key] = KbWait(9)
EmReg_task
sca
EmReg_task
sca
EmReg_task
sca
```

```

load([root_dir 'emotions_table.mat']);
emotions = myExp;
emotions{1,1}
EmReg_task
sca
emotions{1,2}
load('emotions_table.mat')
url_emoscale = '/scales/emotion_scale.png';
myExp{1:13,2} = url_emoscale;
myExp{1,2}
myExp{1,2} = url_emoscale
myExp1{1,2} = url_emoscale
clear myExp
clear myExp1
clear emotions
clear all
scale = imread(myExp{1,2});
emotions = table2cell(myExp)
scale = imread(emotions{1,2});
emotions{1,2}
scale = imread(string(emotions{1,2}));
scale = imread([root_dir string(emotions{1,2})]);
root_dir = '/Volumes/RESEARCH_HD/LifespanProjectTasks/EmoReg/'
scale = imread([root_dir string(emotions{1,2})]);
[root_dir string(emotions{1,2})];
string(emotions{1,2})
cellstr(emotions{1,2})
[root_dir cellstr(emotions{1,2})];
help nominal
[root_dir char(emotions{1,2})];
scale = imread([root_dir char(emotions{1,2})]);
EmReg_task
clear all
[a,b,c] = GetKeyboardIndices
clear PsychHID
[a,b,c] = GetKeyboardIndices
EmReg_task
sca
clear all
save('emotions_table.mat', 'myExp')
[root_dir myExp{1,2}]
root_dir = '/Volumes/RESEARCH_HD/LifespanProjectTasks/EmoReg/'
[root_dir myExp{1,2}]
EmReg_task
clear all
[a,b,c] = GetKeyboardIndices
EmReg_task
scale = Screen('MakeTexture', window, scale_img);
scale_img = imread([root_dir emotions_table{emo,2}]);
scale = Screen('MakeTexture', window, scale_img);
scale_img = imread([root_dir emotions_table{1,2}]);
EmReg_task
help CenterRectOnPointd
EmReg_task
output
EmReg_task
Screen('DrawText', 10, 'Enter a number, then press RETURN', 'center', 'center', [0,0,0],↵
[1,1,1])
window
EmReg_task

```

```

Screen('DrawText', 2, 'Enter a number, then press RETURN', 'center', 'center', [0,0,0], ↵
[1,1,1])
Screen('DrawText', 2, 'Enter a number, then press RETURN', [], [], [0,0,0], [1,1,1])
GetEchoString
EmReg_task
Screen('DrawText', 2, 'Enter a number, then press RETURN', [], [], [0,0,0], [1,1,1])
EmReg_task
6
sca
EmReg_task
7
7
sca
cd /Users/dianaperez/Desktop
var_freq_size_net
cd '/Users/dianaperez/Documents/GitHub/PerezEtAl_HemAsymmetries'
var_freq_size_net
cell2mat(out_str{g})
cell2mat(out_str(g))
var_freq_size_net
perms_var_freq_size
permute_net_assignment
min_ax = min(good_nets(:,:))-(std(good_nets)/2);
min_ax = min(good_nets(:,:))-(std(good_nets(:,:))/2);
min(good_nets(:,:))
min_ax = min(min(good_nets(:,:)))-(min(std(good_nets)/2));
max_ax = max(max(good_nets(:,:)))+(min(std(good_nets)/2));
axis([0, 15, min_ax, max_ax])
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.9, 0.7]);
print(gcf,[output_dir out_str '_PermutationTesting_networkAssignment.jpg'], '-dpng', '-↵
r300');
close gcf
permute_net_assignment
operc_cluster
FCPROCESS_GrattonLab_forced_data('/Users/dianaperez/Desktop/Lifespan_datalist1.xlsx', ↵
'/Users/dianaperez/Desktop/', 'defaults2')
Variant_overlap
unique(varmap_1.data)
Variant_overlap
unique(overlap_map)
template = varmap_1;
template.data = overlap_map;
ft_write_cifti_mod('/Users/dianaperez/Desktop/LS03_longitudinal_variant_overlap.dseries. ↵
nii', template);
template.data = overlap_map';
ft_write_cifti_mod('/Users/dianaperez/Desktop/LS03_longitudinal_variant_overlap.dseries. ↵
nii', template);
operc_cluster
indiv_net_size_by_hems
diff_surf(s,net) = left_hem_surf(s,net) - right_hem_surf(s,net);
indiv_net_size_by_hems
save([output_dir '/HCP384_indiv_network_size_by_hems.mat'], indiv_nets_size)
save↵
('/Users/dianaperez/Desktop/lateralization_code/testing_output/indiv_net_symm/HCP384_indi ↵
v_network_size_by_hems.mat', indiv_nets_size)
save↵
('/Users/dianaperez/Desktop/lateralization_code/testing_output/indiv_net_symm/HCP384_indi ↵
v_network_size_by_hems.mat', 'indiv_nets_size')
clear all
load↵

```

```
( '/Users/dianaperez/Desktop/lateralization_code/testing_output/operc_nets_lefthemcluster_
lefthem.mat' )
load
( '/Users/dianaperez/Desktop/lateralization_code/testing_output/operc_nets_lefthemcluster_
righthem.mat' )
load
( '/Users/dianaperez/Desktop/lateralization_code/testing_output/operc_nets_righthemcluster
_lefthem.mat' )
load
( '/Users/dianaperez/Desktop/lateralization_code/testing_output/operc_nets_righthemcluster
_righthem.mat' )
clear all
load
( '/Users/dianaperez/Desktop/lateralization_code/testing_output/operc_nets_righthemcluster
_lefthem.mat' )
clusterA_LH_nets = RH_cluster;
load
( '/Users/dianaperez/Desktop/lateralization_code/testing_output/operc_nets_righthemcluster
_righthem.mat' )
clear all
load
( '/Users/dianaperez/Desktop/lateralization_code/testing_output/operc_nets_righthemcluster
_lefthem.mat' )
load
( '/Users/dianaperez/Desktop/lateralization_code/testing_output/operc_nets_righthemcluster
_righthem.mat' )
clusterA_RH_nets = RH_cluster;
load
( '/Users/dianaperez/Desktop/lateralization_code/testing_output/operc_nets_righthemcluster
_lefthem.mat' )
clusterA_LH_nets = RH_cluster;
load
( '/Users/dianaperez/Desktop/lateralization_code/testing_output/operc_nets_lefthemcluster_
righthem.mat' )
clusterB_RH_nets = LH_cluster;
load
( '/Users/dianaperez/Desktop/lateralization_code/testing_output/operc_nets_lefthemcluster_
lefthem.mat' )
clusterB_LH_nets = LH_cluster;
clear LH_cluster RH_cluster
rgb_colors = [1 0 0; %DMN
0 0 .6; %Vis
1 1 0; %FP
0 .8 0; %DAN
0 .6 .6; %VAN
0 0 0; %SAL
.3 0 .6; %CON
.2 1 1; %SMd
1 .5 0; %SML
.6 .2 1; %Aud
.2 1 .2; %Tpole
0 .2 .4; %MTL
0 0 1; %PMN
.8 .8 .6]; %PON
scatter(clusterA_LH_nets.number, 80, rgb_colors, 'filled')
scatter(16,clusterA_LH_nets.number, 80, rgb_colors, 'filled')
scatter(16,sum(clusterA_LH_nets.number), 80, rgb_colors, 'filled')
scatter(16,sum(clusterA_LH_nets.number,2), 80, rgb_colors, 'filled')
sum(clusterA_LH_nets.number,2)
sum(clusterA_LH_nets.number,1);
```



```
scatter(16,sum(clusterA_LH_nets.number,1), 80, rgb_colors, 'filled')
scatter(1:16,sum(clusterA_LH_nets.number,1), 80, rgb_colors, 'filled')
rgb = [1 0 0; %DMN
0 0 .6; %Vis
1 1 0; %FP
.67 .67 .67; %Unassigned
0 .8 0; %DAN
.67 .67 .67; %Unassigned2
0 .6 .6; %VAN
0 0 0; % Sal
.3 0 .6; %CON
.2 1 1; %SMd
1 .5 0; % SMI
.6 .2 1; %Aud
.2 1 .2; %Tpole
0 .2 .4; %MTL
0 0 1; %PMN
.8 .8 .6]; %PON
scatter(1:16,sum(clusterA_LH_nets.number,1), 80, rgb, 'filled')
figure
scatter(1:16,sum(clusterA_LH_nets.number,1), 80, rgb, 'filled')
close(gcf)
figure
scatter(1:16,sum(clusterA_LH_nets.number,1), 80, rgb, 'filled')
%-- 1/28/22, 8:41 PM --%
load
('/Users/dianaperez/Desktop/lateralization_code/testing_output/operc_nets_righthemcluster_
_righthem.mat')
clusterA_RH_nets = RH_cluster;
load
('/Users/dianaperez/Desktop/lateralization_code/testing_output/operc_nets_righthemcluster_
_lefthem.mat')
clusterA_LH_nets = LH_cluster;
load
('/Users/dianaperez/Desktop/lateralization_code/testing_output/operc_nets_lefthemcluster_
_righthem.mat')
clusterB_RH_nets = LH_cluster;
load
('/Users/dianaperez/Desktop/lateralization_code/testing_output/operc_nets_lefthemcluster_
_lefthem.mat')
clusterB_LH_nets = LH_cluster;
clear LH_cluster RH_cluster
rgb = [1 0 0; %DMN
0 0 .6; %Vis
1 1 0; %FP
.67 .67 .67; %Unassigned
0 .8 0; %DAN
.67 .67 .67; %Unassigned2
0 .6 .6; %VAN
0 0 0; % Sal
.3 0 .6; %CON
.2 1 1; %SMd
1 .5 0; % SMI
.6 .2 1; %Aud
.2 1 .2; %Tpole
0 .2 .4; %MTL
0 0 1; %PMN
.8 .8 .6]; %PON
figure;
scatter(1:16, RH_cluster.number, 80, rgb, 'filled')
```

```

scatter(1:16, sum(clusterA_LH_nets.number,1), 80, rgb, 'filled')
hold on
scatter(1:16, sum(clusterA_RH_nets.number,1), 80, rgb, 'filled')
close(gcf)
scatter((.5:1:15.5), sum(clusterA_LH_nets.number,1), 80, rgb, 'filled')
scatter((1.5:1:16.5), sum(clusterA_RH_nets.number,1), 80, rgb, 'filled')
hold on
scatter((.5:1:15.5), sum(clusterA_LH_nets.number,1), 80, rgb, 'filled')
close(gcf)
scatter((.5:1:15.5), sum(clusterA_LH_nets.number,1), 80, rgb, 'filled')
hold on
scatter((.5:1:15.5), sum(clusterA_LH_nets.number,1), 80, rgb, 'filled')
scatter((1.5:1:16.5), sum(clusterA_RH_nets.number,1), 80, rgb, 'filled')
close(gcf)
scatter((.75:1:15.75), sum(clusterA_LH_nets.number,1), 80, rgb, 'filled')
hold on
scatter((1.25:1:16.25), sum(clusterA_RH_nets.number,1), 80, rgb, 'filled')
network_names = {'DMN','Vis','FP',' ','DAN','↙',
',','VAN','Sal','CO','SMd','SML','Aud','Tpole','MTL','PMN','PON'};
xticklabels(network_names)
close(gcf)
rgb = [1 0 0; %DMN
0 0 .6; %Vis
1 1 0; %FP
.67 .67 .67; %Unassigned
0 .8 0; %DAN
.67 .67 .67; %Unassigned2
0 .6 .6; %VAN
0 0 0; % Sal
.3 0 .6; %CON
.2 1 1; %SMd
1 .5 0; % SML
.6 .2 1; %Aud
.2 1 .2; %Tpole
0 .2 .4; %MTL
0 0 1; %PMN
.8 .8 .6]; %PON
network_names = {'DMN','Vis','FP',' ','DAN','↙',
',','VAN','Sal','CO','SMd','SML','Aud','Tpole','MTL','PMN','PON'};
scatter((.75:1:15.75), sum(clusterA_LH_nets.number,1), 80, rgb, 'filled')
hold on
scatter((1.25:1:16.25), sum(clusterA_RH_nets.number,1), 80, rgb, 'filled')
xticks([1:16])
xticklabels(network_names)
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.3, 0.9]); %first and second↙
control position on screen, third controls width, and fourth controls height
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.9, 0.5]); %first and second↙
control position on screen, third controls width, and fourth controls height
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.7, 0.5]); %first and second↙
control position on screen, third controls width, and fourth controls height
axis([0, 250, 0, 16.5])
axis([0, 16.5, 0, 250])
axis([0.5, 16.5, 0, 250])
print(gcf, '/Users/dianaperez/Desktop/HCP384_high_asymm_clusterA_nets.jpg', '-dpng', '-↙
r300')
close(gcf)
scatter((.75:1:15.75), sum(clusterB_LH_nets.number,1), 80, rgb, 'filled')
hold on
scatter((1.25:1:16.25), sum(clusterB_RH_nets.number,1), 80, rgb, 'filled')
xticks([1:16])

```

```
xticklabels(network_names)
axis([0.5, 16.5, 0, 250])
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.7, 0.5]); %first and second
control position on screen, third controls width, and fourth controls height
print(gcf, '/Users/dianaperez/Desktop/HCP384_high_asymm_clusterB_nets.jpg', '-dpng', '-r300')
clear all
output_dir =
'/Users/dianaperez/Desktop/lateralization_code/testing_output/indiv_net_symm/';
load([output_dir '/HCP384_indiv_network_size_by_hems.mat'])
load
('/Users/dianaperez/Desktop/lateralization_code/testing_output/HCP384_LhemvsRhem_num_vert
s.mat')
load
('/Users/dianaperez/Desktop/lateralization_code/testing_output/HCP384_new_split_variants_
info.mat')
clear output_mat
numVerts = variants_info.left_hem.group_avg(:,2);
LH_numVerts = numVerts;
RH_numVerts = variants_info.right_hem.group_avg(:,2);
clear numVerts
load
('/Users/dianaperez/Desktop/lateralization_code/testing_output/HCP384_new_split_networksx
Hem.mat')
diff_numvars = networksxHem.clustersLH(:,1) - networksxHem.clustersRH(:,1);
networksxHem.clustersRH(:,1)
networksxHem.clustersRH(:,1)
diff_numvars = networksxHem.clustersLH(:,1) - networksxHem.clustersRH(:,1);
indiv_nets_size.left_hem.surf_area(:,1)
for net = 1:16
diff_numvars = networksxHem.clustersLH(:,net) - networksxHem.clustersRH(:,net);
diff_size_nets = indiv_nets_size.left_hem.surf_area(:,net) - indiv_nets_size.right_hem.
surf_area(:,net);
r = corr(diff_numvars, diff_size_nets);
end
for net = 1:16
diff_numvars = networksxHem.clustersLH(:,net) - networksxHem.clustersRH(:,net);
diff_size_nets = indiv_nets_size.left_hem.surf_area(:,net) - indiv_nets_size.right_hem.
surf_area(:,net);
r(net) = corr(diff_numvars, diff_size_nets);
end
for net = 1:16
diff_numvars = networksxHem.clustersLH(:,net) - networksxHem.clustersRH(:,net);
diff_size_nets = indiv_nets_size.left_hem.surf_area(:,net) - indiv_nets_size.right_hem.
surf_area(:,net);
r(net) = corrcoef(diff_numvars, diff_size_nets);
end
corrcoef(diff_numvars, diff_size_nets)
help corrcoef
for net = 1:16
diff_numvars = networksxHem.clustersLH(:,net) - networksxHem.clustersRH(:,net);
diff_size_nets = indiv_nets_size.left_hem.surf_area(:,net) - indiv_nets_size.right_hem.
surf_area(:,net);
[r p] = corrcoef(diff_numvars, diff_size_nets);
r_nets(net) = r;
p_nets(net) = p;
end
[r p] = corrcoef(diff_numvars, diff_size_nets);
for net = 1:16
diff_numvars = networksxHem.clustersLH(:,net) - networksxHem.clustersRH(:,net);
```

```
diff_size_nets = indiv_nets_size.left_hem.surf_area(:,net) - indiv_nets_size.right_hem.↵
surf_area(:,net);
[r p] = corrcoef(diff_numvars, diff_size_nets);
r_nets(net) = r(1,2);
p_nets(net) = p(1,2);
end
for net = 1:16
%diff_numvars = networksxHem.clustersLH(:,net) - networksxHem.clustersRH(:,net);
diff_numverts = networksxHem.verticesLH(:,net) - networksxHem.verticesRH(:,net);
diff_size_nets = indiv_nets_size.left_hem.surf_area(:,net) - indiv_nets_size.right_hem.↵
surf_area(:,net);
[r p] = corrcoef(diff_numvars, diff_size_nets);
r_nets(net) = r(1,2);
p_nets(net) = p(1,2);
end
%-- 1/29/22, 12:30 PM --%
output_dir =↵
'/Users/dianaperez/Desktop/lateralization_code/testing_output/indiv_net_symm/';
load([output_dir '/HCP384_indiv_network_size_by_hems.mat'])
load↵
('/Users/dianaperez/Desktop/lateralization_code/testing_output/HCP384_new_split_networksx↵
Hem.mat')
names = {'DMN','Vis','FP',' ','DAN','↵
','VAN','Sal','CO','SMd','SML','Aud','Tpole','MTL','PMN','PON'};
rgb = [1 0 0; %DMN
0 0 .6; %Vis
1 1 0; %FP
.67 .67 .67; %Unassigned
0 .8 0; %DAN
.67 .67 .67; %Unassigned2
0 .6 .6; %VAN
0 0 0; % Sal
.3 0 .6; %CON
.2 1 1; %SMd
1 .5 0; % SML
.6 .2 1; %Aud
.2 1 .2; %Tpole
0 .2 .4; %MTL
0 0 1; %PMN
.8 .8 .6]; %PON
scatter(1:16, indiv_nets_size.diff.surf_area, 80, rgb)
diff_surf_area = indiv_nets_size.diff.surf_area
scatter(1:16, diff_surf_area, 80, rgb)
scatter(1:16, diff_surf_area(:,,:), 80, rgb)
avg_diff = mean(diff_surf_area);
diff_surf_area(384,:) = avg_diff;
diff_surf_area = indiv_nets_size.diff.surf_area
diff_surf_area(385,:) = avg_diff;
ind = zeros(385,1);
ind(385) = 1;
net_inds = [ind ind ind ind ind ind ind ind ind ind ind ind ind ind ind];
handles = plotSpread(diff_surf_area, 'categoryMarkers', {'x', '.'}, 'categoryLabels',↵
{'Indiv_diff_scores','Average'}, 'distributionColors', rgb, 'categoryIdx', net_inds,↵
'xNames', names)
for n = 1:numel(network_names)
rgb_for_plot{n} = rgb_colors(n,:);
end
for n = 1:numel(names)
rgb_for_plot{n} = rgb(n,:);
end
```

```
handles = plotSpread(diff_surf_area, 'categoryMarkers', {'x', '.'}, 'categoryLabels',  
{'Indiv_diff_scores','Average'}, 'distributionColors', rgb_for_plot, 'categoryIdx',  
net_inds, 'xNames', names)  
hold on  
scatter(1:16, avg_diff, 'MarkerEdgeColor', 'k', 'MarkerFaceColor', 'k', 'SizeData', 50)  
scatter(8,avg_diff(1,6), 'MarkerEdgeColor', 'w', 'MarkerFaceColor', 'w', 'SizeData', 50)  
ylabel('Difference in surface area of individual-specific networks')  
load  
( '/Users/dianaperez/Desktop/lateralization_code/testing_output/HCP384_new_split_networksx  
Hem.mat')  
diff_numverts = networksxHem.verticesLH(:,net) - networksxHem.verticesRH(:,net);  
for net = 1:16  
diff_numvars = networksxHem.clustersLH(:,net) - networksxHem.clustersRH(:,net);  
%diff_numverts = networksxHem.verticesLH(:,net) - networksxHem.verticesRH(:,net);  
diff_size_nets = indiv_nets_size.left_hem.surf_area(:,net) - indiv_nets_size.right_hem.  
surf_area(:,net);  
[r p] = corrcoef(diff_numvars, diff_surf_area(384,:));  
r_nets(net) = r(1,2);  
p_nets(net) = p(1,2);  
end  
indiv_net_size_by_hems  
for net = 1:16  
diff_numvars = networksxHem.clustersLH(:,net) - networksxHem.clustersRH(:,net);  
%diff_numverts = networksxHem.verticesLH(:,net) - networksxHem.verticesRH(:,net);  
diff_size_nets = indiv_nets_size.left_hem.surf_area(:,net) - indiv_nets_size.right_hem.  
surf_area(:,net);  
[r p] = corrcoef(diff_numvars, diff_size_nets);  
r_nets(net) = r(1,2);  
p_nets(net) = p(1,2);  
end  
load  
( '/Users/dianaperez/Desktop/lateralization_code/testing_output/HCP384_new_split_networksx  
Hem.mat')  
for net = 1:16  
diff_numvars = networksxHem.clustersLH(:,net) - networksxHem.clustersRH(:,net);  
%diff_numverts = networksxHem.verticesLH(:,net) - networksxHem.verticesRH(:,net);  
diff_size_nets = indiv_nets_size.left_hem.surf_area(:,net) - indiv_nets_size.right_hem.  
surf_area(:,net);  
[r p] = corrcoef(diff_numvars, diff_size_nets);  
r_nets(net) = r(1,2);  
p_nets(net) = p(1,2);  
end  
load([output_dir ' /HCP384_indiv_network_size_by_hems.mat'])  
for net = 1:16  
diff_numvars = networksxHem.clustersLH(:,net) - networksxHem.clustersRH(:,net);  
%diff_numverts = networksxHem.verticesLH(:,net) - networksxHem.verticesRH(:,net);  
diff_size_nets = indiv_nets_size.left_hem.surf_area(:,net) - indiv_nets_size.right_hem.  
surf_area(:,net);  
[r p] = corrcoef(diff_numvars, diff_size_nets);  
r_nets(net) = r(1,2);  
p_nets(net) = p(1,2);  
end  
plot(diff_numvars, diff_size_nets)  
close gcf  
plot(diff_numvars, diff_size_nets)  
scatter(diff_numvars, diff_size_nets)  
help polyfit  
hold on  
p = polyfit(diff_numvars, diff_size_nets,1);  
px = [min(diff_numvars) max(diff_numvars)];
```

```

py = polyval(p, px);
plot(px, py, 'LineWidth', 2);
help corr
[r p] = corr(diff_numvars, diff_size_nets)
indiv_net_size_by_hems
title(names{net});
indiv_net_size_by_hems
%-- 1/29/22, 3:48 PM --%
indiv_net_size_by_hems
high_asymmetry_clusters
load('HCP384_variant_maps.mat')
unique(variant_map_mat)
gifti = insert_nonbrain(variant_map_mat, 'both', template);
gifti = insert_nonbrain(variant_map_mat(:, :), 'both', template);
open insert_nonbrain
gifti = insert_nonbrain(variant_map_mat(:, :), 'both', template);
gifti = insert_nonbrain(variant_map_mat, 'both', template);
high_asymmetry_clusters
[data_dir num2str(subs(sub)) netvarmap_str]
high_asymmetry_clusters
unique_IDs_LH = unique_IDs_mat(1:32492, :);
unique_IDs_RH = unique_IDs_mat(32493:end, :);
net_assign_LH = net_assign_mat(1:32492, :);
net_assign_RH = net_assign_mat(32493:end, :);
length(unique_IDs_LH)
size(unique_IDs_LH, 2)
clusterA = [];
clusterA.left_hem.proportion = length(unique_IDs_LH);
clusterA.left_hem.proportion(1) = length(unique_IDs_LH);
high_asymmetry_clusters
for sub = 1:size(unique_IDs_LH, 2)
%% LEFT HEM
[subInfo, clusterA.left_hem.proportion(sub, :), clusterA.left_hem.number(sub, :)] = \
extract_nets(unique_IDs_LH, net_assign_LH, clusterA_seed);
clusterA_LH = [clusterA_LH; subInfo];
[subInfo, clusterA.right_hem.proportion(sub, :), clusterA.right_hem.number(sub, :)] = \
extract_nets(unique_IDs_RH, net_assign_RH, clusterA_seed);
clusterA_RH = [clusterA_RH; subInfo];
[subInfo, clusterB.left_hem.proportion(sub, :), clusterB.left_hem.number(sub, :)] = \
extract_nets(unique_IDs_LH, net_assign_LH, clusterB_seed);
clusterB_LH = [clusterB_LH; subInfo];
[subInfo, clusterB.right_hem.proportion(sub, :), clusterB.right_hem.number(sub, :)] = \
extract_nets(unique_IDs_RH, net_assign_RH, clusterB_seed);
clusterB_RH = [clusterB_RH; subInfo];
end
high_asymmetry_clusters
close gcf
network_names = {'DMN', 'Vis', 'FP', ' ', 'DAN', ' \
', 'VAN', 'Sal', 'CO', 'SMd', 'SMl', 'Aud', 'Tpole', 'MTL', 'PMN', 'PON'};
scatter((.75:1:15.75), sum(clusterA.left_hem.number, 1), 80, rgb, 'filled')
hold on
scatter((1.25:1:16.25), sum(clusterA.right_hem.number, 1), 80, rgb, 'filled')
xticks([1:16])
xticklabels(network_names)
axis([0.5, 16.5, 0, 250])
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.7, 0.5]); %first and second \
control position on screen, third controls width, and fourth controls height
print(gcf, '/Users/dianaperez/Desktop/HCP384_high_asymm_clusterA_nets.jpg', '-dpng', '-\
r300')
close gcf

```

```
scatter((.75:1:15.75), sum(clusterB.left_hem.number,1), 80, rgb, 'filled')
hold on
scatter((1.25:1:16.25), sum(clusterB.right_hem.number,1), 80, rgb, 'filled')
xticks([1:16])
xticklabels(network_names)
axis([0.5, 16.5, 0, 250])
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.7, 0.5]); %first and second
control position on screen, third controls width, and fourth controls height
print(gcf, '/Users/dianaperez/Desktop/HCP384_high_asymm_clusterB_nets.jpg', '-dpng', '-r300')
close gcf
scatter((.75:1:15.75), sum(clusterA.left_hem.number,1), 80, rgb, 'filled')
hold on
scatter((1.25:1:16.25), sum(clusterA.right_hem.number,1), 80, rgb, 'filled')
xticks([1:16])
xticklabels(network_names)
max_ax = max(clusterA.left_hem.number)+(std(clusterA.left_hem.number)/2);
min_ax = min(clusterA.left_hem.number)-(std(clusterA.left_hem.number)/2);
max_ax = max(max(clusterA.left_hem.number)+(min((std(clusterA.left_hem.number)/2)));
max_ax = max(max(clusterA.left_hem.number)+(min((std(clusterA.left_hem.number)/2)));
max_ax = max(max(sum(clusterA.left_hem.number,1))+(min((std(sum(clusterA.left_hem.
number))/2)));
max_ax = max(max(sum(clusterA.left_hem.number,1))+(min((std(sum(clusterA.left_hem.
number))/2)));
min_ax = min(min(sum(clusterA.left_hem.number,1))-(min((std(sum(clusterA.left_hem.
number))/2)));
axis([0.5, 16.5, min_ax, max_ax])
close gcf
scatter((.75:1:15.75), sum(clusterA.left_hem.number,1), 80, rgb, 'filled')
hold on
scatter((1.25:1:16.25), sum(clusterA.right_hem.number,1), 80, rgb, 'filled')
xticks([1:16])
xticklabels(network_names)
max_ax = max(max(sum(clusterA.left_hem.number,1))+(min((std(sum(clusterA.left_hem.
number))/2)));
min_ax = min(min(sum(clusterA.left_hem.number,1))-(min((std(sum(clusterA.left_hem.
number))/2)));
axis([0.5, 16.5, min_ax, max_ax])
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.7, 0.5]); %first and second
control position on screen, third controls width, and fourth controls height
print(gcf, '/Users/dianaperez/Desktop/HCP384_high_asymm_clusterA_nets.jpg', '-dpng', '-r300')
close gcf
scatter((.75:1:15.75), sum(clusterA.left_hem.number,1), 80, rgb)
hold on
scatter((1.25:1:16.25), sum(clusterA.right_hem.number,1), 80, rgb, 'filled')
xticks([1:16])
xticklabels(network_names)
max_ax = max(max(sum(clusterA.left_hem.number,1))+(min((std(sum(clusterA.left_hem.
number))/2)));
min_ax = min(min(sum(clusterA.left_hem.number,1))-(min((std(sum(clusterA.left_hem.
number))/2)));
axis([0.5, 16.5, min_ax, max_ax])
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.7, 0.5]); %first and second
control position on screen, third controls width, and fourth controls height
print(gcf, '/Users/dianaperez/Desktop/HCP384_high_asymm_clusterA_nets.jpg', '-dpng', '-r300')
```



```
scatter((.75:1:15.75), sum(clusterA.left_hem.number,1), 80, rgb, 'd', 'filled')
scatter((.75:1:15.75), sum(clusterA.left_hem.number,1), 80, rgb, 'd', 'filled')
hold on
scatter((1.25:1:16.25), sum(clusterA.right_hem.number,1), 80, rgb, 'filled')
xticks([1:16])
xlabel(network_names)
max_ax = max(max(sum(clusterA.left_hem.number,1)))+(min((std(sum(clusterA.left_hem.
number))/2)));
min_ax = min(min(sum(clusterA.left_hem.number,1)))-(min((std(sum(clusterA.left_hem.
number))/2)));
axis([0.5, 16.5, min_ax, max_ax])
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.7, 0.5]); %first and second
control position on screen, third controls width, and fourth controls height
print(gcf, '/Users/dianaperez/Desktop/HCP384_high_asymm_clusterA_nets.jpg', '-dpng', '-
r300')
close gcf
scatter((.75:1:15.75), sum(clusterB.left_hem.number,1), 80, rgb, 'd', 'filled')
hold on
scatter((1.25:1:16.25), sum(clusterB.right_hem.number,1), 80, rgb, 'filled')
xticks([1:16])
xlabel(network_names)
max_ax = max(max(sum(clusterB.left_hem.number,1)))+(min((std(sum(clusterB.left_hem.
number))/2)));
min_ax = min(min(sum(clusterB.left_hem.number,1)))-(min((std(sum(clusterB.left_hem.
number))/2)));
axis([0.5, 16.5, 0, 250])
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.7, 0.5]); %first and second
control position on screen, third controls width, and fourth controls height
print(gcf, '/Users/dianaperez/Desktop/HCP384_high_asymm_clusterB_nets.jpg', '-dpng', '-
r300')
max_ax = max(max(sum(clusterB.left_hem.number,1)))+(min((std(sum(clusterB.left_hem.
number))/2)));
min_ax = min(min(sum(clusterB.left_hem.number,1)))-(min((std(sum(clusterB.left_hem.
number))/2)));
axis([0.5, 16.5, 0, 250])
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.7, 0.5]); %first and second
control position on screen, third controls width, and fourth controls height
print(gcf, '/Users/dianaperez/Desktop/HCP384_high_asymm_clusterB_nets.jpg', '-dpng', '-
r300')
axis([0.5, 16.5, min_ax, max_ax])
print(gcf, '/Users/dianaperez/Desktop/HCP384_high_asymm_clusterB_nets.jpg', '-dpng', '-
r300')
close gcf
scatter((.75:1:15.75), sum(clusterB.left_hem.number,1), 80, rgb, 'd', 'filled')
hold on
scatter((1.25:1:16.25), sum(clusterB.right_hem.number,1), 80, rgb, 'filled')
xticks([1:16])
xlabel(network_names)
max_ax = max(max(sum(clusterB.left_hem.number,1)))+(min((std(sum(clusterB.left_hem.
number))/2)));
min_ax = min(min(sum(clusterB.left_hem.number,1)))-(min((std(sum(clusterB.left_hem.
number))/2)));
axis([0.5, 16.5, min_ax, max_ax])
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.7, 0.5]); %first and second
control position on screen, third controls width, and fourth controls height
print(gcf, '/Users/dianaperez/Desktop/HCP384_high_asymm_clusterB_nets.jpg', '-dpng', '-
r300')
scatter((.75:1:15.75), sum(clusterA.left_hem.proportion,1), 80, rgb, 'd', 'filled')
hold on
scatter((1.25:1:16.25), sum(clusterA.right_hem.proportion,1), 80, rgb, 'filled')
```



```

xticks([1:16])
xticklabels(network_names)
max_ax = max(max(sum(clusterA.left_hem.proportion,1)))+(min((std(sum(clusterA.left_hem.
proportion))/2)));
min_ax = min(min(sum(clusterA.left_hem.proportion,1)))-(min((std(sum(clusterA.left_hem.
proportion))/2)));
axis([0.5, 16.5, min_ax, max_ax])
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.7, 0.5]); %first and second
control position on screen, third controls width, and fourth controls height
print(gcf, '/Users/dianaperez/Desktop/HCP384_high_asymm_clusterA_nets_proportion.jpg', '-
dpng', '-r300')
var_freq_size_net
surf_areas_LHem(verts)
sum(surf_areas_LHem(verts))
sum(surf_areas_LHem)
sum(surf_areas_LHem(verts))/sum(surf_areas_LHem)
272.1685/6.3782e+04
var_freq_size_net
perms_var_freq_size
permute_net_assignment
high_asymmetry_clusters
scatter((.75:1:13.75), sum(good_nets_LH,1), 150, rgb, 'd', 'filled')
hold on
scatter((1.25:1:14.25), sum(good_nets_RH,1), 150, rgb, 'filled')
xticks([1:16])
xticklabels(network_names)
max_ax = max(max(sum(good_nets_LH,1)))+(min((std(sum(good_nets_LH))/2)));
min_ax = min(min(sum(good_nets_LH,1)))-(min((std(sum(good_nets_LH))/2)));
axis([0.5, 16.5, min_ax, max_ax])
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.7, 0.5]); %first and second
control position on screen, third controls width, and fourth controls height
print(gcf, '/Users/dianaperez/Desktop/HCP384_high_asymm_clusterA_nets.jpg', '-dpng', '-
r300')
xticks([1:14])
axis([0.5, 14.5, min_ax, max_ax])
print(gcf, '/Users/dianaperez/Desktop/HCP384_high_asymm_clusterA_nets.jpg', '-dpng', '-
r300')
close gcf
good_nets_LH = [clusterB.left_hem.number(:,1:3) clusterB.left_hem.number(:,5) clusterB.
left_hem.number(:,7:end)];
good_nets_RH = [clusterB.right_hem.number(:,1:3) clusterB.right_hem.number(:,5) clusterB.
right_hem.number(:,7:end)];
scatter((.75:1:13.75), sum(good_nets_LH,1), 150, rgb, 'd', 'filled')
hold on
scatter((1.25:1:14.25), sum(good_nets_RH,1), 150, rgb, 'filled')
xticks([1:16])
xticklabels(network_names)
max_ax = max(max(sum(good_nets_LH,1)))+(min((std(sum(good_nets_LH))/2)));
min_ax = min(min(sum(good_nets_LH,1)))-(min((std(sum(good_nets_LH))/2)));
axis([0.5, 16.5, min_ax, max_ax])
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.7, 0.5]); %first and second
control position on screen, third controls width, and fourth controls height
print(gcf, '/Users/dianaperez/Desktop/HCP384_high_asymm_clusterB_nets.jpg', '-dpng', '-
r300')
xticks([1:14])
axis([0.5, 14.5, min_ax, max_ax])
max_ax = max(max(sum(good_nets_RH,1)))+(min((std(sum(good_nets_RH))/2)));
min_ax = min(min(sum(good_nets_RH,1)))-(min((std(sum(good_nets_RH))/2)));
axis([0.5, 14.5, min_ax, max_ax])
max_ax = max(max(sum(good_nets_RH,1)))+(min((std(sum(good_nets_RH))/2)));

```

```
min_ax = min(min(sum(good_nets_RH,1)))-(min((std(sum(good_nets_RH))/2)));
axis([0.5, 14.5, min_ax, max_ax])
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.7, 0.5]); %first and second
control position on screen, third controls width, and fourth controls height
print(gcf, '/Users/dianaperez/Desktop/HCP384_high_asymm_clusterB_nets.jpg', '-dpng', '-r300')
ax = gca;
ax.FontSize = 24;
print(gcf, '/Users/dianaperez/Desktop/HCP384_high_asymm_clusterB_nets.jpg', '-dpng', '-r300')
close gcf
network_names =
{'DMN', 'Vis', 'FP', 'DAN', 'VAN', 'Sal', 'CO', 'SMd', 'SMl', 'Aud', 'Tpole', 'MTL', 'PMN', 'PON'};
good_nets_LH = [clusterA.left_hem.number(:,1:3) clusterA.left_hem.number(:,5) clusterA.
left_hem.number(:,7:end)];
good_nets_RH = [clusterA.right_hem.number(:,1:3) clusterA.right_hem.number(:,5) clusterA.
right_hem.number(:,7:end)];
scatter((.75:1:13.75), sum(good_nets_LH,1), 150, rgb, 'd', 'filled')
hold on
scatter((1.25:1:14.25), sum(good_nets_RH,1), 150, rgb, 'filled')
xticks([1:14])
xticklabels(network_names)
max_ax = max(max(sum(good_nets_LH,1)))+(min((std(sum(good_nets_LH))/2)));
min_ax = min(min(sum(good_nets_LH,1)))-(min((std(sum(good_nets_LH))/2)));
axis([0.5, 14.5, min_ax, max_ax])
ax = gca;
ax.FontSize = 24;
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.7, 0.5]); %first and second
control position on screen, third controls width, and fourth controls height
print(gcf, '/Users/dianaperez/Desktop/HCP384_high_asymm_clusterA_nets.jpg', '-dpng', '-r300')
close gcf
good_nets_LH = [clusterB.left_hem.number(:,1:3) clusterB.left_hem.number(:,5) clusterB.
left_hem.number(:,7:end)];
good_nets_RH = [clusterB.right_hem.number(:,1:3) clusterB.right_hem.number(:,5) clusterB.
right_hem.number(:,7:end)];
scatter((.75:1:13.75), sum(good_nets_LH,1), 150, rgb, 'd', 'filled')
hold on
scatter((1.25:1:14.25), sum(good_nets_RH,1), 150, rgb, 'filled')
xticks([1:14])
xticklabels(network_names)
max_ax = max(max(sum(good_nets_RH,1)))+(min((std(sum(good_nets_RH))/2)));
min_ax = min(min(sum(good_nets_RH,1)))-(min((std(sum(good_nets_RH))/2)));
axis([0.5, 14.5, min_ax, max_ax])
ax = gca;
ax.FontSize = 24;
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.7, 0.5]); %first and second
control position on screen, third controls width, and fourth controls height
print(gcf, '/Users/dianaperez/Desktop/HCP384_high_asymm_clusterB_nets.jpg', '-dpng', '-r300')
Overlap_VANvars
permute_net_assignment
var_freq_size_net
permute_net_assignment
indiv_net_size_by_hems
diff_numvars_all = [];
diff_sizenets_all = [];
for net = 1:16
%diff_numvars = networksxHem.clustersLH(:,net) - networksxHem.clustersRH(:,net);
diff_numvars = networksxHem.verticesLH(:,net) - networksxHem.verticesRH(:,net);
```

```

diff_size_nets = indiv_nets_size.left_hem.surf_area(:,net) - indiv_nets_size.right_hem.↵
surf_area(:,net);
diff_numvars_all = [diff_numvars_all; diff_numvars];
diff_sizenets_all = [diff_sizenets_all; diff_size_nets];
[r p] = corr(diff_numvars, diff_size_nets);
rvals(net) = r;
pvals(net) = p;
end
[p_fdr, p_masked] = FDR(pvals, 0.025)
[r_all, p_all] = corr(diff_numvars_all, diff_sizenets_all);
%-- 1/31/22, 6:08 PM --%
indiv_net_size_by_hems
find(diff_numvars==nan)
find(diff_size_nets==nan)
unique
unique(diff_numvars)
unique(diff_size_nets)
unique(diff_surf_area)
find(isnan(diff_numvars))
find(isnan(diff_size_nets))
[r p] = corrcoef(diff_numvars, diff_size_nets);
[r p] = corr(diff_numvars, diff_size_nets, 'rows', 'complete')
doc corr
unique(diff_numvars)
permute_net_assignment
[p_fdr p_masked] = FDR(p_vals, .025);
permute_net_assignment
load↵
('/Users/dianaperez/Desktop/Lateralization/Variant_Properties/HCP752_new_split_RH_network↵
sxHem.mat')
load↵
('/Users/dianaperez/Desktop/Lateralization/Variant_Properties/HCP752_new_split_LH_network↵
sxHem.mat')
LHand_info = load↵
('/Users/dianaperez/Desktop/Lateralization/Variant_Properties/HCP752_new_split_LH_network↵
sxHem.mat')
RHand_info = load↵
('/Users/dianaperez/Desktop/Lateralization/Variant_Properties/HCP752_new_split_RH_network↵
sxHem.mat')
permute_net_assignment
LHand_info = load↵
('/Users/dianaperez/Desktop/Lateralization/Variant_Properties/HCP752_new_split_LH_network↵
sxHem.mat')
RHand_info = load↵
('/Users/dianaperez/Desktop/Lateralization/Variant_Properties/HCP752_new_split_RH_network↵
sxHem.mat')
%-- 1/31/22, 8:19 PM --%
permute_net_assignment
[p_fdr p_masked] = FDR(p_vals, .025);
permute_net_assignment
%-- 2/2/22, 10:05 PM --%
net_specific_diffmap
[varmap_loc num2str(LH(n,1)) net_assign_str]
net_specific_diffmap
size(LHand_net_assign)
net_specific_diffmap
[output_dir group_str{group} '_' network_names{network_id} '_Variants_Overlap.dtseries.↵
nii']
net_specific_diffmap
unique(LH_overlap_map)

```

```

unique(Rh_overlap_map)
unique(RH_overlap_map)
unique9
unique(diff_map)
unique(LH_overlap_map)
unique(RH_overlap_map)
permute_net_assignment
addpath '/Users/dianaperez/Documents/GitHub/PerezEtAl_HemAsymmetries/needed_files'
[p_fdr p_masked] = FDR(p_vals, .025);
network_names = ␣
{'DMN', 'Vis', 'FP', 'DAN', 'VAN', 'Sal', 'CO', 'SMd', 'SML', 'Aud', 'Tpole', 'MTL', 'PMN', 'PON'};
rgb_colors = [1 0 0; %DMN
0 0 .6; %Vis
1 1 0; %FP
0 .8 0; %DAN
0 .6 .6; %VAN
0 0 0; % Sal
.3 0 .6; %CON
.2 1 1; %SMd
1 .5 0; % SML
.6 .2 1; %Aud
.2 1 .2; %Tpole
0 .2 .4; %MTL
0 0 1; %PMN
.8 .8 .6]; %PON
good_nets = [output_mat(:,1:3) output_mat(:,5) output_mat(:,7:end)];
rgb_for_plot = {};
for n = 1:numel(network_names)
    rgb_for_plot{n} = rgb_colors(n,:);
end
ind = zeros(numperms+1,1);
ind(1,1) = 1;
net_inds = [ind ind ind ind ind ind ind ind ind ind ind ind ind ind];
handles = plotSpread(good_nets, 'categoryMarkers', {'x', '.'}, 'categoryLabels', ␣
{'Permuted Differences', 'True Difference'}, 'distributionColors', rgb_for_plot', ␣
'categoryIdx', net_inds, 'xNames', network_names)
ax = gca;
ax.FontSize = 24;
scatter([1:14],good_nets(1,:), 'MarkerEdgeColor', 'k', 'MarkerFaceColor', 'k', ␣
'SizeData', 50)
scatter(6,good_nets(1,6), 'MarkerEdgeColor', 'w', 'MarkerFaceColor', 'w', 'SizeData', 50)
ylabel('Difference in Number of Variants')
xlabel('Assigned Network')
min_ax = min(min(good_nets(:,:)))-(min(std(good_nets)/2));
max_ax = max(max(good_nets(:,:)))+(min(std(good_nets)/2));
axis([0, 15, min_ax, max_ax])
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.9, 0.7]);
print(gcf,[output_dir out_str '_PermutationTesting_networkAssignment.jpg'],'-dpng','-␣
r300');
perms_var_freq_size
permute_net_assignment
indiv_net_size_by_hems
net_specific_diffmap
numsubs = size(LHand_net_assign/1);
numsubs = size(LHand_net_assign,1);
numsubs = size(LHand_net_assign,2);
net_specific_diffmap
indiv_net_size_by_hems
p = polyfit(diff_numvars, diff_size_nets,1);
px = [min(diff_numvars) max(diff_numvars)];

```

```
py = polyval(p, px);
plot(px, py, 'LineWidth', 2);
title(names{net});
scatter(diff_numvars, diff_size_nets)
hold on
p = polyfit(diff_numvars, diff_size_nets,1);
px = [min(diff_numvars) max(diff_numvars)];
py = polyval(p, px);
plot(px, py, 'LineWidth', 2);
title(names{net});
indiv_net_size_by_hems
diff_numvars = networksxHem.verticesLH(:,net) - networksxHem.verticesRH(:,net);
diff_size_nets = indiv_nets_size.left_hem.surf_area(:,net) - indiv_nets_size.right_hem.
surf_area(:,net);
diff_numvars_all = [diff_numvars_all; diff_numvars];
diff_sizenets_all = [diff_sizenets_all; diff_size_nets];
[r p] = corr(diff_numvars, diff_size_nets);
rvals(net) = r;
pvals(net) = p;
scatter(diff_numvars, diff_size_nets)
hold on
p = polyfit(diff_numvars, diff_size_nets,1);
px = [min(diff_numvars) max(diff_numvars)];
py = polyval(p, px);
plot(px, py, 'LineWidth', 2);
title(names{net});
close(gcf)
diff_numvars = networksxHem.verticesLH(:,net) - networksxHem.verticesRH(:,net);
diff_size_nets = indiv_nets_size.left_hem.surf_area(:,net) - indiv_nets_size.right_hem.
surf_area(:,net);
diff_numvars_all = [diff_numvars_all; diff_numvars];
diff_sizenets_all = [diff_sizenets_all; diff_size_nets];
[r p] = corr(diff_numvars, diff_size_nets);
rvals(net) = r;
pvals(net) = p;
scatter(diff_numvars, diff_size_nets)
hold on
p = polyfit(diff_numvars, diff_size_nets,1);
px = [min(diff_numvars) max(diff_numvars)];
py = polyval(p, px);
plot(px, py, 'LineWidth', 2);
title(names{net});
xlabel('Difference in number of Variant Vertices (Left-Right Hem)');
ylabel('Difference in Network Surface Area in mm^2 (Left-Right Hem)');
diff_numvars = networksxHem.clustersLH(:,net) - networksxHem.clustersRH(:,net);
%diff_numvars = networksxHem.verticesLH(:,net) - networksxHem.verticesRH(:,net);
diff_size_nets = indiv_nets_size.left_hem.surf_area(:,net) - indiv_nets_size.right_hem.
surf_area(:,net);
diff_numvars_all = [diff_numvars_all; diff_numvars];
diff_sizenets_all = [diff_sizenets_all; diff_size_nets];
[r p] = corr(diff_numvars, diff_size_nets);
rvals(net) = r;
pvals(net) = p;
scatter(diff_numvars, diff_size_nets)
hold on
p = polyfit(diff_numvars, diff_size_nets,1);
px = [min(diff_numvars) max(diff_numvars)];
py = polyval(p, px);
plot(px, py, 'LineWidth', 2);
title(names{net});
```

```

xlabel('Difference in number of Variant Regions (Left-Right Hem)');
ylabel('Difference in Network Surface Area in mm^2(Left-Right Hem)');
close(gcf)
diff_numvars = networksxHem.clustersLH(:,net) - networksxHem.clustersRH(:,net);
%diff_numvars = networksxHem.verticesLH(:,net) - networksxHem.verticesRH(:,net);
diff_size_nets = indiv_nets_size.left_hem.surf_area(:,net) - indiv_nets_size.right_hem.
surf_area(:,net);
diff_numvars_all = [diff_numvars_all; diff_numvars];
diff_sizenets_all = [diff_sizenets_all; diff_size_nets];
[r p] = corr(diff_numvars, diff_size_nets);
rvals(net) = r;
pvals(net) = p;
scatter(diff_numvars, diff_size_nets)
hold on
p = polyfit(diff_numvars, diff_size_nets,1);
px = [min(diff_numvars) max(diff_numvars)];
py = polyval(p, px);
plot(px, py, 'LineWidth', 2);
title(names{net});
xlabel('Difference in number of Variant Regions (Left-Right Hem)');
ylabel('Difference in Network Surface Area in mm^2(Left-Right Hem)');
indiv_net_size_by_hems
[r,p] = corr(diff_numvars_all, diff_sizenets_all)
indiv_net_size_by_hems
al_goodplot(diff_surf_area, [1:16], .5, rgb_for_plot, 'bilateral', [], [])
al_goodplot(diff_surf_area, [1:16], .5, rgb_for_plot, 'bilateral')
al_goodplot(diff_surf_area, [1:16], .5, rgb, 'bilateral')
close(gcf)
al_goodplot(diff_surf_area, [1:2:32], .5, rgb, 'bilateral')
ylabel('Difference in surface area of individual-specific networks')
close(gcf)
al_goodplot(diff_surf_area, [1:2:32], .5, rgb, 'bilateral')
close(gcf)
al_goodplot(diff_surf_area(:,1), [ ], .5, rgb(1,:), 'bilateral')
al_goodplot
permute_net_assignment
DMN_verts = allSubs_info.networksxHem.verticesLH - allSubs_info.networksxHem.verticesRH;
close(gcf)
al_goodplot(DMN_verts(:,1), [ ], .5, rgb(1,:), 'bilateral')
rgb = [1 0 0; %DMN
0 0 .6; %Vis
1 1 0; %FP
.67 .67 .67; %Unassigned
0 .8 0; %DAN
.67 .67 .67; %Unassigned2
0 .6 .6; %VAN
0 0 0; % Sal
.3 0 .6; %CON
.2 1 1; %SMd
1 .5 0; % SMI
.6 .2 1; %Aud
.2 1 .2; %Tpole
0 .2 .4; %MTL
0 0 1; %PMN
.8 .8 .6]; %PON
al_goodplot(DMN_verts(:,1), [ ], .5, rgb(1,:), 'bilateral')
indiv_net_size_by_hems
[r,p] = corr(diff_numvars_all, diff_sizenets_all)
permute_net_assignment
diff = left - right;

```

```

mean(diff)
permute_net_assignment
var_freq_size_net
sessions = [4:8, 6:10, 6:10, 1:5, 1:5, 1:5];
sessions(2)
sessions = [4:8; 6:10; 6:10; 1:5; 1:5; 1:5];
length(sessions(2))
length(sessions(2))
length(sessions(2,:))
similarity_analysis_forced_same_amt_data
%-- 2/4/22, 3:49 PM --%
similarity_analysis_forced_same_amt_data
figure('Position',[1 1 1000 800]);
imagesc(simmat,[0 1]); colormap('jet');
hline_new([5,10,15,20,25,30]+0.5,'k',2);
vline_new([5,10,15,20,25,30]+0.5,'k',2);
set(gca,'XTick',[4,13.5,19,24.5,30.5,34,37,40,45], 'YTick',↵
[4,13.5,19,24.5,30.5,34,37,40,45], 'XTickLabel',...
{'LS02', 'LS03', 'LS04', 'LS05', 'LS07', 'LS08', 'LS10', 'LS11', 'LS14'}, 'YTickLabel',↵
{'LS02', 'LS03', 'LS04', 'LS05', 'LS07', 'LS08', 'LS10', 'LS11', 'LS14'});
set(gca,'XTick',[3, 8, 13, 18, 23, 28], 'YTick', [3, 8, 13, 18, 23, 28], 'XTickLabel',...
{'LS02', 'LS03', 'LS05', 'LS08', 'LS11', 'LS14'}, 'YTickLabel', {'LS02', 'LS03', 'LS05',↵
'LS08', 'LS11', 'LS14'});
axis square;
colorbar;
title('Correlation Matrix Similarity');
saveas(gcf,[outDir 'SimilarityMat_rest_matched_data.tiff'],'tiff');
outDir = '/Users/dianaperez/Desktop/';
saveas(gcf,[outDir 'SimilarityMat_rest_matched_data.tiff'],'tiff');
close all
%-- 2/8/22, 10:01 AM --%
indiv_net_size_by_hems
numvars = networksxHem.clustersLH(:,net) + networksxHem.clustersRH(:,net);
size_nets = indiv_nets_size.left_hem.surf_area(:,net) + indiv_nets_size.right_hem.↵
surf_area(:,net);
indiv_net_size_by_hems
[r p] = corr(diff_numvars_all, diff_size_nets_all)
[r p] = corr(diff_numvars_all, diff_sizenets_all)
scatter(diff_numvars_all, diff_sizenets_all)
hold on
p = polyfit(diff_numvars_all, diff_sizenets_all,1);
px = [min(diff_numvars_all) max(diff_numvars_all)];
py = polyval(p, px);
plot(px, py, 'LineWidth', 2);
indiv_net_size_by_hems
diff_surf_area(:,6) = [];
indiv_net_size_by_hems
close gcf
indiv_net_size_by_hems
title('All networks (Left Hem)');
title('All networks (Left Hem)');
indiv_net_size_by_hems
EmReg_task
PsychHID('Devices')
EmReg_task
sca
PsychHID('Devices')
EmReg_task
sca
%-- 2/8/22, 1:01 PM --%

```

```

EmReg_task
sca
%-- 2/8/22, 1:02 PM --%
EmReg_task
sca
clear PsychHID
clear KbCheck
EmReg_task
sca
%-- 2/8/22, 1:06 PM --%
EmReg_task
DrawFormattedText(window, emotions_table{emo,1}, xCenter/2, (yCenter-(yCenter/3)), -1);
DrawFormattedText(window, emotions_table{emo,1}, xCenter/4, (yCenter-(yCenter/3)), -1);
DrawFormattedText(window, emotions_table{emo,1}, xCenter/2, (yCenter-(yCenter/3)), -1);
Screen('Flip', window);
Screen('Close');
Screen('Close');
Screen('Close', window);
DrawFormattedText(window, emotions_table{emo,1}, xCenter/4, (yCenter-(yCenter/3)), -1);
EmReg_task
DrawFormattedText(window, emotions_table{emo,1}, xCenter/2, (yCenter-(yCenter/3)), -1);
DrawFormattedText(window, emotions_table{emo,3}, xCenter/2, yCenter, 0);
Screen('Flip', window);
DrawFormattedText(window, emotions_table{emo,1}, 'center', 'center', -1);
Screen('Flip', window);
DrawFormattedText(window, emotions_table{emo,1}, 'center', 'center'/3, -1);
DrawFormattedText(window, emotions_table{emo,1}, 'center', (yCenter-(yCenter/3)), -1);
DrawFormattedText(window, emotions_table{emo,3}, 'center', yCenter, 0);
Screen('Flip', window);
number = GetEchoNumber(window, 'Enter a number, then press RETURN', 'center', yCenter, \
[0,0,0], [], -1)
EmReg_task
GetEchoNumber
EmReg_task
double(output)
x = 0;
Screen('DrawText', 10, double(output), 0, 540, [0,0,0], [1,1,1])
Screen('Flip', 10, 0, 1)
Screen('DrawText', 10, double(output), 540, 540, [0,0,0], [1,1,1])
Screen('Flip', 10, 0, 1)
Screen('DrawText', 10, double(output), 550, 540, [0,0,0], [1,1,1])
Screen('Flip', 10, 0, 1)
Screen('DrawText', 10, double(output), 580, 540, [0,0,0], [1,1,1])
Screen('Flip', 10, 0, 1)
Screen('DrawText', 10, double(output), 600, 540, [0,0,0], [1,1,1])
Screen('Flip', 10, 0, 1)
x = 630;
bgColor = [1,1,1];
78
EmReg_task
7
8
4
3
4
5
6
7
sca
EmReg_task

```



```
6
4
6
3
2
5
6
3
4
5
6
3
4
5
6
sca
EmReg_task
Screen('DrawTexture', window, scale, [], [xCenter (yCenter+(yCenter/3))], 0); %change↵
location to bottom
EmReg_task
%-- 2/9/22, 12:01 PM --%
cd /Users/dianaperez/Desktop/Mentoring/
%-- 2/9/22, 1:55 PM --%
network_seg_new
help sprintf
network_seg_new
%-- 2/10/22, 10:20 AM --%
EmReg_task
sca
%-- 2/10/22, 11:08 AM --%
EmReg_task
Condition = conditions(Condition,:);
sca
EmReg_task
%-- 2/10/22, 1:22 PM --%
EmReg_task
7
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-2
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4
sca
yCenter-(yCenter/3)
yCenter-(yCenter/2)
yCenter-(yCenter/8)
yCenter-(yCenter/4)
yCenter-(yCenter*(3/4))
EmReg_task
sca
```

EmReg_task

sca

EmReg_task

sca

EmReg_task

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sca

EmReg_task

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numel(emotions_table)

length(emotions_table)

EmReg_task

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EmReg_task

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EmReg_task
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-2
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[droppedframes] = Screen('PlayMovie', [clips_dir clips{conditions(1)}], 1);
EmReg_task
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EmReg_task

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%-- 2/10/22, 1:47 PM --%

PsychoToolbox/PsychDemos/MovieDemo.m

Psychtoolbox/PsychDemos/MovieDemo.m

```
cd /Users/dianaperez/Documents/MATLAB/
```

MovieDemo

sca

%-- 2/10/22, 2:14 PM --%

```
LoadMovieIntoTexturesDemo( '/Volumes/RESEARCH_HD/LifespanProjectTasks/EmoReg/clips/Champ.✓  
mp4' )
```

```
ShowCursor;
```

%-- 2/10/22, 3:10 PM --%

```
LoadMovieIntoTexturesDemo( '/Volumes/RESEARCH_HD/LifespanProjectTasks/EmoReg/clips/Champ.mp4' )
```

```
ShowCursor;
```

%-- 2/10/22, 3:33 PM --%

pathtool

```
cd ~/Downloads
```

DownloadPsychtoolbox

%-- 2/10/22, 3:41 PM --%

EmReg_task

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sca

EmReg_task

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EmReg_task
sca
EmReg_task
Screen('Flip', window);
Screen('PlayMovie', moviePtr, 1);
Screen('Flip', window);
Screen('PlayMovie', moviePtr, 1);
[moviePtr] = Screen('OpenMovie', window, [clips_dir clips{conditions(1)}]);
Screen('PlayMovie', moviePtr, 1);
sca
EmReg_task
[moviePtr] = Screen('OpenMovie', window, [clips_dir clips{conditions(1)}]);
Screen('Flip', window);
[moviePtr] = Screen('OpenMovie', window, [clips_dir clips{conditions(1)}]);
[clips_dir clips{conditions(1)}]
[moviePtr] = Screen('OpenMovie', window, ↵
'/Volumes/RESEARCH_HD/LifespanProjectTasks/EmoReg/clips/21Grams_converted.mov');
Screen('PlayMovie', moviePtr, 1)
Screen('PlayMovie', ↵
'/Volumes/RESEARCH_HD/LifespanProjectTasks/EmoReg/clips/21Grams_converted.mov', 1)
Screen('CloseAll')
sca
EmReg_task
help tic
toc
tic
toc
sca
EmReg_task
sca
EmReg_task
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sca
EmReg_task
sca
EmReg_task
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sca
KbName('spaceBar')
KbName('space')
KbName('leftarrow')
KbName('q')
Condition = 0;
conditions(Condition)
EmReg_task
sca
EmReg_task
sca
EmReg_task
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EmReg_task
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EmReg_task
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EmReg_task
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EmReg_task
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sca
EmReg_task
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sca
EmReg_task
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q 2
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76
EmReg_task
2
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sca
%-- 3/5/22, 3:24 PM --%
EmReg_task
%-- 3/5/22, 3:30 PM --%
EmReg_task
help strfind
sca
EmReg_task
addpath '/Volumes/RESEARCH_HD/LifespanProjectTasks/EmoReg'
EmReg_task
```



```
sca
EmReg_task
7
EmReg_task
7
5
EmReg_task
7
7
7
Screen('Flip', window);
7
help DrawFormattedText
help Screen
EmReg_task
7
EmReg_task
7
EmReg_task
%-- 3/8/22, 10:53 AM --%
net_size_asymmetry
addpath(genpath('/Volumes/Back_Up/Box/Dependencies/cifti-matlab-master/'))
net_size_asymmetry
print(gcf,[output_dir 'HCP384_indiv_nets_asymm_surfArea.jpg'],'-dpng','-r300');
%-- 3/8/22, 2:38 PM --%
net_size_asymmetry
scatter(8,avg_diff(1,6), 'MarkerEdgeColor', 'w', 'MarkerFaceColor', 'w', 'SizeData', 50)
load
('/Users/dianaperez/Desktop/Research/lateralization_code/testing_output/HCP384_new_split_
networksxHem.mat')
diff_numvars_all = [];
asymm_numvars_all = [];
diff_sizenets_all = [];
net_size_asymmetry
length(networksxHem.verticesLH(net))
size(networksxHem.verticesLH(net))
size(networksxHem.verticesLH)
length(networksxHem.verticesLH)
net_size_asymmetry
diff_numvars(s)
max([networksxHem.verticesLH(s,net), networksxHem.verticesRH(s,net)])
max([networksxHem.verticesLH(s,net), networksxHem.verticesRH(s,net)])
networksxHem.verticesLH(s,net)
networksxHem.verticesRH(s,net)
net_size_asymmetry
diff_numvars_all = asymm_numvars_all;
[r p] = corr(diff_numvars_all, diff_sizenets_all)
rvals(17) = r;
pvals(17) = p;
scatter(diff_numvars_all, diff_sizenets_all)
hold on
p = polyfit(diff_numvars_all, diff_sizenets_all,1);
px = [min(diff_numvars_all) max(diff_numvars_all)];
py = polyval(p, px);
plot(px, py, 'LineWidth', 2);
title('All networks');
xlabel('Number of Variant Vertices (right hem)');
ylabel('Network Surface Area in mm^2 (right hem)');
nonzero_numvars = diff_numvars_all(diff_numvars_all~=0);
find(nonzero_numvars==0)
```

```

nonzero_sizenets = diff_sizenets_all(diff_sizenets_all~=0);
[r p] = corr(nonzero_numvars, nonzero_sizenets)
rvals(17) = r;
pvals(17) = p;
nonzero_numvars = find(diff_numvars_all~=0);
diff_sizenets_all=diff_sizenets_all(nonzero_numvars);
diff_numvars_all=diff_numvars_all(nonzero_numvars);
find(diff_sizenets_all==0)
find(diff_numvars_all==0)
[r p] = corr(diff_numvars_all, diff_sizenets_all)
rvals(17) = r;
pvals(17) = p;
scatter(diff_numvars_all, diff_sizenets_all)
hold on
p = polyfit(diff_numvars_all, diff_sizenets_all,1);
px = [min(diff_numvars_all) max(diff_numvars_all)];
py = polyval(p, px);
plot(px, py, 'LineWidth', 2);
title('All networks');
xlabel('Number of Variant Vertices (right hem)');
ylabel('Network Surface Area in mm^2 (right hem)');
net_size_asymmetry
help mean
averages = mean(asymm_numvars_all,2);
averages = mean(asymm_numvars_all,1);
names = ␣
{'DMN','Vis','FP','DAN','VAN','Sal','CO','SMd','SML','Aud','Tpole','MTL','PMN','PON'};
rgb = [1 0 0; %DMN
0 0 .6; %Vis
1 1 0; %FP
% .67 .67 .67; %Unassigned
0 .8 0; %DAN
% .67 .67 .67; %Unassigned2
0 .6 .6;%VAN
0 0 0; % Sal
.3 0 .6; %CON
.2 1 1; %SMd
1 .5 0; % SML
.6 .2 1; %Aud
.2 1 .2; %Tpole
0 .2 .4; %MTL
0 0 1; %PMN
.8 .8 .6]; %PON
for n = 1:numel(names)
rgb_for_plot{n} = rgb(n,:);
end
to_plot = averages;
to_plot = asymm_numvars_all;
to_plot(385,:) = averages;
ind = zeros(385,1);
ind(385) = 1;
net_inds = [ind ind ind ind ind ind ind ind ind ind ind ind ind];
handles = plotSpread(to_plot, 'categoryMarkers', {'x', '.'}, 'categoryLabels', ␣
{'Indiv_asymm_scores','Average'}, 'distributionColors', rgb_for_plot, 'categoryIdx', ␣
net_inds, 'xNames', names)
for n = 1:numel(names)
rgb_for_plot{n} = rgb(n,:);
end
names = ␣
{'DMN','Vis','FP','DAN','VAN','Sal','CO','SMd','SML','Aud','Tpole','MTL','PMN','PON'};

```

```

for n = 1:numel(names)
    rgb_for_plot{n} = rgb(n,:);
end
clear rgb_for_plot
for n = 1:numel(names)
    rgb_for_plot{n} = rgb(n,:);
end
handles = plotSpread(to_plot, 'categoryMarkers', {'x', '.'}, 'categoryLabels', ↵
{'Indiv_asymm_scores', 'Average'}, 'distributionColors', rgb_for_plot, 'categoryIdx', ↵
net_inds, 'xNames', names)
close gcf
hist(asym_numvars_all(:,1))
hist(asymm_numvars_all(:,1))
right_lateralized_DMN = find(asymm_numvars_all(:,1)<0);
figure;
net_size_asymmetry
hist(asymm_numvars_all(:,1))
right_lateralized_DMN = find(asymm_numvars_all(:,1)<0);
figure;
scatter(asymm_numvars_all(right_lateralized_DMN,1), diff_sizenets_all↵
(right_lateralized_DMN,1))
left_lateralized_DMN = find(asymm_numvars_all(:,1)>0);
figure;
scatter(asymm_numvars_all(left_lateralized_DMN,1), diff_sizenets_all↵
(left_lateralized_DMN,1))
right_lateralized_DMN = find(diff_sizenets_all(:,1)<0);
figure;
scatter(asymm_numvars_all(right_lateralized_DMN,1), diff_sizenets_all↵
(right_lateralized_DMN,1))
left_lateralized_DMN = find(diff_sizenets_all(:,1)>0);
figure;
scatter(asymm_numvars_all(left_lateralized_DMN,1), diff_sizenets_all↵
(left_lateralized_DMN,1))
hist(asymm_numvars_all(:,1))
right_lateralized_DMN = find(asymm_numvars_all(:,1)<0);
mean(diff_sizenets_all(right_lateralized_DMN,1))
left_lateralized_DMN = find(asymm_numvars_all(:,1)>0);
mean(diff_sizenets_all(left_lateralized_DMN,1))
high_asymmetry_clusters
cd '/Users/dianaperez/Documents/GitHub/PerezEtAl_HemAsymmetries'
high_asymmetry_clusters
rgb = [1 0 0; %DMN
0 0 .6; %Vis
1 1 0; %FP
%.67 .67 .67; %Unassigned
0 .8 0; %DAN
%.67 .67 .67; %Unassigned2
0 .6 .6; %VAN
0 0 0; % Sal
.3 0 .6; %CON
.2 1 1; %SMd
1 .5 0; % SML
.6 .2 1; %Aud
.2 1 .2; %Tpole
0 .2 .4; %MTL
0 0 1; %PMN
.8 .8 .6]; %PON
network_names = ↵
{'DMN', 'Vis', 'FP', 'DAN', 'VAN', 'Sal', 'CO', 'SMd', 'SML', 'Aud', 'Tpole', 'MTL', 'PMN', 'PON'};
good_nets_LH = [clusterA.left_hem.number(:,1:3) clusterA.left_hem.number(:,5) clusterA.↵

```

```
left_hem.number(:,7:end));
good_nets_RH = [clusterA.right_hem.number(:,1:3) clusterA.right_hem.number(:,5) clusterA.
right_hem.number(:,7:end)];
sum_good_nets = sum(good_nets_LH,1);
sum_good_nets_RH = sum(good_nets_RH,1);
for net = 1:numel(network_names)
    asymm(net) = (sum_good_nets(net) - sum_good_nets_RH(net))/max([sum_good_nets(net),
sum_good_nets_RH(net)]);
end
plot(asymm)
scatter(asymm)
scatter(sum_good_nets,sum_good_nets_RH, 80, rgb_colors, 'filled')
scatter(sum_good_nets,sum_good_nets_RH, 80, rgb, 'filled')
line = reffline(1,0);
line.Color = 'black';
sum_variants = sum_good_nets + sum_good_nets_RH;
sum_good_nets/sum_variants
sum_good_nets./sum_variants
scatter((sum_good_nets./sum_variants),(sum_good_nets_RH./sum_variants), 80, rgb,
'filled')
close gcf
scatter((sum_good_nets./sum_variants),(sum_good_nets_RH./sum_variants), 80, rgb,
'filled')
(sum_good_nets./sum_variants)
(sum_good_nets_RH./sum_variants)
prop_vars_LH = ~isnan(sum_good_nets./sum_variants)
prop_vars_LH = sum_good_nets./sum_variants;
prop_vars_LH = prop_vars_LH(~isnan(prop_vars_LH))
prop_vars_LH = prop_vars_LH(~isnan(prop_vars_LH))
prop_vars_RH = sum_good_nets_RH./sum_variants;
prop_vars_RH = prop_vars_RH(~isnan(prop_vars_RH))
scatter(prop_vars_LH,prop_vars_RH, 80, rgb, 'filled')
rgb = rgb(~isnan(sum_good_nets_RH./sum_variants),:);
scatter(prop_vars_LH,prop_vars_RH, 80, rgb, 'filled')
close gcf
scatter(prop_vars_LH,prop_vars_RH, 80, rgb, 'filled')
ax_max = max([x;y])+(std([x;y])/2)
axis([0 ax_max 0 ax_max]);
line = reffline(1,0);
line.Color = 'black';
xlabel('Left Hemisphere')
ylabel('Right Hemisphere')
ax = gca;
ax.FontSize = 24;
line = reffline(1,0);
line.Color = 'black';
xlabel('Left Hemisphere')
ylabel('Right Hemisphere')
ax = gca;
ax.FontSize = 24;
load
('/Users/dianaperez/Desktop/Research/lateralization_code/testing_output/HCP384_variants_i
nfo.mat')
clear all
load
('/Users/dianaperez/Desktop/Research/lateralization_code/testing_output/HCP384_variants_i
nfo.mat')
numVars = variants_info.left_hem.group_avg{:,1};
numVars_LH = numVars;
numVars_RH = variants_info.right_hem.group_avg{:,1};
```

```
diff_numVars = numVars_LH - numVars_RH;
for s = 1:384
    asymm_numVars(s,1) = diff_numVars(s)/max([numVars_LH(s), numVars_RH(s)]);
end
mean(asymm_numVars)
plot(asymm_numVars)
scatter(asymm_numVars)
hist(asymm_numVars)
close gcf
numVerts = variants_info.left_hem.group_avg(:,2);
numVerts_LH = variants_info.left_hem.group_avg(:,2);
numVerts_RH = variants_info.right_hem.group_avg(:,2);
diff_numVerts = numVerts_LH - numVerts_RH;
for s = 1:384
    asymm_numVerts(s,1) = diff_numVerts(s)/max([numVerts_LH(s), numVerts_RH(s)]);
end
diff_numVerts = numVerts_LH - numVerts_RH;
for s = 1:384
    asymm_numVerts(s,1) = diff_numVerts(s)/max([numVerts_LH(s), numVerts_RH(s)]);
end
mean(asymm_numVerts)
avg_surf_LH = variants_info.left_hem.group_avg(:,4);
avg_surf_RH = variants_info.right_hem.group_avg(:,4);
diff_avgSurf = avg_surf_LH - avg_surf_RH;
for s = 1:384
    asymm_avg_surf(s,1) = diff_avgSurf(s)/max([avg_surf_LH(s), avg_surf_RH(s)]);
end
mean(asymm_avg_surf)
load
('/Users/dianaperez/Desktop/Research/lateralization_code/testing_output/HCP384_networksxH
em.mat')
clear all
load
('/Users/dianaperez/Desktop/Research/lateralization_code/testing_output/HCP384_networksxH
em.mat')
diff_clusters = networksxHem.clustersLH - networksxHem.clustersRH;
for net = 1:16
    for s = 1:384
        asymm_net(s,net) = diff_clusters(s,net)/max([networksxHem.clustersLH(s,net),
networksxHem.clustersRH(s,net)]);
    end
end
mean(asymm_net,1)
for net = 1:16
    for s = 1:384
        asymm_net(s,net) = diff_clusters(s,net)/max([networksxHem.clustersLH(s,net),
networksxHem.clustersRH(s,net)]);
    end
    if isnan(asymm_net(s,net))
        asymm_net(s,net) = 0;
    end
end
end
mean(asymm_net,1)
bar(asymm)
bar(ans)
load
('/Users/dianaperez/Desktop/Research/lateralization_code/testing_output/network_symmetry/
WashU120_network_sizes_across_hems.mat')
clear all
load
```

```

('/Users/dianaperez/Desktop/Research/lateralization_code/testing_output/network_symmetry/
WashU120_network_sizes_across_hems.mat')
diff_surf = net_size(:,3) - net_size(:,7);
for net = 1:16
asymm(net) = diff_surf/max([net_size(net,3), net_size(net,7)]);
if isnan(asymm(net))
asymm(net) = 0;
end
end
for net = 1:16
asymm(net) = diff_surf/max([net_size(net,3), net_size(net,7)]);
asymm(net) = diff_surf(net)/max([net_size(net,3), net_size(net,7)]);
end
for net = 1:16
asymm(net) = diff_surf(net)/max([net_size(net,3), net_size(net,7)]);
if isnan(asymm(net))
asymm(net) = 0;
end
end
rgb = [1 0 0; %DMN
0 0 .6; %Vis
1 1 0; %FP
%.67 .67 .67; %Unassigned
0 .8 0; %DAN
% .67 .67 .67; %Unassigned2
0 .6 .6; %VAN
0 0 0; % Sal
.3 0 .6; %CON
.2 1 1; %SMd
1 .5 0; % SML
.6 .2 1; %Aud
.2 1 .2; %Tpole
0 .2 .4; %MTL
0 0 1; %PMN
.8 .8 .6]; %PON
asymm(4) = [];
asymm(5) = [];
for net = 1:14
bar_series(net) = bar(asymm(net), 'BarWidth', 0.9);
set(bar_series(net), 'FaceColor', rgb(net,:));
hold on;
end
close(gcf)
count = 1;
for net = 1:14
bar_series(net) = bar(count, asymm(net), 'BarWidth', 0.9);
set(bar_series(net), 'FaceColor', rgb(net,:));
count = count + 1;
hold on;
end
network_names =
{'DMN', 'Vis', 'FP', 'DAN', 'VAN', 'Sal', 'CO', 'SMd', 'SML', 'Aud', 'Tpole', 'MTL', 'PMN', 'PON'};
xticklabels(network_names)
clear all
load
('/Users/dianaperez/Desktop/Research/lateralization_code/testing_output/HCP384_networksxH
em.mat')
diff_numvars = networksxHem.clustersLH - networksxHem.clustersRH;
for net = 1:16
average_diff(net) = mean(diff_numvars(:,net));

```

```

asymm_netvars(net) = average_diff/max([mean(networksxHem.clustersLH(:,net)), mean(
(networksxHem.clustersRH(:,net))]);
if isnan(asymm_netvars(net))
asymm_netvars(net) = 0;
end
end
for net = 1:16
average_diff(net) = mean(diff_numvars(:,net));
asymm_netvars(net) = average_diff(net)/max([mean(networksxHem.clustersLH(:,net)), mean(
(networksxHem.clustersRH(:,net))]);
if isnan(asymm_netvars(net))
asymm_netvars(net) = 0;
end
end
for net = 1:16
for s = 1:384
asymm_netvars_ind(s,net) = (networksxHem.clustersLH(s,net)-networksxHem.clustersRH(s,
net))/max([networksxHem.clustersLH(:,net), networksxHem.clustersRH(:,net)]);
end
end
for net = 1:16
for s = 1:384
asymm_netvars_ind(s,net) = (networksxHem.clustersLH(s,net)-networksxHem.clustersRH(s,
net))/max([networksxHem.clustersLH(s,net), networksxHem.clustersRH(s,net)]);
if isnan(asymm_netvars_ind(s,net))
asymm_netvars_ind(s,net) = 0;
end
end
end
mean(asymm_netvars_ind,1)
count = 1;
avg_asymm = ans;
avg_asymm(6) = [];
avg_asymm(4) = [];
for net = 1:14
bar_series(net) = bar(count, avg_asymm(net), 'BarWidth', 0.9);
set(bar_series(net), 'FaceColor', rgb(net,:));
count = count + 1;
hold on;
end
rgb = [1 0 0; %DMN
0 0 .6; %Vis
1 1 0; %FP
%.67 .67 .67; %Unassigned
0 .8 0; %DAN
% .67 .67 .67; %Unassigned2
0 .6 .6; %VAN
0 0 0; % Sal
.3 0 .6; %CON
.2 1 1; %SMd
1 .5 0; % SMI
.6 .2 1; %Aud
.2 1 .2; %Tpole
0 .2 .4; %MTL
0 0 1; %PMN
.8 .8 .6]; %PON
count = 1;
for net = 1:14
bar_series(net) = bar(count, avg_asymm(net), 'BarWidth', 0.9);
set(bar_series(net), 'FaceColor', rgb(net,:));

```

```

count = count + 1;
hold on;
end
close(gcf)
count = 1;
for net = 1:14
bar_series(net) = bar(count, avg_asymm(net), 'BarWidth', 0.9);
set(bar_series(net), 'FaceColor', rgb(net,:));
count = count + 1;
hold on;
end
close(gcf)
net_size_asymmetry
axis([1 1 1.25 1.25])
axis([1, 1, 1.25, 1.25,])
axis([-1, 1, -1.25, 1.25,])
p = polyfit(diff_numvars_all, diff_sizenets_all,1);
px = [min(diff_numvars_all) max(diff_numvars_all)];
py = polyval(p, px);
plot(px, py, 'b', 'LineWidth', 2);
plot(px, py, 'k', 'LineWidth', 2);
axis([-1.25, 1.25,-1,1])
axis([-1, 1, -1.25, 1.25,])
net_size_asymmetry
axis([-1, 1, -1.25, 1.25,])
p = polyfit(diff_numvars_all, diff_sizenets_all,1);
px = [min(diff_numvars_all) max(diff_numvars_all)];
py = polyval(p, px);
plot(px, py, 'k','LineWidth', 2);
axis([-1, 1, -1, 1])
[p_fdr, p_masked] = FDR(p_vals, 0.025)
permute_net_assignment
[true_diff perm_diffs] = network_comparisons(LH_info.networksxHem.clustersLH(:,3),
LHand_info.networksxHem.clustersRH(:,3), flip_switch, numperms, 0, 1, net);
permute_net_assignment
[p_fdr p_masked] = FDR(p_vals, .025);
permute_net_assignment
flip_switch = zeros(670,1);
flip_switch(1:floor(670/2)) = 1;
[p_fdr p_masked] = FDR(p_vals, .025);
clear all
net_size_across_hems
net_size_asymmetry
help dicomhdr
help dicominfo
%-- 3/13/22, 3:28 PM --%
net_size_across_hems
load
('/Users/dianaperez/Desktop/Research/lateralization_code/testing_output/HCP384_new_split_
variants_info.mat')
load
('/Users/dianaperez/Desktop/Research/lateralization_code/testing_output/HCP384_networksxH
em.mat')
lh_numvars = mean(networksxHem.clustersLH,1);
rh_numvars = mean(networksxHem.clustersRH,1);
scatter(lh_numvars, rh_numvars, 80, rgb_colors, 'filled')
scatter(lh_numvars, rh_numvars, 80, 'k')
axis([0 2.5 0 2.5])
line = reffline(1,0);
line.Color = 'black';

```



```

xlabel('Left Hemisphere')
ax = gca;
ax.FontSize = 24;
for net = 1:16
se_lh(net) = std(networksxHem.clustersLH(:,net))/sqrt(384);
se_rh(net) = std(networksxHem.clustersRH(:,net))/sqrt(384);
end
errorbar(lh_numvars, rh_numvars, se_rh, se_rh, se_lh, se_lh, 'o')
rgb_colors = [1 0 0; %DMN
0 0 .6; %Vis
1 1 0; %FP
.67 .67 .67; %Unassigned
0 .8 0; %DAN
.67 .67 .67; %Unassigned2
0 .6 .6; %VAN
0 0 0; % Sal
.3 0 .6; %CON
.2 1 1; %SMd
1 .5 0; % SMI
.6 .2 1; %Aud
.2 1 .2; %Tpole
0 .2 .4; %MTL
0 0 1; %PMN
.8 .8 .6]; %PON
scatter(lh_numvars, rh_numvars, 80, rgb_colors, 'filled')
scatter(lh_numvars, rh_numvars, 80, 'k')
axis([0 2.5 0 2.5])
line = reffline(1,0);
line.Color = 'black';
xlabel('Left Hemisphere')
ax = gca;
ax.FontSize = 24;
for net = 1:16
se_lh(net) = std(networksxHem.clustersLH(:,net))/sqrt(384);
se_rh(net) = std(networksxHem.clustersRH(:,net))/sqrt(384);
end
errorbar(lh_numvars, rh_numvars, se_rh, se_rh, se_lh, se_lh, 'o')
close(gcf)
scatter(lh_numvars, rh_numvars, 80, rgb_colors, 'filled')
hold on
scatter(lh_numvars, rh_numvars, 80, 'k')
axis([0 2.5 0 2.5])
line = reffline(1,0);
line.Color = 'black';
xlabel('Left Hemisphere')
ax = gca;
ax.FontSize = 24;
for net = 1:16
se_lh(net) = std(networksxHem.clustersLH(:,net))/sqrt(384);
se_rh(net) = std(networksxHem.clustersRH(:,net))/sqrt(384);
end
errorbar(lh_numvars, rh_numvars, se_rh, se_rh, se_lh, se_lh, 'o')
errorbar.Color
errorbar.Colors
close(gcf)
scatter(lh_numvars, rh_numvars, 80, rgb_colors, 'filled')
hold on
scatter(lh_numvars, rh_numvars, 80, 'k')
axis([0 2.5 0 2.5])
line = reffline(1,0);

```

```

line.Color = 'black';
xlabel('Left Hemisphere')
ax = gca;
ax.FontSize = 24;
for net = 1:16
se_lh(net) = std(networksxHem.clustersLH(:,net))/sqrt(384);
se_rh(net) = std(networksxHem.clustersRH(:,net))/sqrt(384);
end
h = errorbar(lh_numvars, rh_numvars, se_rh, se_rh, se_lh, se_lh, 'o')
close(gcf)
scatter(lh_numvars, rh_numvars, 80, rgb_colors, 'filled')
hold on
scatter(lh_numvars, rh_numvars, 80, 'k')
axis([0 2.5 0 2.5])
line = reline(1,0);
line.Color = 'black';
xlabel('Left Hemisphere')
ax = gca;
ax.FontSize = 24;
for net = 1:16
se_lh(net) = std(networksxHem.clustersLH(:,net))/sqrt(384);
se_rh(net) = std(networksxHem.clustersRH(:,net))/sqrt(384);
end
h = errorbar(lh_numvars, rh_numvars, se_rh, se_rh, se_lh, se_lh, 'o')
h.Color = [0,0,0];
scatter(lh_numvars, rh_numvars, 80, rgb_colors, 'filled')
scatter(lh_numvars, rh_numvars, 80, 'k')
print(gcf,
'/Users/dianaperez/Desktop/Research/lateralization_code/HCP384_numvars_scatterplot.jpg',
'-dpng', '-r300')
%-- 3/13/22, 8:17 PM --%
net_size_asymmetry
addpath
'/Users/dianaperez/Documents/GitHub/PerezEtAl_HemAsymmetries/needed_files/plotSpread/plot
Spread'
net_size_asymmetry
mean(asymm_numvars_all,1)
mean(diff_sizenets_all,1)
permute_net_assignment
addpath '/Users/dianaperez/Documents/GitHub/PerezEtAl_HemAsymmetries/needed_files'
permute_net_assignment
network_names =
{'DMN', 'Vis', 'FP', 'DAN', 'Lang', 'Sal', 'CO', 'SMd', 'SML', 'Aud', 'Tpole', 'MTL', 'PMN', 'PON'};
rgb_colors = [1 0 0; %DMN
0 0 .6; %Vis
1 1 0; %FP
0 .8 0; %DAN
0 .6 .6; %VAN
0 0 0; % Sal
.3 0 .6; %CON
.2 1 1; %SMd
1 .5 0; % SML
.6 .2 1; %Aud
.2 1 .2; %Tpole
0 .2 .4; %MTL
0 0 1; %PMN
.8 .8 .6]; %PON
good_nets = [output_mat(:,1:3) output_mat(:,5) output_mat(:,7:end)];
rgb_for_plot = {};
for n = 1:numel(network_names)

```

```

rgb_for_plot{n} = rgb_colors(n,:);
end
ind = zeros(numperms+1,1);
ind(1,1) = 1;
net_inds = [ind ind ind ind ind ind ind ind ind ind ind ind ind ind ind];
handles = plotSpread(good_nets, 'categoryMarkers', {'x', '.'}, 'categoryLabels', ↵
{'Permuted Differences', 'True Difference'}, 'distributionColors', rgb_for_plot', ↵
'categoryIdx', net_inds, 'xNames', network_names)
ax = gca;
ax.FontSize = 24;
scatter([1:14], good_nets(1,:), 'MarkerEdgeColor', 'k', 'MarkerFaceColor', 'k', ↵
'SizeData', 50)
scatter(6, good_nets(1,6), 'MarkerEdgeColor', 'w', 'MarkerFaceColor', 'w', 'SizeData', 50)
ylabel('Difference in Number of Variants')
xlabel('Assigned Network')
min_ax = min(min(good_nets(:,:)))-(min(std(good_nets)/2));
max_ax = max(max(good_nets(:,:)))+(min(std(good_nets)/2));
axis([0, 15, min_ax, max_ax])
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.9, 0.7]);
print(gcf, [output_dir out_str '_PermutationTesting_networkAssignment.jpg'], '-dpng', '-↵
r300');
network_names = ↵
{'DMN', 'Vis', 'FP', 'DAN', 'Lang', 'Sal', 'CO', 'SMd', 'SML', 'Aud', 'Tpole', 'MTL', 'PMN', 'PON'};
rgb_colors = [1 0 0; %DMN
0 0 .6; %Vis
1 1 0; %FP
0 .8 0; %DAN
0 .6 .6; %VAN
0 0 0; % Sal
.3 0 .6; %CON
.2 1 1; %SMd
1 .5 0; % SML
.6 .2 1; %Aud
.2 1 .2; %Tpole
0 .2 .4; %MTL
0 0 1; %PMN
.8 .8 .6]; %PON
good_nets = [output_mat(:,1:3) output_mat(:,5) output_mat(:,7:end)];
rgb_for_plot = {};
for n = 1:numel(network_names)
rgb_for_plot{n} = rgb_colors(n,:);
end
ind = zeros(numperms+1,1);
ind(1,1) = 1;
net_inds = [ind ind ind ind ind ind ind ind ind ind ind ind ind ind ind];
handles = plotSpread(good_nets, 'categoryMarkers', {'x', '.'}, 'categoryLabels', ↵
{'Permuted Differences', 'True Difference'}, 'distributionColors', rgb_for_plot', ↵
'categoryIdx', net_inds, 'xNames', network_names)
ax = gca;
ax.FontSize = 24;
scatter([1:14], good_nets(1,:), 'MarkerEdgeColor', 'k', 'MarkerFaceColor', 'k', ↵
'SizeData', 50)
scatter(6, good_nets(1,6), 'MarkerEdgeColor', 'w', 'MarkerFaceColor', 'w', 'SizeData', 50)
ylabel('Difference in Number of Variants')
xlabel('Assigned Network')
min_ax = min(min(good_nets(:,:)))-(min(std(good_nets)/2));
max_ax = max(max(good_nets(:,:)))+(min(std(good_nets)/2));
axis([0, 15, min_ax, max_ax])
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.9, 0.7]);
print(gcf, [output_dir out_str '_PermutationTesting_networkAssignment.jpg'], '-dpng', '-↵

```

```

r300');
permute_net_assignment
var_freq_size_net
addpath '/Users/dianaperez/Documents/GitHub/PerezEtAl_HemAsymmetries'
var_freq_size_net
%-- 3/15/22, 1:24 PM --%
net_size_across_hems
root_dir = '/Users/dianaperez/Desktop/Research/lateralization_code/'; % location of code
directory
%root_dir = '/projects/p31161/lateralization_code/';
data_location = [root_dir 'testing_output/'];
LHand_info = load([data_location 'HCP752_LH_networksxHem.mat']);
RHand_info = load([data_location 'HCP752_RH_networksxHem.mat']);
clear all
root_dir = '/Users/dianaperez/Desktop/Research/lateralization_code/'; % location of code
directory
%root_dir = '/projects/p31161/lateralization_code/';
data_location = [root_dir 'testing_output/'];
LHand_info = load([data_location 'HCP752_LH_networksxHem.mat']);
RHand_info = load([data_location 'HCP752_RH_networksxHem.mat']);
lh_numvars = mean(LHand_info.networksxHem.clustersLH,1);
rh_numvars = mean(RHand_info.networksxHem.clustersRH,1);
rgb_colors = [1 0 0; %DMN
0 0 .6; %Vis
1 1 0; %FP
.67 .67 .67; %Unassigned
0 .8 0; %DAN
.67 .67 .67; %Unassigned2
0 .6 .6; %VAN
0 0 0; % Sal
.3 0 .6; %CON
.2 1 1; %SMd
1 .5 0; % SMI
.6 .2 1; %Aud
.2 1 .2; %Tpole
0 .2 .4; %MTL
0 0 1; %PMN
.8 .8 .6]; %PON
scatter(lh_numvars, rh_numvars, 80, rgb_colors, 'filled')
hold on
scatter(lh_numvars, rh_numvars, 80, 'k')
lh_numvars = mean(RHand_info.networksxHem.clustersLH,1);
rh_numvars = mean(RHand_info.networksxHem.clustersRH,1);
scatter(lh_numvars, rh_numvars, 80, rgb_colors, 'd', 'filled')
hold on
scatter(lh_numvars, rh_numvars, 80, 'd', 'k')
line = reffline(1,0);
line.Color = 'black';
xlabel('Left Hemisphere')
ax = gca;
ax.FontSize = 24;
ylabel('Right Hemisphere')
close gcf
% load data
root_dir = '/Users/dianaperez/Desktop/Research/lateralization_code/'; % location of code
directory
data_location = [root_dir 'testing_output/'];
LHand_info = load([data_location 'HCP752_LH_networksxHem.mat']);
RHand_info = load([data_location 'HCP752_RH_networksxHem.mat']);
%plot left-handers

```

```
lh_numvars = mean(LHand_info.networksxHem.clustersLH,1);
rh_numvars = mean(RHand_info.networksxHem.clustersRH,1);
scatter(lh_numvars, rh_numvars, 80, rgb_colors, 'filled')
hold on
scatter(lh_numvars, rh_numvars, 80, 'k')
for net = 1:16
    se_lh(net) = std(LHand_info.networksxHem.clustersLH(:,net))/sqrt(40);
    se_rh(net) = std(RHand_info.networksxHem.clustersRH(:,net))/sqrt(40);
end
h = errorbar(lh_numvars, rh_numvars, se_rh, se_rh, se_lh, se_lh, 'o')
h.Color = [0,0,0];
% plot right-handers
lh_numvars = mean(RHand_info.networksxHem.clustersLH,1);
rh_numvars = mean(RHand_info.networksxHem.clustersRH,1);
scatter(lh_numvars, rh_numvars, 80, rgb_colors, 'd', 'filled')
hold on
scatter(lh_numvars, rh_numvars, 80, 'd','k')
axis([0 2.5 0 2.5])
line = refline(1,0);
line.Color = 'black';
xlabel('Left Hemisphere')
ylabel('Right Hemisphere')
ax = gca;
ax.FontSize = 24;
for net = 1:16
    se_lh(net) = std(RHand_info.networksxHem.clustersLH(:,net))/sqrt(670);
    se_rh(net) = std(RHand_info.networksxHem.clustersRH(:,net))/sqrt(670);
end
h = errorbar(lh_numvars, rh_numvars, se_rh, se_rh, se_lh, se_lh, 'o')
h.Color = [0,0,0];
close(gcf)
root_dir = '/Users/dianaperez/Desktop/Research/lateralization_code/'; % location of code
directory
data_location = [root_dir 'testing_output/'];
LHand_info = load([data_location 'HCP752_LH_networksxHem.mat']);
RHand_info = load([data_location 'HCP752_RH_networksxHem.mat']);
%plot left-handers
lh_numvars = mean(LHand_info.networksxHem.clustersLH,1);
rh_numvars = mean(LHand_info.networksxHem.clustersRH,1);
for net = 1:16
    se_lh(net) = std(LHand_info.networksxHem.clustersLH(:,net))/sqrt(40);
    se_rh(net) = std(LHand_info.networksxHem.clustersRH(:,net))/sqrt(40);
end
h = errorbar(lh_numvars, rh_numvars, se_rh, se_rh, se_lh, se_lh, 'o')
h.Color = [0,0,0];
scatter(lh_numvars, rh_numvars, 80, rgb_colors, 'filled')
hold on
scatter(lh_numvars, rh_numvars, 80, 'k')
% plot right-handers
lh_numvars = mean(RHand_info.networksxHem.clustersLH,1);
rh_numvars = mean(RHand_info.networksxHem.clustersRH,1);
for net = 1:16
    se_lh(net) = std(RHand_info.networksxHem.clustersLH(:,net))/sqrt(670);
    se_rh(net) = std(RHand_info.networksxHem.clustersRH(:,net))/sqrt(670);
end
h = errorbar(lh_numvars, rh_numvars, se_rh, se_rh, se_lh, se_lh, 'o')
h.Color = [0,0,0];
scatter(lh_numvars, rh_numvars, 80, rgb_colors, 'd', 'filled')
hold on
scatter(lh_numvars, rh_numvars, 80, 'd','k')
```

```
axis([0 2.5 0 2.5])
line = reffline(1,0);
line.Color = 'black';
xlabel('Left Hemisphere')
ylabel('Right Hemisphere')
ax = gca;
ax.FontSize = 24;
root_dir = '/Users/dianaperez/Desktop/Research/lateralization_code/'; % location of code
directory
data_location = [root_dir 'testing_output/'];
LHand_info = load([data_location 'HCP752_LH_networksxHem.mat']);
RHand_info = load([data_location 'HCP752_RH_networksxHem.mat']);
%plot left-handers
lh_numvars = mean(LHand_info.networksxHem.clustersLH,1);
rh_numvars = mean(LHand_info.networksxHem.clustersRH,1);
for net = 1:16
se_lh(net) = std(LHand_info.networksxHem.clustersLH(:,net))/sqrt(40);
se_rh(net) = std(LHand_info.networksxHem.clustersRH(:,net))/sqrt(40);
end
h = errorbar(lh_numvars, rh_numvars, se_rh, se_rh, se_lh, se_lh, 'o')
h.Color = [0,0,0];
hold on
scatter(lh_numvars, rh_numvars, 80, rgb_colors, 'filled')
hold on
scatter(lh_numvars, rh_numvars, 80, 'k')
% plot right-handers
lh_numvars = mean(RHand_info.networksxHem.clustersLH,1);
rh_numvars = mean(RHand_info.networksxHem.clustersRH,1);
for net = 1:16
se_lh(net) = std(RHand_info.networksxHem.clustersLH(:,net))/sqrt(670);
se_rh(net) = std(RHand_info.networksxHem.clustersRH(:,net))/sqrt(670);
end
h = errorbar(lh_numvars, rh_numvars, se_rh, se_rh, se_lh, se_lh, 'o')
h.Color = [0,0,0];
scatter(lh_numvars, rh_numvars, 80, rgb_colors, 'd', 'filled')
hold on
scatter(lh_numvars, rh_numvars, 80, 'd','k')
axis([0 2.5 0 2.5])
line = reffline(1,0);
line.Color = 'black';
xlabel('Left Hemisphere')
ylabel('Right Hemisphere')
ax = gca;
ax.FontSize = 24;
print(gcf,
'/Users/dianaperez/Desktop/Research/lateralization_code/HCP752_numvars_LhandvRhand_scatter
rplot.jpg', '-dpng', '-r300')
close gcf
lh_numvars = mean(LHand_info.networksxHem.clustersLH,1);
rh_numvars = mean(LHand_info.networksxHem.clustersRH,1);
for net = 1:16
se_lh(net) = std(LHand_info.networksxHem.clustersLH(:,net))/sqrt(40);
se_rh(net) = std(LHand_info.networksxHem.clustersRH(:,net))/sqrt(40);
end
h = errorbar(lh_numvars, rh_numvars, se_rh, se_rh, se_lh, se_lh, 'o')
h.Color = [0,0,0];
hold on
scatter(lh_numvars, rh_numvars, 80, rgb_colors, 'filled')
hold on
scatter(lh_numvars, rh_numvars, 80, 'k')
```

```

line = reffline(1,0);
line.Color = 'black';
xlabel('Left Hemisphere')
ylabel('Right Hemisphere')
ax = gca;
ax.FontSize = 24;
print(gcf,
'/Users/dianaperez/Desktop/Research/lateralization_code/HCP752_numvars_LhandvRhand_scatte
rplot.jpg', '-dpng', '-r300')
print(gcf,
'/Users/dianaperez/Desktop/Research/lateralization_code/HCP752_numvars_Lhand_scatterplot.
jpg', '-dpng', '-r300')
close gcf
lh_numvars = mean(RHand_info.networksxHem.clustersLH,1);
rh_numvars = mean(RHand_info.networksxHem.clustersRH,1);
for net = 1:16
se_lh(net) = std(RHand_info.networksxHem.clustersLH(:,net))/sqrt(670);
se_rh(net) = std(RHand_info.networksxHem.clustersRH(:,net))/sqrt(670);
end
h = errorbar(lh_numvars, rh_numvars, se_rh, se_rh, se_lh, se_lh, 'o')
h.Color = [0,0,0];
scatter(lh_numvars, rh_numvars, 80, rgb_colors, 'd', 'filled')
hold on
scatter(lh_numvars, rh_numvars, 80, 'd','k')
axis([0 2.5 0 2.5])
line = reffline(1,0);
line.Color = 'black';
xlabel('Left Hemisphere')
ylabel('Right Hemisphere')
ax = gca;
ax.FontSize = 24;
print(gcf,
'/Users/dianaperez/Desktop/Research/lateralization_code/HCP752_numvars_Rhand_scatterplot.
jpg', '-dpng', '-r300')
close gcf
lh_numvars = mean(RHand_info.networksxHem.clustersLH,1);
rh_numvars = mean(RHand_info.networksxHem.clustersRH,1);
for net = 1:16
se_lh(net) = std(RHand_info.networksxHem.clustersLH(:,net))/sqrt(670);
se_rh(net) = std(RHand_info.networksxHem.clustersRH(:,net))/sqrt(670);
end
h = errorbar(lh_numvars, rh_numvars, se_rh, se_rh, se_lh, se_lh, 'o')
h.Color = [0,0,0];
hold on
scatter(lh_numvars, rh_numvars, 80, rgb_colors, 'd', 'filled')
hold on
scatter(lh_numvars, rh_numvars, 80, 'd','k')
axis([0 2.5 0 2.5])
line = reffline(1,0);
line.Color = 'black';
xlabel('Left Hemisphere')
ylabel('Right Hemisphere')
ax = gca;
ax.FontSize = 24;
print(gcf,
'/Users/dianaperez/Desktop/Research/lateralization_code/HCP752_numvars_Rhand_scatterplot.
jpg', '-dpng', '-r300')
% load data
root_dir = '/Users/dianaperez/Desktop/Research/lateralization_code/'; % location of code
directory

```

```
data_location = [root_dir 'testing_output/'];
LHand_info = load([data_location 'HCP752_LH_networksxHem.mat']);
RHand_info = load([data_location 'HCP752_RH_networksxHem.mat']);
%plot left-handers
lh_numvars = mean(LHand_info.networksxHem.clustersLH,1);
rh_numvars = mean(LHand_info.networksxHem.clustersRH,1);
for net = 1:16
se_lh(net) = std(LHand_info.networksxHem.clustersLH(:,net))/sqrt(40);
se_rh(net) = std(LHand_info.networksxHem.clustersRH(:,net))/sqrt(40);
end
h = errorbar(lh_numvars, rh_numvars, se_rh, se_rh, se_lh, se_lh, 'o')
h.Color = [0,0,0];
hold on
scatter(lh_numvars, rh_numvars, 80, rgb_colors, 'filled')
hold on
scatter(lh_numvars, rh_numvars, 80, 'k')
% plot right-handers
lh_numvars = mean(RHand_info.networksxHem.clustersLH,1);
rh_numvars = mean(RHand_info.networksxHem.clustersRH,1);
for net = 1:16
se_lh(net) = std(RHand_info.networksxHem.clustersLH(:,net))/sqrt(670);
se_rh(net) = std(RHand_info.networksxHem.clustersRH(:,net))/sqrt(670);
end
h = errorbar(lh_numvars, rh_numvars, se_rh, se_rh, se_lh, se_lh, 'o')
h.Color = [0,0,0];
hold on
scatter(lh_numvars, rh_numvars, 80, rgb_colors, 'd', 'filled')
hold on
scatter(lh_numvars, rh_numvars, 80, 'd','k')
axis([0 2.5 0 2.5])
line = reffline(1,0);
line.Color = 'black';
xlabel('Left Hemisphere')
ylabel('Right Hemisphere')
ax = gca;
ax.FontSize = 24;
print(gcf,
'/Users/dianaperez/Desktop/Research/lateralization_code/HCP752_numvars_LhandvRhand_scatter
rplot.jpg', '-dpng', '-r300')
%-- 3/17/22, 11:12 AM --%
wb_dir = '/Applications/workbench/bin_macosx64/';
group_avg =
'/Users/dianaperez/Desktop/Research/lateralization_code/group_avgs/120_LR_minsize400_reco
lored_manualconsensus_LR.dlabel.nii';
borderlabel_file = '/Users/dianaperez/Downloads/wbBorderLabelFile.txt';
output_dlabel = '/Users/dianaperez/Desktop/WashU120_labelfile_fornetoutlines.dlabel.nii';
system([wb_dir 'wb_command -cifti-label-import ' group_avg ' ' borderlabel_file ' '
output_dlabel])
dir = '/Applications/workbench/bin_macosx64/';
group_avg =
'/Users/dianaperez/Desktop/Research/lateralization_code/group_avgs/120_LR_minsize400_reco
lored_manualconsensus_LR.dlabel.nii';
borderlabel_file = '/Users/dianaperez/Downloads/wbBorderLabelFiles.txt';
output_dlabel = '/Users/dianaperez/Desktop/WashU120_labelfile_fornetoutlines.dlabel.nii';
system([wb_dir 'wb_command -cifti-label-import ' group_avg ' ' borderlabel_file ' '
output_dlabel])
wb_dir = '/Applications/workbench/bin_macosx64/';
group_avg =
'/Users/dianaperez/Desktop/Research/lateralization_code/group_avgs/120_LR_minsize400_reco
lored_manualconsensus_LR.dlabel.nii';
```



```
borderlabel_file = '/Users/dianaperez/Downloads/wbBorderLabelFile.txt';
output_dlabel = '/Users/dianaperez/Desktop/WashU120_labelfile_fornetoutlines.dlabel.nii';
system([wb_dir 'wb_command -cifti-label-import ' group_avg ' ' borderlabel_file ' '↵
output_dlabel])
output_lefthem = '/Users/dianaperez/Desktop/WashU120_labelfile_fornetoutlines_left.label.↵
gii';
output_righthem = '/Users/dianaperez/Desktop/WashU120_labelfile_fornetoutlines_right.↵
label.gii';
system([wb_dir 'wb_command -cifti-separate ' output_dlabel ' COLUMN -label CORTEX_LEFT '↵
output_lefthem ' -label CORTEX_RIGHT ' output_righthem])
system([wb_dir 'wb_command -label-to-border↵
/Users/dianaperez/Documents/Dependencies/32k_ConteAtlas_v2_distribute/Conte69_atlas-v2.↵
LR.32k_fs_LR.wb/Conte69.L.midthickness.32k_fs_LR.surf.gii ' output_lefthem↵
WashU120_netborders_L.border])
system([wb_dir 'wb_command -label-to-border↵
/Users/dianaperez/Documents/Dependencies/32k_ConteAtlas_v2_distribute/Conte69_atlas-v2.↵
LR.32k_fs_LR.wb/Conte69.R.midthickness.32k_fs_LR.surf.gii ' output_righthem↵
WashU120_netborders_R.border])
system([wb_dir 'wb_command -label-to-border↵
/Users/dianaperez/Documents/Dependencies/32k_ConteAtlas_v2_distribute/Conte69_atlas-v2.↵
LR.32k_fs_LR.wb/Conte69.L.midthickness.32k_fs_LR.surf.gii ' output_lefthem '↵
WashU120_netborders_L.border'])
system([wb_dir 'wb_command -label-to-border↵
/Users/dianaperez/Documents/Dependencies/32k_ConteAtlas_v2_distribute/Conte69_atlas-v2.↵
LR.32k_fs_LR.wb/Conte69.R.midthickness.32k_fs_LR.surf.gii ' output_righthem '↵
WashU120_netborders_R.border'])
cd /Users/dianaperez/Documents
cd Dependencies
cd 32k_ConteAtlas_v2_distribute/
system([wb_dir 'wb_command -label-to-border↵
/Users/dianaperez/Documents/Dependencies/Resources/Conte69_atlas-v2.LR.32k_fs_LR.↵
wb/Conte69.L.midthickness.32k_fs_LR.surf.gii ' output_lefthem ' WashU120_netborders_L.↵
border'])
system([wb_dir 'wb_command -label-to-border↵
/Users/dianaperez/Documents/Dependencies/Resources/Conte69_atlas-v2.LR.32k_fs_LR.↵
wb/Conte69.R.midthickness.32k_fs_LR.surf.gii ' output_righthem ' WashU120_netborders_R.↵
border'])
load('/Users/dianaperez/Desktop/Research/lateralization_code/HCP384_net_assignments.mat')
net_specific_diffmap
addpath '/Users/dianaperez/Documents/GitHub/PerezEtAl_HemAsymmetries'
net_specific_diffmap
template.brainstructure>0
whole_map = [diff_map; diff_map];
diff_map = whole_map(template.brainstructure>0);
template.data = diff_map;
ft_write_cifti_mod([output_dir out_str '_' network_names{network_id} '_Variants_DiffMap.↵
dtseries.nii'], template);
net_specific_diffmap
var_freq_size_net
clear all
load↵
('/Users/dianaperez/Desktop/Research/lateralization_code/testing_output/HCP752_new_split_↵
LH_networksxHem.mat')
network_names = {'DMN' 'Vis' 'FP' 'DAN' 'Lang' 'Sal' 'CO' 'SMd' 'SMl'↵
'Aud' 'Tpole' 'MTL' 'PMN' 'PON'};
good_nets_LH = [networksxHem.verticesLH(:,1:3) networksxHem.verticesLH(:,5) networksxHem.↵
verticesLH(:,7:end)];
good_nets_LH = [networksxHem.clustersLH(:,1:3) networksxHem.clustersLH(:,5) networksxHem.↵
clustersLH(:,7:end)];
good_nets_RH = [networksxHem.clustersRH(:,1:3) networksxHem.clustersRH(:,5) networksxHem.↵
```

```
clustersRH(:,7:end)];
good_nets_LH = mean(good_nets_LH);
good_nets_RH = mean(good_nets_RH);
means = [good_nets_LH' good_nets_RH'];
bar(1:14, means)
legend('Left Hem', 'Right Hem')
xticks(1:14)
xticklabels(network_names)
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.9, 0.7]);
ylabel('Average Number of Variant Regions')
xlabel('Network');
title('Variant Regions Assigned to Each Network Across Hemispheres')
ax = gca;
ax.FontSize = 24;
for net = 1:16
se_lh(net) = std(networksxHem.clustersLH(:,net))/sqrt(40);
se_rh(net) = std(networksxHem.clustersRH(:,net))/sqrt(40);
end
se_lh(4) = [];
se_lh(5) = [];
se_rh(4) = [];
se_rh(5) = [];
hold on
h = errorbar(1:14, means, se_lh, se_lh, se_rh, se_rh, 'o')
h = errorbar(means, se_lh, se_lh, se_rh, se_rh, 'o')
h = errorbar(1:14, means(:,1), se_lh, se_lh, 'o')
bar(1:14, means)
legend('Left Hem', 'Right Hem')
xticks(1:14)
xticklabels(network_names)
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.9, 0.7]);
ylabel('Average Number of Variant Regions')
xlabel('Network');
title('Variant Regions Assigned to Each Network Across Hemispheres')
ax = gca;
ax.FontSize = 24;
h = errorbar(.5:1:13.5, means(:,1), se_lh, se_lh)
close gcf
bar(1:14, means)
legend('Left Hem', 'Right Hem')
xticks(1:14)
xticklabels(network_names)
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.9, 0.7]);
ylabel('Average Number of Variant Regions')
xlabel('Network');
title('Variant Regions Assigned to Each Network Across Hemispheres')
ax = gca;
ax.FontSize = 24;
hold on
h = errorbar(.5:1:13.5, means(:,1), se_lh, se_lh)
close gcf
bar(1:14, means)
legend('Left Hem', 'Right Hem')
xticks(1:14)
xticklabels(network_names)
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.9, 0.7]);
ylabel('Average Number of Variant Regions')
xlabel('Network');
title('Variant Regions Assigned to Each Network Across Hemispheres')
ax = gca;
```

```
ax.FontSize = 24;
hold on
h = errorbar(.75:1:13.75, means(:,1), se_lh, se_lh)
h = errorbar(.95:1:13.95, means(:,1), se_lh, se_lh)
h = errorbar(.90:1:13.90, means(:,1), se_lh, se_lh)
h = errorbar(.85:1:13.85, means(:,1), se_lh, se_lh)
close(gcf)
bar(1:14, means)
legend('Left Hem', 'Right Hem')
xticks(1:14)
xticklabels(network_names)
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.9, 0.7]);
ylabel('Average Number of Variant Regions')
xlabel('Network');
title('Variant Regions Assigned to Each Network Across Hemispheres')
ax = gca;
ax.FontSize = 24;
hold on
h = errorbar([1:14 1:14], means, se_lh, se_lh, se_rh, se_rh)
h = errorbar([1:14 1:14], means, se_lh, se_lh, se_rh, se_rh)
h = errorbar([1:14; 1:14], means, se_lh, se_lh, se_rh, se_rh)
h = errorbar([1:14; 1:14], means, [se_lh; se_rh], [se_lh; se_rh])
[1:14; 1:14]
h = errorbar([1:14; 1:14]', means, [se_lh; se_rh]', [se_lh; se_rh]')
bar(1:14, means)
legend('Left Hem', 'Right Hem')
xticks(1:14)
xticklabels(network_names)
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.9, 0.7]);
ylabel('Average Number of Variant Regions')
xlabel('Network');
title('Variant Regions Assigned to Each Network Across Hemispheres')
ax = gca;
ax.FontSize = 24;
hold on
h = errorbar(.85:1:13.85, means(:,1), se_lh, se_lh)
h.Color = [0,0,0];
g = errorbar(1.15:1:14.15, means(:,1), se_lh, se_lh)
g.Color = [0,0,0];
close(gcf)
bar(1:14, means)
legend('Left Hem', 'Right Hem')
xticks(1:14)
xticklabels(network_names)
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.9, 0.7]);
ylabel('Average Number of Variant Regions')
xlabel('Network');
title('Variant Regions Assigned to Each Network Across Hemispheres')
ax = gca;
ax.FontSize = 24;
hold on
h = errorbar(.85:1:13.85, means(:,1), se_lh, se_lh)
h.Color = [0,0,0];
g = errorbar(1.15:1:14.15, means(:,1), se_rh, se_rh)
g.Color = [0,0,0];
h.LineStyle = 'none';
g.LineStyle = 'none';
close(gcf)
bar(1:14, means)
legend('Left Hem', 'Right Hem')
```

```

xticks(1:14)
xticklabels(network_names)
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.9, 0.7]);
ylabel('Average Number of Variant Regions')
xlabel('Network');
title('Variant Regions Assigned to Each Network Across Hemispheres')
ax = gca;
ax.FontSize = 24;
hold on
h = errorbar(.85:1:13.85, means(:,1), se_lh, se_lh)
h.Color = [0,0,0];
g = errorbar(1.15:1:14.15, means(:,2), se_rh, se_rh)
g.Color = [0,0,0];
h.LineStyle = 'none';
g.LineStyle = 'none';
h.LineWidth = 2;
g.LineWidth = 2;
legend('Left Hem', 'Right Hem')
print(gcf, '
/Users/dianaperez/Desktop/Research/lateralization_code/HCP752_numvars_Lhand_leftvrighthe
m.jpg', '-dpng', '-r300')
left_handlers = networksxHem;
load
('/Users/dianaperez/Desktop/Research/lateralization_code/testing_output/HCP752_new_split_
RH_networksxHem.mat')
network_names = {'DMN' 'Vis' 'FP' 'DAN' 'Lang' 'Sal' 'CO' 'SMD' 'SML'
'Aud' 'Tpole' 'MTL' 'PMN' 'PON'};
good_nets_LH = [networksxHem.verticesLH(:,1:3) networksxHem.verticesLH(:,5) networksxHem.
verticesLH(:,7:end)];
good_nets_LH = [networksxHem.clustersLH(:,1:3) networksxHem.clustersLH(:,5) networksxHem.
clustersLH(:,7:end)];
good_nets_RH = [networksxHem.clustersRH(:,1:3) networksxHem.clustersRH(:,5) networksxHem.
clustersRH(:,7:end)];
good_nets_LH = mean(good_nets_LH);
good_nets_RH = mean(good_nets_RH);
means = [good_nets_LH' good_nets_RH'];
for net = 1:16
se_lh(net) = std(networksxHem.clustersLH(:,net))/sqrt(670);
se_rh(net) = std(networksxHem.clustersRH(:,net))/sqrt(670);
end
se_lh(4) = [];
se_lh(5) = [];
se_rh(4) = [];
se_rh(5) = [];
bar(1:14, means)
legend('Left Hem', 'Right Hem')
xticks(1:14)
xticklabels(network_names)
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.9, 0.7]);
ylabel('Average Number of Variant Regions')
xlabel('Network');
title('Variant Regions Assigned to Each Network Across Hemispheres')
ax = gca;
ax.FontSize = 24;
hold on
h = errorbar(.85:1:13.85, means(:,1), se_lh, se_lh)
h.Color = [0,0,0];
g = errorbar(1.15:1:14.15, means(:,2), se_rh, se_rh)
g.Color = [0,0,0];
h.LineStyle = 'none';

```

```

g.LineStyle = 'none';
h.LineWidth = 2;
g.LineWidth = 2;
legend('Left Hem', 'Right Hem')
close(gcf)
load
('/Users/dianaperez/Desktop/Research/lateralization_code/testing_output/HCP752_new_split_
RH_networksxHem.mat')
network_names = {'DMN' 'Vis' 'FP' 'DAN' 'Lang' 'Sal' 'CO' 'SMd' 'SMl'
'Aud' 'Tpole' 'MTL' 'PMN' 'PON'};
good_nets_LH = [networksxHem.verticesLH(:,1:3) networksxHem.verticesLH(:,5) networksxHem.
verticesLH(:,7:end)];
good_nets_LH = [networksxHem.clustersLH(:,1:3) networksxHem.clustersLH(:,5) networksxHem.
clustersLH(:,7:end)];
good_nets_RH = [networksxHem.clustersRH(:,1:3) networksxHem.clustersRH(:,5) networksxHem.
clustersRH(:,7:end)];
good_nets_LH = mean(good_nets_LH);
good_nets_RH = mean(good_nets_RH);
means = [good_nets_LH' good_nets_RH'];
for net = 1:16
se_lh(net) = std(networksxHem.clustersLH(:,net))/sqrt(670);
se_rh(net) = std(networksxHem.clustersRH(:,net))/sqrt(670);
end
se_lh(4) = [];
se_lh(5) = [];
se_rh(4) = [];
se_rh(5) = [];
bar(1:14, means)
legend('Left Hem', 'Right Hem')
xticks(1:14)
xticklabels(network_names)
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.9, 0.7]);
ylabel('Average Number of Variant Regions')
xlabel('Network');
title('Variant Regions Assigned to Each Network Across Hemispheres')
ax = gca;
ax.FontSize = 24;
hold on
h = errorbar(.85:1:13.85, means(:,1), se_lh, se_lh)
h.Color = [0,0,0];
g = errorbar(1.15:1:14.15, means(:,2), se_rh, se_rh)
g.Color = [0,0,0];
h.LineStyle = 'none';
g.LineStyle = 'none';
h.LineWidth = 2;
g.LineWidth = 2;
legend('Left Hem', 'Right Hem')
print(gcf,
'/Users/dianaperez/Desktop/Research/lateralization_code/HCP752_numvars_Rhand_leftvrighthe
m.jpg', '-dpng', '-r300')
close(gcf)
right_handlers = networksxHem;
mean_lefthanders_LH = mean(left_handlers.clustersLH);
mean_lefthanders_RH = mean(left_handlers.clustersRH);
mean_righthanders_LH = mean(right_handlers.clustersLH);
mean_righthanders_RH = mean(right_handlers.clustersRH);
diff_lefthanders = mean_lefthanders_LH - mean_lefthanders_RH;
diff_righthanders = mean_righthanders_LH - mean_righthanders_RH;
diff_lefthanders = left_handlers.clustersLH - left_handlers.clustersRH;
mean_diff_lefthanders = mean(diff_lefthanders, 1);

```

```

diff_righthanders = right_handers.clustersLH - right_handers.clustersRH;
mean_diff_righthanders = mean(diff_righthanders, 1);
diff_lefthanders(4) = [], diff_lefthanders(5) = [];
diff_righthanders(4) = [], diff_righthanders(5) = [];
diff_lefthanders = mean_lefthanders_LH - mean_lefthanders_RH;
diff_righthanders = mean_righthanders_LH - mean_righthanders_RH;
diff_lefthanders(4) = [];
diff_lefthanders(5) = [];
diff_righthanders(4) = []; diff_righthanders(5) = [];
bar(1:14, diffs)
legend('Left Handers', 'Right Handers')
xticks(1:14)
xticklabels(network_names)
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.9, 0.7]);
ylabel('Difference in Average Number of Variant Regions (left - right hem)')
xlabel('Network');
diffs = [diff_lefthanders; diff_righthanders];
bar(1:14, diffs)
legend('Left Handers', 'Right Handers')
xticks(1:14)
xticklabels(network_names)
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.9, 0.7]);
ylabel('Difference in Average Number of Variant Regions (left - right hem)')
xlabel('Network');
ax = gca;
ax.FontSize = 24;
hold on
diff_lefthanders = left_handers.clustersLH - left_handers.clustersRH;
mean_diff_lefthanders = mean(diff_lefthanders, 1);
diff_righthanders = right_handers.clustersLH - right_handers.clustersRH;
mean_diff_righthanders = mean(diff_righthanders, 1);
load
('/Users/dianaperez/Desktop/Research/lateralization_code/testing_output/HCP752_new_split_
RH_networksxHem.mat')
M
load
('/Users/dianaperez/Desktop/Research/lateralization_code/testing_output/HCP752_new_split_
MH_networksxHem.mat')
load
('/Users/dianaperez/Desktop/Research/lateralization_code/testing_output/HCP752_new_split_
RH_networksxHem.mat')
load
('/Users/dianaperez/Desktop/Research/lateralization_code/testing_output/HCP752_new_split_
RH_matched_networksxHem.mat')
load
('/Users/dianaperez/Desktop/Research/lateralization_code/testing_output/HCP752_new_split_
RH_networksxHem.mat')
load
('/Users/dianaperez/Desktop/Research/lateralization_code/testing_output/HCP752_new_split_
MH_networksxHem.mat')
load
('/Users/dianaperez/Desktop/Research/lateralization_code/testing_output/HCP752_new_split_
RH_matched_networksxHem.mat')
load
('/Users/dianaperez/Desktop/Research/lateralization_code/testing_output/HCP752_new_split_
RH_networksxHem.mat')
load
('/Users/dianaperez/Desktop/Research/lateralization_code/testing_output/HCP752_RH_network
sxHem.mat')
clear all

```

```
load
('/Users/dianaperez/Desktop/Research/lateralization_code/testing_output/HCP752_RH_network
sxHem.mat')
network_names = {'DMN' 'Vis' 'FP' 'DAN' 'Lang' 'Sal' 'CO' 'SMd' 'SML'
'Aud' 'Tpole' 'MTL' 'PMN' 'PON'};
good_nets_LH = [networksxHem.verticesLH(:,1:3) networksxHem.verticesLH(:,5) networksxHem.
verticesLH(:,7:end)];
good_nets_LH = [networksxHem.clustersLH(:,1:3) networksxHem.clustersLH(:,5) networksxHem.
clustersLH(:,7:end)];
good_nets_RH = [networksxHem.clustersRH(:,1:3) networksxHem.clustersRH(:,5) networksxHem.
clustersRH(:,7:end)];
good_nets_LH = mean(good_nets_LH);
good_nets_RH = mean(good_nets_RH);
means = [good_nets_LH' good_nets_RH'];
for net = 1:16
se_lh(net) = std(networksxHem.clustersLH(:,net))/sqrt(670);
se_rh(net) = std(networksxHem.clustersRH(:,net))/sqrt(670);
end
se_lh(4) = [];
se_lh(5) = [];
se_rh(4) = [];
se_rh(5) = [];
bar(1:14, means)
legend('Left Hem', 'Right Hem')
xticks(1:14)
xticklabels(network_names)
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.9, 0.7]);
ylabel('Average Number of Variant Regions')
xlabel('Network');
title('Variant Regions Assigned to Each Network Across Hemispheres')
ax = gca;
ax.FontSize = 24;
hold on
h = errorbar(.85:1:13.85, means(:,1), se_lh, se_lh)
h.Color = [0,0,0];
g = errorbar(1.15:1:14.15, means(:,2), se_rh, se_rh)
g.Color = [0,0,0];
h.LineStyle = 'none';
g.LineStyle = 'none';
h.LineWidth = 2;
g.LineWidth = 2;
legend('Left Hem', 'Right Hem')
print(gcf,
'/Users/dianaperez/Desktop/Research/lateralization_code/HCP752_numvars_Rhand_leftvrighthe
m.jpg', '-dpng', '-r300')
close gcf
load
('/Users/dianaperez/Desktop/Research/lateralization_code/testing_output/HCP752_RH_network
sxHem.mat')
network_names = {'DMN' 'Vis' 'FP' 'DAN' 'Lang' 'Sal' 'CO' 'SMd' 'SML'
'Aud' 'Tpole' 'MTL' 'PMN' 'PON'};
good_nets_LH = [networksxHem.verticesLH(:,1:3) networksxHem.verticesLH(:,5) networksxHem.
verticesLH(:,7:end)];
good_nets_LH = [networksxHem.clustersLH(:,1:3) networksxHem.clustersLH(:,5) networksxHem.
clustersLH(:,7:end)];
good_nets_RH = [networksxHem.clustersRH(:,1:3) networksxHem.clustersRH(:,5) networksxHem.
clustersRH(:,7:end)];
good_nets_LH = mean(good_nets_LH);
good_nets_RH = mean(good_nets_RH);
means = [good_nets_LH' good_nets_RH'];
```

```

for net = 1:16
se_lh(net) = std(networksxHem.clustersLH(:,net))/sqrt(670);
se_rh(net) = std(networksxHem.clustersRH(:,net))/sqrt(670);
end
se_lh(4) = [];
se_lh(5) = [];
se_rh(4) = [];
se_rh(5) = [];
bar(1:14, means)
legend('Left Hem', 'Right Hem')
xticks(1:14)
xticklabels(network_names)
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.9, 0.7]);
ylabel('Average Number of Variant Regions')
xlabel('Network');
title('Variant Regions Assigned to Each Network Across Hemispheres')
ax = gca;
ax.FontSize = 24;
hold on
h = errorbar(.85:1:13.85, means(:,1), se_lh, se_lh)
h.Color = [0,0,0];
g = errorbar(1.15:1:14.15, means(:,2), se_rh, se_rh)
g.Color = [0,0,0];
h.LineStyle = 'none';
g.LineStyle = 'none';
h.LineWidth = 2;
g.LineWidth = 2;
legend('Left Hem', 'Right Hem')
print(gcf, '
/Users/dianaperez/Desktop/Research/lateralization_code/HCP752_numvars_Rhand_leftvrighthe
m.jpg', '-dpng', '-r300')
help axis
axis([0,2.5, 0,15])
axis([0,15,0,2.5])
load
('/Users/dianaperez/Desktop/Research/lateralization_code/testing_output/HCP752_RH_network
sxHem.mat')
network_names = {'DMN' 'Vis' 'FP' 'DAN' 'Lang' 'Sal' 'CO' 'SMd' 'SMl'
'Aud' 'Tpole' 'MTL' 'PMN' 'PON'};
good_nets_LH = [networksxHem.verticesLH(:,1:3) networksxHem.verticesLH(:,5) networksxHem.
verticesLH(:,7:end)];
good_nets_LH = [networksxHem.clustersLH(:,1:3) networksxHem.clustersLH(:,5) networksxHem.
clustersLH(:,7:end)];
good_nets_RH = [networksxHem.clustersRH(:,1:3) networksxHem.clustersRH(:,5) networksxHem.
clustersRH(:,7:end)];
good_nets_LH = mean(good_nets_LH);
good_nets_RH = mean(good_nets_RH);
means = [good_nets_LH' good_nets_RH'];
for net = 1:16
se_lh(net) = std(networksxHem.clustersLH(:,net))/sqrt(670);
se_rh(net) = std(networksxHem.clustersRH(:,net))/sqrt(670);
end
se_lh(4) = [];
se_lh(5) = [];
se_rh(4) = [];
se_rh(5) = [];
bar(1:14, means)
legend('Left Hem', 'Right Hem')
xticks(1:14)
xticklabels(network_names)

```



```
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.9, 0.7]);
ylabel('Average Number of Variant Regions')
xlabel('Network');
title('Variant Regions Assigned to Each Network Across Hemispheres')
ax = gca;
ax.FontSize = 24;
hold on
h = errorbar(.85:1:13.85, means(:,1), se_lh, se_lh)
h.Color = [0,0,0];
g = errorbar(1.15:1:14.15, means(:,2), se_rh, se_rh)
g.Color = [0,0,0];
h.LineStyle = 'none';
g.LineStyle = 'none';
h.LineWidth = 2;
g.LineWidth = 2;
legend('Left Hem', 'Right Hem')
print(gcf, '
/Users/dianaperez/Desktop/Research/lateralization_code/HCP752_numvars_Rhand_leftvrighthe
m.jpg', '-dpng', '-r300')
axis([0,15,0,2.5])
print(gcf, '
/Users/dianaperez/Desktop/Research/lateralization_code/HCP752_numvars_Rhand_leftvrighthe
m.jpg', '-dpng', '-r300')
right_handlers = networksxHem;
close gcf
load
('/Users/dianaperez/Desktop/Research/lateralization_code/testing_output/HCP752_new_split_
LH_networksxHem.mat')
network_names = {'DMN' 'Vis' 'FP' 'DAN' 'Lang' 'Sal' 'CO' 'SMd' 'SMl'
'Aud' 'Tpole' 'MTL' 'PMN' 'PON'};
good_nets_LH = [networksxHem.verticesLH(:,1:3) networksxHem.verticesLH(:,5) networksxHem.
verticesLH(:,7:end)];
good_nets_LH = [networksxHem.clustersLH(:,1:3) networksxHem.clustersLH(:,5) networksxHem.
clustersLH(:,7:end)];
good_nets_RH = [networksxHem.clustersRH(:,1:3) networksxHem.clustersRH(:,5) networksxHem.
clustersRH(:,7:end)];
good_nets_LH = mean(good_nets_LH);
good_nets_RH = mean(good_nets_RH);
means = [good_nets_LH' good_nets_RH'];
for net = 1:16
se_lh(net) = std(networksxHem.clustersLH(:,net))/sqrt(40);
se_rh(net) = std(networksxHem.clustersRH(:,net))/sqrt(40);
end
se_lh(4) = [];
se_lh(5) = [];
se_rh(4) = [];
se_rh(5) = [];
bar(1:14, means)
legend('Left Hem', 'Right Hem')
xticks(1:14)
xticklabels(network_names)
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.9, 0.7]);
ylabel('Average Number of Variant Regions')
xlabel('Network');
title('Variant Regions Assigned to Each Network Across Hemispheres')
ax = gca;
ax.FontSize = 24;
hold on
h = errorbar(.85:1:13.85, means(:,1), se_lh, se_lh)
h.Color = [0,0,0];
```

```

g = errorbar(1.15:1:14.15, means(:,2), se_rh, se_rh)
g.Color = [0,0,0];
h.LineStyle = 'none';
g.LineStyle = 'none';
h.LineWidth = 2;
g.LineWidth = 2;
legend('Left Hem', 'Right Hem')
print(gcf, '
/Users/dianaperez/Desktop/Research/lateralization_code/HCP752_numvars_Lhand_leftvrighthe
m.jpg', '-dpng', '-r300')
close gcf
left_handlers = networksxHem;
diff_lefthanders = left_handlers.clustersLH - left_handlers.clustersRH;
mean_diff_lefthanders = mean(diff_lefthanders, 1);
diff_righthanders = right_handlers.clustersLH - right_handlers.clustersRH;
mean_diff_righthanders = mean(diff_righthanders, 1);
for net = 1:16
se_lh(net) = std(diff_lefthanders(:,net))/sqrt(40);
se_rh(net) = std(diff_righthanders(:,net))/sqrt(670);
end
se_lh(4) = [];
se_lh(5) = [];
se_rh(4) = [];
se_rh(5) = [];
diffs = [mean_diff_lefthanders; mean_diff_righthanders];
bar(1:14, diffs)
diffs(:,4) = []; diff(:,5) = [];
diffs(:,5) = []
bar(1:14, diffs)
legend('Left Handers', 'Right Handers')
xticks(1:14)
xticklabels(network_names)
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.9, 0.7]);
ylabel('Difference in Average Number of Variant Regions (left - right hem)')
xlabel('Network');
ax = gca;
ax.FontSize = 24;
hold on
h = errorbar(.85:1:13.85, diffs(:,1), se_lh, se_lh)
h.Color = [0,0,0];
g = errorbar(1.15:1:14.15, diffs(:,2), se_rh, se_rh)
g.Color = [0,0,0];
h.LineStyle = 'none';
g.LineStyle = 'none';
h.LineWidth = 2;
g.LineWidth = 2;
h = errorbar(.85:1:13.85, diffs(1,:), se_lh, se_lh)
h.Color = [0,0,0];
g = errorbar(1.15:1:14.15, diffs(2,:), se_rh, se_rh)
g.Color = [0,0,0];
h.LineStyle = 'none';
g.LineStyle = 'none';
h.LineWidth = 2;
g.LineWidth = 2;
legend('Left Handers', 'Right Handers')
print(gcf, '
/Users/dianaperez/Desktop/Research/lateralization_code/HCP752_numvars_LvRhand_L-Rhem.
jpg', '-dpng', '-r300')
close gcf
load

```

```

('/Users/dianaperez/Desktop/Research/lateralization_code/testing_output/HCP752_new_split_
LH_networksxHem.mat')
network_names = {'DMN' 'Vis' 'FP' 'DAN' 'Lang' 'Sal' 'CO' 'SMd' 'SML'
'Aud' 'Tpole' 'MTL' 'PMN' 'PON'};
good_nets_LH = [networksxHem.verticesLH(:,1:3) networksxHem.verticesLH(:,5) networksxHem.
verticesLH(:,7:end)];
good_nets_LH = [networksxHem.clustersLH(:,1:3) networksxHem.clustersLH(:,5) networksxHem.
clustersLH(:,7:end)];
good_nets_RH = [networksxHem.clustersRH(:,1:3) networksxHem.clustersRH(:,5) networksxHem.
clustersRH(:,7:end)];
good_nets_LH = mean(good_nets_LH);
good_nets_RH = mean(good_nets_RH);
means = [good_nets_LH' good_nets_RH'];
for net = 1:16
se_lh(net) = std(networksxHem.clustersLH(:,net))/sqrt(40);
se_rh(net) = std(networksxHem.clustersRH(:,net))/sqrt(40);
end
se_lh(4) = [];
se_lh(5) = [];
se_rh(4) = [];
se_rh(5) = [];
bar(1:14, means)
legend('Left Hem', 'Right Hem')
xticks(1:14)
xticklabels(network_names)
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.9, 0.7]);
ylabel('Average Number of Variant Regions')
xlabel('Network');
title('Variant Regions Assigned to Each Network Across Hemispheres')
ax = gca;
ax.FontSize = 24;
hold on
h = errorbar(.85:1:13.85, means(:,1), se_lh, se_lh)
h.Color = [0,0,0];
g = errorbar(1.15:1:14.15, means(:,2), se_rh, se_rh)
g.Color = [0,0,0];
h.LineStyle = 'none';
g.LineStyle = 'none';
h.LineWidth = 2;
g.LineWidth = 2;
legend('Left Hem', 'Right Hem')
print(gcf,
'/Users/dianaperez/Desktop/Research/lateralization_code/HCP752_numvars_Lhand_leftvrighthe
m.jpg', '-dpng', '-r300')
close gcf
left_handlers = networksxHem;
%% right handlers
load
('/Users/dianaperez/Desktop/Research/lateralization_code/testing_output/HCP752_RH_network
sxHem.mat')
network_names = {'DMN' 'Vis' 'FP' 'DAN' 'Lang' 'Sal' 'CO' 'SMd' 'SML'
'Aud' 'Tpole' 'MTL' 'PMN' 'PON'};
good_nets_LH = [networksxHem.verticesLH(:,1:3) networksxHem.verticesLH(:,5) networksxHem.
verticesLH(:,7:end)];
good_nets_LH = [networksxHem.clustersLH(:,1:3) networksxHem.clustersLH(:,5) networksxHem.
clustersLH(:,7:end)];
good_nets_RH = [networksxHem.clustersRH(:,1:3) networksxHem.clustersRH(:,5) networksxHem.
clustersRH(:,7:end)];
good_nets_LH = mean(good_nets_LH);
good_nets_RH = mean(good_nets_RH);

```

```

means = [good_nets_LH' good_nets_RH'];
for net = 1:16
se_lh(net) = std(networksxHem.clustersLH(:,net))/sqrt(670);
se_rh(net) = std(networksxHem.clustersRH(:,net))/sqrt(670);
end
se_lh(4) = [];
se_lh(5) = [];
se_rh(4) = [];
se_rh(5) = [];
bar(1:14, means)
legend('Left Hem', 'Right Hem')
xticks(1:14)
xticklabels(network_names)
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.9, 0.7]);
ylabel('Average Number of Variant Regions')
xlabel('Network');
title('Variant Regions Assigned to Each Network Across Hemispheres')
ax = gca;
ax.FontSize = 24;
hold on
h = errorbar(.85:1:13.85, means(:,1), se_lh, se_lh)
h.Color = [0,0,0];
g = errorbar(1.15:1:14.15, means(:,2), se_rh, se_rh)
g.Color = [0,0,0];
h.LineStyle = 'none';
g.LineStyle = 'none';
h.LineWidth = 2;
g.LineWidth = 2;
legend('Left Hem', 'Right Hem')
axis([0,15,0,2.5])
print(gcf,
'/Users/dianaperez/Desktop/Research/lateralization_code/HCP752_numvars_Rhand_leftvrighthem.jpg', '-dpng', '-r300')
close gcf
right_handlers = networksxHem;
%% difference (left - right)
% mean_lefthanders_LH = mean(left_handlers.clustersLH);
% mean_lefthanders_RH = mean(left_handlers.clustersRH);
% mean_righthanders_LH = mean(right_handlers.clustersLH);
% mean_righthanders_RH = mean(right_handlers.clustersRH);
% diff_lefthanders = mean_lefthanders_LH - mean_lefthanders_RH;
% diff_righthanders = mean_righthanders_LH - mean_righthanders_RH;
% diff_lefthanders(4) = []; diff_lefthanders(5) = [];
% diff_righthanders(4) = []; diff_righthanders(5) = [];
% diffs = [diff_lefthanders; diff_righthanders];
%% OR
diff_lefthanders = left_handlers.clustersLH - left_handlers.clustersRH;
mean_diff_lefthanders = mean(diff_lefthanders, 1);
diff_righthanders = right_handlers.clustersLH - right_handlers.clustersRH;
mean_diff_righthanders = mean(diff_righthanders, 1);
for net = 1:16
se_lh(net) = std(diff_lefthanders(:,net))/sqrt(40);
se_rh(net) = std(diff_righthanders(:,net))/sqrt(670);
end
se_lh(4) = [];
se_lh(5) = [];
se_rh(4) = [];
se_rh(5) = [];
diffs = [mean_diff_lefthanders; mean_diff_righthanders];
diffs(:,4) = []; diffs(:,5) = [];

```

```

bar(1:14, diffs)
legend('Left Handers', 'Right Handers')
xticks(1:14)
xticklabels(network_names)
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.9, 0.7]);
ylabel('Difference in Average Number of Variant Regions (left - right hem)')
xlabel('Network');
ax = gca;
ax.FontSize = 24;
hold on
h = errorbar(.85:1:13.85, diffs(1,:), se_lh, se_lh)
h.Color = [0,0,0];
g = errorbar(1.15:1:14.15, diffs(2,:), se_rh, se_rh)
g.Color = [0,0,0];
h.LineStyle = 'none';
g.LineStyle = 'none';
h.LineWidth = 2;
g.LineWidth = 2;
legend('Left Handers', 'Right Handers')
print(gcf, '
/Users/dianaperez/Desktop/Research/lateralization_code/HCP752_numvars_LvRhand_L-Rhem.
jpg', '-dpng', '-r300')
close gcf
diff_lefthanders = left_handers.clustersLH - left_handers.clustersRH;
mean_diff_lefthanders = mean(diff_lefthanders, 1);
diff_righthanders = right_handers.clustersLH - right_handers.clustersRH;
mean_diff_righthanders = mean(diff_righthanders, 1);
for net = 1:16
se_lh(net) = std(diff_lefthanders(:,net))/sqrt(40);
se_rh(net) = std(diff_righthanders(:,net))/sqrt(670);
end
se_lh(4) = [];
se_lh(5) = [];
se_rh(4) = [];
se_rh(5) = [];
diffs = [mean_diff_lefthanders; mean_diff_righthanders];
diffs(:,4) = []; diffs(:,5) = [];
b = bar(1:14, diffs)
legend('Left Handers', 'Right Handers')
b(1).FaceColor = [.2 .6 .5];
b(2).FaceColor = [0.4940 0.1840 0.5560];
xticks(1:14)
xticklabels(network_names)
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.9, 0.7]);
ylabel('Difference in Average Number of Variant Regions (left - right hem)')
xlabel('Network');
ax = gca;
ax.FontSize = 24;
hold on
h = errorbar(.85:1:13.85, diffs(1,:), se_lh, se_lh)
h.Color = [0,0,0];
g = errorbar(1.15:1:14.15, diffs(2,:), se_rh, se_rh)
g.Color = [0,0,0];
h.LineStyle = 'none';
g.LineStyle = 'none';
h.LineWidth = 2;
g.LineWidth = 2;
legend('Left Handers', 'Right Handers')
print(gcf, '
/Users/dianaperez/Desktop/Research/lateralization_code/HCP752_numvars_LvRhand_L-Rhem.

```

[illegible]

[illegible]

%-- 3/21/22, 9:54 AM --%

EmReg_task

```
input_condition{1}>4
```

```
isnum(input_condition{1})
```

```
isenum(input_condition{1})
```

```
help ischar
```

```
isnumeric(input_condition{1})
```

help inputdlg

```
cell2mat(input_condition{1})
```

```
cell2mat(input_condition)
```

```
isnumeric(ans)
```

```
num=cell2mat(input_condition)
```

```
ischar(num
```

```
ischar(num)
```

```
num=str2num(input_condition)
```

```
num=str2num(input_condition{1})
```

EmReg_task

%-- 3/21/22, 10:16 AM --%

EmReg_task

%-- 3/21/22, 10:21 AM --%

EmReg_task

```
cd /Volumes/RESEARCH_HD
```

EmReg_task

1
1
1
1
1
1
1
1
1
1
1
1
1
1
1

sca

EmReg_task

```
0
0
0
0
0
0
0
0
0
0
0
0
0
0
0
0
load('EmoRegBaseline_LStrst.mat')
EmReg_task
1
1
2
3
4
5
4
3
4
5
6
4
4
-1
f3
4
5
3
53
3
4
3
2
3
4
5
4
-3
2
cd /Volumes/RESEARCH_HD
q 3
4
3
5
5 5
4
5
6
5
4
5
6
4
```



```
3
4
q 2
3
4
3
2
3
4
5
4
5
4
5
3
-2
3
q 3
2
3
4
5
3
4
5
4
3
2
3
4
1
3
2
3
4
5
4
3
2
3
4
5
6
4
3
2
3
%-- 3/28/22, 2:02 PM --%
network_seg_new
addpath(genpath('/Users/dianaperez/Documents/Dependencies/Resources/'))
network_seg_new
within(within<0) = 0;
network_seg_new
mean(within>0)
network_seg_new
between(between<0) = 0;
network_seg_new
mean(within_network)
mean(within_network{:, :, :})
network_seg_new
```

```
network_z.within{sub,ses}(net)
permute_net_assignment
net_size_across_hems
cd '/Users/dianaperez/Documents/GitHub/PerezEtAl_HemAsymmetries'
net_size_across_hems
group_avg.data(26895)
group_avg.data(26764)
newcifti = zeros(1:59412);
newcifti = zeros(1,1:59412);
newcifti = zeros(1,59412);
old_cifti = group_avg.data(brainstructure==1 || brainstructure==2);
old_cifti = group_avg.data(group_avg.brainstructure==1 || group_avg.brainstructure==2);
old_cifti_LH = group_avg.data(group_avg.brainstructure==1);
old_cifti_RH = group_avg.data(group_avg.brainstructure==2);
old_cifti = [old_cifti_LH; old_cifti_RH];
clear old_cifti_LH old_cifti_RH
unique(old_cifti)
newcifti(old_cifti==14) = 14;
template = ft_read_cifti_mod(
('/Users/dianaperez/Desktop/Research/Lateralization/Spatial_Location/HCP/HCP_newsplitvars
_diffMap_afterclustercorrect_allSubs.dtseries.nii');
template.data = newcifti;
ft_write_cifti_mod('/Users/dianaperez/Desktop/Network14only.dtseries.nii', template)
template.data = newcifti;
ft_write_cifti_mod('/Users/dianaperez/Desktop/Network14only.dtseries.nii', template)
old_cifti = group_avg.data(group_avg.brainstructure>0);
old_cifti = [];
old_cifti = group_avg.data(group_avg.brainstructure>0);
group_avg.brainstructure>0
old_cifti = group_avg.data(group_avg.brainstructure>0);
newcifti(group_avg.data==14) = 14;
template.data = newcifti;
ft_write_cifti_mod('/Users/dianaperez/Desktop/Network14only.dtseries.nii', template)
newcifti(group_avg.data==13) = 13;
template.data = newcifti;
ft_write_cifti_mod('/Users/dianaperez/Desktop/Network13only.dtseries.nii', template)
newcifti = zeros(59412,1);
newcifti(group_avg.data==0) = 1;
ft_write_cifti_mod('/Users/dianaperez/Desktop/NonNetworksOnly.dtseries.nii', template)
template.data = newcifti;
ft_write_cifti_mod('/Users/dianaperez/Desktop/NonNetworksOnly.dtseries.nii', template)
template.data = newcifti(1:59412);
ft_write_cifti_mod('/Users/dianaperez/Desktop/NonNetworksOnly.dtseries.nii', template)
newcifti = zeros(59412,1);
newcifti(group_avg.data==0) = 1;
template.data = newcifti(1:59412);
ft_write_cifti_mod('/Users/dianaperez/Desktop/NonNetworksOnly.dtseries.nii', template)
newcifti = zeros(59412,1);
newcifti(group_avg.data==4) = 4;
template.data = newcifti(1:59412);
ft_write_cifti_mod('/Users/dianaperez/Desktop/Network40only.dtseries.nii', template)
newcifti = zeros(59412,1);
newcifti(group_avg.data==6) = 6;
template.data = newcifti(1:59412);
ft_write_cifti_mod('/Users/dianaperez/Desktop/Network60only.dtseries.nii', template)
permute_net_assignment
net_size_across_hems
[out_dir out_str '_groupavg_nets_surfarea_scatter.jpg']
%-- 3/31/22, 10:24 AM --%
cd /Users/dianaperez/Documents/GitHub/
```

```
var_freq_size_net
addpath(genpath('/Users/dianaperez/Documents/Dependencies/cifti-matlab-master'))
var_freq_size_net
good_nets_RH = sum(good_nets_RH);
good_nets_LH = sum(good_nets_LH);
means = [good_nets_LH' good_nets_RH'];
bar(1:14, means)
legend('Left Hem', 'Right Hem')
xticks(1:14)
xticklabels(network_names)
clear all
network_seg_new
addpath '/Users/dianaperez/Documents/GitHub/GrattonLab-General-Repo/FCProcess'
network_seg_new
untitled
addpath(genpath('/Users/dianaperez/Documents/Dependencies/cifti-matlab-master'))
untitled
addpath(genpath('/Users/dianaperez/Documents/Dependencies/Resources'))
untitled
size_nets(net)/288
untitled
mean(node_SI)
untitled
rois(1)
rois(end)
test = matrix(rois)
test = matrix(rois,rois)
untitled
mean(node_SI)
untitled
mean(node_SI)
node_SI = [0.57379448,0.42760423,0.40719646,-0.0084462678,-2.2896698,0.39838830,↵
-0.31563443,-1.0662855,-0.026975855,-0.13974182,-1.0860931]
mean(node_SI)
untitled
figure(matrix_orig)
imagesc(matrix_orig)
imagesc(matrix_sorted)
imagesc(matrix_orig)
imagesc(matrix_sorted)
imagesc(matrix_orig,[0 1]); colormap('jet');
imagesc(matrix_sorted,[0 1]); colormap('jet');
figure
imagesc(matrix_orig,[0 1]); colormap('jet');
imagesc(matrix_orig,[0.4 1]); colormap('jet');
imagesc(matrix_orig,[-0.4 1]); colormap('jet');
figure
imagesc(matrix_sorted,[-0.4 1]); colormap('jet');
untitled
skip = net_size(1);
net_size(1) = [];
weights = net_size./(300-skip)
atlas_params.networks{2:e14}
atlas_params.networks{2:14}
untitled
atlas_params.networks{2:14}
network_z.networks{1:13} = atlas_params.networks{2:14}
untitled
maskmat = ones(size(tmp_between));
rois(1):rois(end)
```

```
maskmat(:,rois(1):rois(end)) = 0;
untitled
help reshape
between = reshape(between,length(rois), 300-length(rois));
untitled
clear all
untitled
node_between = between(roi,:);
untitled
mean(within)
untitled
mean(between)
untitled
net_size(1)
net_size(5)
untitled
net
untitled
node_within = tmp_within(roi,:);
node_within(roi) = [];
node_within(node_within<0) = [];
node_mean_within = mean(node_within);
untitled
%-- 4/4/22, 1:24 PM --%
segregation_ind_by_node
addpath '/Users/dianaperez/Documents/GitHub/GrattonLab-General-Repo/FCProcess'
segregation_ind_by_node
mean(within_corrs)
mean(means_within)
mean(between_corrs)
mean(means_between)
segregation_ind_by_node
[output_dir subs(sub) '_seg_index_node.mat']
[output_dir subs{sub} '_seg_index_node.mat']
save([output_dir subs{sub} '_seg_index_node.mat'], 'sub_struct')
segregation_ind_by_node
load('/Users/dianaperez/Desktop/Segregation_Analyses/LS03_seg_index_node.mat')
clear all
load('/Users/dianaperez/Desktop/Segregation_Analyses/LS03_seg_index_node.mat')
ft_read_cifti_mod('/Users/dianaperez/Desktop/sub-LS16_vs_120_avg_corr_LR_cortex_corr.
dtseries.nii')
ls16 = ans;
cd /Volumes/Back_Up/
ls
cd Lifespan
cd /Users/dianaperez/Desktop
cd Research/CNS_analyses/
template = ft_read_cifti_mod('sub-LS02_spCorrMap.dtseries.nii');
template.data = ls16.data;
ft_write_cifti_mod('sub-LS16_spCorrMap.dtseries.nii', template);
%-- 4/7/22, 12:53 PM --%
CreateVariantFiles_LS03specific
addpath(genpath('/Users/dianaperez/Documents/Dependencies/'))
CreateVariantFiles_LS03specific
SNRpath
ft_read_cifti_mod(SNRpath)
CreateVariantFiles_LS03specific
addpath '/Users/dianaperez/Documents/GitHub/Lifespan-Analysis'
CreateVariantFiles_LS03specific
%-- 4/9/22, 11:31 AM --%
```

```

cd /Volumes/RESEARCH_HD
cd /Users/dianaperez/Documents/
reliability_analysis
mean = mean(corr,1);
times =[5:5:100];
figure;
plot(times(1:20),corr(1,1:20),'Color',[1, 0, 0],'LineWidth', 3) %LS02
hold on
plot(times(1:20),corr(2,1:20),'Color',[0, 1, 0],'LineWidth', 3) %LS03
hold on
plot(times(1:20),corr(3,1:20),'Color',[0, 0, 1],'LineWidth', 3)%LS05
hold on
plot(times(1:3),corr(4,1:3),'Color',[0, 1, 1],'LineWidth', 3)%LS08
hold on
plot(times(1:20),corr(5,1:20),'Color',[1, 0, 1],'LineWidth', 3)%LS11
hold on
plot(times(1:20),corr(6,1:20),'Color',[0.4660 0.6740 0.1880],'LineWidth', 3)%LS14
hold on
plot(times(1:20),corr(7,1:9),'Color',[0.9290 0.6940 0.1250],'LineWidth', 3)%LS14
hold on
plot(times(1:20),corr(8,1:14),'Color',[0.4940 0.1840 0.5560],'LineWidth', 3)%LS14
hold on
plot(times(1:20),mean, ':', 'Color', [0,0,0], 'LineWidth',3) %average
close gcf
times =[5:5:100];
figure;
plot(times(1:20),corr(1,1:20),'Color',[1, 0, 0],'LineWidth', 3) %LS02
hold on
plot(times(1:20),corr(2,1:20),'Color',[0, 1, 0],'LineWidth', 3) %LS03
hold on
plot(times(1:20),corr(3,1:20),'Color',[0, 0, 1],'LineWidth', 3)%LS05
hold on
plot(times(1:3),corr(4,1:3),'Color',[0, 1, 1],'LineWidth', 3)%LS08
hold on
plot(times(1:20),corr(5,1:20),'Color',[1, 0, 1],'LineWidth', 3)%LS11
hold on
plot(times(1:20),corr(6,1:20),'Color',[0.4660 0.6740 0.1880],'LineWidth', 3)%LS14
hold on
plot(times(1:20),corr(7,1:9),'Color',[0.9290 0.6940 0.1250],'LineWidth', 3)%LS14
hold on
plot(times(1:20),corr(8,1:14),'Color',[0.4940 0.1840 0.5560],'LineWidth', 3)%LS14
hold on
plot(times(1:20),mean(1:20), ':', 'Color', [0,0,0], 'LineWidth',3) %average
plot(times(1:20),mean(1:20), ':', 'Color', [0,0,0], 'LineWidth',3)
close gcf
times =[5:5:100];
figure;
plot(times(1:20),corr(1,1:20),'Color',[1, 0, 0],'LineWidth', 3) %LS02
hold on
plot(times(1:20),corr(2,1:20),'Color',[0, 1, 0],'LineWidth', 3) %LS03
hold on
plot(times(1:20),corr(3,1:20),'Color',[0, 0, 1],'LineWidth', 3)%LS05
hold on
plot(times(1:3),corr(4,1:3),'Color',[0, 1, 1],'LineWidth', 3)%LS08
hold on
plot(times(1:20),corr(5,1:20),'Color',[1, 0, 1],'LineWidth', 3)%LS11
hold on
plot(times(1:20),corr(6,1:20),'Color',[0.4660 0.6740 0.1880],'LineWidth', 3)%LS14
hold on
plot(times(1:9),corr(7,1:9),'Color',[0.9290 0.6940 0.1250],'LineWidth', 3)%LS14

```

```
hold on
plot(times(1:14),corr(8,1:14),'Color',[0.4940 0.1840 0.5560],'LineWidth', 3)%LS14
hold on
plot(times(1:20),mean(1:20), ':', 'Color', [0,0,0], 'LineWidth',3) %average
mean = [0.580319941 0.671566779 0.723272658 0.757708593 0.778572007 0.788029527↵
0.79889902 0.808729104 0.818726873 0.821323197 0.827748935 0.836636393 0.843602278↵
0.847735413 0.852112674 0.856295884 0.861091436 0.862434778 0.865438688 0.867434122]
close gcf
figure;
plot(times(1:20),corr(1,1:20),'Color',[1, 0, 0],'LineWidth', 3) %LS02
hold on
plot(times(1:20),corr(2,1:20),'Color',[0, 1, 0],'LineWidth', 3) %LS03
hold on
plot(times(1:20),corr(3,1:20),'Color',[0, 0, 1],'LineWidth', 3)%LS05
hold on
plot(times(1:3),corr(4,1:3),'Color',[0, 1, 1],'LineWidth', 3)%LS08
hold on
plot(times(1:20),corr(5,1:20),'Color',[1, 0, 1],'LineWidth', 3)%LS11
hold on
plot(times(1:20),corr(6,1:20),'Color',[0.4660 0.6740 0.1880],'LineWidth', 3)%LS14
hold on
plot(times(1:9),corr(7,1:9),'Color',[0.9290 0.6940 0.1250],'LineWidth', 3)%LS14
hold on
plot(times(1:14),corr(8,1:14),'Color',[0.4940 0.1840 0.5560],'LineWidth', 3)%LS14
hold on
plot(times(1:20),mean(1:20), ':', 'Color', [0,0,0], 'LineWidth',3) %average
ylabel('Pearson Correlation (r)');
xlabel('Time (Minutes)');
m = findobj(gca,'Type','line');
hleg1 = legend(m(1:7), 'Average', 'LS14', 'LS11', 'LS08', 'LS05', 'LS03', 'LS02',↵
'Location', 'SouthEast');
hleg1.FontSize = 14;
ax = gca;
ax.FontSize = 17;
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.5, 0.7]);
close gcf
figure;
plot(times(1:20),corr(1,1:20),'Color',[1, 0, 0],'LineWidth', 3) %LS02
hold on
plot(times(1:20),corr(2,1:20),'Color',[0, 1, 0],'LineWidth', 3) %LS03
hold on
plot(times(1:20),corr(3,1:20),'Color',[0, 0, 1],'LineWidth', 3)%LS05
hold on
plot(times(1:3),corr(4,1:3),'Color',[0, 1, 1],'LineWidth', 3)%LS08
hold on
plot(times(1:20),corr(5,1:20),'Color',[1, 0, 1],'LineWidth', 3)%LS11
hold on
plot(times(1:20),corr(6,1:20),'Color',[0.4660 0.6740 0.1880],'LineWidth', 3)%LS14
hold on
plot(times(1:9),corr(7,1:9),'Color',[0.9290 0.6940 0.1250],'LineWidth', 3)%LS14
hold on
plot(times(1:14),corr(8,1:14),'Color',[0.4940 0.1840 0.5560],'LineWidth', 3)%LS14
hold on
plot(times(1:20),mean(1:20), ':', 'Color', [0,0,0], 'LineWidth',3) %average
ylabel('Pearson Correlation (r)');
xlabel('Time (Minutes)');
m = findobj(gca,'Type','line');
hleg1 = legend(m(1:9), 'Average', 'LS14', 'LS11', 'LS08', 'LS05', 'LS03', 'LS02',↵
'Location', 'SouthEast');
hleg1.FontSize = 14;
```

```

ax = gca;
ax.FontSize = 17;
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.5, 0.7]);
close gcf
times = [5:5:100];
figure;
plot(times(1:20), corr(1,1:20), 'Color', [1, 0, 0], 'LineWidth', 3) %LS02
hold on
plot(times(1:20), corr(2,1:20), 'Color', [0, 1, 0], 'LineWidth', 3) %LS03
hold on
plot(times(1:20), corr(3,1:20), 'Color', [0, 0, 1], 'LineWidth', 3) %LS05
hold on
plot(times(1:3), corr(4,1:3), 'Color', [0, 1, 1], 'LineWidth', 3) %LS08
hold on
plot(times(1:20), corr(5,1:20), 'Color', [1, 0, 1], 'LineWidth', 3) %LS11
hold on
plot(times(1:20), corr(6,1:20), 'Color', [0.4660 0.6740 0.1880], 'LineWidth', 3) %LS14
hold on
plot(times(1:9), corr(7,1:9), 'Color', [0.9290 0.6940 0.1250], 'LineWidth', 3) %LS14
hold on
plot(times(1:14), corr(8,1:14), 'Color', [0.4940 0.1840 0.5560], 'LineWidth', 3) %LS14
hold on
plot(times(1:20), mean(1:20), ':', 'Color', [0,0,0], 'LineWidth', 3) %average
ylabel('Pearson Correlation (r)');
xlabel('Time (Minutes)');
m = findobj(gca, 'Type', 'line');
hleg1 = legend(m(1:9), 'Average', 'LS17', 'LS17', 'LS14', 'LS11', 'LS08', 'LS05', 'LS03',
'LS02', 'Location', 'SouthEast');
hleg1.FontSize = 14;
ax = gca;
ax.FontSize = 17;
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.5, 0.7]);
close gcf
figure;
plot(times(1:20), corr(1,1:20), 'Color', [1, 0, 0], 'LineWidth', 3) %LS02
hold on
plot(times(1:20), corr(2,1:20), 'Color', [0, 1, 0], 'LineWidth', 3) %LS03
hold on
plot(times(1:20), corr(3,1:20), 'Color', [0, 0, 1], 'LineWidth', 3) %LS05
hold on
plot(times(1:3), corr(4,1:3), 'Color', [0, 1, 1], 'LineWidth', 3) %LS08
hold on
plot(times(1:20), corr(5,1:20), 'Color', [1, 0, 1], 'LineWidth', 3) %LS11
hold on
plot(times(1:20), corr(6,1:20), 'Color', [0.4660 0.6740 0.1880], 'LineWidth', 3) %LS14
hold on
plot(times(1:9), corr(7,1:9), 'Color', [0.9290 0.6940 0.1250], 'LineWidth', 3) %LS14
hold on
plot(times(1:14), corr(8,1:14), 'Color', [0.4940 0.1840 0.5560], 'LineWidth', 3) %LS14
hold on
plot(times(1:20), mean(1:20), ':', 'Color', [0,0,0], 'LineWidth', 3) %average
ylabel('Pearson Correlation (r)');
xlabel('Time (Minutes)');
m = findobj(gca, 'Type', 'line');
hleg1 = legend(m(1:9), 'Average', 'LS17', 'LS16', 'LS14', 'LS11', 'LS08', 'LS05', 'LS03',
'LS02', 'Location', 'SouthEast');
hleg1.FontSize = 14;
ax = gca;
ax.FontSize = 17;
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.5, 0.7]);

```

```
print(gcf,↵
['/Volumes/RESEARCH_HD/Lifespan/CNS_Analyses/ReliabilityLifespanRestDatatruhalf' num2str↵
(pts2sample) '.jpg'], '-dpng', '-r300');
similarity_analysis_forced_same_amt_data
matched_data = datasample(masked_data,min_data,2,'Replace', false);
similarity_analysis_forced_same_amt_data
addpath(genpath('/Users/dianaperez/Documents/Dependencies/'))
matcheddata_corrlin(count,:) = single(FisherTransform(corrmat_matched_data(maskmat)));
similarity_analysis_forced_same_amt_data
figure('Position',[1 1 1000 800]);
imagesc(simmat,[0 1]); colormap('jet');
hline_new([5,10,15,20,25,30,35,40]+0.5,'k',2);
hline_new(0,'k',2);
hline_new(0.5,'k',2);
vline_new([0,5,10,15,20,25,30,35,40]+0.5,'k',2);
set(gca,'XTick',[4,13.5,19,24.5,30.5,34,37,40], 'YTick',↵
[4,13.5,19,24.5,30.5,34,37,40,45], 'XTickLabel',...
{'LS02', 'LS03', 'LS04', 'LS05', 'LS07', 'LS08', 'LS10', 'LS11', 'LS14', 'LS16', 'LS17'},↵
'YTickLabel', {'LS02', 'LS03', 'LS04', 'LS05', 'LS07', 'LS08', 'LS10', 'LS11', 'LS14'});
set(gca,'XTick',[2.5,7.5,12.5,17.5,22.5,27.5,32.5,37.5], 'YTick',↵
[2.5,7.5,12.5,17.5,22.5,27.5,32.5,37.5], 'XTickLabel',...
{'LS02', 'LS03', 'LS04', 'LS05', 'LS07', 'LS08', 'LS10', 'LS11', 'LS14', 'LS16', 'LS17'},↵
'YTickLabel', {'LS02', 'LS03', 'LS04', 'LS05', 'LS07', 'LS08', 'LS10', 'LS11', 'LS14'});
set(gca,'XTick',[3,8,13,18,23,28,33,38], 'YTick', [3,8,13,18,23,28,33,38],↵
'XTickLabel',...
{'LS02', 'LS03', 'LS04', 'LS05', 'LS07', 'LS08', 'LS10', 'LS11', 'LS14', 'LS16', 'LS17'},↵
'YTickLabel', {'LS02', 'LS03', 'LS04', 'LS05', 'LS07', 'LS08', 'LS10', 'LS11', 'LS14'});
axis square;
colorbar;
set(gca,'XTick',[3,8,13,18,23,28,33,38], 'YTick', [3,8,13,18,23,28,33,38],↵
'XTickLabel',...
{'LS02', 'LS03', 'LS05', 'LS08', 'LS11', 'LS14', 'LS16', 'LS17'}, 'YTickLabel', {'LS02',↵
'LS03', 'LS05', 'LS08', 'LS11', 'LS14', 'LS16', 'LS17'});
axis square;
colorbar;
title('Correlation Matrix Similarity');
saveas(gcf,[outdir 'SimilarityMat_rest_matched_data.tiff'],'tiff');
close('all');
figure('Position',[1 1 1000 800]);
imagesc(simmat,[0 1]); colormap('jet');
hline_new([0,5,10,15,20,25,30,35,40]+0.5,'k',2);
vline_new([0,5,10,15,20,25,30,35,40]+0.5,'k',2);
set(gca,'XTick',[3,8,13,18,23,28,33,38], 'YTick', [3,8,13,18,23,28,33,38],↵
'XTickLabel',...
{'LS02', 'LS03', 'LS05', 'LS08', 'LS11', 'LS14', 'LS16', 'LS17'}, 'YTickLabel', {'LS02',↵
'LS03', 'LS05', 'LS08', 'LS11', 'LS14', 'LS16', 'LS17'});
axis square;
colorbar;
ax = gca;
ax.FontSize = 24;
saveas(gcf,[outdir 'SimilarityMat_rest_matched_data.tiff'],'tiff');
reliability_analysis
mean = [0.580319941 0.671566779 0.723272658 0.757708593 0.778572007 0.788029527↵
0.79889902 0.808729104 0.818726873 0.821323197 0.827748935 0.836636393 0.843602278↵
0.847735413 0.852112674 0.856295884 0.861091436 0.862434778 0.865438688 0.867434122];
hleg1.FontSize = 20;
ax = gca;
ax.FontSize = 24;
hleg1.FontSize = 20;
hleg1.FontSize = 15;
```



```

hleg1.FontSize = 18;
hleg1.FontSize = 16;
hleg1 = legend(m(1:9), 'Mean', 'LS17', 'LS16', 'LS14', 'LS11', 'LS08', 'LS05', 'LS03',
'LS02', 'Location', 'SouthEast');
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.5, 0.7]);
hleg1.FontSize = 20;
ax.FontSize = 24;
hleg1.FontSize = 24;
hleg1.FontSize = 23;
print(gcf,
['/Volumes/RESEARCH_HD/Lifespan/CNS_Analyses/ReliabilityLifespanRestDatatruhalf' num2str
(pts2sample) '.jpg'], '-dpng', '-r300');
similarity_analysis_wLongitudinal
hline_new([8,18,19,29,31,36,37,42,47,53]+0.5, 'k', 2);
hline_new([8,18,19,29,31,36,37,42,47,52]+0.5, 'k', 2);
close(gcf
figure('Position', [1 1 1000 800]);
imagesc(simmat, [0 1]); colormap('jet');
hline_new([8,18,19,29,31,36,37,42,47,52]+0.5, 'k', 2);
hline_new([3,13,24]+0.5, 'k', .5);
vline_new([8,18,19,29,31,36,37,42, 47,52]+0.5, 'k', 2);
vline_new([3,13,24]+0.5, 'k', .5);
set(gca, 'XTick', [4,13.5,19,24.5,30.5,34,37,40,45, 50], 'YTick',
[4,13.5,19,24.5,30.5,34,37,40,45,50], 'XTickLabel',...
{'LS02', 'LS03', 'LS04', 'LS05', 'LS07', 'LS08', 'LS10', 'LS11', 'LS14', 'LS16', 'LS17'},
'YTickLabel', {'LS02', 'LS03', 'LS04', 'LS05', 'LS07', 'LS08', 'LS10', 'LS11', 'LS14',
'LS16', 'LS17'});
axis square;
colorbar;
set(gca, 'XTick', [2,13.5,19,24.5,30.5,34,37,40,45, 50, 55], 'YTick',
[4,13.5,19,24.5,30.5,34,37,40,45,50, 55], 'XTickLabel',...
{'LS02', 'LS03', 'LS04', 'LS05', 'LS07', 'LS08', 'LS10', 'LS11', 'LS14', 'LS16', 'LS17'},
'YTickLabel', {'LS02', 'LS03', 'LS04', 'LS05', 'LS07', 'LS08', 'LS10', 'LS11', 'LS14',
'LS16', 'LS17'});
axis square;
colorbar;
ax = gca;
ax.FontSize = 24;
ax.FontSize = 20;
subs = {'LS02', 'LS03', 'LS05'};
sessions = [8,10,10];
clear corrmatrix
count = 1;
for sub = 1:numel(subs)
for ses = 1:sessions(sub)
data = load(sprintf('%s/sub-%s/sub-%s_sess-%d_task-rest_corrmatrix_Seitzman300.mat',
datadir, subs{sub}, subs{sub}, ses));
corrmatrix(count, :, :, :) = data.corrmatrix;
% sub_corrmatrix(ses, :, :, :) = data.corrmatrix;
count = count + 1;
end
end
clear corrmatrix
maskmat = ones(atlas_params.num_rois);
maskmat = logical(triu(maskmat,1));
count = 1;
for sub = 1:numel(subs)
for ses = 1:sessions(sub)
tmp = corrmatrix(count, :, :, :);
if fisher

```

```
corrln(count,:) = single(FisherTransform(tmp(maskmat)));
else
corrln(count,:) = single(tmp(maskmat));
end
count = count + 1;
end
end
simmat = corr(corrln');
figure('Position',[1 1 1000 800]);
imagesc(simmat,[0 1]); colormap('jet');
hline_new([8,18]+0.5,'k',2);
hline_new([3,13,23]+0.5,'k',.5);
vline_new([8,18]+0.5,'k',2);
vline_new([3,13,23]+0.5,'k',.5);
set(gca,'XTick',[4,13.5,23.5], 'YTick', [4,13.5,23.5], 'XTickLabel',...
{'LS02', 'LS03', 'LS05'}, 'YTickLabel', {'LS02', 'LS03', 'LS05'});
axis square;
colorbar;
ax = gca;
ax.FontSize = 20;
ax.FontSize = 24;
saveas(gcf,[outdir 'SimilarityMat_longitudinalSubs.tiff'],'tiff');
close('all');
ses_pre = [3,5,5];
ses_post = [5,5,5];
count = 1;
clear corrmat
for sub = 1:numel(subs)
for ses = 1:ses_pre(sub)
data = load(sprintf('%s/sub-%s/sub-%s_sess-%d_task-rest_corrmat_Seitzman300.mat',↵
datadir, subs{sub}, subs{sub}, ses));
corrmat(count,:,:) = data.corrmat;
sub_corrmat(ses,:,:) = data.corrmat;
count = count + 1;
end
if ses > 1
mean_pre(sub,:,:) = squeeze(mean(sub_corrmat));
clear sub_corrmat
else
mean_pre(sub,:,:) = squeeze(sub_corrmat);
end
end
count = 1;
for sub = 1:numel(subs)
for ses = ses_pre(sub)+1:ses_pre(sub)+ses_post(sub)
data = load(sprintf('%s/sub-%s/sub-%s_sess-%d_task-rest_corrmat_Seitzman300.mat',↵
datadir, subs{sub}, subs{sub}, ses));
corrmat(count,:,:) = data.corrmat;
sub_corrmat(ses,:,:) = data.corrmat;
count = count + 1;
end
if ses > 1
mean_post(sub,:,:) = squeeze(mean(sub_corrmat));
clear sub_corrmat
else
mean_post(sub,:,:) = squeeze(sub_corrmat);
end
end
maskmat = ones(atlas_params.num_rois);
maskmat = logical(triu(maskmat,1));
```

```
clear corrlin
count = 1;
for sub = 1:numel(subs)
tmp = mean_pre(count,:,:,:);
corrlin_pre(count,:) = single(tmp(maskmat));
tmp = mean_post(count,:,:,:);
corrlin_post(count,:) = single(tmp(maskmat));
count = count + 1;
end
simmat = corr(corrlin_pre', corrlin_post');
figure('Position',[1 1 1000 800]);
imagesc(simmat,[0 1]); colormap('jet');
hline_new([1:2]+0.5,'k',2);
vline_new([1:2]+0.5,'k',2);
set(gca,'XTick',[1:3], 'YTick', [1:3], 'XTickLabel',...
{'LS02', 'LS03', 'LS05'}, 'YTickLabel', {'LS02', 'LS03', 'LS05'});
axis square;
ax = gca;
ax.FontSize = 24;
colorbar;
ax.FontSize = 24;
ax = gca;
ax.FontSize = 24;
xtitle('Timepoint A')
title('Similarity Across Sessions ~1 year apart')
xlabel('Timepoint A')
xlabel('Timepoint B')
ylabel('Timepoint A')
saveas(gcf,[outDir 'SimilarityMat_longitudinal_avg.tiff'],'tiff');
segregation_ind_by_net
help ones
segregation_ind_by_net
help sum
segregation_ind_by_net
weights(net)
seg_index_net(net,ses)
weights(net)
seg_index_net(net,ses)
weights(net)*seg_index_net(net,ses)
segregation_ind_by_net
seg_index_net(15,:) = sum(weighted_seg_ind_net);
segregation_ind_by_net
segregation_ind_by_ses
segregation_ind_by_sub
mean_matrix = mean_matrix(:,300,300);
segregation_ind_by_sub
mean = mean_matrix(:,:,:);
mean = mean_matrix(1,:,:);
mean = squeeze(mean_matrix);
mean_matrix(:,:,:)= squeeze(mean(sub_corrmat));
mean_matrix = squeeze(mean(sub_corrmat));
segregation_ind_by_sub
segregation_ind_by_net
load('/Users/dianaperez/Desktop/Segregation_Analyses/allsubs_seg_index_sub.mat')
clear all
load('/Users/dianaperez/Desktop/Segregation_Analyses/allsubs_seg_index_sub.mat')
segregation_ind_by_sub
load('/Users/dianaperez/Desktop/Segregation_Analyses/allsubs_seg_index_ses.mat')
clear all
load('/Users/dianaperez/Desktop/Segregation_Analyses/allsubs_seg_index_ses.mat')
```

```
segregation_ind_by_sub
segregation_ind_by_ses
load('/Users/dianaperez/Desktop/Segregation_Analyses/LS02_seg_index_net.mat')
clear all
load('/Users/dianaperez/Desktop/Segregation_Analyses/LS02_seg_index_net.mat')
load('/Users/dianaperez/Desktop/Segregation_Analyses/LS03_seg_index_net.mat')
load('/Users/dianaperez/Desktop/Segregation_Analyses/LS05_seg_index_net.mat')
load('/Users/dianaperez/Desktop/Segregation_Analyses/LS08_seg_index_net.mat')
load('/Users/dianaperez/Desktop/Segregation_Analyses/LS11_seg_index_net.mat')
load('/Users/dianaperez/Desktop/Segregation_Analyses/LS14_seg_index_net.mat')
load('/Users/dianaperez/Desktop/Segregation_Analyses/LS16_seg_index_net.mat')
load('/Users/dianaperez/Desktop/Segregation_Analyses/LS17_seg_index_net.mat')
segregation_ind_by_node
load('/Users/dianaperez/Desktop/Segregation_Analyses/allsubs_seg_index_ses.mat')
clear all
load('/Users/dianaperez/Desktop/Segregation_Analyses/allsubs_seg_index_ses.mat')
similarity_analysis
.3855-.3762
help reshape
similarity_analysis
40*40
similarity_analysis
help reshape
similarity_analysis
figure('Position',[1 1 1000 800]);
imagesc(matrix,[0 1]); colormap('jet');
min(matrix)
min(min(matrix))
min(min(matrix))
max(max(matrix))
min(diff_score)
max(diff_score)
close all
figure('Position',[1 1 1000 800]);
imagesc(matrix,[0 .2]); colormap('jet');
colorbar
imagesc(matrix,[0 .01]); colormap('jet');
imagesc(matrix,[0 .1]); colormap('jet');
imagesc(matrix,[-.2 .2]); colormap('jet');
imagesc(matrix,[-.15 .15]); colormap('jet');
imagesc(matrix,[-.14 .14]); colormap('jet');
abs(matrix)
close all
figure('Position',[1 1 1000 800]);
imagesc(matrix,[-.14 .14]); colormap('jet');
matrix = abs(diff_score);
matrix = reshape(matrix, [40,40]);
figure('Position',[1 1 1000 800]);
imagesc(matrix,[0 .14]); colormap('jet');
colorbar
imagesc(matrix,[0 .1]); colormap('jet');
imagesc(matrix,[0 .14]); colormap('jet');
colorbar
hline_new([0,5,10,15,20,25,30,35,40]+0.5,'k',2);
vline_new([0,5,10,15,20,25,30,35,40]+0.5,'k',2);
set(gca,'XTick',[3,8,13,18,23,28,33,38], 'YTick', [3,8,13,18,23,28,33,38], \
'XTickLabel',...
{'LS02', 'LS03', 'LS05', 'LS08', 'LS11', 'LS14', 'LS16', 'LS17'}, 'YTickLabel', {'LS02', \
'LS03', 'LS05', 'LS08', 'LS11', 'LS14', 'LS16', 'LS17'});
axis square;
```

```

FT_matrix = FisherTransform(matrix);
max(FT_matrix)
max(max(FT_matrix))
figure('Position',[1 1 1000 800]);
imagesc(matrix,[0 .14]); colormap('jet');
imagesc(FT_matrix,[0 .14]); colormap('jet');
hline_new([0,5,10,15,20,25,30,35,40]+0.5,'k',2);
vline_new([0,5,10,15,20,25,30,35,40]+0.5,'k',2);
set(gca,'XTick',[3,8,13,18,23,28,33,38], 'YTick', [3,8,13,18,23,28,33,38], 'XTickLabel',...
{'LS02', 'LS03', 'LS05', 'LS08', 'LS11', 'LS14', 'LS16', 'LS17'}, 'YTickLabel', {'LS02', 'LS03', 'LS05', 'LS08', 'LS11', 'LS14', 'LS16', 'LS17'});
axis square;
colorbar;
close all
similarity_analysis
run('/Users/dianaperez/Documents/GitHub/Lifespan-Analysis/Segregation_Index/segregation_ind_by_node.m')
%-- 4/10/22, 4:51 PM --%
CreateVariantFiles_LS03specific
addpath '/Users/dianaperez/Documents/Dependencies/cifti-matlab-master'
CreateVariantFiles_LS03specific
ft_read_cifti_mod('/Volumes/RESEARCH_HD/Lifespan/CNS_analyses/spCor_Maps/sub-LS17_first-half_vs_120_avg_corr_LR_cortex_corr.dtseries.nii')
ft_read_cifti_mod('/Volumes/RESEARCH_HD/Lifespan/CNS_analyses/spCor_Maps/sub-LS03_vs_120_avg_corr_LR_cortex_corr.dtseries.nii')
readtable('/Volumes/fsmresfiles/PBS/Gratton_Lab/Lifespan/Post-COVID/BIDS/derivatives/preproc_fmriprip-20.2.0/fmriprip/sub-LS02/ses-5/func/FD_outputs/sub-LS02_ses-5_task-rest_desc-framenums_fFD.txt')
clear all
readtable('/Volumes/fsmresfiles/PBS/Gratton_Lab/Lifespan/Post-COVID/BIDS/derivatives/preproc_fmriprip-20.2.0/fmriprip/sub-LS02/ses-5/func/FD_outputs/sub-LS02_ses-5_task-rest_desc-framenums_fFD.txt')
mean(ans)
Var1 = ans{:,1};
mean(Var1);
get_FDnums
CreateVariantFiles_LS03specific
similarity_analysis
hline_new([0,5,10,15,20,25,30,35,40]+0.5,'k',2);
vline_new([0,5,10,15,20,25,30,35,40]+0.5,'k',2);
set(gca,'XTick',[3,8,13,18,23,28,33,38], 'YTick', [3,8,13,18,23,28,33,38], 'XTickLabel',...
{'LS02', 'LS03', 'LS05', 'LS08', 'LS11', 'LS14', 'LS16', 'LS17'}, 'YTickLabel', {'LS02', 'LS03', 'LS05', 'LS08', 'LS11', 'LS14', 'LS16', 'LS17'});
axis square;
colorbar;
addpath '/Users/dianaperez/Documents/GitHub/Lifespan-Analysis'
hline_new([0,5,10,15,20,25,30,35,40]+0.5,'k',2);
vline_new([0,5,10,15,20,25,30,35,40]+0.5,'k',2);
set(gca,'XTick',[3,8,13,18,23,28,33,38], 'YTick', [3,8,13,18,23,28,33,38], 'XTickLabel',...
{'LS02', 'LS03', 'LS05', 'LS08', 'LS11', 'LS14', 'LS16', 'LS17'}, 'YTickLabel', {'LS02', 'LS03', 'LS05', 'LS08', 'LS11', 'LS14', 'LS16', 'LS17'});
axis square;
colorbar;
saveas(gcf,[outdir 'SimilarityMat_SegregationIndex.tiff'],'tiff');
outdir = '/Users/dianaperez/Desktop/Segregation_Analyses';
saveas(gcf,[outdir 'SimilarityMat_SegregationIndex.tiff'],'tiff');
CreateVariantFiles_LS03specific

```

```

variant_overlap2
ft_write_cifti([out_dir subs{sub} '_variant_overlap_split-half.dtseries.nii'], template)
variant_overlap2
ft_write_cifti_mod(out_fname, template)
variant_overlap2
mean(corr)
subs = {'LS02', 'LS03', 'LS05', 'LS08', 'LS11', 'LS14', 'LS16', 'LS17'};
data_dir = '/Volumes/RESEARCH_HD/Lifespan/CNS_analyses/spCor_Maps/split-half/';
out_dir = '/Volumes/RESEARCH_HD/Lifespan/CNS_analyses/';
corr = [];
for sub = 1:numel(subs)
map1 = ft_read_cifti_mod([data_dir 'sub-' subs{sub} '_first-✓
half_vs_120_avg_corr_LR_cortex_corr.dtseries.nii']);
map2 = ft_read_cifti_mod([data_dir 'sub-' subs{sub} '_second-✓
half_vs_120_avg_corr_LR_cortex_corr.dtseries.nii']);
corr(sub) = corr(map1.data, map2.data);
end
subs = {'LS02', 'LS03', 'LS05', 'LS08', 'LS11', 'LS14', 'LS16', 'LS17'};
data_dir = '/Volumes/RESEARCH_HD/Lifespan/CNS_analyses/spCor_Maps/split-half/';
out_dir = '/Volumes/RESEARCH_HD/Lifespan/CNS_analyses/';
corr = [];
for sub1 = 1:numel(subs)
for sub2 = 1: numel(subs)
map1 = ft_read_cifti_mod([data_dir 'sub-' subs{sub1} '_first-✓
half_vs_120_avg_corr_LR_cortex_corr.dtseries.nii']);
map2 = ft_read_cifti_mod([data_dir 'sub-' subs{sub2} '_second-✓
half_vs_120_avg_corr_LR_cortex_corr.dtseries.nii']);
corr(sub1,sub2) = corr(map1.data, map2.data);
end
splithalf_spcorrmap_simmat
figure('Position',[1 1 1000 800]);
imagesc(corr,[0 1]); colormap('jet');
%hline_new([0,5,10,15,20,25,30,35,40]+0.5,'k',2);
%vline_new([0,5,10,15,20,25,30,35,40]+0.5,'k',2);
set(gca,'XTick',[1:8], 'YTick', [1:8], 'XTickLabel',...
{'LS02', 'LS03', 'LS05', 'LS08', 'LS11', 'LS14', 'LS16', 'LS17'}, 'YTickLabel', {'LS02',✓
'LS03', 'LS05', 'LS08', 'LS11', 'LS14', 'LS16', 'LS17'});
axis square;
colorbar;
hline_new([1:8]+0.5,'k',2);
vline_new([1:8]+0.5,'k',2);
hline_new([0:8]+0.5,'k',2);
vline_new([0:8]+0.5,'k',2);
ax =gca;
ax.FontSize = 24
saveas(gcf,[out_dir 'SimilarityMat_SpCorrMap.tiff'],'tiff');
similarity_analysis_forced_same_amt_data
for s = 1:numel(subject)
lines = [count:(count+sessions-1)]
sub_vals = simmat(lines,:);
maskmat = zeros(sessions,sessions)'
count = count+sessions;
end
similarity_analysis_forced_same_amt_data
maskmat = logical(triu(maskmat, 1));
maskmat = ones(sessions,sessions);
maskmat = logical(triu(maskmat, 1));
count = 1;
within = [];
between = [];

```

```
for s = 1:numel(subject)
lines = [count:(count+sessions-1)];
sub_vals = simmat(lines,:);
maskmat = ones(sessions,sessions);
maskmat = logical(triu(maskmat, 1));
within_sub = sub_vals(lines,lines);
within = [within; within_sub(maskmat)];
maskmat = ones(size(sub_vals));
maskmat(lines,lines) = 0;
between = [between; sub_vals(maskmat==1)];
count = count+sessions;
end
delete_this
mean(within)
mean(between)
similarity_analysis_wLongitudinal
clear all
segregation_ind_by_sub
segregation_ind_by_node
CreateVariantFiles_LS03specific
overlapmap
unique(group_map)
ft_write_cifti_mod([out_dir '/Lifespan_Variant_Overlap_Map.dtseries.nii'], overlap_map);
segregation_ind_by_ses_longitudinal
%-- 4/12/22, 2:49 PM --%
CreateVariantFiles_LS03specific
addpath '/Users/dianaperez/Documents/Dependencies/cifti-matlab-master'
CreateVariantFiles_LS03specific
rest_file
CreateVariantFiles_LS03specific
rest_file
outfilerest
CreateVariantFiles_LS03specific
variant_overlap2
[data_dir subs{sub} '_allsess_uniqueIDs_variants_sizeExcluded_thresh-5_smooth_2.55.
dtseries.nii']
['/Users/dianaperez/Desktop/' subs{sub}
'_allsess_timeA_uniqueIDs_variants_sizeExcluded_thresh-5_smooth_2.55.dtseries.nii']
out_fname
mean([0.292185810351499,0.567052791491455,0.388237057287416])
clear all
load('/Volumes/RESEARCH_HD/Lifespan/CNS_analyses/allsubs_seg_index_ses_longitudinal.mat')
LS02_pre = ses_SI(1,1:3);
LS03_pre = ses_SI(2,1:5);
LS05_pre = ses_SI(3,1:5);
LS02_post = ses_SI(1,4:8);
LS03_post = ses_SI(2,6:10);
LS05_post = ses_SI(3,6:10);
vals = [LS02_pre LS02_post LS03_pre LS03_post LS05_pre LS05_post];
corr(vals)
for val1 = 1:length(vals)
for val2 = 1:length(vals)
diff = abs(val1-val2);
end
end
for val1 = 1:length(vals)
for val2 = 1:length(vals)
diff(val1,val2) = abs(val1-val2);
end
end
```

```
for val1 = 1:length(vals)
for val2 = 1:length(vals)
diff(val1,val2) = abs(vals(val1)-vals(val2));
end
end
figure('Position',[1 1 1000 800]);
imagesc(diff,[0 .14]); colormap('jet');
max(diff)
max(max(diff))
max(max(diff))
imagesc(diff,[0 .12]); colormap('jet');
colorbar;
axis square;
hline_new([0,8,18,28]+0.5,'k',2);
vline_new([0,8,18,28]+0.5,'k',2);
set(gca,'XTick',[4, 13, 23], 'YTick',[4, 13, 23], 'XTickLabel',...
{'LS02', 'LS03', 'LS05'}, 'YTickLabel', {'LS02', 'LS03', 'LS05'});
ax = gca;
ax.FontSize = 24;
hline_new([3,13,23]+0.5,'k',.5);
vline_new([3,13,23]+0.5,'k',.5);
saveas(gcf,[output_dir 'SimilarityMat_SegregationIndex_Longitudinal.tiff'],'tiff');
output_dir = '/Volumes/RESEARCH_HD/Lifespan/CNS_analyses/';
saveas(gcf,[output_dir 'SimilarityMat_SegregationIndex_Longitudinal.tiff'],'tiff');
help scatter
scatter([1,1,1,1,1;2,2,2,2,2], [.1,.2,.6,.2,.1;.5,.6,.5,.6,.5])
scatter([1,1,1,1,1;2,2,2,2,2], [.1,.2,.6,.2,.1;.5,.6,.5,.6,.5])
scatter([1,1,1,1,1;2,2,2,2,2], [.1,.2,.6,.2,.1;.5,.6,.5,.6,.5])
scatter([1,1,1,1,1], [.1,.2,.6,.2,.1])
scatter([1,1,1,1,1], [.1,.2,.6,.2,.1], 'filled')
clear all
rgb_colors = [1, 0, 0;0, 1, 0;...
0, 0, 1,...
0, 1, 1,...
1, 0, 1,...
0.4660 0.6740 0.188,...
0.9290 0.6940 0.1250,...
0.4940 0.1840 0.5560];
rgb_colors = [1 0 0;
0, 1, 0;
0, 0, 1,
0, 1, 1,
1, 0, 1,
0.4660 0.6740 0.188,
0.9290 0.6940 0.1250,
0.4940 0.1840 0.5560];
load('/Users/dianaperez/Desktop/Segregation_Analyses/allsubs_seg_index_ses.mat')
segregation_ind_by_ses
addpath '/Users/dianaperez/Documents/GitHub/GrattonLab-General-Repo/FCProcess'
segregation_ind_by_ses
within_sub_reliability
test = randperm(catData);
test = randperm(1,catData);
help randperm
test = randperm(1:size(catData));
test = randperm(1:size(catData,1));
size(catData,1)
size(catData,2)
test = 1:size(catData,2);
test = randperm(size(catData,2));
```



```

within_sub_reliability
sampledData{s} = datasample(rest_of_data,sampStep*t,2);
within_sub_reliability
size(allsubs_seg_ind{1})
size(allsubs_seg_ind{1},2)
allsubs_seg_ind{1,1}
size(allsubs_seg_ind{1},2)
times = [2.5:2.5:100];
figure;
for s = 1:numel(subs)
plot(times(1:size(allsubs_seg_ind{1},2)),allsubs_seg_ind{1,1},'Color',rgb_color\
(s,:), 'LineWidth', 3)
hold on
end
times = [2.5:2.5:100];
figure;
for s = 1:numel(subject)
plot(times(1:size(allsubs_seg_ind{1},2)),allsubs_seg_ind{1,1},'Color',rgb_color\
(s,:), 'LineWidth', 3)
hold on
end
within_sub_reliability
plot(times(1:size(allsubs_seg_ind{1},2)),allsubs_seg_ind{1,1},'Color',rgb_color\
(s,:), 'LineWidth', 3)
times(1:size(allsubs_seg_ind{1},2))
size(allsubs_seg_ind{1},2)
times(s, 1:size(allsubs_seg_ind{1},2))
times_all(s, 1:size(allsubs_seg_ind{1},2))
plot(times_all(s,1:size(allsubs_seg_ind{1},2)),allsubs_seg_ind{1,1},'Color',rgb_color\
(s,:), 'LineWidth', 3)
plot(times_all(s,1:size(allsubs_seg_ind{1},2)),allsubs_seg_ind{1,1},'Color',rgb_colors\
(s,:), 'LineWidth', 3)
figure;
for s = 1:numel(subject)
plot(times_all(s,1:size(allsubs_seg_ind{1},2)),allsubs_seg_ind{1,1},'Color',rgb_colors\
(s,:), 'LineWidth', 3)
hold on
end
within_sub_reliability
xlabel('Time (Minutes)');
ylabel('Segregation Index');
xlabel('Time (Minutes)');
ylabel('Segregation Index');
xlabel('Time (Minutes)');
m = findobj(gca,'Type','line');
hleg1 = legend(m(1:8), 'LS17', 'LS16', 'LS14', 'LS11', 'LS08', 'LS05', 'LS03', 'LS02',\
'Location', 'SouthEast');
hleg1.FontSize = 20;
ax = gca;
ax.FontSize = 24;
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.5, 0.7]);
%
figure;
for s = 1:numel(subject)
plot(times_all(s,1:size(allsubs_seg_ind{s},2)),smooth(allsubs_seg_ind{1,s}),'Color',\
rgb_colors(s,:), 'LineWidth', 3)
hold on
end
%-- 4/12/22, 9:31 PM --%
within_sub_reliability

```

```

addpath '/Users/dianaperez/Documents/GitHub/GrattonLab-General-Repo/FCProcess'
within_sub_reliability
addpath '/Users/dianaperez/Documents/GitHub/Lifespan-Analysis'
within_sub_reliability
%-- 4/12/22, 10:24 PM --%
within_sub_reliability
addpath '/Users/dianaperez/Documents/GitHub/GrattonLab-General-Repo/FCProcess'
within_sub_reliability
help smooth
within_sub_reliability
help axis
axis([0 100 0 .15])
print(gcf,['/Volumes/RESEARCH_HD/Lifespan/CNS_Analyses/ReliabilityLifespanSegIndex.
jpg'],'-dpng','-r300');
similarity_analysis_forced_same_amt_data
ax = gca;
ax.FontSize = 24;
saveas(gcf,[outdir 'SimilarityMat_rest.tiff'],'tiff');
close('all');
count = 1;
within = [];
between = [];
for s = 1:numel(subject)
lines = [count:(count+sessions-1)];
sub_vals = simmat(lines,:);
maskmat = ones(sessions,sessions);
maskmat = logical(triu(maskmat, 1));
within_sub = sub_vals(:,lines);
within = [within; within_sub(maskmat)];
maskmat = ones(size(sub_vals));
maskmat(:,lines) = 0;
between = [between; sub_vals(maskmat==1)];
count = count+sessions;
end
mean(between)
mean(within)
similarity_analysis_forced_same_amt_data
rgb_colors = [1 0 0;%LS02
0, 1, 0;%LS03
0, 0, 1;%LS05
0, 1, 1;%LS08
1, 0, 1;%LS11
0.4660 0.6740 0.188;%LS14
0.9290 0.6940 0.1250;%LS16
0.4940 0.1840 0.5560;%LS17
0 0 0;
0 0 0;
0 0 0;
0 0 0;
0 0 0;
0 0 0;
0 0 0;
0 0 0;
0 0 0;
0 0 0;
0 0 0;
0 0 0];
subs = {'LS02', 'LS03', 'LS05', 'LS08', 'LS11', 'LS14',
'LS16', 'LS17', 'INET001', 'INET002',
'INET003', 'INET005', 'INET006', 'INET010', 'INET016', 'INET018', 'INET019', 'INET030'};
lifespan_SI = load('/Users/dianaperez/Desktop/Segregation_Analyses/allsubs_seg_index_ses.
mat');

```

```
iNet_SI = load(
('/Users/dianaperez/Desktop/Segregation_Analyses/INET_allsubs_seg_index_ses.mat');
handles = plotSpread([Lifespan_SI iNet_SI], 'distributionColors', rgb_for_plot', ↵
'xNames', subs)
handles = plotSpread([lifespan_SI iNet_SI], 'distributionColors', rgb_for_plot', ↵
'xNames', subs)
handles = plotSpread([lifespan_SI iNet_SI], 'distributionColors', rgb_colors', 'xNames', ↵
subs)
for n = 1:numel(network_names)
rgb_for_plot{n} = rgb_colors(n,:);
end
rgb_colors = [1 0 0;%LS02
0, 1, 0;%LS03
0, 0, 1;%LS05
0, 1, 1;%LS08
1, 0, 1;%LS11
0.4660 0.6740 0.188;%LS14
0.9290 0.6940 0.1250;%LS16
0.4940 0.1840 0.5560;%LS17
0 0 0;
0 0 0;
0 0 0;
0 0 0;
0 0 0;
0 0 0;
0 0 0;
0 0 0;
0 0 0;
0 0 0];
subs = {'LS02', 'LS03', 'LS05', 'LS08', 'LS11', 'LS14', ↵
'LS16', 'LS17', 'INET001', 'INET002', ↵
'INET003', 'INET005', 'INET006', 'INET010', 'INET016', 'INET018', 'INET019', 'INET030'};
for n = 1:numel(subs)
rgb_for_plot{n} = rgb_colors(n,:);
end
handles = plotSpread([lifespan_SI iNet_SI], 'distributionColors', rgb_for_plot', ↵
'xNames', subs)
rgb_colors = [1 0 0;%LS02
0, 1, 0;%LS03
0, 0, 1;%LS05
0, 1, 1;%LS08
1, 0, 1;%LS11
0.4660 0.6740 0.188;%LS14
0.9290 0.6940 0.1250;%LS16
0.4940 0.1840 0.5560;%LS17
0 0 0;
0 0 0;
0 0 0;
0 0 0;
0 0 0;
0 0 0;
0 0 0;
0 0 0;
0 0 0;
0 0 0];
subs = {'LS02', 'LS03', 'LS05', 'LS08', 'LS11', 'LS14', ↵
'LS16', 'LS17', 'INET001', 'INET002', ↵
'INET003', 'INET005', 'INET006', 'INET010', 'INET016', 'INET018', 'INET019', 'INET030'};
for n = 1:numel(subs)
rgb_for_plot{n} = rgb_colors(n,:);
```

```

end
load('/Users/dianaperez/Desktop/Segregation_Analyses/allsubs_seg_index_ses.mat');
lifespan_SI = ses_SI;
load('/Users/dianaperez/Desktop/Segregation_Analyses/INET_allsubs_seg_index_ses.mat');
iNet_SI = ses_SI;
handles = plotSpread([lifespan_SI iNet_SI], 'distributionColors', rgb_for_plot', ↵
'xNames', subs)
handles = plotSpread([lifespan_SI' iNet_SI'], 'distributionColors', rgb_for_plot', ↵
'xNames', subs)
[lifespan_SI' iNet_SI']
make_figures
handles = plotSpread([lifespan_SI'], 'distributionColors', rgb_for_plot', 'xNames', subs)
make_figures
clear rgb_for_plot
rgb_colors = [1 0 0;%LS02
0, 1, 0;%LS03
0, 0, 1;%LS05
0, 1, 1;%LS08
1, 0, 1;%LS11
0.4660 0.6740 0.188;%LS14
0.9290 0.6940 0.1250;%LS16
0.4940 0.1840 0.5560];%LS17
%      0 0 0;
%      0 0 0;
%      0 0 0;
%      0 0 0;
%      0 0 0;
%      0 0 0;
%      0 0 0;
%      0 0 0;
%      0 0 0;
%      0 0 0;
subs = {'LS02', 'LS03', 'LS05', 'LS08', 'LS11', 'LS14', ↵
'LS16', 'LS17'};%,'INET001','INET002', ↵
'INET003','INET005','INET006','INET010','INET016','INET018','INET019','INET030'};
for n = 1:numel(subs)
    rgb_for_plot{n} = rgb_colors(n,:);
end
make_figures
ax = gca;
ax.FontSize = 24;
handles = plotSpread([lifespan_SI'], 'categoryMarkers', '.', 'distributionColors', ↵
rgb_for_plot', 'xNames', subs)
help plot
handles = plotSpread([lifespan_SI'], 'categoryMarkers', '.', 'distributionColors', ↵
rgb_for_plot', 'xNames', subs)
iNet_SI = ses_SI;
iNet_SI(5,:) = -1;
iNet_SI = ses_SI';
iNet_SI(5,:) = -1;
SIs = [lifespan_SI' iNetSI];
SIs = [lifespan_SI' iNet_SI];
make_figures
axis([0 18 .25 .6])
axis([0 19 .25 .6])
make_figures
ax = gca;
ax.FontSize = 20;
make_figures
axis([0 19 .25 .6])

```

```
ax = gca;
ax.FontSize = 20;
make_figures
help plot
handles = plotSpread(SIs, 'distributionMarkers', 'o', 'distributionColors', ↵
rgb_for_plot', 'xNames', subs)
make_figures
plotSpread
make_figures
rgb_colors = [1 0 0;%LS02
0, 1, 0;%LS03
0, 0, 1;%LS05
0, 1, 1;%LS08
1, 0, 1;%LS11
0.4660 0.6740 0.188;%LS14
0.9290 0.6940 0.1250;%LS16
0.4940 0.1840 0.5560;%LS17
0 0 0;
0 0 0;
0 0 0;
0 0 0;
0 0 0;
0 0 0;
0 0 0;
0 0 0;
0 0 0;
0 0 0];
make_figures
load('/Volumes/RESEARCH_HD/Lifespan/CNS_analyses/allsubs_seg_index_ses_longitudinal.mat')
make_figures
axis([0 19 .25 .6])
handles.MarkerFaceColor
handles
make_figures
axis([0 19 .25 .6])
ax = gca;
ax.FontSize = 20;
load('/Volumes/RESEARCH_HD/Lifespan/CNS_analyses/allsubs_seg_index_ses_longitudinal.mat')
longitudinal_SI = ses_SI;
rgb_colors = [1 0 0;%LS02
0, 1, 0;%LS03
0, 0, 1;%LS05
for n = 1:3
rgb_for_plot{n} = rgb_colors(n,:);
end
handles = plotSpread([lifespan_SI' iNet_SI], 'distributionMarkers', {'x'}, ↵
'distributionColors', rgb_for_plot')
handles = plotSpread(longitudinal_SI', 'distributionMarkers', {'x'}, ↵
'distributionColors', rgb_for_plot')
clear rgb_for_plot
rgb_colors = [1 0 0;%LS02
0, 1, 0;%LS03
0, 0, 1;%LS05
for n = 1:3
rgb_for_plot{n} = rgb_colors(n,:);
end
handles = plotSpread(longitudinal_SI', 'distributionMarkers', {'x'}, ↵
'distributionColors', rgb_for_plot')
close all
make_figures
```

```

clear rgb_for_plot
make_figures
longitudinal_SI(:,6:10) = [];
longitudinal_SI(1,4:5) = -1;
make_figures
plotSpread
make_figures
%-- 4/13/22, 9:50 AM --%
make_figures
axis([0 19 .3 .55])
axis([0 19 .31 .54])
axis([0 19 .32 .53])
axis([0 19 .3 .55])
similarity_analysis
addpath '/Users/dianaperez/Documents/GitHub/Lifespan-Analysis'
similarity_analysis
hline_new([0,4,8,12,16,20,24,28,32]+0.5, 'k',2);
vline_new([0,4,8,12,16,20,24,28,32]+0.5, 'k',2);
vline_new([0,4,8,12,16,20,24,28,32,36,40]+0.5, 'k',2);
hline_new([0,4,8,12,16,20,24,28,32,36,40]+0.5, 'k',2);
axis square;
colorbar;
SIs = [0.40465483 0.47502443 0.42045888 0.35866937 0.35903123 0.39794636 0.39622128↵
0.37635228 0.48386648 0.45387810 0.50429422 0.49092054 0.35866398 0.49722296 0.45307758↵
0.44976237 0.40819031 0.39566854];
make_figures
lifespan_SI(:,1) = [];
iNet_SI(1,:) = [];
pre_first_ses = [0.40465483 0.47502443 0.42045888];
post_first_ses = [0.35866937 0.35903123 0.39794636 0.39622128 0.37635228 0.48386648↵
0.45387810 0.50429422 0.49092054 0.35866398 0.49722296 0.45307758 0.44976237 0.40819031↵
0.39566854];
handles = plotSpread([pre_first_ses post_first_ses], 'distributionMarkers', {'o', '*'},↵
'distributionColors', rgb_for_plot, 'xNames', subs)
handles = plotSpread([pre_first_ses post_first_ses], 'distributionMarkers',↵
{'o','o','o','*','*','*','*','*','*','*','*','*','*','*','*','*'},↵
'distributionColors', rgb_for_plot, 'xNames', subs)
close all
make_figures
ph
ph.MarkerFaceColor
help plot
ah
ah.MarkerFaceColor
ah.MarkerFaceColor = plotColors{iData,iCategory};
fh.MarkerFaceColor = plotColors{iData,iCategory};
parserObj.MarkerFaceColor = plotColors{iData,iCategory};
close all
p = plot([1 2 3 4 5 6],[0 3 1 6 4 10],'-o','LineWidth',3);
clear all
p = plot([1 2 3 4 5 6],[0 3 1 6 4 10],'-o','LineWidth',3);
clear all
make_figures
'MarkerFaceColor', plotColors{iData,iCategory},...
make_figures
handles = plotSpread([lifespan_SI iNet_SI], 'distributionMarkers', {'o'},↵
'distributionColors', rgb_for_plot, 'xNames', subs)
close all
make_figures
close all

```

```

make_figures
handles = plotSpread([lifespan_SI' iNet_SI], 'distributionMarkers', {'o'},↵
'distributionColors', rgb_for_plot', 'xNames', subs)
make_figures
handles = plotSpread([lifespan_SI' iNet_SI], 'distributionMarkers', {'o'},↵
'distributionColors', rgb_for_plot', 'xNames', subs, 'binWidth', .5)
make_figures
handles = plotSpread([lifespan_SI' iNet_SI], 'distributionMarkers', {'o'},↵
'distributionColors', rgb_for_plot', 'xNames', subs, 'binWidth', 1)
rgb_colors = [0 0 0;
0 0 0;
0 0 0;
0 0 0;
0 0 0;
0 0 0;
0 0 0;
0 0 0;
1 1 1;
1 1 1;
1 1 1;
1 1 1;
1 1 1;
1 1 1;
1 1 1;
1 1 1;
1 1 1;
1 1 1;
1 1 1];
for n = 1:numel(subs)
rgb_for_plot{n} = rgb_colors(n,:);
end
handles = plotSpread([lifespan_SI' iNet_SI], 'distributionMarkers', {'o'},↵
'distributionColors', rgb_for_plot', 'xNames', subs, 'binWidth', 1)
pre_first_ses = [0.40465483 0.47502443 0.42045888];
post_first_ses = [0.35866937 0.35903123 0.39794636 0.39622128 0.37635228 0.48386648↵
0.45387810 0.50429422 0.49092054 0.35866398 0.49722296 0.45307758 0.44976237 0.40819031↵
0.39566854];
handles = plotSpread([pre_first_ses post_first_ses], 'distributionMarkers', {'^'},↵
'distributionColors', rgb_for_plot', 'xNames', subs)
rgb_colors = [1 0 0;%LS02
0, 1, 0;%LS03
0, 0, 1;%LS05
0, 1, 1;%LS08
1, 0, 1;%LS11
0.4660 0.6740 0.188;%LS14
0.9290 0.6940 0.1250;%LS16
0.4940 0.1840 0.5560;%LS17
0 0 0;
0 0 0;
0 0 0;
0 0 0;
0 0 0;
0 0 0;
0 0 0;
0 0 0;
0 0 0;
0 0 0];
subs = {'LS02', 'LS03', 'LS05', 'LS08', 'LS11', 'LS14',↵
'LS16', 'LS17', 'INET001', 'INET002',↵
'INET003', 'INET005', 'INET006', 'INET010', 'INET016', 'INET018', 'INET019', 'INET030'};
for n = 1:numel(subs)

```

```

rgb_for_plot{n} = rgb_colors(n,:);
end
handles = plotSpread([pre_first_ses post_first_ses], 'distributionMarkers', {'^'}, ↵
'distributionColors', rgb_for_plot', 'xNames', subs)
pre_first_ses = [-1 -1 -1];
handles = plotSpread([pre_first_ses post_first_ses], 'distributionMarkers', {'^'}, ↵
'distributionColors', rgb_for_plot', 'xNames', subs)
handles = plotSpread([pre_first_ses post_first_ses], 'distributionMarkers', {'d'}, ↵
'distributionColors', rgb_for_plot', 'xNames', subs)
close all
make_figures
handles = plotSpread([lifespan_SI' iNet_SI], 'distributionMarkers', {'o'}, ↵
'distributionColors', rgb_for_plot', 'xNames', subs, 'binWidth', 1)
close all
make_figures
handles = plotSpread([lifespan_SI' iNet_SI], 'distributionMarkers', {'o'}, ↵
'distributionColors', rgb_for_plot', 'xNames', subs, 'binWidth', 1)
make_figures
handles = plotSpread([lifespan_SI' iNet_SI], 'distributionMarkers', {'o'}, ↵
'distributionColors', rgb_for_plot', 'xNames', subs, 'binWidth', 1)
rgb_colors = [0 0 0;
0 0 0;
1 1 1;
0 0 0;
0 0 0;
0 0 0;
0 0 0;
0 0 0;
0 0 0;
1 1 1;
1 1 1;
1 1 1;
1 1 1;
1 1 1;
1 1 1;
1 1 1;
1 1 1;
1 1 1;
1 1 1;
1 1 1;
1 1 1;
1 1 1;
1 1 1;
1 1 1;
for n = 1:numel(subs)
rgb_for_plot{n} = rgb_colors(n,:);
end
handles = plotSpread([lifespan_SI' iNet_SI], 'distributionMarkers', {'o'}, ↵
'distributionColors', rgb_for_plot', 'xNames', subs, 'binWidth', 1)
within_sub_reliability
addpath '/Users/dianaperez/Documents/GitHub/GrattonLab-General-Repo/FCProcess'
within_sub_reliability
mean(allsubs_seg_ind{1,1},2);
mean(allsubs_seg_ind{1,1},1);
plot(times_all(s,1:size(allsubs_seg_ind{s},2)),mean(allsubs_seg_ind{1,s},1),'Color',↵
rgb_colors(s,:), 'LineWidth', 3)
hold on
plot(times_all(s,1:size(allsubs_seg_ind{s},2)),mean(allsubs_seg_ind{1,s},1),'Color',↵
rgb_colors(s,:), 'LineWidth', 3)
within_sub_reliability
clear all
within_sub_reliability
mean_seg_inds{s} = mean(sub_seg_index,1);
mean_abs_diff{s} = mean(abs_diffs,1);
within_sub_reliability
times=[2.5:2.5:100];

```



```

figure;
for s = 1:numel(subject)
plot(times_all(s,1:size(mean_seg_inds{s},2)),mean_abs_diffs(s,1:size(allsubs_seg_ind{s},2)), 'Color',rgb_colors(s,:), 'LineWidth', 3)
hold on
end
for s = 1:numel(subject)
plot(times_all(s,1:size(mean_seg_inds{s},2)),mean_abs_diff(s,1:size(allsubs_seg_ind{s},2)), 'Color',rgb_colors(s,:), 'LineWidth', 3)
hold on
end
mean_abs_diff(s,1:size(allsubs_seg_ind{s},2))
allsubs_seg_ind{s}
size(mean_abs_diff{s},2)
mean_abs_diff(s,1:size(mean_abs_diff{s},2))
mean_abs_diff{1,s}
for s = 1:numel(subject)
plot(times_all(s,1:size(mean_seg_inds{s},2)),mean_abs_diff{1,s}, 'Color',rgb_colors(s,:), 'LineWidth', 3)
hold on
end
size(mean_seg_inds{s},2)
times_all(s,1:size(mean_seg_inds{s},2))
mean_abs_diff{1,s}
within_sub_reliability
mean_seg_inds{s} = mean(sub_seg_index,1);
within_sub_reliability
mean(allsubs_abs_diffs{1,s},2)
mean(allsubs_abs_diffs{1,s},1)
times = [2.5:2.5:100];
figure;
for s = 1:numel(subject)
plot(times_all(s,1:size(allsubs_seg_inds{s},2)),mean(allsubs_abs_diffs{1,s},1), 'Color',rgb_colors(s,:), 'LineWidth', 3)
hold on
end
times = [2.5:2.5:100];
figure;
for s = 1:numel(subject)
plot(times_all(s,1:size(allsubs_seg_ind{s},2)),mean(allsubs_abs_diffs{1,s},1), 'Color',rgb_colors(s,:), 'LineWidth', 3)
hold on
end
mean(allsubs_abs_diffs{1,s}(:,1:size(allsubs_seg_ind{s},2))
mean(allsubs_abs_diffs{1,s}(:,1:size(allsubs_seg_ind{s},2)))
times = [2.5:2.5:100];
figure;
for s = 1:numel(subject)
plot(times_all(s,1:size(allsubs_seg_ind{s},2)),mean(allsubs_abs_diffs{1,s}(:,1:size(allsubs_seg_ind{s},2))),1), 'Color',rgb_colors(s,:), 'LineWidth', 3)
hold on
end
times = [2.5:2.5:100];
figure;
for s = 1:numel(subject)
plot(times_all(s,1:size(allsubs_seg_ind{s},2)),mean(allsubs_abs_diffs{1,s}(:,1:size(allsubs_seg_ind{s},2))),1), 'Color',rgb_colors(s,:), 'LineWidth', 3)
hold on
end
times_all(s,1:size(allsubs_seg_ind{s},2))

```

```

mean(allsubs_abs_diffs{1,s}(:,1:size(allsubs_seg_ind{s}),1),1)
mean(allsubs_abs_diffs{1,s}(:,1:size(allsubs_seg_ind{s}),2),2)
mean(allsubs_abs_diffs{1,s}(:,1:size(allsubs_seg_ind{s}),1),1)
allsubs_abs_diffs{1,s}(:,1:size(allsubs_seg_ind{s},2))
mean(allsubs_abs_diffs{1,s}(:,1:size(allsubs_seg_ind{s},2)),1)
times = [2.5:2.5:100];
figure;
for s = 1:numel(subject)
plot(times_all(s,1:size(allsubs_seg_ind{s},2)),mean(allsubs_abs_diffs{1,s}(:,1:size(
allsubs_seg_ind{s},2)),1),'Color',rgb_colors(s,:), 'LineWidth', 3)
hold on
end
figure;
for s = 1:numel(subject)
plot(times_all(s,1:size(allsubs_seg_ind{s},2)),mean(allsubs_seg_ind{1,s}(:,1:size(
allsubs_seg_ind{s},2)),1),'Color',rgb_colors(s,:), 'LineWidth', 3)
hold on
end
within_sub_reliability
ylabel('% Difference');
xlabel('Time (Minutes)');
axis([0 100 0 .3])
ylabel('% Difference');
xlabel('Time (Minutes)');
m = findobj(gca,'Type','line');
hleg1 = legend(m(1:8), 'LS17', 'LS16', 'LS14', 'LS11', 'LS08', 'LS05', 'LS03', 'LS02',
'Location', 'NorthEast');
hleg1.FontSize = 20;
ax = gca;
ax.FontSize = 24;
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.5, 0.7]);
%
times = [2.5:2.5:100];
figure;
for s = 1:numel(subject)
plot(times_all(s,1:size(allsubs_seg_ind{s},2)),mean(allsubs_abs_diffs{1,s}(:,1:size(
allsubs_seg_ind{s},2)),1),'Color',rgb_colors(s,:), 'LineWidth', 3)
hold on
end
ylabel('% Difference');
xlabel('Time (Minutes)');
m = findobj(gca,'Type','line');
hleg1 = legend(m(1:8), 'LS17', 'LS16', 'LS14', 'LS11', 'LS08', 'LS05', 'LS03', 'LS02',
'Location', 'NorthEast');
hleg1.FontSize = 20;
hleg1.FontSize = 24;
hleg1.FontSize = 20;
ax = gca;
ax.FontSize = 24;
ax.FontSize = 15;
ax.FontSize = 10;
xticks([0:10:100])
print(gcf,['/Volumes/RESEARCH_HD/Lifespan/CNS_Analyses/ReliabilityLifespanSegIndex_Diff.
jpg'],'-dpng','-r300');
figure;
for s = 1:numel(subject)
plot(times_all(s,1:size(allsubs_seg_ind{s},2)),mean(allsubs_seg_ind{1,s}(:,1:size(
allsubs_seg_ind{s},2)),1),'Color',rgb_colors(s,:), 'LineWidth', 3)
hold on
end

```

```

ylabel('Expectation Value');
xlabel('Time (Minutes)');
m = findobj(gca,'Type','line');
ax = gca;
ax.FontSize = 24;
xticks([0:10:100])
axis([0 100 .3 .55])
print(gcf,['/Volumes/RESEARCH_HD/Lifespan/CNS_Analyses/ReliabilityLifespanSegIndex.
jpg'],'-dpng','-r300');
similarity_analysis_wLongitudinal
hline_new([0,8,18]+0.5,'k',2);
hline_new([0,3,13,23]+0.5,'k',.5);
vline_new([0,8,18]+0.5,'k',2);
vline_new([0,3,13,23]+0.5,'k',.5);
hline_new([0,8,18,28]+0.5,'k',2);
hline_new([0,3,13,23]+0.5,'k',.5);
vline_new([0,8,18,28]+0.5,'k',2);
vline_new([0,3,13,23]+0.5,'k',.5);
set(gca,'XTick',[4,13.5,23.5], 'YTick', [4,13.5,23.5], 'XTickLabel',...
{'LS02', 'LS03', 'LS05'}, 'YTickLabel', {'LS02', 'LS03', 'LS05'});
axis square;
colorbar;
ax = gca;
ax.FontSize = 24;
xticklabel([])
xticklabels([])
yticklabels([])
ses_pre = [3,5,5];
ses_post = [5,5,5];
count = 1;
within = [];
between = [];
s = 1;
ses_pre = [3,5,5];
ses_post = [5,5,5];
count = 1;
within = [];
between = [];
lines = [count:(ses_pre(s)+ses_post(s))-1];
lines = [count:(ses_pre(s)+ses_post(s))];
sub_vals = simmat(lines,:);
maskmat = ones(sessions,sessions);
maskmat = logical(triu(maskmat, 1));
sub_vals = simmat(lines,:);
maskmat = ones(length(lines),length(lines));
sub_vals = simmat(lines,:);
within_sub = sub_vals(:,lines);
within = [within; within_sub(maskmat)];
maskmat = ones(size(sub_vals));
maskmat(:,lines) = 0;
between = [between; sub_vals(maskmat==1)];
count = lines(end)+1;
s = 2;
lines = [count:(ses_pre(s)+ses_post(s))];
(ses_pre(s)+ses_post(s))
lines = [count:count+(ses_pre(s)+ses_post(s))];
lines = [count:count+(ses_pre(s)+ses_post(s))-1];
sub_vals = simmat(lines,:);
maskmat = ones(sessions,sessions);
maskmat = logical(triu(maskmat, 1));

```

```
within_sub = sub_vals(:,lines);
within = [within; within_sub(maskmat)];
maskmat = ones(size(sub_vals));
maskmat(:,lines) = 0;
between = [between; sub_vals(maskmat==1)];
count = lines(end)+1;
sub_vals = simmat(lines,:);
maskmat = ones(length(lines),length(lines));
maskmat = logical(triu(maskmat, 1));
within_sub = sub_vals(:,lines);
within = [within; within_sub(maskmat)];
maskmat = ones(size(sub_vals));
maskmat(:,lines) = 0;
between = [between; sub_vals(maskmat==1)];
count = lines(end)+1;
s = 1;
count = 1;
within = [];
between = [];
lines = [count:count+(ses_pre(s)+ses_post(s))-1];
sub_vals = simmat(lines,:);
maskmat = ones(length(lines),length(lines));
maskmat = logical(triu(maskmat, 1));
within_sub = sub_vals(:,lines);
within_sub(maskmat)
within = [within; within_sub(maskmat)];
maskmat = ones(size(sub_vals));
maskmat(:,lines) = 0;
between = [between; sub_vals(maskmat==1)];
count = lines(end)+1;
s = 2;
lines = [count:count+(ses_pre(s)+ses_post(s))-1];
sub_vals = simmat(lines,:);
maskmat = ones(length(lines),length(lines));
maskmat = logical(triu(maskmat, 1));
within_sub = sub_vals(:,lines);
within = [within; within_sub(maskmat)];
maskmat = ones(size(sub_vals));
maskmat(:,lines) = 0;
between = [between; sub_vals(maskmat==1)];
count = lines(end)+1;
s = 3;
lines = [count:count+(ses_pre(s)+ses_post(s))-1];
sub_vals = simmat(lines,:);
maskmat = ones(length(lines),length(lines));
maskmat = logical(triu(maskmat, 1));
within_sub = sub_vals(:,lines);
within = [within; within_sub(maskmat)];
maskmat = ones(size(sub_vals));
maskmat(:,lines) = 0;
between = [between; sub_vals(maskmat==1)];
count = lines(end)+1;
mean(between)
mean(within)
reliability_analysis
clear all
reliability_analysis
%-- 4/14/22, 5:29 PM --%
3818/136
3818/270
```

```

reliability_analysis_v2
70/2.5
reliability_analysis_v2
count+sampStep
reliability_analysis_v2
indices_for_data(set,:) = count:sampStep;
count:sampStep;
count
count+sampStep
reliability_analysis_v2
start_at_pt = datasample(num_sets,1);
count = start_at_pt
num_sets
datasample(num_sets,1)
rng('shuffle')
datasample(num_sets,1)
datasample(1:num_sets,1)
count = datasample(1:num_sets,1);
true_half_inds = [];
while length(true_half_inds) < pts2sample
if count <= num_sets
true_half_inds = [true_half_inds; indices_for_sets(count,:)]';
count = count + 1;
end
end
while length(true_half_inds) < pts2sample
if count <= num_sets
true_half_inds = [true_half_inds; indices_for_sets(count,:)]';
count = count + 1;
end
end
while length(true_half_inds) < pts2sample
if count <= num_sets
true_half_inds = [true_half_inds; indices_for_data(count,:)]';
count = count + 1;
end
end
3944*1.1
/60
(3944*1.1)/60
3818/136
136*28
(3808*1.1)/60
reliability_analysis_v2
indices_for_data(end-num_true_half_sets:end);
indices_for_data(end-num_true_half_sets-1:end);
indices_for_data(end-(num_true_half_sets-1):end);
indices_for_rest_of_data = indices_for_data;
indices_for_rest_of_data(end-(num_true_half_sets-1):end) = [];
indices_for_rest_of_data = indices_for_data;
indices_for_rest_of_data((end-(num_true_half_sets-1)):end,:) = [];
reliability_analysis_v2
size(indices_for_rest_of_data,1)
times = [2.5:2.5:((size(indices_for_rest_of_data,1))*2.5)];
datasample(1:size(indices_for_rest_of_data,1),1);
reliability_analysis_v2
corr_means = mean(allsubs_corrs);
reliability_analysis_v2
mean = [0.467687886 0.559761435 0.632339679 0.66006551 0.690802314 0.718423844
0.734759345 0.732076428 0.752826409 0.764049345 0.772971331 0.781086781 0.792362846

```

```
0.790422743 0.798111731 0.809298493 0.802306023 0.81222891 0.81431187 0.816426924✓
0.822798413 0.828645946 0.830524224 0.83257417 0.83115552 0.837167681 0.840971793✓
0.842499929 0.839556975 0.841703753 0.84395807 0.846296928 0.850515097 0.850531645✓
0.8533126 0.850525042 0.853731175 0.85390665 0.854444922 0.85268042 0.857144836✓
0.858457492 0.858360196 0.859544358 0.861655104 0.861233008 0.862832248 0.863479792✓
0.862828184 0.857708573 0.87665832 0.877845915 0.87690863 0.878025055];
s = 3;
means{1,s}
figure;
for s = 1:numel(subject)
plot(times_all(s,1:size(allsubs_seg_ind{s},2)),means{1,s},'Color',rgb_colors✓
(s,:),'LineWidth', 3)
hold on
end
for s = 1:numel(subject)
plot(times_all(s,1:size(means{1,s},2)),means{1,s},'Color',rgb_colors(s,:),'LineWidth', 3)
hold on
end
plot(times(1:20),mean(1:20), ':', 'Color', [0,0,0], 'LineWidth',3) %average
plot(times_all(1,1:20),mean(1:20), ':', 'Color', [0,0,0], 'LineWidth',3) %average
plot(times_all(1,1:40),mean(1:40), ':', 'Color', [0,0,0], 'LineWidth',3) %average
plot(times_all(1,1:4),mean(1:48), ':', 'Color', [0,0,0], 'LineWidth',3) %average
plot(times_all(1,1:48),mean(1:48), ':', 'Color', [0,0,0], 'LineWidth',3) %average
reliability_analysis_v2
mean = [0.466963624 0.568742573 0.62424519 0.662057459 0.689187726 0.709678761✓
0.726053023 0.74064818 0.751189469 0.762092324 0.771169663 0.778562048 0.785648176✓
0.791579589 0.797392614 0.802157453 0.806759886 0.810889354 0.814636111 0.818249976✓
0.821505513 0.824875033 0.828085976 0.83154268 0.83431512 0.836681223 0.838774137✓
0.841217936 0.840038058 0.841879417 0.84428506 0.846331487 0.847734422 0.849486222✓
0.851167548 0.852191338 0.853726477 0.854901912 0.85528071 0.856386732 0.857766222✓
0.858754624 0.859717988 0.86083224 0.861552202 0.862601614 0.863284792 0.86391976✓
0.864501404 0.8579562 0.8746753 0.875439495 0.87587905 0.87634486];
plot(times_all(1,1:4),mean(1:48), ':', 'Color', [0,0,0], 'LineWidth',3) %average
plot(times_all(1,1:48),mean(1:48), ':', 'Color', [0,0,0], 'LineWidth',3) %average
ylabel('Pearson Correlation (r)');
xlabel('Time (Minutes)');
m = findobj(gca,'Type','line');
hleg1 = legend(m(1:9), 'Mean', 'LS17', 'LS16', 'LS14', 'LS11', 'LS08', 'LS05', 'LS03',✓
'LS02', 'Location', 'SouthEast');
hleg1.FontSize = 20;
ax = gca;
ax.FontSize = 24;
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.5, 0.7]);
ylabel('Pearson Correlation (r)');
xlabel('Time (Minutes)');
m = findobj(gca,'Type','line');
hleg1 = legend(m(1:9), 'Mean', 'LS17', 'LS16', 'LS14', 'LS11', 'LS08', 'LS05', 'LS03',✓
'LS02', 'Location', 'SouthEast');
hleg1.FontSize = 20;
ax = gca;
ax.FontSize = 24;
clear set
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.5, 0.7]);
xticks([5:5:120])
axis([5 120 .3 1])
xticks([10:10:120])
print(gcf,✓
['/Volumes/RESEARCH_HD/Lifespan/CNS_Analyses/ReliabilityLifespanRestDatatruhalf' num2str✓
(pts2sample) '.jpg'],'-png','-r300');
print(gcf,✓
```

```
['/Users/dianaperez/Desktop/Lifespan/CNS_Analyses/ReliabilityLifespanRestDatatruhal'↵
num2str(pts2sample) '.jpg'], '-dpng', '-r300');
print(gcf,['/Users/dianaperez/Desktop/Lifespan/ReliabilityLifespanRestDatatruhal'↵
num2str(pts2sample) '.jpg'], '-dpng', '-r300');
print(gcf,['/Users/dianaperez/Desktop/Lifespan/ReliabilityLifespanRestData.jpg'], '-↵
dpng', '-r300');
within_sub_reliability_v2
addpath '/Users/dianaperez/Documents/GitHub/GrattonLab-General-Repo/FCProcess'
within_sub_reliability_v2
figure;
for s = 1:numel(subject)
plot(times_all(s,1:size(allsubs_seg_ind{s},2)),mean(allsubs_seg_ind{1,s}(:,1:size↵
(allsubs_seg_ind{s},2)),1), 'Color',rgb_colors(s,:), 'LineWidth', 3)
hold on
end
ylabel('Expectation Value');
xlabel('Time (Minutes)');
m = findobj(gca,'Type','line');
ax = gca;
ax.FontSize = 24;
print(gcf,['/Volumes/RESEARCH_HD/Lifespan/CNS_Analyses/ReliabilityLifespanSegIndex.↵
jpg'], '-dpng', '-r300');
%-- 4/15/22, 2:51 PM --%
within_sub_reliability_v2
addpath '/Users/dianaperez/Documents/GitHub/GrattonLab-General-Repo/FCProcess'
within_sub_reliability_v2
s = 1
mean(allsubs_seg_ind{1,s}(:,1:size(allsubs_seg_ind{s},2)),1)
s = 2
[0.17513525,0.23543307,0.26925221,0.29455280,0.31462324,0.32750398,0.33855036,0.34856731,↵
0.35695845,0.36352569,0.36956799,0.37590152,0.38027725,0.38332370,0.38977948,0.39222571,0↵
.39717701,0.39897895,0.40176255,0.40565437,0.40783823,0.40995008,0.41150612,0.41320971,0.↵
41662928,0.41748437,0.41961083,0.42092493,0.42268413,0.42396396,0.42558959,0.42672858,0.4↵
2872712,0.42987919,0.43015543,0.43192738,0.43253067,0.43378815,0.43526950,0.43594587,0.43↵
680459,0.43797994,0.43884566,0.43988848,0.44037372,0.44093323,0.44164586,0.44217351,0.442↵
47830]
mean(allsubs_seg_ind{1,s}(:,1:size(allsubs_seg_ind{s},2)),1)
s = 3
mean(allsubs_seg_ind{1,s}(:,1:size(allsubs_seg_ind{s},2)),1)
s = 4
mean(allsubs_seg_ind{1,s}(:,1:size(allsubs_seg_ind{s},2)),1)
s = 5
mean(allsubs_seg_ind{1,s}(:,1:size(allsubs_seg_ind{s},2)),1)
s = 6
mean(allsubs_seg_ind{1,s}(:,1:size(allsubs_seg_ind{s},2)),1)
s = 7
mean(allsubs_seg_ind{1,s}(:,1:size(allsubs_seg_ind{s},2)),1)
s = 8
mean(allsubs_seg_ind{1,s}(:,1:size(allsubs_seg_ind{s},2)),1)
means = [0.228121374 0.282854956 0.311825923 0.330394855 0.344075468 0.3540615↵
0.362085299 0.368580906 0.374353669 0.378886496 0.38303392 0.386997298 0.39018919↵
0.392779018 0.395671349 0.3921087 0.394652817 0.396349383 0.398223 0.39998822↵
0.401678269 0.402921549 0.404130596 0.405287927 0.406779144 0.407857626 0.408819803↵
0.40779906 0.4087539 0.409494202 0.41042813 0.411371703 0.4121878 0.412706817↵
0.413360333 0.414032785 0.41465492 0.419794232 0.420634614 0.420983464 0.421506702↵
0.421954376 0.422355276 0.42278862 0.423175282 0.42350195 0.423740618 0.42402081↵
0.434813233 0.42669529 0.42691267 0.42712188 0.42740375];
plot(times_all(1,1:48),means(1:48), ':', 'Color', [0,0,0], 'LineWidth',3)
ylabel('Segregation Index');
xlabel('Time (Minutes)');
```



```

m = findobj(gca,'Type','line');
hleg1 = legend(m(1:9), 'Mean', 'LS17', 'LS16', 'LS14', 'LS11', 'LS08', 'LS05', 'LS03', 'LS02', 'Location', 'NorthEast');
hleg1.FontSize = 20;
ax = gca;
ax.FontSize = 24;
m = findobj(gca,'Type','line');
hleg1 = legend(m(1:9), 'Mean', 'LS17', 'LS16', 'LS14', 'LS11', 'LS08', 'LS05', 'LS03', 'LS02', 'Location', 'SouthEast');
hleg1.FontSize = 20;
ax = gca;
ax.FontSize = 24;
axis([0 120 .1 .5])
times = [2.5:2.5:100];
figure;
for s = 1:numel(subject)
plot(times_all(s,1:size(allsubs_seg_ind{s},2)),mean(allsubs_abs_diffs{1,s}(:,1:size(allsubs_seg_ind{s},2)),1), 'Color', rgb_colors(s,:), 'LineWidth', 3)
hold on
end
figure;
for s = 1:numel(subject)
plot(times_all(s,1:size(allsubs_seg_ind{s},2)),mean(allsubs_abs_diffs{1,s}(:,1:size(allsubs_seg_ind{s},2)),1), 'Color', rgb_colors(s,:), 'LineWidth', 3)
hold on
means_diff(s,:) = mean(allsubs_abs_diffs{1,s}(:,1:size(allsubs_seg_ind{s},2)),1);
end
figure;
for s = 1:numel(subject)
plot(times_all(s,1:size(allsubs_seg_ind{s},2)),mean(allsubs_abs_diffs{1,s}(:,1:size(allsubs_seg_ind{s},2)),1), 'Color', rgb_colors(s,:), 'LineWidth', 3)
hold on
means_diff{s} = mean(allsubs_abs_diffs{1,s}(:,1:size(allsubs_seg_ind{s},2)),1);
end
means_diff = [];
figure;
for s = 1:numel(subject)
plot(times_all(s,1:size(allsubs_seg_ind{s},2)),mean(allsubs_abs_diffs{1,s}(:,1:size(allsubs_seg_ind{s},2)),1), 'Color', rgb_colors(s,:), 'LineWidth', 3)
hold on
means_diff{s} = mean(allsubs_abs_diffs{1,s}(:,1:size(allsubs_seg_ind{s},2)),1);
end
mean_diff_all = [0.451054915    0.31958749    0.250078328    0.205589374    0.173270193
0.149834654    0.130898036    0.116096011    0.103579279    0.09357542    0.085252215    0.077510628
0.071893195    0.067765274    0.063014982    0.0609474    0.057319998    0.055589204    0.053587631
0.051688755    0.049855826    0.049189074    0.047716066    0.047485058    0.046371319    0.046092851
0.045026644    0.049051036    0.048147125    0.04798667    0.047803287    0.048079218    0.047927852
0.047976362    0.048676489    0.048785449    0.048605224    0.048932758    0.049221036];
plot(times_all(1,1:48),mean_diff_all(1:48), ':', 'Color', [0,0,0], 'LineWidth', 3)
mean_diff_all = [0.451054915    0.31958749    0.250078328    0.205589374    0.173270193
0.149834654    0.130898036    0.116096011    0.103579279    0.09357542    0.085252215    0.077510628
0.071893195    0.067765274    0.063014982    0.0609474    0.057319998    0.055589204    0.053587631
0.051688755    0.049855826    0.049189074    0.047716066    0.047485058    0.046371319    0.046092851
0.045026644    0.049051036    0.048147125    0.04798667    0.047803287    0.048079218    0.047927852
0.047976362    0.048676489    0.048785449    0.048605224    0.048932758    0.049221036    0.049052662
0.049501161    0.049540971    0.049734258    0.04993298    0.050447245    0.050401381    0.0504621
0.050506165    0.048969311    0.042753616    0.042789566    0.043138017    0.043480223    0.026972895];
plot(times_all(1,1:48),mean_diff_all(1:48), ':', 'Color', [0,0,0], 'LineWidth', 3)
axis([0 120 0 .6])
ylabel('% Difference');

```



```
xlabel('Time (Minutes)');
m = findobj(gca,'Type','line');
hleg1 = legend(m(1:9), 'Mean', 'LS17', 'LS16', 'LS14', 'LS11', 'LS08', 'LS05', 'LS03',
'LS02', 'Location', 'NorthEast');
hleg1.FontSize = 20;
ax = gca;
ax.FontSize = 24;
print(gcf,[output_dir '/ReliabilityLifespanSegIndex_Diff.jpg'],'-dpng','-r300');
figure;
for s = 1:numel(subject)
plot(times_all(s,1:size(allsubs_seg_ind{s},2)),mean(allsubs_seg_ind{1,s}(:,1:size(
allsubs_seg_ind{s},2)),1), 'Color',rgb_colors(s,:), 'LineWidth', 3)
hold on
end
means = [0.228121374    0.282854956 0.311825923 0.330394855 0.344075468 0.3540615
0.362085299 0.368580906 0.374353669 0.378886496 0.38303392 0.386997298 0.39018919
0.392779018 0.395671349 0.3921087 0.394652817 0.396349383 0.398223 0.39998822
0.401678269 0.402921549 0.404130596 0.405287927 0.406779144 0.407857626 0.408819803
0.40779906 0.4087539 0.409494202 0.41042813 0.411371703 0.4121878 0.412706817
0.413360333 0.414032785 0.41465492 0.419794232 0.420634614 0.420983464 0.421506702
0.421954376 0.422355276 0.42278862 0.423175282 0.42350195 0.423740618 0.42402081
0.434813233 0.42669529 0.42691267 0.42712188 0.42740375];
plot(times_all(1,1:48),means(1:48), ':', 'Color', [0,0,0], 'LineWidth',3)
ylabel('Segregation Index');
xlabel('Time (Minutes)');
% m = findobj(gca,'Type','line');
% hleg1 = legend(m(1:9), 'Mean', 'LS17', 'LS16', 'LS14', 'LS11', 'LS08', 'LS05', 'LS03',
'LS02', 'Location', 'SouthEast');
% hleg1.FontSize = 20;
ax = gca;
ax.FontSize = 24;
axis([0 120 0 .6])
axis([0 120 0.1 .5])
axis([0 120 0.15 .5])
xticks(0:10:120)
ax = gca;
ax.FontSize = 20;
print(gcf,[output_dir '/ReliabilityLifespanSegIndex.jpg'],'-dpng','-r300');
figure;
for s = 1:numel(subject)
plot(times_all(s,1:size(allsubs_seg_ind{s},2)),mean(allsubs_abs_diffs{1,s}(:,1:size(
allsubs_seg_ind{s},2)),1), 'Color',rgb_colors(s,:), 'LineWidth', 3)
hold on
means_diff{s} = mean(allsubs_abs_diffs{1,s}(:,1:size(allsubs_seg_ind{s},2)),1);
end
mean_diff_all = [0.451054915    0.31958749 0.250078328 0.205589374 0.173270193
0.149834654 0.130898036 0.116096011 0.103579279 0.09357542 0.085252215 0.077510628
0.071893195 0.067765274 0.063014982 0.0609474 0.057319998 0.055589204 0.053587631
0.051688755 0.049855826 0.049189074 0.047716066 0.047485058 0.046371319 0.046092851
0.045026644 0.049051036 0.048147125 0.04798667 0.047803287 0.048079218 0.047927852
0.047976362 0.048676489 0.048785449 0.048605224 0.048932758 0.049221036 0.049052662
0.049501161 0.049540971 0.049734258 0.04993298 0.050447245 0.050401381 0.0504621
0.050506165 0.048969311 0.042753616 0.042789566 0.043138017 0.043480223 0.026972895];
plot(times_all(1,1:48),mean_diff_all(1:48), ':', 'Color', [0,0,0], 'LineWidth',3)
axis([0 120 0 .6])
ylabel('% Difference');
xlabel('Time (Minutes)');
m = findobj(gca,'Type','line');
hleg1 = legend(m(1:9), 'Mean', 'LS17', 'LS16', 'LS14', 'LS11', 'LS08', 'LS05', 'LS03',
'LS02', 'Location', 'NorthEast');
```

```

hleg1.FontSize = 20;
ax = gca;
ax.FontSize = 20;
xticks(0:10:120)
print(gcf,[output_dir '/ReliabilityLifespanSegIndex_Diff.jpg'],'-dpng','-r300');
similarity_analysis_wLongitudinal
%-- 4/19/22, 3:47 PM --%
within_sub_reliability_v2
addpath '/Users/dianaperez/Documents/GitHub/GrattonLab-General-Repo/FCProcess'
within_sub_reliability_v2
addpath '/Users/dianaperez/Documents/GitHub/Lifespan-Analysis'
within_sub_reliability_v2
rgb_colors = [1 0 0;%LS02 red
0, 1, 0;%LS03 green
0, 0, 1;%LS05 blue
0, 1, 1;%LS08 cyan
1, 0, 1;%LS11 magenta
0.4660 0.6740 0.188;%LS14 other green
0.9290 0.6940 0.1250;%LS16 yellow orange-y
0.4940 0.1840 0.5560;%LS17 purple
0.4660 0.6740 0.1880;
0.4660 0.6740 0.1880];
rgb_colors = [1 0 0;%LS02 red
0, 1, 0;%LS03 green
0, 0, 1;%LS05 blue
0, 1, 1;%LS08 cyan
1, 0, 1;%LS11 magenta
0.4660 0.6740 0.188;%LS14 other green
0.9290 0.6940 0.1250;%LS16 yellow orange-y
0.4940 0.1840 0.5560; 0.3010 0.7450 0.9330; 0.8500 0.3250 0.0980];%LS17 purple
axis([0 120 0 .6])
ylabel('% Difference');
xlabel('Time (Minutes)');
m = findobj(gca,'Type','line');
hleg1 = legend(m(1:9), 'Mean', 'LS17', 'LS16', 'LS14', 'LS11', 'LS08', 'LS05', 'LS03',↵
'LS02', 'Location', 'NorthEast');
hleg1.FontSize = 20;
ax = gca;
ax.FontSize = 24;
%
print(gcf,[output_dir '/ReliabilityiNETSegIndex_Diff.jpg'],'-dpng','-r300');
axis([0 80 0 .6])
axis([0 80 0 .5])
axis([0 70 0 .5])
hleg1 = legend(m(1:9), 'Mean', 'LS17', 'LS16', 'LS14', 'LS11', 'LS08', 'LS05', 'LS03',↵
'LS02', 'Location', 'NorthEast');
hleg1 = legend(m(1:10), subject, 'Location', 'NorthEast');
print(gcf,[output_dir '/ReliabilityLifespanSegIndex_Diff.jpg'],'-dpng','-r300');
figure;
for s = 1:numel(subject)
plot(times_all(s,1:size(allsubs_seg_ind{s},2)),mean(allsubs_seg_ind{1,s}(:,1:size↵
(allsubs_seg_ind{s},2)),1),'Color',rgb_colors(s,:), 'LineWidth', 3)
hold on
end
endax = gca;
ax.FontSize = 20;
ax = gca;
ax.FontSize = 20;
ylabel('Segregation Index');
xlabel('Time (Minutes)');

```

```

print(gcf,[output_dir '/ReliabilityiNETSegIndex.jpg'],'-dpng','-r300')
load('/Users/dianaperez/Desktop/Segregation_Analyses/INET_allsubs_seg_index_ses.mat')
clear all
load('/Users/dianaperez/Desktop/Segregation_Analyses/INET_allsubs_seg_index_ses.mat')
load('/Users/dianaperez/Desktop/lifespan_allsubs_seg_index_sub_by_net.mat')
run('/Users/dianaperez/Desktop/segregation_ind_by_sub_by_net.m')
%-- 4/29/22, 3:41 PM --%
hist(logx)
hist(log)
hist(log(x))
rng('default')
rng('default')
x = random(pd,10000,1);
pd = makedist('Lognormal','mu',5,'sigma',2)
x = random(pd,10000,1);
logx = log(x);
hist(logx)
histfit(logx)
cmap = colormap(jet);
H = histfit(logx);
H2 = histcounts(logx);
b = bar(H2, 'facecolor', 'flat');
b.CData = cmap;
H2 = histcounts(logx,256);
b = bar(H2, 'facecolor', 'flat');
b.CData = cmap;
rng('default')
pd = makedist('Lognormal','mu',5,'sigma',2)
x = random(pd,10000,1);
logx = log(x);
cmap = colormap(jet);
nbins = 256;
data = histcounts(logx, nbins);
b = bar(data, 'facecolor', 'flat');
b.CData = cmap;
%-- 5/8/22, 1:31 PM --%
reliability_analysis_v2
mean = [0.515268013 0.622747069 0.681274594 0.718237485 0.745798623 0.765701861✓
0.781850887 0.794715224 0.80590723 0.814858538 0.822555385 0.829795586 0.835691422✓
0.840888584 0.84614858 0.850029338 0.854318895 0.857281215 0.85914574 0.861973127✓
0.864639569 0.867157803 0.869307432 0.871292319 0.872455141 0.864466596 0.85001729✓
0.91664582 0.91766846];
figure;
for s = 1:numel(subject)
plot(times_all(s,1:size(means{1,s},2)),means{1,s},'LineWidth', 3)
hold on
end
ylabel('Pearson Correlation (r)');
xlabel('Time (Minutes)');
plot(times_all(1,1:48),mean(1:48), ':', 'Color', [0,0,0], 'LineWidth',3) %average
plot(times_all(1,1:29),mean(1:29), ':', 'Color', [0,0,0], 'LineWidth',3) %average
clear gcf
figure;
for s = 1:numel(subject)
plot(times_all(s,1:size(means{1,s},2)),means{1,s},'Color',rgb_colors(s,:), 'LineWidth', 3)
hold on
end
plot(times_all(1,1:48),mean(1:48), ':', 'Color', [0,0,0], 'LineWidth',3) %average
ylabel('Pearson Correlation (r)');
xlabel('Time (Minutes)');

```

```

figure;
for s = 1:numel(subject)
plot(times_all(s,1:size(means{1,s},2)),means{1,s},'LineWidth', 3)
hold on
end
plot(times_all(1,1:48),mean(1:48), ':', 'Color', [0,0,0], 'LineWidth',3) %average
ylabel('Pearson Correlation (r)');
xlabel('Time (Minutes)');
plot(times_all(1,1:29),mean(1:29), ':', 'Color', [0,0,0], 'LineWidth',3) %average
ylabel('Pearson Correlation (r)');
xlabel('Time (Minutes)');
clear gcf
figure;
for s = 1:numel(subject)
plot(times_all(s,1:size(means{1,s},2)),means{1,s},'LineWidth', 3)
hold on
end
plot(times_all(4,1:29),mean(1:29), ':', 'Color', [0,0,0], 'LineWidth',3)
plot(times_all(4,1:26),mean(1:26), ':', 'Color', [0,0,0], 'LineWidth',3)
clear gcf
figure;
for s = 1:numel(subject)
plot(times_all(s,1:size(means{1,s},2)),means{1,s},'LineWidth', 3)
hold on
end
plot(times_all(4,1:26),mean(1:26), ':', 'Color', [0,0,0], 'LineWidth',3)
close gcf
figure;
for s = 1:numel(subject)
plot(times_all(s,1:size(means{1,s},2)),means{1,s},'LineWidth', 3)
hold on
end
plot(times_all(4,1:25),mean(1:25), ':', 'Color', [0,0,0], 'LineWidth',3)
ylabel('Pearson Correlation (r)');
xlabel('Time (Minutes)');
ax = gca;
ax.FontSize = 24;
print(gcf,['/Volumes/RESEARCH_HD/Lifespan/CNS_Analyses/ReliabilityiNETRestDatatruhalf' num2str(pts2sample) '.jpg'],'-dpng','-r300');
print(gcf,['/Users/dianaperez/Desktop/ReliabilityiNETRestDatatruhalf' num2str(pts2sample) '.jpg'],'-dpng','-r300');
similarity_analysis_forced_same_amt_data
hline_new([0,4,8,12,16,20,24,28,32]+0.5,'k',2);
hline_new([0,5,10,15,20,25,30,35,40,44,48]+0.5,'k',2);
figure('Position',[1 1 1000 800]);
imagesc(simmat,[0 1]); colormap('jet');
hline_new([0,4,8,12,16,20,24,28,32,36,40]+0.5,'k',2);
vline_new([0,4,8,12,16,20,24,28,32,36,40]+0.5,'k',2);
axis square;
colorbar;
title('Correlation Matrix Similarity');
set(gca,'XTick',[2,6,10,14,18,22,26,30,34,38], 'YTick', [2,6,10,14,18,22,26,30,34,38], 'XTickLabel',...
{'INET001','INET002'},
{'INET003','INET005','INET006','INET010','INET016','INET018','INET019','INET030'},
'YTickLabel', {'INET001','INET002'},
{'INET003','INET005','INET006','INET010','INET016','INET018','INET019','INET030'})
ax = gca;
ax.FontSize = 20
ax.FontSize = 18

```

```

ax.FontSize = 16
ax.FontSize = 14
saveas(gcf, '/Users/dianaperez/Desktop/iNet_SimilarityMat_rest.tiff', 'tiff')
%-- 5/21/22, 7:34 PM --%
var_freq_size_net
addpath '/Users/dianaperez/Documents/Dependencies/cifti-matlab-master'
var_freq_size_net
[data_location num2str(subs{g,1}(n)) netAssign_str]
var_freq_size_net
network_names = {'DMN' 'Vis' 'FP' 'DAN' 'Lang' 'Sal' 'CO' 'SMd' 'SMl'
'Aud' 'Tpole' 'MTL' 'PMN' 'PON'};
good_nets_LH = [networksxHem.verticesLH(:,1:3) networksxHem.verticesLH(:,5) networksxHem.
verticesLH(:,7:end)];
good_nets_RH = [networksxHem.verticesRH(:,1:3) networksxHem.verticesRH(:,5) networksxHem.
verticesRH(:,7:end)];
good_nets_LH = mean(good_nets_LH);
good_nets_RH = mean(good_nets_RH);
means = [good_nets_LH' good_nets_RH'];
bar(1:14, means)
network_names = {'DMN' 'Vis' 'FP' 'DAN' 'Lang' 'Sal' 'CO' 'SMd' 'SMl'
'Aud' 'Tpole' 'MTL' 'PMN' 'PON'};
good_nets_LH = [networksxHem.verticesLH(:,1:3) networksxHem.verticesLH(:,5) networksxHem.
verticesLH(:,7:end)];
good_nets_RH = [networksxHem.verticesRH(:,1:3) networksxHem.verticesRH(:,5) networksxHem.
verticesRH(:,7:end)];
good_nets_LH = mean(good_nets_LH);
good_nets_RH = mean(good_nets_RH);
means = [good_nets_LH' good_nets_RH'];
b = bar(1:14, means)
b(1) = [1,1,1];
b(1).FaceColor = [1,1,1];
b(1).FaceColor = [0,0,0];
b(2).FaceColor = [17,17,17];
b(2).FaceColor = [.5,.5,.5];
legend('Left Handers', 'Right Handers')
xticks(1:14)
xticklabels(network_names)
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.9, 0.7]);
ylabel('Average Number of Variant vertices');
xlabel('Network');
perms_var_freq_size
permute_net_assignment
root_dir = '/Users/dianaperez/Desktop/Research/Lateralization/lateralization_code/'; %
location of code directory
data_location = [root_dir 'testing_output/'];
LHand_info = load([data_location 'HCP752_LH_networksxHem.mat']);
RHand_info = load([data_location 'HCP752_RH_networksxHem.mat']);
%plot left-handers
clear all
root_dir = '/Users/dianaperez/Desktop/Research/Lateralization/lateralization_code/'; %
location of code directory
data_location = [root_dir 'testing_output/'];
LHand_info = load([data_location 'HCP752_LH_networksxHem.mat']);
RHand_info = load([data_location 'HCP752_RH_networksxHem.mat']);
%plot left-handers
lh_numvars = mean(LHand_info.networksxHem.clustersLH,1);
rh_numvars = mean(RHand_info.networksxHem.clustersRH,1);
for net = 1:16
se_lh(net) = std(LHand_info.networksxHem.clustersLH(:,net))/sqrt(40);
se_rh(net) = std(RHand_info.networksxHem.clustersRH(:,net))/sqrt(40);

```

```

end
h = errorbar(lh_numvars, rh_numvars, se_rh, se_rh, se_lh, se_lh, 'o')
h.Color = [0,0,0];
clear al
clear all
root_dir = '/Users/dianaperez/Desktop/Research/Lateralization/lateralization_code/'; %
location of code directory
data_location = [root_dir 'testing_output/'];
LHand_info = load([data_location 'HCP752_LH_networksxHem.mat']);
RHand_info = load([data_location 'HCP752_RH_networksxHem.mat']);
%plot left-handers
Lhand_lh_numvars = mean(LHand_info.networksxHem.clustersLH,1);
Lhand_rh_numvars = mean(LHand_info.networksxHem.clustersRH,1);
Lhand_diff_numvars = Lhand_lh_numvars-Lhand_rh_numvars;
clear all
load
('Users/dianaperez/Desktop/Research/lateralization_code/testing_output/HCP752_new_split_
LH_networksxHem.mat')
network_names = {'DMN' 'Vis' 'FP' 'DAN' 'Lang' 'Sal' 'CO' 'SMd' 'SMl'
'Aud' 'Tpole' 'MTL' 'PMN' 'PON'};
load
('Users/dianaperez/Desktop/Research/Lateralization/lateralization_code/testing_output/HCP
752_new_split_LH_networksxHem.mat')
network_names = {'DMN' 'Vis' 'FP' 'DAN' 'Lang' 'Sal' 'CO' 'SMd' 'SMl'
'Aud' 'Tpole' 'MTL' 'PMN' 'PON'};
good_nets_LH = [networksxHem.verticesLH(:,1:3) networksxHem.verticesLH(:,5) networksxHem.
verticesLH(:,7:end)];
good_nets_LH = [networksxHem.clustersLH(:,1:3) networksxHem.clustersLH(:,5) networksxHem.
clustersLH(:,7:end)];
good_nets_RH = [networksxHem.clustersRH(:,1:3) networksxHem.clustersRH(:,5) networksxHem.
clustersRH(:,7:end)];
good_nets_LH = mean(good_nets_LH);
good_nets_RH = mean(good_nets_RH);
means = [good_nets_LH' good_nets_RH'];
for net = 1:16
se_lh(net) = std(networksxHem.clustersLH(:,net))/sqrt(40);
se_rh(net) = std(networksxHem.clustersRH(:,net))/sqrt(40);
end
se_lh(4) = [];
se_lh(5) = [];
se_rh(4) = [];
se_rh(5) = [];
bar(1:14, means)
legend('Left Hem', 'Right Hem')
xticks(1:14)
xticklabels(network_names)
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.9, 0.7]);
ylabel('Average Number of Variant Regions')
xlabel('Network');
title('Variant Regions Assigned to Each Network Across Hemispheres')
ax = gca;
ax.FontSize = 24;
hold on
h = errorbar(.85:1:13.85, means(:,1), se_lh, se_lh)
h.Color = [0,0,0];
g = errorbar(1.15:1:14.15, means(:,2), se_rh, se_rh)
g.Color = [0,0,0];
h.LineStyle = 'none';
g.LineStyle = 'none';
h.LineWidth = 2;

```

```

g.LineWidth = 2;
legend('Left Hem', 'Right Hem')
close(gcf)
left_handlers = networksxHem;
load
('/Users/dianaperez/Desktop/Research/Lateralization/lateralization_code/testing_output/HCP752_RH_networksxHem.mat')
network_names = {'DMN' 'Vis' 'FP' 'DAN' 'Lang' 'Sal' 'CO' 'SMd' 'SMl'
'Aud' 'Tpole' 'MTL' 'PMN' 'PON'};
good_nets_LH = [networksxHem.verticesLH(:,1:3) networksxHem.verticesLH(:,5) networksxHem.
verticesLH(:,7:end)];
good_nets_LH = [networksxHem.clustersLH(:,1:3) networksxHem.clustersLH(:,5) networksxHem.
clustersLH(:,7:end)];
good_nets_RH = [networksxHem.clustersRH(:,1:3) networksxHem.clustersRH(:,5) networksxHem.
clustersRH(:,7:end)];
good_nets_LH = mean(good_nets_LH);
good_nets_RH = mean(good_nets_RH);
means = [good_nets_LH' good_nets_RH'];
for net = 1:16
se_lh(net) = std(networksxHem.clustersLH(:,net))/sqrt(670);
se_rh(net) = std(networksxHem.clustersRH(:,net))/sqrt(670);
end
se_lh(4) = [];
se_lh(5) = [];
se_rh(4) = [];
se_rh(5) = [];
bar(1:14, means)
legend('Left Hem', 'Right Hem')
xticks(1:14)
xticklabels(network_names)
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.9, 0.7]);
ylabel('Average Number of Variant Regions')
xlabel('Network');
title('Variant Regions Assigned to Each Network Across Hemispheres')
ax = gca;
ax.FontSize = 24;
hold on
h = errorbar(.85:1:13.85, means(:,1), se_lh, se_lh)
h.Color = [0,0,0];
g = errorbar(1.15:1:14.15, means(:,2), se_rh, se_rh)
g.Color = [0,0,0];
h.LineStyle = 'none';
g.LineStyle = 'none';
h.LineWidth = 2;
g.LineWidth = 2;
legend('Left Hem', 'Right Hem')
axis([0,15,0,2.5])
load
('/Users/dianaperez/Desktop/Research/Lateralization/lateralization_code/testing_output/HCP752_RH_networksxHem.mat')
network_names = {'DMN' 'Vis' 'FP' 'DAN' 'Lang' 'Sal' 'CO' 'SMd' 'SMl'
'Aud' 'Tpole' 'MTL' 'PMN' 'PON'};
good_nets_LH = [networksxHem.verticesLH(:,1:3) networksxHem.verticesLH(:,5) networksxHem.
verticesLH(:,7:end)];
good_nets_LH = [networksxHem.clustersLH(:,1:3) networksxHem.clustersLH(:,5) networksxHem.
clustersLH(:,7:end)];
good_nets_RH = [networksxHem.clustersRH(:,1:3) networksxHem.clustersRH(:,5) networksxHem.
clustersRH(:,7:end)];
good_nets_LH = mean(good_nets_LH);
good_nets_RH = mean(good_nets_RH);

```



```

means = [good_nets_LH' good_nets_RH'];
for net = 1:16
se_lh(net) = std(networksxHem.clustersLH(:,net))/sqrt(670);
se_rh(net) = std(networksxHem.clustersRH(:,net))/sqrt(670);
end
se_lh(4) = [];
se_lh(5) = [];
se_rh(4) = [];
se_rh(5) = [];
bar(1:14, means)
legend('Left Hem', 'Right Hem')
xticks(1:14)
xticklabels(network_names)
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.9, 0.7]);
ylabel('Average Number of Variant Regions')
xlabel('Network');
title('Variant Regions Assigned to Each Network Across Hemispheres')
ax = gca;
ax.FontSize = 24;
hold on
h = errorbar(.85:1:13.85, means(:,1), se_lh, se_lh)
h.Color = [0,0,0];
g = errorbar(1.15:1:14.15, means(:,2), se_rh, se_rh)
g.Color = [0,0,0];
h.LineStyle = 'none';
g.LineStyle = 'none';
h.LineWidth = 2;
g.LineWidth = 2;
legend('Left Hem', 'Right Hem')
axis([0,15,0,2.5])
close gcf
right_handlers = networksxHem;
diff_lefthanders = left_handlers.clustersLH - left_handlers.clustersRH;
mean_diff_lefthanders = mean(diff_lefthanders, 1);
diff_righthanders = right_handlers.clustersLH - right_handlers.clustersRH;
mean_diff_righthanders = mean(diff_righthanders, 1);
for net = 1:16
se_lh(net) = std(diff_lefthanders(:,net))/sqrt(40);
se_rh(net) = std(diff_righthanders(:,net))/sqrt(670);
end
se_lh(4) = [];
se_lh(5) = [];
se_rh(4) = [];
se_rh(5) = [];
diffs = [mean_diff_lefthanders; mean_diff_righthanders];
diffs(:,4) = []; diffs(:,5) = [];
b = bar(1:14, diffs)
b(1).FaceColor = [0,0,0];%[.2 .6 .5];
b(2).FaceColor = [.5,.5,.5];%[0.4940 0.1840 0.5560];
legend('Left Handers', 'Right Handers')
xticks(1:14)
xticklabels(network_names)
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.9, 0.7]);
ylabel(' ')
xlabel(' ');
ax = gca;
ax.FontSize = 24;
hold on
h = errorbar(.85:1:13.85, diffs(1,:), se_lh, se_lh)
h.Color = [0,0,0];

```



```

g = errorbar(1.15:1:14.15, diffs(2,:), se_rh, se_rh)
g.Color = [0,0,0];
h.LineStyle = 'none';
g.LineStyle = 'none';
h.LineWidth = 2;
g.LineWidth = 2;
legend('Left Handers', 'Right Handers')
h = errorbar(.85:1:13.85, diffs(1,:), [], se_lh)
h.Color = [1,1,1];
close(gcf)
diff_lefthanders = left_handers.clustersLH - left_handers.clustersRH;
mean_diff_lefthanders = mean(diff_lefthanders, 1);
diff_righthanders = right_handers.clustersLH - right_handers.clustersRH;
mean_diff_righthanders = mean(diff_righthanders, 1);
for net = 1:16
se_lh(net) = std(diff_lefthanders(:,net))/sqrt(40);
se_rh(net) = std(diff_righthanders(:,net))/sqrt(670);
end
se_lh(4) = [];
se_lh(5) = [];
se_rh(4) = [];
se_rh(5) = [];
diffs = [mean_diff_lefthanders; mean_diff_righthanders];
diffs(:,4) = []; diffs(:,5) = [];
b = bar(1:14, diffs)
b(1).FaceColor = [0,0,0];%[.2 .6 .5];
b(2).FaceColor = [.5,.5,.5];%[0.4940 0.1840 0.5560];
legend('Left Handers', 'Right Handers')
xticks(1:14)
xticklabels(network_names)
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.9, 0.7]);
ylabel(' ')
xlabel(' ');
ax = gca;
ax.FontSize = 24;
hold on
h = errorbar(.85:1:13.85, diffs(1,:), se_lh, se_lh)
h.Color = [0,0,0];
g = errorbar(1.15:1:14.15, diffs(2,:), se_rh, se_rh)
g.Color = [0,0,0];
h.LineStyle = 'none';
g.LineStyle = 'none';
h.LineWidth = 2;
g.LineWidth = 2;
legend('Left Handers', 'Right Handers')
h = errorbar(.85:1:13.85, diffs(1,:), se_lh, se_lh)
h.Color = [1,1,1];
close(gcf)
diff_lefthanders = left_handers.clustersLH - left_handers.clustersRH;
mean_diff_lefthanders = mean(diff_lefthanders, 1);
diff_righthanders = right_handers.clustersLH - right_handers.clustersRH;
mean_diff_righthanders = mean(diff_righthanders, 1);
for net = 1:16
se_lh(net) = std(diff_lefthanders(:,net))/sqrt(40);
se_rh(net) = std(diff_righthanders(:,net))/sqrt(670);
end
se_lh(4) = [];
se_lh(5) = [];
se_rh(4) = [];
se_rh(5) = [];

```

```

diffs = [mean_diff_lefthanders; mean_diff_righthanders];
diffs(:,4) = []; diffs(:,5) = [];
b = bar(1:14, diffs)
b(1).FaceColor = [0,0,0];%[.2 .6 .5];
b(2).FaceColor = [.5,.5,.5];%[0.4940 0.1840 0.5560];
legend('Left Handers', 'Right Handers')
xticks(1:14)
xticklabels(network_names)
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.9, 0.7]);
ylabel(' ')
xlabel(' ');
ax = gca;
ax.FontSize = 24;
hold on
h = errorbar(.85:1:13.85, diffs(1,:), se_lh, se_lh)
h.Color = [0,0,0];
g = errorbar(1.15:1:14.15, diffs(2,:), se_rh, se_rh)
g.Color = [0,0,0];
h.LineStyle = 'none';
g.LineStyle = 'none';
h.LineWidth = 2;
g.LineWidth = 2;
legend('Left Handers', 'Right Handers')
close gcf
diff_lefthanders = left_handers.clustersLH - left_handers.clustersRH;
mean_diff_lefthanders = mean(diff_lefthanders, 1);
diff_righthanders = right_handers.clustersLH - right_handers.clustersRH;
mean_diff_righthanders = mean(diff_righthanders, 1);
for net = 1:16
se_lh(net) = std(diff_lefthanders(:,net))/sqrt(40);
se_rh(net) = std(diff_righthanders(:,net))/sqrt(670);
end
se_lh(4) = [];
se_lh(5) = [];
se_rh(4) = [];
se_rh(5) = [];
diffs = [mean_diff_lefthanders; mean_diff_righthanders];
diffs(:,4) = []; diffs(:,5) = [];
b = bar(1:14, diffs)
b(1).FaceColor = [0,0,0];%[.2 .6 .5];
b(2).FaceColor = [.5,.5,.5];%[0.4940 0.1840 0.5560];
legend('Left Handers', 'Right Handers')
xticks(1:14)
xticklabels(network_names)
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.9, 0.7]);
ylabel(' ')
xlabel(' ');
ax = gca;
ax.FontSize = 24;
hold on
h = errorbar(.85:1:13.85, diffs(1,:), se_lh, se_lh)
h.Color = [0.5,0.5,0.5];
g = errorbar(1.15:1:14.15, diffs(2,:), se_rh, se_rh)
g.Color = [0,0,0];
h.LineStyle = 'none';
g.LineStyle = 'none';
h.LineWidth = 2;
g.LineWidth = 2;
legend('Left Handers', 'Right Handers')
print(gcf, '/Users/dianaperez/Desktop/HCP752_numvars_LvRhand_L-Rhem.jpg', '-dpng', '-<

```

```

r300')
%-- 5/26/22, 6:43 PM --%
help rand
help datasample
randperm
help randperm
for i = 1:1000000000000000
mat(i) = randperm([0,1], 1);
end
for i = 1:1000000000000000
mat(i) = randn([0,1], 1);
end
help randn
for i = 1:10
mat(i) = datasample([0,1], 1);
end
sum(mat)/length(mat)
for i = 1:10
mat(i) = datasample([0,1], 1);
end
for i = 1:10000000000000000000
mat(i) = datasample([0,1], 1);
end
sum(mat)/length(mat)
for i = 1:1000000000
mat(i) = datasample([0,1], 1);
end
sum(mat)/length(mat)
sum(mat)/length(mat)
%-- 6/6/22, 3:23 PM --%
ismac
ispc
islinux
IsLinux
isunix
path1 = '/Users/dianaperez/Desktop/';
if ismac
path2 = replace('\', path1, '/');
elseif ispc
path2 = replace('/', path1, '\');
end
if ispc
path2 = replace('\', path1, '/');
elseif ismac
path2 = replace('/', path1, '\');
end
if ispc
path2 = replace(path1, '\', '/');
elseif ismac
path2 = replace(path1, '/', '\');
end
perms_var_freq_size
permute_net_assignment
[p_fdr, p_masked] = fdr([0.00004373 0.00000249 0.7522832 0.03489360 0.00000000
0.26316159 0.00762987 0.41333184 0.00000029 0.11045880 0.02515492 0.03078876
0.63493542], 0.025)
[p_fdr, p_masked] = FDR([0.00004373 0.00000249 0.7522832 0.03489360 0.00000000
0.26316159 0.00762987 0.41333184 0.00000029 0.11045880 0.02515492 0.03078876
0.63493542], 0.025)
[p_fdr, p_masked] = FDR([0.004405384 0.0000000004045834881115

```

```
0.0000000000000000], 0.025)
cd /Users/dianaperez/Desktop
cd MSC_variants
ft_read_cifti_mod('Templatematch_spCorr_bysubject_IDs_reassigned.dtseries.nii')
addpath '/Users/dianaperez/Documents/Dependencies/cifti-matlab-master'
ft_read_cifti_mod('Templatematch_spCorr_bysubject_IDs_reassigned.dtseries.nii')
clear all
ft_read_cifti_mod('Templatematch_spCorr_bysubject_IDs_reassigned.dtseries.nii')
MSC_vars = ans;
unique(MSC_vars.data(:,1))
unique(MSC_vars.data(:,2))
unique(MSC_vars.data(:,3))
unique(MSC_vars.data(:,4))
unique(MSC_vars.data(:,7))
template = ft_read_cifti_mod('MSC08_topVariants_binary_corr<0.3.dtseries.nii')
template.data = MSC_vars.data(:,1);
unique(template.data)
ft_write_cifti_mod('MSC01_variants_reassigned.dtseries.nii', template)
template.data = MSC_vars.data(:,2);
ft_write_cifti_mod('MSC02_variants_reassigned.dtseries.nii', template)
unique(template.data)
template.data = MSC_vars.data(:,3);
unique(template.data)
ft_write_cifti_mod('MSC03_variants_reassigned.dtseries.nii', template)
template.data = MSC_vars.data(:,4);
unique(template.data)
ft_write_cifti_mod('MSC04_variants_reassigned.dtseries.nii', template)
template.data = MSC_vars.data(:,5);
unique(template.data)
ft_write_cifti_mod('MSC05_variants_reassigned.dtseries.nii', template)
template.data = MSC_vars.data(:,6);
unique(template.data)
ft_write_cifti_mod('MSC06_variants_reassigned.dtseries.nii', template)
template.data = MSC_vars.data(:,7);
ft_write_cifti_mod('MSC07_variants_reassigned.dtseries.nii', template)
template.data = MSC_vars.data(:,8);
ft_write_cifti_mod('MSC09_variants_reassigned.dtseries.nii', template)
template.data = MSC_vars.data(:,9);
ft_write_cifti_mod('MSC10_variants_reassigned.dtseries.nii', template)
clear all
segregation_ind_by_net
cd '/Users/dianaperez/Documents/GitHub/GrattonLab-General-Repo/FCProcess'
addpath '/Users/dianaperez/Documents/GitHub/GrattonLab-General-Repo/FCProcess'
segregation_ind_by_net
output_dir = '/Users/dianaperez/Desktop/Research/Segregation_Analyses/';
save([output_dir subs{sub} '_seg_index_net_withNaNnegs.mat'], 'sub_struct')
segregation_ind_by_net
segregation_ind_by_ses
segregation_ind_by_sub
segregation_ind_by_ses
segregation_ind_by_net
clear all
load␣
('/Users/dianaperez/Desktop/Research/Segregation_Analyses/INET019_seg_index_net_withNaNne
gs.mat')
load␣
('/Users/dianaperez/Desktop/Research/Segregation_Analyses/INET018_seg_index_net_withNaNne
gs.mat')
load␣
('/Users/dianaperez/Desktop/Research/Segregation_Analyses/INET016_seg_index_net_withNaNne
```

```
gs.mat')
load
('/Users/dianaperez/Desktop/Research/Segregation_Analyses/INET010_seg_index_net_withNaNne
gs.mat')
load
('/Users/dianaperez/Desktop/Research/Segregation_Analyses/INET006_seg_index_net_withNaNne
gs.mat')
load
('/Users/dianaperez/Desktop/Research/Segregation_Analyses/INET005_seg_index_net_withNaNne
gs.mat')
load
('/Users/dianaperez/Desktop/Research/Segregation_Analyses/INET003_seg_index_net_withNaNne
gs.mat')
load
('/Users/dianaperez/Desktop/Research/Segregation_Analyses/INET002_seg_index_net_withNaNne
gs.mat')
load
('/Users/dianaperez/Desktop/Research/Segregation_Analyses/INET001_seg_index_net_withNaNne
gs.mat')
segregation_ind_by_net
load
('/Users/dianaperez/Desktop/Research/Segregation_Analyses/LS17_seg_index_net_withNaNnegs.
mat')
load
('/Users/dianaperez/Desktop/Research/Segregation_Analyses/LS16_seg_index_net_withNaNnegs.
mat')
load
('/Users/dianaperez/Desktop/Research/Segregation_Analyses/LS14_seg_index_net_withNaNnegs.
mat')
load
('/Users/dianaperez/Desktop/Research/Segregation_Analyses/LS11_seg_index_net_withNaNnegs.
mat')
load
('/Users/dianaperez/Desktop/Research/Segregation_Analyses/LS08_seg_index_net_withNaNnegs.
mat')
load
('/Users/dianaperez/Desktop/Research/Segregation_Analyses/LS05_seg_index_net_withNaNnegs.
mat')
load
('/Users/dianaperez/Desktop/Research/Segregation_Analyses/LS03_seg_index_net_withNaNnegs.
mat')
load
('/Users/dianaperez/Desktop/Research/Segregation_Analyses/LS02_seg_index_net_withNaNnegs.
mat')
load
('/Users/dianaperez/Desktop/Research/Segregation_Analyses/INET030_seg_index_net_with0negs
.mat')
load
('/Users/dianaperez/Desktop/Research/Segregation_Analyses/INET019_seg_index_net_with0negs
.mat')
load
('/Users/dianaperez/Desktop/Research/Segregation_Analyses/INET018_seg_index_net_with0negs
.mat')
load
('/Users/dianaperez/Desktop/Research/Segregation_Analyses/INET016_seg_index_net_with0negs
.mat')
load
('/Users/dianaperez/Desktop/Research/Segregation_Analyses/INET010_seg_index_net_with0negs
.mat')
load
```

```
( '/Users/dianaperez/Desktop/Research/Segregation_Analyses/INET006_seg_index_net_with0negs
.mat')
load✓
( '/Users/dianaperez/Desktop/Research/Segregation_Analyses/INET005_seg_index_net_with0negs
.mat')
load✓
( '/Users/dianaperez/Desktop/Research/Segregation_Analyses/INET003_seg_index_net_with0negs
.mat')
load✓
( '/Users/dianaperez/Desktop/Research/Segregation_Analyses/INET002_seg_index_net_with0negs
.mat')
load✓
( '/Users/dianaperez/Desktop/Research/Segregation_Analyses/INET001_seg_index_net_with0negs
.mat')
load✓
( '/Users/dianaperez/Desktop/Research/Segregation_Analyses/LS17_seg_index_net_with0negs.
mat')
load✓
( '/Users/dianaperez/Desktop/Research/Segregation_Analyses/LS16_seg_index_net_with0negs.
mat')
load✓
( '/Users/dianaperez/Desktop/Research/Segregation_Analyses/LS14_seg_index_net_with0negs.
mat')
load✓
( '/Users/dianaperez/Desktop/Research/Segregation_Analyses/LS11_seg_index_net_with0negs.
mat')
load✓
( '/Users/dianaperez/Desktop/Research/Segregation_Analyses/LS08_seg_index_net_with0negs.
mat')
load✓
( '/Users/dianaperez/Desktop/Research/Segregation_Analyses/LS05_seg_index_net_with0negs.
mat')
load✓
( '/Users/dianaperez/Desktop/Research/Segregation_Analyses/LS03_seg_index_net_with0negs.
mat')
load✓
( '/Users/dianaperez/Desktop/Research/Segregation_Analyses/LS02_seg_index_net_with0negs.
mat')
load✓
( '/Users/dianaperez/Desktop/Research/Segregation_Analyses/INET_allsubs_seg_index_ses_with
NaNnegs.mat')
load✓
( '/Users/dianaperez/Desktop/Research/Segregation_Analyses/INET_allsubs_seg_index_ses_with
0negs.mat')
clear all
segregation_ind_by_ses
load('/Users/dianaperez/Desktop/Research/Segregation_Analyses/allsubs_seg_index_ses.mat')
segregation_ind_by_ses
load✓
( '/Users/dianaperez/Desktop/Research/Segregation_Analyses/LS_allsubs_seg_index_sub_with0n
eg.mat')
load✓
( '/Users/dianaperez/Desktop/Research/Segregation_Analyses/LS_allsubs_seg_index_sub_withNa
Nneg.mat')
load✓
( '/Users/dianaperez/Desktop/Research/Segregation_Analyses/INET_allsubs_seg_index_sub_with
0neg.mat')
load✓
( '/Users/dianaperez/Desktop/Research/Segregation_Analyses/INET_allsubs_seg_index_sub_with
NaNneg.mat')
```

```

scatter(sub_SI)
scatter(sub_SI, length(sub_SI))
length(sub_SI)
help scatter
scatter(length(sub_SI), sub_SI)
sub_SI
scatter(length(sub_SI), sub_SI')
scatter(10, sub_SI')
scatter(10, sub_SI)
plot(10, sub_SI)
plot(1:10, sub_SI)
scatter(1:10, sub_SI)
clear all
load
('Users/dianaperez/Desktop/Research/Segregation_Analyses/INET_allsubs_seg_index_sub_with
NaNneg.mat')
iNet_SI = sub_SI';
load
('Users/dianaperez/Desktop/Research/Segregation_Analyses/LS_allsubs_seg_index_sub_withNa
Nneg.mat')
lifespan_SI = sub_SI';
iNet_SI(5,:) = -1;
load
('Users/dianaperez/Desktop/Research/Segregation_Analyses/INET_allsubs_seg_index_sub_with
NaNneg.mat')
iNet_SI = sub_SI';
lifespan_SI(9) = -1;
lifespan_SI(10) = -1;
handles = plotSpread([lifespan_SI iNet_SI], 'distributionMarkers', {'o'}, 'xNames',
{'YA', 'OA'}, 'binWidth', 1)
handles = plotSpread([lifespan_SI iNet_SI], 'xNames', {'YA', 'OA'}, 'binWidth', 1)
load
('Users/dianaperez/Desktop/Research/Segregation_Analyses/INET_allsubs_seg_index_sub_with
NaNneg.mat')
iNet_SI = sub_SI';
load
('Users/dianaperez/Desktop/Research/Segregation_Analyses/LS_allsubs_seg_index_sub_withNa
Nneg.mat')
lifespan_SI = sub_SI';
% lifespan_SI(9) = -1;
% lifespan_SI(10) = -1;
ind=[1,1,1,1,1,1,1,1,1,1,0,0,0,0,0,0,0,0];
handles = plotSpread([iNet_SI; lifespan_SI], 'xNames', {'YA', 'OA'}, 'binWidth', 1)
handles = plotSpread([iNet_SI; lifespan_SI], 'xNames', {'YA', 'OA'},
'distributionColors', {'b', 'r'}, 'binWidth', 1)
ind=[1,1,1,1,1,1,1,1,1,1,2,2,2,2,2,2,2,2];
handles = plotSpread([iNet_SI; lifespan_SI], 'distributionIdx', ind, 'xNames', {'YA',
'OA'}, 'distributionColors', {'b', 'r'}, 'binWidth', 1)
handles = plotSpread([iNet_SI; lifespan_SI], 'distributionIdx',
ind, 'distributionMarkers', {'o'}, 'xNames', {'YA', 'OA'}, 'distributionColors', {'b',
'r'}, 'binWidth', 1)
handles = plotSpread([iNet_SI; lifespan_SI], 'distributionIdx',
ind, 'distributionMarkers', {'x'}, 'xNames', {'YA', 'OA'}, 'distributionColors', {'b',
'r'}, 'binWidth', 1)
load
('Users/dianaperez/Desktop/Research/Segregation_Analyses/INET_allsubs_seg_index_sub_with
0neg.mat')
iNet_SI = sub_SI';
load
('Users/dianaperez/Desktop/Research/Segregation_Analyses/LS_allsubs_seg_index_sub_with0n

```

```

eg.mat')
lifespan_SI = sub_SI';
handles = plotSpread([iNet_SI; lifespan_SI], 'distributionIdx',
ind, 'distributionMarkers', {'o'}, 'xNames', {'YA', 'OA'}, 'distributionColors', {'b',
'r'}, 'binWidth', 1)
handles = plotSpread([iNet_SI; lifespan_SI], 'distributionIdx',
ind, 'distributionMarkers', {'o'}, 'xNames', {'YA', 'OA'}, 'distributionColors', {'b',
'r'}, 'binWidth', 1)
ax = gca;
ax.FontSize = 24;
m = findobj(gca, 'Type', 'Scatter')
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.3, 0.7]); %first and second
control position on screen, third controls width, and fourth controls height
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.3, 0.8]); %first and second
control position on screen, third controls width, and fourth controls height
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0, 0.8]); %first and second
control position on screen, third controls width, and fourth controls height
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.8, 0.8]); %first and second
control position on screen, third controls width, and fourth controls height
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.3, 0.7]); %first and second
control position on screen, third controls width, and fourth controls height
axis([0, 3, 0, 1])
axis([0, 3, 0.3, .8])
print(gcf, '/Users/dianaperez/Desktop/SI_by_sub_NanVs0.jpg', '-dpng', '-r300')
load('/Users/dianaperez/Desktop/Research/Segregation_Analyses/allsubs_seg_index_ses.mat')
lifespan_SI = ses_SI;
load('/Users/dianaperez/Desktop/Research/Segregation_Analyses/INET_allsubs_seg_index_ses.
mat')
INET_SI(:,5) = -1;
INET_SI = ses_SI;
INET_SI(:,5) = -1;
ind=[1,1,1,1,1,1,1,1,1,1,1,2,2,2,2,2,2,2,2];
subs = {'LS02', 'LS03', 'LS05', 'LS08', 'LS11', 'LS14',
'LS16', 'LS17', 'INET001', 'INET002',
'INET003', 'INET005', 'INET006', 'INET010', 'INET016', 'INET018', 'INET019', 'INET030'};
subs = {'INET001', 'INET002',
'INET003', 'INET005', 'INET006', 'INET010', 'INET016', 'INET018', 'INET019', 'INET030', 'LS02',
'LS03', 'LS05', 'LS08', 'LS11', 'LS14', 'LS16', 'LS17'};
handles = plotSpread([iNet_SI; lifespan_SI], 'distributionIdx',
ind, 'distributionMarkers', {'o'}, 'xNames', {'YA', 'OA'}, 'distributionColors', {'b',
'r'}, 'binWidth', 1)
load('/Users/dianaperez/Desktop/Research/Segregation_Analyses/allsubs_seg_index_ses.mat')
lifespan_SI = ses_SI;
load('/Users/dianaperez/Desktop/Research/Segregation_Analyses/INET_allsubs_seg_index_ses.
mat')
iNet_SI = ses_SI;
iNet_SI(:,5) = -1;
[iNet_SI; lifespan_SI]
handles = plotSpread([iNet_SI; lifespan_SI], 'distributionIdx',
ind, 'distributionMarkers', {'x'}, 'xNames', subs, 'distributionColors', {'b',
'r'}, 'binWidth', 1)
handles = plotSpread([iNet_SI; lifespan_SI])
figure; handles = plotSpread([iNet_SI; lifespan_SI])
figure; handles = plotSpread([iNet_SI; lifespan_SI], 'distributionIdx', ind)
ind
figure; handles = plotSpread([iNet_SI; lifespan_SI], 'distributionIdx', [ind;ind;ind;
ind;ind])
[ind;ind;ind;ind;ind]
[iNet_SI; lifespan_SI]
figure; handles = plotSpread([iNet_SI; lifespan_SI])

```



```
figure; handles = plotSpread([iNet_SI; lifespan_SI]', 'categoryIdx', ↵
ind, 'categorynMarkers', {'x'}, 'xNames', subs, 'categoryColors', {'b', 'r'}, 'binWidth', 1)
handles = plotSpread([iNet_SI; lifespan_SI]', 'categoryIdx', ind, 'categoryMarkers', ↵
{'x'}, 'xNames', subs, 'categoryColors', {'b', 'r'}, 'binWidth', 1)
handles = plotSpread([iNet_SI; lifespan_SI]', 'categoryIdx', ind, 'categoryMarkers', ↵
{'x', 'x'}, 'xNames', subs, 'categoryColors', {'b', 'r'}, 'binWidth', 1)
handles = plotSpread([iNet_SI; lifespan_SI]', 'categoryIdx', [ind;ind;ind;ind;↵
ind], 'categoryMarkers', {'x', 'x'}, 'xNames', subs, 'categoryColors', {'b', ↵
'r'}, 'binWidth', 1)
subs = {'INET001', 'INET002', ↵
'INET003', 'INET005', 'INET006', 'INET010', 'INET016', 'INET018', 'INET019', 'INET030', 'LS02', ↵
'LS03', 'LS05', 'LS08', 'LS11', 'LS14', 'LS16', 'LS17'};
handles = plotSpread([iNet_SI; lifespan_SI]', 'categoryIdx', [ind;ind;ind;ind;↵
ind], 'categoryMarkers', {'x'}, 'xNames', subs, 'categoryColors', {'b', 'r'}, 'binWidth', 1)
ax = gca;
ax.FontSize = 24;
axis([0, 3, 0.3, .8])
subs = {'INET001', 'INET002', ↵
'INET003', 'INET005', 'INET006', 'INET010', 'INET016', 'INET018', 'INET019', 'INET030', 'LS02', ↵
'LS03', 'LS05', 'LS08', 'LS11', 'LS14', 'LS16', 'LS17'};
handles = plotSpread([iNet_SI; lifespan_SI]', 'categoryIdx', [ind;ind;ind;ind;↵
ind], 'categoryMarkers', {'x', 'x'}, 'xNames', subs, 'categoryColors', {'b', ↵
'r'}, 'binWidth', 1)
ax = gca;
ax.FontSize = 24;
axis([0, 3, 0.3, .8])
axis([0, 3, 20, .8])
handles = plotSpread([iNet_SI; lifespan_SI]', 'categoryIdx', [ind;ind;ind;ind;↵
ind], 'categoryMarkers', {'x', 'x'}, 'xNames', subs, 'categoryColors', {'b', ↵
'r'}, 'binWidth', 1)
ax = gca;
ax.FontSize = 24;
axis([0, 20, .3, .8])
axis([0, 19, .3, .8])
axis([0, 18, .3, .8])
axis([0, 18.5, .3, .8])
axis([0.5, 18.5, .3, .8])
axis([0.5, 18.5, .1, .8])
axis([0.5, 18.5, .2, .6])
axis([0.5, 18.5, .3, .6])
axis([0.5, 18.5, .3, .55])
ax.FontSize = 20;
ax.FontSize = 24;
load↵
('/Users/dianaperez/Desktop/Research/Segregation_Analyses/LS_allsubs_seg_index_ses_with0n↵
egs.mat')
load↵
('/Users/dianaperez/Desktop/Research/Segregation_Analyses/INET_allsubs_seg_index_ses_with↵
0negs.mat')
load↵
('/Users/dianaperez/Desktop/Research/Segregation_Analyses/LS_allsubs_seg_index_ses_with0n↵
egs.mat')
lifespan_SI = ses_SI;
load↵
('/Users/dianaperez/Desktop/Research/Segregation_Analyses/INET_allsubs_seg_index_ses_with↵
0negs.mat')
iNet_SI = ses_SI;
iNet_SI(:,5) = -1;
handles = plotSpread([iNet_SI; lifespan_SI]', 'categoryIdx', [ind;ind;ind;ind;↵
ind], 'categoryMarkers', {'o', 'o'}, 'xNames', subs, 'categoryColors', {'b', ↵
```

```
'r'}, 'binWidth', 1)
axis([0.5, 18.5, .3, .8])
axis([0.5, 18.5, .3, .7])
print(gcf, '/Users/dianaperez/Desktop/SI_by_ses_NanVs0.jpg', '-dpng', '-r300')
load
(' /Users/dianaperez/Desktop/Research/Segregation_Analyses/LS17_seg_index_net_withNaNnegs.
mat')
help contains
data = [];
for sub = 1:numel(subs)
load([subs(sub) '_seg_index_net_withNaNnegs.mat'])
sub_struct(15,:) = [];
if contains(subs(sub), 'INET')
sub_struct(:,5) = -1
end
data = [data; sub_struct];
end
data = [];
for sub = 1:numel(subs)
load([subs{sub} '_seg_index_net_withNaNnegs.mat'])
sub_struct(15,:) = [];
if contains(subs{sub}, 'INET')
sub_struct(:,5) = -1
end
data = [data; sub_struct];
end
data = [];
for sub = 1:numel(subs)
load([' /Users/dianaperez/Desktop/Research/Segregation_Analyses/' subs{sub}
'_seg_index_net_withNaNnegs.mat'])
sub_struct(15,:) = [];
if contains(subs{sub}, 'INET')
sub_struct(:,5) = -1
end
data = [data; sub_struct];
end
data = [];
for sub = 1:numel(subs)
load([' /Users/dianaperez/Desktop/Research/Segregation_Analyses/' subs{sub}
'_seg_index_net_withNaNnegs.mat'])
sub_struct.seg_ind(15,:) = [];
if contains(subs{sub}, 'INET')
sub_struct.seg_ind(:,5) = -1
end
data = [data; sub_struct.seg_ind];
end
data'
[1:14]*18
ind = [1:14];
ind = [ind;ind;ind;ind;ind;ind;ind;ind;ind;ind;ind;ind;ind;ind;ind;ind;ind;ind];
ind = [1:14]';
ind = [ind;ind;ind;ind;ind;ind;ind;ind;ind;ind;ind;ind;ind;ind;ind;ind;ind;ind];
help repmat
ind = repmat([1:14]',252,1);
ind = repmat([1:14]',18,1);
handles = plotSpread(data)
data = [];
for sub = 1:numel(subs)
load([' /Users/dianaperez/Desktop/Research/Segregation_Analyses/' subs{sub}
'_seg_index_net_withNaNnegs.mat'])
```

```
sub_struct.seg_ind(15,:) = [];
if contains(subs{sub}, 'INET')
sub_struct.seg_ind(:,5) = -1
end
data = [data sub_struct.seg_ind];
end
handles = plotSpread(data)
handles = plotSpread(data')
cx_ind(1:50) = 1;
cx_ind(51:90) = 2;
data = data';
cx_ind2 = repmat(cx_ind, 18,1);
cx_ind2 = repmat(cx_ind4, 18,1);
cx_ind2 = repmat(cx_ind, 14,1);
atlas_dir = '/Volumes/fsmresfiles/PBS/Gratton_Lab/Atlases/';
atlas = 'Seitzman300';
atlas_params = atlas_parameters_GrattonLab(atlas,atlas_dir);% load atlas that contains
roi info (including which rois belong to each network)
nets = {'unassigned' 'SMd' 'SML' 'CO' 'Aud' 'DMN' 'PMN' 'Vis' 'FP' 'Sal' 'Lang' 'DAN'
'MTL' 'PON'};
handles = plotSpread(data, 'categoryIdx', cx_ind, 'categoryMarkers', {'o','o'}, 'xNames',
nets, 'categoryColors', {'b', 'r'}, 'binWidth', 1)
handles = plotSpread(data, 'categoryIdx', cx_ind, 'categoryMarkers', {'o','o'}, 'xNames',
nets, 'categoryColors', {'b', 'r'}, 'binWidth', 1)
data = [];
nets = {'unassigned' 'SMd' 'SML' 'CO' 'Aud' 'DMN' 'PMN' 'Vis' 'FP' 'Sal' 'Lang' 'DAN'
'MTL' 'PON'};
for sub = 1:numel(subs)
load(['/Users/dianaperez/Desktop/Research/Segregation_Analyses/' subs{sub}
'_seg_index_net_withNaNegs.mat'])
sub_struct.seg_ind(15,:) = [];
if contains(subs{sub}, 'INET')
sub_struct.seg_ind(:,5) = -1
end
data = [data sub_struct.seg_ind];
end
data = data';
cx_ind(1:50) = 1;
cx_ind(51:90) = 2;
cx_ind = repmat(cx_ind, 14,1);
%ind = repmat([1:14]',18,1);
handles = plotSpread(data, 'categoryIdx', cx_ind, 'categoryMarkers', {'o','o'}, 'xNames',
nets, 'categoryColors', {'b', 'r'}, 'binWidth', 1)
handles = plotSpread(data, 'categoryIdx', cx_ind, 'categoryMarkers', {'.', '.'}, 'xNames',
nets, 'categoryColors', {'b', 'r'}, 'binWidth', 1)
close(gcf)
data = [];
nets = {'unassigned' 'SMd' 'SML' 'CO' 'Aud' 'DMN' 'PMN' 'Vis' 'FP' 'Sal' 'Lang' 'DAN'
'MTL' 'PON'};
for sub = 1:numel(subs)
load(['/Users/dianaperez/Desktop/Research/Segregation_Analyses/' subs{sub}
'_seg_index_net_withNaNegs.mat'])
sub_struct.seg_ind(15,:) = [];
if contains(subs{sub}, 'INET')
sub_struct.seg_ind(:,5) = -1
end
data = [data sub_struct.seg_ind];
end
data = data';
cx_ind(1:50) = 1;
```

```
cx_ind(51:90) = 2;
cx_ind = repmat(cx_ind, 14,1);
%ind = repmat([1:14]',18,1);
handles = plotSpread(data, 'categoryIdx', cx_ind, 'categoryMarkers', {'x','x'}, 'xNames', ↵
nets, 'categoryColors', {'b', 'r'}, 'binWidth', 1)
hold on
% 0
data = [];
for sub = 1:numel(subs)
load([' /Users/dianaperez/Desktop/Research/Segregation_Analyses/' subs{sub} ↵
'_seg_index_net_with0negs.mat'])
sub_struct.seg_ind(15,:) = [];
if contains(subs{sub}, 'INET')
sub_struct.seg_ind(:,5) = -1
end
data = [data sub_struct.seg_ind];
end
data = data';
handles = plotSpread(data, 'categoryIdx', cx_ind, 'categoryMarkers', {'.', '.'}, 'xNames', ↵
nets, 'categoryColors', {'b', 'r'}, 'binWidth', 1)
clear cx_ind
data = [];
nets = {'unassigned' 'SMd' 'SML' 'CO' 'Aud' 'DMN' 'PMN' 'Vis' 'FP' 'Sal' 'Lang' 'DAN' ↵
'MTL' 'PON'};
for sub = 1:numel(subs)
load([' /Users/dianaperez/Desktop/Research/Segregation_Analyses/' subs{sub} ↵
'_seg_index_net_withNaNegs.mat'])
sub_struct.seg_ind(15,:) = [];
if contains(subs{sub}, 'INET')
sub_struct.seg_ind(:,5) = -1
end
data = [data sub_struct.seg_ind];
end
data = data';
cx_ind(1:50) = 1;
cx_ind(51:90) = 2;
cx_ind = repmat(cx_ind, 14,1);
%ind = repmat([1:14]',18,1);
handles = plotSpread(data, 'categoryIdx', cx_ind, 'categoryMarkers', {'x','x'}, 'xNames', ↵
nets, 'categoryColors', {'b', 'r'}, 'binWidth', 1)
hold on
% 0
data = [];
for sub = 1:numel(subs)
load([' /Users/dianaperez/Desktop/Research/Segregation_Analyses/' subs{sub} ↵
'_seg_index_net_with0negs.mat'])
sub_struct.seg_ind(15,:) = [];
if contains(subs{sub}, 'INET')
sub_struct.seg_ind(:,5) = -1
end
data = [data sub_struct.seg_ind];
end
data = data';
handles = plotSpread(data, 'categoryIdx', cx_ind, 'categoryMarkers', {'.', '.'}, 'xNames', ↵
nets, 'categoryColors', {'b', 'r'}, 'binWidth', 1)
ax = gca;
ax.FontSize = 24;
axis([0.5, 15.5, .4, .9])
axis([0.5, 15.5, .3, .9])
axis([0.5, 15.5, .2, .9])
```

```
axis([0.5, 15.5, 0, .9])
axis([0.5, 15, 0, .9])
axis([0, 15, 0, .9])
axis([0, 15, -0.2, .9])
axis([0, 15, -0.3, .9])
data = [];
nets = {'unassign' 'SMd' 'SML' 'CO' 'Aud' 'DMN' 'PMN' 'Vis' 'FP' 'Sal' 'Lang' 'DAN'
'MTL' 'PON'};
for sub = 1:numel(subs)
load(['\Users\dianaperez\Desktop\Research\Segregation_Analyses\' subs{sub}
'_seg_index_net_withNaNegs.mat'])
sub_struct.seg_ind(15,:) = [];
if contains(subs{sub}, 'INET')
sub_struct.seg_ind(:,5) = -1
end
data = [data sub_struct.seg_ind];
end
data = data';
cx_ind(1:50) = 1;
cx_ind(51:90) = 2;
cx_ind = repmat(cx_ind, 14,1);
%ind = repmat([1:14]',18,1);
handles = plotSpread(data, 'categoryIdx', cx_ind, 'categoryMarkers', {'x','x'}, 'xNames',
nets, 'categoryColors', {'b', 'r'}, 'binWidth', 1)
hold on
% 0
data = [];
for sub = 1:numel(subs)
load(['\Users\dianaperez\Desktop\Research\Segregation_Analyses\' subs{sub}
'_seg_index_net_with0negs.mat'])
sub_struct.seg_ind(15,:) = [];
if contains(subs{sub}, 'INET')
sub_struct.seg_ind(:,5) = -1
end
data = [data sub_struct.seg_ind];
end
data = data';
handles = plotSpread(data, 'categoryIdx', cx_ind, 'categoryMarkers', {'.', '.'}, 'xNames',
nets, 'categoryColors', {'b', 'r'}, 'binWidth', 1)
ax = gca;
ax.FontSize = 24;
axis([0.5, 15.5, .4, .9])
data = [];
nets = {'unassign' 'SMd' 'SML' 'CO' 'Aud' 'DMN' 'PMN' 'Vis' 'FP' 'Sal' 'Lang' 'DAN'
'MTL' 'PON'};
for sub = 1:numel(subs)
load(['\Users\dianaperez\Desktop\Research\Segregation_Analyses\' subs{sub}
'_seg_index_net_withNaNegs.mat'])
sub_struct.seg_ind(15,:) = [];
if contains(subs{sub}, 'INET')
sub_struct.seg_ind(:,5) = -1
end
data = [data sub_struct.seg_ind];
end
data = data';
cx_ind = [];
cx_ind(1:50) = 1;
cx_ind(51:90) = 2;
cx_ind = repmat(cx_ind, 14,1);
%ind = repmat([1:14]',18,1);
```

```
handles = plotSpread(data, 'categoryIdx', cx_ind, 'categoryMarkers', {'x','x'}, 'xNames', ↵
nets, 'categoryColors', {'b', 'r'}, 'binWidth', 1)
hold on
% 0
data = [];
for sub = 1:numel(subs)
load([' /Users/dianaperez/Desktop/Research/Segregation_Analyses/' subs{sub} ↵
'_seg_index_net_with0negs.mat'])
sub_struct.seg_ind(15,:) = [];
if contains(subs{sub}, 'INET')
sub_struct.seg_ind(:,5) = -1
end
data = [data sub_struct.seg_ind];
end
data = data';
handles = plotSpread(data, 'categoryIdx', cx_ind, 'categoryMarkers', {'.', '.'}, 'xNames', ↵
nets, 'categoryColors', {'b', 'r'}, 'binWidth', 1)
ax = gca;
ax.FontSize = 24;
axis([0, 15, -0.3, .9])
print(gcf, '/Users/dianaperez/Desktop/SI_by_net_allsess_NanVs0.jpg', '-dpng', '-r300')
print(gcf, '/Users/dianaperez/Desktop/SI_by_net_allsess_NanVs0.jpg', '-dpng', '-r300')
help mean
data = [];
avg = [];
nets = {'unassign' 'SMd' 'SML' 'CO' 'Aud' 'DMN' 'PMN' 'Vis' 'FP' 'Sal' 'Lang' 'DAN' ↵
'MTL' 'PON'};
for sub = 1:numel(subs)
load([' /Users/dianaperez/Desktop/Research/Segregation_Analyses/' subs{sub} ↵
'_seg_index_net_withNaNegs.mat'])
sub_struct.seg_ind(15,:) = [];
if contains(subs{sub}, 'INET')
sub_struct.seg_ind(:,5) = -1
end
avg = [avg; mean(sub_struct.seg_ind, 1)];
data = [data sub_struct.seg_ind];
end
data = [];
avg = [];
nets = {'unassign' 'SMd' 'SML' 'CO' 'Aud' 'DMN' 'PMN' 'Vis' 'FP' 'Sal' 'Lang' 'DAN' ↵
'MTL' 'PON'};
for sub = 1:numel(subs)
load([' /Users/dianaperez/Desktop/Research/Segregation_Analyses/' subs{sub} ↵
'_seg_index_net_withNaNegs.mat'])
sub_struct.seg_ind(15,:) = [];
avg = [avg; mean(sub_struct.seg_ind, 1)];
if contains(subs{sub}, 'INET')
sub_struct.seg_ind(:,5) = -1
end
data = [data sub_struct.seg_ind];
end
mean(sub_struct.seg_ind, 1)
avg = [avg; mean(sub_struct.seg_ind, 2)];
mean(sub_struct.seg_ind, 2)
data = [];
avg = [];
nets = {'unassign' 'SMd' 'SML' 'CO' 'Aud' 'DMN' 'PMN' 'Vis' 'FP' 'Sal' 'Lang' 'DAN' ↵
'MTL' 'PON'};
for sub = 1:numel(subs)
load([' /Users/dianaperez/Desktop/Research/Segregation_Analyses/' subs{sub} ↵
```

```

'_seg_index_net_withNaNegs.mat'])
sub_struct.seg_ind(15,:) = [];
avg = [avg mean(sub_struct.seg_ind, 2)];
if contains(subs{sub}, 'INET')
sub_struct.seg_ind(:,5) = -1
end
data = [data sub_struct.seg_ind];
end
data = data';
cx_ind = [1,1,1,1,1,1,1,1,1,1,2,2,2,2,2,2,2,2];
cx_ind = repmat(cx_ind, 14,1);
cx_ind = repmat(cx_ind, 1,14);
cx_ind = [1,1,1,1,1,1,1,1,1,1,2,2,2,2,2,2,2,2];
cx_ind = repmat(cx_ind, 1,14);
cx_ind = [1,1,1,1,1,1,1,1,1,1,2,2,2,2,2,2,2,2];
cx_ind = repmat(cx_ind, 1,14);
cx_ind = [1,1,1,1,1,1,1,1,1,1,2,2,2,2,2,2,2,2];
cx_ind = repmat(cx_ind, 14,1);
data = [];
avg = [];
nets = {'unassign' 'SMd' 'SML' 'CO' 'Aud' 'DMN' 'PMN' 'Vis' 'FP' 'Sal' 'Lang' 'DAN'
'MTL' 'PON'};
for sub = 1:numel(subs)
load(['/Users/dianaperez/Desktop/Research/Segregation_Analyses/' subs{sub}
'_seg_index_net_withNaNegs.mat'])
sub_struct.seg_ind(15,:) = [];
avg = [avg mean(sub_struct.seg_ind, 2)];
if contains(subs{sub}, 'INET')
sub_struct.seg_ind(:,5) = -1
end
data = [data sub_struct.seg_ind];
end
data = data';
avg_nan = avg';
cx_ind = [];
cx_ind(1:50) = 1;
cx_ind(51:90) = 2;
cx_ind = repmat(cx_ind, 14,1);
%ind = repmat([1:14]',18,1);
handles = plotSpread(data, 'categoryIdx', cx_ind, 'categoryMarkers', {'x','x'}, 'xNames',
nets, 'categoryColors', {'b', 'r'}, 'binWidth', 1)
hold on
% 0
data = [];
avg = [];
for sub = 1:numel(subs)
load(['/Users/dianaperez/Desktop/Research/Segregation_Analyses/' subs{sub}
'_seg_index_net_with0negs.mat'])
sub_struct.seg_ind(15,:) = [];
avg = [avg mean(sub_struct.seg_ind, 2)];
if contains(subs{sub}, 'INET')
sub_struct.seg_ind(:,5) = -1
end
data = [data sub_struct.seg_ind];
end
avg_zero = avg';
data = data';
handles = plotSpread(data, 'categoryIdx', cx_ind, 'categoryMarkers', {'.', '.'}, 'xNames',
nets, 'categoryColors', {'b', 'r'}, 'binWidth', 1)
ax = gca;

```

```

ax.FontSize = 24;
axis([0, 15, -0.3, .9])
%print(gcf, '/Users/dianaperez/Desktop/SI_by_net_allsess_NanVs0.jpg', '-dpng', '-r300')
close(gcf)
cx_ind = [1,1,1,1,1,1,1,1,1,1,1,2,2,2,2,2,2,2,2];
cx_ind = repmat(cx_ind, 14,1);
handles = plotSpread(avg_nan, 'categoryIdx', cx_ind, 'categoryMarkers', 'x', 'x', 'xNames', nets, 'categoryColors', {'b', 'r'}, 'binWidth', 1)
hold on
handles = plotSpread(avg_zero, 'categoryIdx', cx_ind, 'categoryMarkers', '.', '.', 'xNames', nets, 'categoryColors', {'b', 'r'}, 'binWidth', 1)
cx_ind = [1,1,1,1,1,1,1,1,1,1,1,2,2,2,2,2,2,2,2];
cx_ind = [1,1,1,1,1,1,1,1,1,1,1,2,2,2,2,2,2,2,2]';
cx_ind = repmat(cx_ind, 14,1);
cx_ind = [1,1,1,1,1,1,1,1,1,1,1,2,2,2,2,2,2,2,2]';
cx_ind = repmat(cx_ind, 1,14);
close(gcf)
cx_ind = [1,1,1,1,1,1,1,1,1,1,1,2,2,2,2,2,2,2,2]';
cx_ind = repmat(cx_ind, 1,14);
handles = plotSpread(avg_nan, 'categoryIdx', cx_ind, 'categoryMarkers', 'x', 'x', 'xNames', nets, 'categoryColors', {'b', 'r'}, 'binWidth', 1)
hold on
handles = plotSpread(avg_zero, 'categoryIdx', cx_ind, 'categoryMarkers', '.', '.', 'xNames', nets, 'categoryColors', {'b', 'r'}, 'binWidth', 1)
cx_ind = [1,1,1,1,1,1,1,1,1,1,1,2,2,2,2,2,2,2,2]';
cx_ind = repmat(cx_ind, 1,14);
handles = plotSpread(avg_nan, 'categoryIdx', cx_ind, 'categoryMarkers', 'x', 'x', 'xNames', nets, 'categoryColors', {'b', 'r'}, 'binWidth', 1)
hold on
handles = plotSpread(avg_zero, 'categoryIdx', cx_ind, 'categoryMarkers', 'o', 'o', 'xNames', nets, 'categoryColors', {'b', 'r'}, 'binWidth', 1)
ax = gca;
ax.FontSize = 24;
axis([0, 15, -0.3, .9])
axis([0, 15, -0.1, .9])
print(gcf, '/Users/dianaperez/Desktop/SI_by_net_NanVs0.jpg', '-dpng', '-r300')
var_freq_size_net
perms_var_freq_size
load
('/Users/dianaperez/Desktop/Research/Lateralization/lateralization_code/testing_output/HCP384_variants_info.mat')
clear all
load
('/Users/dianaperez/Desktop/Research/Lateralization/lateralization_code/testing_output/HCP384_variants_info.mat')
lh_numvars = variants_info.left_hem.group_avg(:,3);
rh_numvars = variants_info.right_hem.group_avg(:,3);
scatter(lh_numvars, rh_numvars, 80, rgb_colors, 'filled')
hold on
scatter(lh_numvars, rh_numvars, 80, 'k')
axis([0 2.5 0 2.5])
line = reline(1,0);
line.Color = 'black';
xlabel('Left Hemisphere')
ax = gca;
ax.FontSize = 24;
for net = 1:16
se_lh(net) = std(networksxHem.clustersLH(:,net))/sqrt(384);
se_rh(net) = std(networksxHem.clustersRH(:,net))/sqrt(384);
end

```



```

h = errorbar(lh_numvars, rh_numvars, se_rh, se_rh, se_lh, se_lh, 'o')
h.Color = [0,0,0];
rgb_colors = [1 0 0; %DMN
0 0 .6; %Vis
1 1 0; %FP
.67 .67 .67; %Unassigned
0 .8 0; %DAN
.67 .67 .67; %Unassigned2
0 .6 .6; %VAN
0 0 0; % Sal
.3 0 .6; %CON
.2 1 1; %SMd
1 .5 0; % SMI
.6 .2 1; %Aud
.2 1 .2; %Tpole
0 .2 .4; %MTL
0 0 1; %PMN
.8 .8 .6]; %PON
scatter(lh_numvars, rh_numvars, 80, rgb_colors, 'filled')
hold on
scatter(lh_numvars, rh_numvars, 80, 'k')
axis([0 2.5 0 2.5])
line = refline(1,0);
line.Color = 'black';
xlabel('Left Hemisphere')
ax = gca;
ax.FontSize = 24;
for net = 1:16
se_lh(net) = std(networksxHem.clustersLH(:,net))/sqrt(384);
se_rh(net) = std(networksxHem.clustersRH(:,net))/sqrt(384);
end
h = errorbar(lh_numvars, rh_numvars, se_rh, se_rh, se_lh, se_lh, 'o')
h.Color = [0,0,0];
avgVarSize = variants_info.left_hem.group_avg(:,3);
lh_numvars = variants_info.left_hem.group_avg(:,3);
rh_numvars = variants_info.right_hem.group_avg(:,3);
scatter(lh_numvars, rh_numvars, 80, rgb_colors, 'filled')
hold on
scatter(lh_numvars, rh_numvars, 80, 'k')
axis([0 2.5 0 2.5])
line = refline(1,0);
line.Color = 'black';
xlabel('Left Hemisphere')
ax = gca;
ax.FontSize = 24;
for net = 1:16
se_lh(net) = std(networksxHem.clustersLH(:,net))/sqrt(384);
se_rh(net) = std(networksxHem.clustersRH(:,net))/sqrt(384);
end
h = errorbar(lh_numvars, rh_numvars, se_rh, se_rh, se_lh, se_lh, 'o')
h.Color = [0,0,0];
load
(' /Users/dianaperez/Desktop/Research/Lateralization/lateralization_code/testing_output/HCP
P384_networksxHem.mat')
var_freq_size_net
[data_location num2str(subs{g,1}(n))
netAssign_str]
[data_location num2str(subs{g,1}(n)) netAssign_str]
var_freq_size_net
load

```

```

('/Users/dianaperez/Desktop/Research/Lateralization/lateralization_code/testing_output/HCP384_networksxHem.mat')
clear all
load
('/Users/dianaperez/Desktop/Research/Lateralization/lateralization_code/testing_output/HCP384_networksxHem.mat')
lh_numvars = [];
rh_numvars = [];
for net = 1:16
vars_LH = []; vars_RH = [];
vars_LH = networksxHem.avgSizeLH(find(networksxHem.avgSizeLH{:,net}>0), net);
lh_numvars(net) = mean(vars_LH);
vars_RH = networksxHem.avgSizeRH(find(networksxHem.avgSizeRH{:,net}>0), net);
rh_numvars(net) = mean(vars_RH);
end
lh_numvars = [];
rh_numvars = [];
for net = 1:16
vars_LH = []; vars_RH = [];
vars_LH = networksxHem.avgSizeLH(find(networksxHem.avgSizeLH(:,net)>0), net);
lh_numvars(net) = mean(vars_LH);
vars_RH = networksxHem.avgSizeRH(find(networksxHem.avgSizeRH(:,net)>0), net);
rh_numvars(net) = mean(vars_RH);
end
lh_numvars = [];
rh_numvars = [];
for net = 1:16
vars_LH = []; vars_RH = [];
vars_LH = networksxHem.avgSizeLH(find(networksxHem.avgSizeLH(:,net)>0), net);
if length(vars_LH)>0
lh_numvars(net) = mean(vars_LH);
else
lh_numvars(net) = 0;
end
vars_RH = networksxHem.avgSizeRH(find(networksxHem.avgSizeRH(:,net)>0), net);
if length(vars_RH)>0
rh_numvars(net) = mean(vars_RH);
else
rh_numvars(net) = 0;
end
end
scatter(lh_numvars, rh_numvars, 80, rgb_colors, 'filled')
hold on
scatter(lh_numvars, rh_numvars, 80, 'k')
rgb_colors = [1 0 0; %DMN
0 0 .6; %Vis
1 1 0; %FP
.67 .67 .67; %Unassigned
0 .8 0; %DAN
.67 .67 .67; %Unassigned2
0 .6 .6; %VAN
0 0 0; % Sal
.3 0 .6; %CON
.2 1 1; %SMd
1 .5 0; % SMI
.6 .2 1; %Aud
.2 1 .2; %Tpole
0 .2 .4; %MTL
0 0 1; %PMN
.8 .8 .6]; %PON

```

```
scatter(lh_numvars, rh_numvars, 80, rgb_colors, 'filled')
hold on
scatter(lh_numvars, rh_numvars, 80, 'k')
axis([0 2.5 0 2.5])
line = reffline(1,0);
line.Color = 'black';
xlabel('Left Hemisphere')
ax = gca;
ax.FontSize = 24;
scatter(lh_numvars, rh_numvars, 80, rgb_colors, 'filled')
hold on
scatter(lh_numvars, rh_numvars, 80, 'k')
axis([150 700 150 700])
line = reffline(1,0);
line.Color = 'black';
xlabel('Left Hemisphere')
ax = gca;
ax.FontSize = 24;
ylabel('Right Hemisphere')
for net = 1:16
se_lh(net) = std(networksxHem.clustersLH(:,net))/sqrt(384);
se_rh(net) = std(networksxHem.clustersRH(:,net))/sqrt(384);
end
h = errorbar(lh_numvars, rh_numvars, se_rh, se_rh, se_lh, se_lh, 'o')
h.Color = [0,0,0];
h = errorbar(lh_numvars, rh_numvars, se_rh, se_rh, se_lh, se_lh, 'o', 'MarkerSize', 80)
h = errorbar(lh_numvars, rh_numvars, se_rh, se_rh, se_lh, se_lh, 'o', 'MarkerSize', 50)
h = errorbar(lh_numvars, rh_numvars, se_rh, se_rh, se_lh, se_lh, 'MarkerSize', 50)
h = errorbar(lh_numvars, rh_numvars, se_rh, se_rh, se_lh, se_lh, 'o', 'MarkerSize', 20)
h = errorbar(lh_numvars, rh_numvars, se_rh, se_rh, se_lh, se_lh, 'o', 'MarkerSize', 10)
scatter(lh_numvars, rh_numvars, 80, rgb_colors, 'filled')
hold on
scatter(lh_numvars, rh_numvars, 80, 'k')
axis([150 700 150 700])
line = reffline(1,0);
line.Color = 'black';
xlabel('Left Hemisphere')
ylabel('Right Hemisphere')
ax = gca;
ax.FontSize = 24;
h = errorbar(lh_numvars, rh_numvars, se_rh, se_rh, se_lh, se_lh, 'o', 'MarkerSize', 10)
h = errorbar(lh_numvars, rh_numvars, se_rh, se_rh, se_lh, se_lh, 'o', 'MarkerSize', 15)
scatter(lh_numvars, rh_numvars, 80, rgb_colors, 'filled')
hold on
scatter(lh_numvars, rh_numvars, 80, 'k')
%axis([0 2.5 0 2.5])
axis([150 700 150 700])
line = reffline(1,0);
line.Color = 'black';
xlabel('Left Hemisphere')
ylabel('Right Hemisphere')
ax = gca;
ax.FontSize = 24;
vars_RH = networksxHem.avgSizeRH(find(networksxHem.avgSizeRH(:,14)>0), 14);
h = errorbar(lh_numvars, rh_numvars, se_rh, se_rh, se_lh, se_lh, 'o', 'MarkerSize', 10)
h.Color = [0,0,0];
print(gcf, '
/Users/dianaperez/Desktop/Research/Lateralization/lateralization_code/HCP384_varsize_sca
tterplot.jpg', '-dpng', '-r300')
help median
```

```

lh_numvars = [];
rh_numvars = [];
for net = 1:16
vars_LH = []; vars_RH = [];
vars_LH = networksxHem.avgSizeLH(find(networksxHem.avgSizeLH(:,net)>0), net);
if length(vars_LH)>0
lh_numvars(net) = median(vars_LH);
else
lh_numvars(net) = 0;
end
vars_RH = networksxHem.avgSizeRH(find(networksxHem.avgSizeRH(:,net)>0), net);
if length(vars_RH)>0
rh_numvars(net) = median(vars_RH);
else
rh_numvars(net) = 0;
end
end
scatter(lh_numvars, rh_numvars, 80, rgb_colors, 'filled')
hold on
scatter(lh_numvars, rh_numvars, 80, 'k')
%axis([0 2.5 0 2.5])
axis([150 700 150 700])
line = reffline(1,0);
line.Color = 'black';
xlabel('Left Hemisphere')
ylabel('Right Hemisphere')
ax = gca;
ax.FontSize = 24;
for net = 1:16
se_lh(net) = std(networksxHem.clustersLH(:,net))/sqrt(384);
se_rh(net) = std(networksxHem.clustersRH(:,net))/sqrt(384);
end
h = errorbar(lh_numvars, rh_numvars, se_rh, se_rh, se_lh, se_lh, 'o', 'MarkerSize', 10)
h.Color = [0,0,0];
rgb_colors = [1 0 0; %DMN
0 0 .6; %Vis
1 1 0; %FP
.67 .67 .67; %Unassigned
0 .8 0; %DAN
.67 .67 .67; %Unassigned2
0 .6 .6; %VAN
0 0 0; % Sal
.3 0 .6; %CON
.2 1 1; %SMd
1 .5 0; % SMI
.6 .2 1; %Aud
0 .2 .4; %Tpole
.2 1 .2; %MTL
0 0 1; %PMN
.8 .8 .6]; %PON
lh_numvars = [];
rh_numvars = [];
for net = 1:16
vars_LH = []; vars_RH = [];
vars_LH = networksxHem.avgSizeLH(find(networksxHem.avgSizeLH(:,net)>0), net);
if length(vars_LH)>0
lh_numvars(net) = median(vars_LH);
else
lh_numvars(net) = 0;
end
end

```

```
vars_RH = networksxHem.avgSizeRH(find(networksxHem.avgSizeRH(:,net)>0), net);
if length(vars_RH)>0
rh_numvars(net) = median(vars_RH);
else
rh_numvars(net) = 0;
end
end
scatter(lh_numvars, rh_numvars, 80, rgb_colors, 'filled')
hold on
scatter(lh_numvars, rh_numvars, 80, 'k')
%axis([0 2.5 0 2.5])
axis([150 700 150 700])
line = reffline(1,0);
line.Color = 'black';
xlabel('Left Hemisphere')
ylabel('Right Hemisphere')
ax = gca;
ax.FontSize = 24;
for net = 1:16
se_lh(net) = std(networksxHem.clustersLH(:,net))/sqrt(384);
se_rh(net) = std(networksxHem.clustersRH(:,net))/sqrt(384);
end
h = errorbar(lh_numvars, rh_numvars, se_rh, se_rh, se_lh, se_lh, 'o', 'MarkerSize', 10)
h.Color = [0,0,0];
clear all
load
(' /Users/dianaperez/Desktop/Research/Lateralization/lateralization_code/testing_output/HCP384_networksxHem.mat')
lh_numvars = [];
rh_numvars = [];
for net = 1:16
vars_LH = []; vars_RH = [];
vars_LH = networksxHem.avgSizeLH(find(networksxHem.avgSizeLH(:,net)>0), net);
if length(vars_LH)>0
lh_numvars(net) = median(vars_LH);
else
lh_numvars(net) = 0;
end
vars_RH = networksxHem.avgSizeRH(find(networksxHem.avgSizeRH(:,net)>0), net);
if length(vars_RH)>0
rh_numvars(net) = median(vars_RH);
else
rh_numvars(net) = 0;
end
end
scatter(lh_numvars, rh_numvars, 80, rgb_colors, 'filled')
hold on
scatter(lh_numvars, rh_numvars, 80, 'k')
%axis([0 2.5 0 2.5])
axis([150 700 150 700])
line = reffline(1,0);
line.Color = 'black';
xlabel('Left Hemisphere')
ylabel('Right Hemisphere')
ax = gca;
ax.FontSize = 24;
for net = 1:16
se_lh(net) = std(networksxHem.clustersLH(:,net))/sqrt(384);
se_rh(net) = std(networksxHem.clustersRH(:,net))/sqrt(384);
end
```

```

h = errorbar(lh_numvars, rh_numvars, se_rh, se_rh, se_lh, se_lh, 'o', 'MarkerSize', 10)
h.Color = [0,0,0];
rgb_colors = [1 0 0; %DMN
0 0 .6; %Vis
1 1 0; %FP
.67 .67 .67; %Unassigned
0 .8 0; %DAN
.67 .67 .67; %Unassigned2
0 .6 .6; %VAN
0 0 0; % Sal
.3 0 .6; %CON
.2 1 1; %SMd
1 .5 0; % SMI
.6 .2 1; %Aud
0 .2 .4; %Tpole
.2 1 .2; %MTL
0 0 1; %PMN
.8 .8 .6]; %PON
lh_numvars = [];
rh_numvars = [];
for net = 1:16
vars_LH = []; vars_RH = [];
vars_LH = networksxHem.avgSizeLH(find(networksxHem.avgSizeLH(:,net)>0), net);
if length(vars_LH)>0
lh_numvars(net) = median(vars_LH);
else
lh_numvars(net) = 0;
end
vars_RH = networksxHem.avgSizeRH(find(networksxHem.avgSizeRH(:,net)>0), net);
if length(vars_RH)>0
rh_numvars(net) = median(vars_RH);
else
rh_numvars(net) = 0;
end
end
scatter(lh_numvars, rh_numvars, 80, rgb_colors, 'filled')
hold on
scatter(lh_numvars, rh_numvars, 80, 'k')
%axis([0 2.5 0 2.5])
axis([150 700 150 700])
line = reffline(1,0);
line.Color = 'black';
xlabel('Left Hemisphere')
ylabel('Right Hemisphere')
ax = gca;
ax.FontSize = 24;
for net = 1:16
se_lh(net) = std(networksxHem.clustersLH(:,net))/sqrt(384);
se_rh(net) = std(networksxHem.clustersRH(:,net))/sqrt(384);
end
h = errorbar(lh_numvars, rh_numvars, se_rh, se_rh, se_lh, se_lh, 'o', 'MarkerSize', 10)
h.Color = [0,0,0];
scatter(lh_numvars, rh_numvars, 80, rgb_colors, 'filled')
hold on
scatter(lh_numvars, rh_numvars, 80, 'k')
for net = 1:16
se_lh(net) = std(networksxHem.avgSizeLH(:,net))/sqrt(384);
se_rh(net) = std(networksxHem.avgSizeRH(:,net))/sqrt(384);
end
h = errorbar(lh_numvars, rh_numvars, se_rh, se_rh, se_lh, se_lh, 'o', 'MarkerSize', 10)

```

```

h.Color = [0,0,0];
scatter(lh_numvars, rh_numvars, 80, rgb_colors, 'filled')
hold on
scatter(lh_numvars, rh_numvars, 80, 'k')
print(gcf,
'/Users/dianaperez/Desktop/Research/Lateralization/lateralization_code/HCP384_varsize_sca
tterplot.jpg', '-dpng', '-r300')
workbenchdir = '/Applications/workbench/bin_macosx64/';
varmap_path = '/Users/dianaperez/Desktop/MS_C_variants/MS_C01_variants_reassigned.dtseries.
nii'
boder_labels = '/Users/dianaperez/Downloads/wbBorderLabelFile.txt';
output_path = '/Users/dianaperez/Desktop/MS_C_variants/MS_C01_variant_borders.dlabel.nii';
system([workbenchdir 'wb_command -cifti-label-import ' varmap_path border_labels
output_path])
system([workbenchdir 'wb_command -cifti-label-import ' varmap_path boder_labels
output_path])
system([workbenchdir 'wb_command -cifti-label-import ' varmap_path ' ' border_labels ' '
output_path])
system([workbenchdir 'wb_command -cifti-label-import ' varmap_path ' ' boder_labels ' '
output_path])
output_path_left = '/Users/dianaperez/Desktop/MS_C_variants/MS_C01_variant_borders_left.
dlabel.gii';
output_path_right = '/Users/dianaperez/Desktop/MS_C_variants/MS_C01_variant_borders_right.
dlabel.gii';
system([workbenchdir 'wb_command -cifti-separate ' output_path ' COLUMN -label
CORTEX_LEFT ' output_path_left ' -label CORTEX_RIGHT ' output_path_right])
output_path_left = '/Users/dianaperez/Desktop/MS_C_variants/MS_C01_variant_borders_left.
label.gii';
output_path_right = '/Users/dianaperez/Desktop/MS_C_variants/MS_C01_variant_borders_right.
label.gii';
system([workbenchdir 'wb_command -cifti-separate ' output_path ' COLUMN -label
CORTEX_LEFT ' output_path_left ' -label CORTEX_RIGHT ' output_path_right])
midthick_path =
'/Users/dianaperez/Desktop/Organization/CNS_analyses/Dependencies/32k_ConteAtlas_v2_distr
ibute/Conte69_atlas-v2.LR.32k_fs_LR.wb/Conte69.L.midthickness.32k_fs_LR.surf.gii';
output_border_L = '/Users/dianaperez/Desktop/MS_C_variants/MS_C01_variants_L.border';
system([workbenchdir 'wb_command -label-to-border ' midthick_path ' ' output_path_left '
' output_border_L])
midthick_path =
'/Users/dianaperez/Desktop/Organization/CNS_analyses/Dependencies/32k_ConteAtlas_v2_distr
ibute/Conte69_atlas-v2.LR.32k_fs_LR.wb/Conte69.R.midthickness.32k_fs_LR.surf.gii';
output_border_R = '/Users/dianaperez/Desktop/MS_C_variants/MS_C01_variants_R.border';
system([workbenchdir 'wb_command -label-to-border ' midthick_path ' ' output_path_right
' ' output_border_R])
midthick_path =
'/Users/dianaperez/Desktop/Organization/CNS_analyses/Dependencies/32k_ConteAtlas_v2_distr
ibute/Conte69.L.midthickness.32k_fs_LR.surf.gii';
output_border_L = '/Users/dianaperez/Desktop/MS_C_variants/MS_C01_variants_L.border';
system([workbenchdir 'wb_command -label-to-border ' midthick_path ' ' output_path_left '
' output_border_L])
midthick_path =
'/Users/dianaperez/Desktop/Organization/CNS_analyses/Dependencies/32k_ConteAtlas_v2_distr
ibute/Conte69.R.midthickness.32k_fs_LR.surf.gii';
output_border_R = '/Users/dianaperez/Desktop/MS_C_variants/MS_C01_variants_R.border';
system([workbenchdir 'wb_command -label-to-border ' midthick_path ' ' output_path_right
' ' output_border_R])
varmap_path = '/Users/dianaperez/Desktop/MS_C_variants/MS_C02_variants_reassigned.dtseries.
nii';
boder_labels = '/Users/dianaperez/Downloads/wbBorderLabelFile.txt';
output_path = '/Users/dianaperez/Desktop/MS_C_variants/MS_C02_variant_borders.dlabel.nii';

```

```

system([workbenchdir 'wb_command -cifti-label-import ' varmap_path ' ' border_labels ' '↵
output_path])
output_path_left = '/Users/dianaperez/Desktop/MS_C_variants/MS_C02_variant_borders_left.↵
label.gii';
output_path_right = '/Users/dianaperez/Desktop/MS_C_variants/MS_C02_variant_borders_right.↵
label.gii';
system([workbenchdir 'wb_command -cifti-separate ' output_path ' COLUMN -label↵
CORTEX_LEFT ' output_path_left ' -label CORTEX_RIGHT ' output_path_right])
midthick_path =↵
'/Users/dianaperez/Desktop/Organization/CNS_analyses/Dependencies/32k_ConteAtlas_v2_distr↵
ibute/Conte69.L.midthickness.32k_fs_LR.surf.gii';
output_border_L = '/Users/dianaperez/Desktop/MS_C_variants/MS_C02_variants_L.border';
system([workbenchdir 'wb_command -label-to-border ' midthick_path ' ' output_path_left '↵
' output_border_L])
midthick_path =↵
'/Users/dianaperez/Desktop/Organization/CNS_analyses/Dependencies/32k_ConteAtlas_v2_distr↵
ibute/Conte69.R.midthickness.32k_fs_LR.surf.gii';
output_border_R = '/Users/dianaperez/Desktop/MS_C_variants/MS_C02_variants_R.border';
system([workbenchdir 'wb_command -label-to-border ' midthick_path ' ' output_path_right↵
' ' output_border_R])
varmap_path = '/Users/dianaperez/Desktop/MS_C_variants/MS_C02_variants_reassigned.dtseries.↵
nii';
border_labels = '/Users/dianaperez/Downloads/wbBorderLabelFile.txt';
output_path = '/Users/dianaperez/Desktop/MS_C_variants/MS_C02_variant_borders.dlabel.nii';
system([workbenchdir 'wb_command -cifti-label-import ' varmap_path ' ' border_labels ' '↵
output_path])
output_path_left = '/Users/dianaperez/Desktop/MS_C_variants/MS_C02_variant_borders_left.↵
label.gii';
output_path_right = '/Users/dianaperez/Desktop/MS_C_variants/MS_C02_variant_borders_right.↵
label.gii';
system([workbenchdir 'wb_command -cifti-separate ' output_path ' COLUMN -label↵
CORTEX_LEFT ' output_path_left ' -label CORTEX_RIGHT ' output_path_right])
midthick_path =↵
'/Users/dianaperez/Desktop/Organization/CNS_analyses/Dependencies/32k_ConteAtlas_v2_distr↵
ibute/Conte69.L.midthickness.32k_fs_LR.surf.gii';
output_border_L = '/Users/dianaperez/Desktop/MS_C_variants/MS_C02_variants_L.border';
system([workbenchdir 'wb_command -label-to-border ' midthick_path ' ' output_path_left '↵
' output_border_L])
midthick_path =↵
'/Users/dianaperez/Desktop/Organization/CNS_analyses/Dependencies/32k_ConteAtlas_v2_distr↵
ibute/Conte69.R.midthickness.32k_fs_LR.surf.gii';
output_border_R = '/Users/dianaperez/Desktop/MS_C_variants/MS_C02_variants_R.border';
system([workbenchdir 'wb_command -label-to-border ' midthick_path ' ' output_path_right↵
' ' output_border_R])
clear all
load↵
('/Users/dianaperez/Desktop/Research/Lateralization/lateralization_code/testing_output/HC↵
P384_networksxHem.mat')
lh_numvars = [];
rh_numvars = [];
for net = 1:16
vars_LH = []; vars_RH = [];
vars_LH = networksxHem.avgSizeLH(find(networksxHem.surfaceSizeLH(:,net)>0), net);
if length(vars_LH)>0
lh_numvars(net) = median(vars_LH);
else
lh_numvars(net) = 0;
end
vars_RH = networksxHem.avgSizeRH(find(networksxHem.surfaceRH(:,net)>0), net);
if length(vars_RH)>0

```



```

rh_numvars(net) = median(vars_RH);
else
rh_numvars(net) = 0;
end
end
lh_numvars = [];
rh_numvars = [];
for net = 1:16
vars_LH = []; vars_RH = [];
vars_LH = networksxHem.avgSizeLH(find(networksxHem.surfaceLH(:,net)>0), net);
if length(vars_LH)>0
lh_numvars(net) = median(vars_LH);
else
lh_numvars(net) = 0;
end
vars_RH = networksxHem.avgSizeRH(find(networksxHem.surfaceRH(:,net)>0), net);
if length(vars_RH)>0
rh_numvars(net) = median(vars_RH);
else
rh_numvars(net) = 0;
end
end
scatter(lh_numvars, rh_numvars, 80, rgb_colors, 'filled')
hold on
scatter(lh_numvars, rh_numvars, 80, 'k')
rgb_colors = [1 0 0; %DMN
0 0 .6; %Vis
1 1 0; %FP
.67 .67 .67; %Unassigned
0 .8 0; %DAN
.67 .67 .67; %Unassigned2
0 .6 .6; %VAN
0 0 0; % Sal
.3 0 .6; %CON
.2 1 1; %SMd
1 .5 0; % SMI
.6 .2 1; %Aud
0 .2 .4; %Tpole
.2 1 .2; %MTL
0 0 1; %PMN
.8 .8 .6]; %PON
scatter(lh_numvars, rh_numvars, 80, rgb_colors, 'filled')
hold on
scatter(lh_numvars, rh_numvars, 80, 'k')
line = reffline(1,0);
line.Color = 'black';
xlabel('Left Hemisphere')
ylabel('Right Hemisphere')
ax = gca;
ax.FontSize = 24;
for net = 1:16
se_lh(net) = std(networksxHem.surfaceLH(:,net))/sqrt(384);
se_rh(net) = std(networksxHem.surfaceRH(:,net))/sqrt(384);
end
h = errorbar(lh_numvars, rh_numvars, se_rh, se_rh, se_lh, se_lh, 'o', 'MarkerSize', 10)
h.Color = [0,0,0];
axis([150 700 150 700])
line = reffline(1,0);
lh_numvars = [];
rh_numvars = [];

```

```

for net = 1:16
vars_LH = []; vars_RH = [];
vars_LH = networksxHem.surfaceLH(find(networksxHem.surfaceLH(:,net)>0), net);
if length(vars_LH)>0
lh_numvars(net) = median(vars_LH);
else
lh_numvars(net) = 0;
end
vars_RH = networksxHem.surfaceRH(find(networksxHem.surfaceRH(:,net)>0), net);
if length(vars_RH)>0
rh_numvars(net) = median(vars_RH);
else
rh_numvars(net) = 0;
end
end
scatter(lh_numvars, rh_numvars, 80, rgb_colors, 'filled')
hold on
scatter(lh_numvars, rh_numvars, 80, 'k')
lh_numvars = [];
rh_numvars = [];
for net = 1:16
vars_LH = []; vars_RH = [];
vars_LH = networksxHem.surfaceLH(find(networksxHem.surfaceLH(:,net)>0), net);
if length(vars_LH)>0
lh_numvars(net) = mean(vars_LH);
else
lh_numvars(net) = 0;
end
vars_RH = networksxHem.surfaceRH(find(networksxHem.surfaceRH(:,net)>0), net);
if length(vars_RH)>0
rh_numvars(net) = mean(vars_RH);
else
rh_numvars(net) = 0;
end
end
scatter(lh_numvars, rh_numvars, 80, rgb_colors, 'filled')
hold on
scatter(lh_numvars, rh_numvars, 80, 'k')
axis([200 700 200 700])
line = reffline(1,0);
line.Color = 'black';
xlabel('Left Hemisphere')
ylabel('Right Hemisphere')
ax = gca;
ax.FontSize = 24;
for net = 1:16
se_lh(net) = std(networksxHem.surfaceLH(:,net))/sqrt(384);
se_rh(net) = std(networksxHem.surfaceRH(:,net))/sqrt(384);
end
h = errorbar(lh_numvars, rh_numvars, se_rh, se_rh, se_lh, se_lh, 'o', 'MarkerSize', 10)
h.Color = [0,0,0];
print(gcf,
'/Users/dianaperez/Desktop/Research/lateralization_code/HCP384_variant_territory_scatterp
lot.jpg', '-dpng', '-r300')
lh_numvars = [];
rh_numvars = [];
for net = 1:16
vars_LH = []; vars_RH = [];
vars_LH = networksxHem.surfaceLH(find(networksxHem.surfaceLH(:,net)>0), net);
if length(vars_LH)>0

```

```

lh_numvars(net) = mean(vars_LH);
else
lh_numvars(net) = 0;
end
vars_RH = networksxHem.surfaceRH(find(networksxHem.surfaceRH(:,net)>0), net);
if length(vars_RH)>0
rh_numvars(net) = mean(vars_RH);
else
rh_numvars(net) = 0;
end
end
scatter(lh_numvars, rh_numvars, 80, rgb_colors, 'filled')
hold on
scatter(lh_numvars, rh_numvars, 80, 'k')
%axis([0 2.5 0 2.5])
axis([200 700 200 700])
line = refline(1,0);
line.Color = 'black';
xlabel('Left Hemisphere')
ylabel('Right Hemisphere')
ax = gca;
ax.FontSize = 24;
for net = 1:16
se_lh(net) = std(networksxHem.surfaceLH(:,net))/sqrt(384);
se_rh(net) = std(networksxHem.surfaceRH(:,net))/sqrt(384);
end
h = errorbar(lh_numvars, rh_numvars, se_rh, se_rh, se_lh, se_lh, 'o', 'MarkerSize', 10)
h.Color = [0,0,0];
print(gcf, '
/Users/dianaperez/Desktop/Research/lateralization_code/HCP384_variant_territory_scatterp
lot.jpg', '-dpng', '-r300')
%-- 6/20/22, 7:38 AM --%
rgb_colors = [1 0 0; %DMN
0 0 .6; %Vis
1 1 0; %FP
.67 .67 .67; %Unassigned
0 .8 0; %DAN
.67 .67 .67; %Unassigned2
0 .6 .6; %VAN
0 0 0; % Sal
.3 0 .6; %CON
.2 1 1; %SMd
1 .5 0; % SMI
.6 .2 1; %Aud
0 .2 .4; %Tpole
.2 1 .2; %MTL
0 0 1; %PMN
.8 .8 .6]; %PON
load
(' /Users/dianaperez/Desktop/Research/Lateralization/lateralization_code/testing_output/HCP
P384_networksxHem.mat')
lh_numvars = [];
rh_numvars = [];
for net = 1:16
vars_LH = []; vars_RH = [];
vars_LH = networksxHem.surfaceLH(find(networksxHem.surfaceLH(:,net)>0), net);
if length(vars_LH)>0
lh_numvars(net) = mean(vars_LH);
else
lh_numvars(net) = 0;

```

```

end
vars_RH = networksxHem.surfaceRH(find(networksxHem.surfaceRH(:,net)>0), net);
if length(vars_RH)>0
rh_numvars(net) = mean(vars_RH);
else
rh_numvars(net) = 0;
end
end
scatter(lh_numvars, rh_numvars, 80, rgb_colors, 'filled')
hold on
scatter(lh_numvars, rh_numvars, 80, 'k')
%axis([0 2.5 0 2.5])
axis([200 700 200 700])
line = reffline(1,0);
line.Color = 'black';
xlabel('Left Hemisphere')
ylabel('Right Hemisphere')
ax = gca;
ax.FontSize = 24;
for net = 1:16
se_lh(net) = std(networksxHem.surfaceLH(:,net))/sqrt(384);
se_rh(net) = std(networksxHem.surfaceRH(:,net))/sqrt(384);
end
h = errorbar(lh_numvars, rh_numvars, se_rh, se_rh, se_lh, se_lh, 'o', 'MarkerSize', 10)
h.Color = [0,0,0];
var_freq_size_net
addpath(genpath('/Users/dianaperez/Documents/Dependencies/'))
var_freq_size_net
addpath(genpath('/Users/dianaperez/Desktop/Organization/CNS_analyses/Dependencies/'))
var_freq_size_net
addpath
addpath help
help addpath
rmpath(genpath('/Users/dianaperez/Desktop/Organization/CNS_analyses/Dependencies/'))
addpath(genpath('/Users/dianaperez/Documents/Dependencies/'))
var_freq_size_net
rgb_colors = [1 0 0; %DMN
0 0 .6; %Vis
1 1 0; %FP
.67 .67 .67; %Unassigned
0 .8 0; %DAN
.67 .67 .67; %Unassigned2
0 .6 .6;%VAN
0 0 0; % Sal
.3 0 .6; %CON
.2 1 1; %SMd
1 .5 0; % SMI
.6 .2 1; %Aud
0 .2 .4; %Tpole
.2 1 .2; %MTL
0 0 1; %PMN
.8 .8 .6]; %PON
load
(' /Users/dianaperez/Desktop/Research/Lateralization/lateralization_code/testing_output/HCP384_networksxHem.mat')
lh_numvars = [];
rh_numvars = [];
for net = 1:16
vars_LH = []; vars_RH = [];
vars_LH = networksxHem.surfaceLH(find(networksxHem.surfaceLH(:,net)>0), net);

```

```
if length(vars_LH)>0
lh_numvars(net) = mean(vars_LH);
else
lh_numvars(net) = 0;
end
vars_RH = networksxHem.surfaceRH(find(networksxHem.surfaceRH(:,net)>0), net);
if length(vars_RH)>0
rh_numvars(net) = mean(vars_RH);
else
rh_numvars(net) = 0;
end
end
scatter(lh_numvars, rh_numvars, 80, rgb_colors, 'filled')
hold on
scatter(lh_numvars, rh_numvars, 80, 'k')
%axis([0 2.5 0 2.5])
axis([200 700 200 700])
line = refline(1,0);
line.Color = 'black';
xlabel('Left Hemisphere')
ylabel('Right Hemisphere')
ax = gca;
ax.FontSize = 24;
for net = 1:16
se_lh(net) = std(networksxHem.surfaceLH(:,net))/sqrt(384);
se_rh(net) = std(networksxHem.surfaceRH(:,net))/sqrt(384);
end
h = errorbar(lh_numvars, rh_numvars, se_rh, se_rh, se_lh, se_lh, 'o', 'MarkerSize', 10)
h.Color = [0,0,0];
print(gcf, '
/Users/dianaperez/Desktop/Research/lateralization_code/HCP384_variant_territory_scatterp
lot.jpg', '-dpng', '-r300')
vars_LH = networksxHem.surfaceLH(find(networksxHem.surfaceLH(:,14)>0), 14);
vars_RH = networksxHem.surfaceRH(find(networksxHem.surfaceRH(:,14)>0), 14);
vars_RH = networksxHem.surfaceRH(find(networksxHem.surfaceRH(:,14)>0), 14);
load
('Users/dianaperez/Desktop/Research/Lateralization/lateralization_code/testing_output/HCP
P384_networksxHem.mat')
lh_numvars = [];
rh_numvars = [];
for net = 1:16
vars_LH = []; vars_RH = [];
if length(vars_LH)>30 || length(vars_RH)>30
vars_LH = networksxHem.surfaceLH(find(networksxHem.surfaceLH(:,net)>0), net);
if length(vars_LH)>0
lh_numvars(net) = mean(vars_LH);
else
lh_numvars(net) = 0;
end
vars_RH = networksxHem.surfaceRH(find(networksxHem.surfaceRH(:,net)>0), net);
if length(vars_RH)>0
rh_numvars(net) = mean(vars_RH);
else
rh_numvars(net) = 0;
end
else
lh_numvars = 0;
rh_numvars = 0;
end
end
```

```

scatter(lh_numvars, rh_numvars, 80, rgb_colors, 'filled')
hold on
scatter(lh_numvars, rh_numvars, 80, 'k')
lh_numvars = [];
rh_numvars = [];
for net = 1:16
vars_LH = []; vars_RH = [];
if length(vars_LH)>30 || length(vars_RH)>30
vars_LH = networksxHem.surfaceLH(find(networksxHem.surfaceLH(:,net)>0), net);
if length(vars_LH)>0
lh_numvars(net) = mean(vars_LH);
else
lh_numvars(net) = 0;
end
vars_RH = networksxHem.surfaceRH(find(networksxHem.surfaceRH(:,net)>0), net);
if length(vars_RH)>0
rh_numvars(net) = mean(vars_RH);
else
rh_numvars(net) = 0;
end
else
lh_numvars(net) = 0;
rh_numvars(net) = 0;
end
end
scatter(lh_numvars, rh_numvars, 80, rgb_colors, 'filled')
hold on
scatter(lh_numvars, rh_numvars, 80, 'k')
var_territory_scatterplot
load
('/Users/dianaperez/Desktop/Research/Lateralization/lateralization_code/testing_output/HCP384_networksxHem.mat')
clear all
load
('/Users/dianaperez/Desktop/Research/Lateralization/lateralization_code/testing_output/HCP384_networksxHem.mat')
var_territory_scatterplot
close(gcf)
var_territory_scatterplot
clear all
var_freq_size_net
find(Lhem(:,1)==Lhem_var_IDs(v))
num_verts = length(verts)
surf_area = sum(surf_areas_LHem(verts))
Lhem(verts,2)
sub_info(v,:) = [Lhem_var_IDs(v) num_verts surf_area net]
var_freq_size_net
clear all
var_freq_size_net
% Network assignment
network_names = {'DMN' 'Vis' 'FP' 'DAN' 'Lang' 'Sal' 'CO' 'SMd' 'SML'
'Aud' 'Tpole' 'MTL' 'PMN' 'PON'};
good_nets_LH = [networksxHem.verticesLH(:,1:3) networksxHem.verticesLH(:,5) networksxHem.verticesLH(:,7:end)];
good_nets_RH = [networksxHem.verticesRH(:,1:3) networksxHem.verticesRH(:,5) networksxHem.verticesRH(:,7:end)];
good_nets_LH = mean(good_nets_LH);
good_nets_RH = mean(good_nets_RH);
means = [good_nets_LH' good_nets_RH'];
bar(1:14, means)

```

```

legend('Left Hem', 'Right Hem')
xticks(1:14)
xticklabels(network_names)
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.9, 0.7]);
ylabel('Average Number of Variant vertices');
xlabel('Network');
title('Variant vertices Assigned to Each Network Across Hemispheres');
ax = gca;
ax.FontSize = 24;
print(gcf,[output_dir cell2mat(out_str(g)) '_netAssignment_verts.jpg'],'-dpng','-r300');
rgb_colors = [1 0 0; %DMN
0 0 .6; %Vis
1 1 0; %FP
.67 .67 .67; %Unassigned
0 .8 0; %DAN
.67 .67 .67; %Unassigned2
0 .6 .6;%VAN
0 0 0; % Sal
.3 0 .6; %CON
.2 1 1; %SMd
1 .5 0; % SML
.6 .2 1; %Aud
0 .2 .4; %Tpole
.2 1 .2; %MTL
0 0 1; %PMN
.8 .8 .6]; %PON
lh_numvars = [];
rh_numvars = [];
for net = 1:16
vars_LH = []; vars_RH = [];
vars_LH = networksxHem.verticesLH(find(networksxHem.verticesLH(:,net)>0), net);
if length(vars_LH)>30 || length(vars_RH)>30
if length(vars_LH)>0
lh_numvars(net) = mean(vars_LH);
else
lh_numvars(net) = 0;
end
vars_RH = networksxHem.verticesRH(find(networksxHem.verticesRH(:,net)>0), net);
if length(vars_RH)>0
rh_numvars(net) = mean(vars_RH);
else
rh_numvars(net) = 0;
end
else
lh_numvars(net) = 0;
rh_numvars(net) = 0;
end
end
scatter(lh_numvars, rh_numvars, 80, rgb_colors, 'filled')
hold on
scatter(lh_numvars, rh_numvars, 80, 'k')
%axis([0 2.5 0 2.5])
%axis([200 700 200 700])
line = reffline(1,0);
line.Color = 'black';
xlabel('Left Hemisphere')
ylabel('Right Hemisphere')
ax = gca;
ax.FontSize = 24;
for net = 1:16

```

```

%se_lh(net) = std(networksxHem.surfaceLH(:,net))/sqrt(384);
se_lh(net) = std(vars_LH)/sqrt(length(vars_LH));
%se_rh(net) = std(networksxHem.surfaceRH(:,net))/sqrt(384);
se_rh(net) = std(vars_RH)/sqrt(length(vars_RH));
end
h = errorbar(lh_numvars, rh_numvars, se_rh, se_rh, se_lh, se_lh, 'o', 'MarkerSize', 10)
h.Color = [0,0,0];
mean(vars_LH)
mean(vars_RH)
mean(networksxHem.verticesLH(find(networksxHem.verticesLH(:,net)>0), net))
mean(networksxHem.verticesLH(:, 7))
mean(networksxHem.verticesRH(:, 7))
mean(networksxHem.verticesLH(find(networksxHem.verticesLH(:,7)>0), 7))
mean(networksxHem.verticesRH(:, 7))
mean(networksxHem.verticesRH(find(networksxHem.verticesRH(:,7)>0), 7))
mean(networksxHem.verticesLH(:, 7)) - mean(networksxHem.verticesRH(:, 7))
mean(networksxHem.verticesLH(find(networksxHem.verticesLH(:,7)>0), 7)) - mean(
networksxHem.verticesRH(find(networksxHem.verticesRH(:,7)>0), 7))
%-- 7/11/22, 12:22 PM --%
var_freq_size_net
leftHem = variants_info.left_hem.group_avg{:,4};
rightHem = variants_info.right_hem.group_avg{:,4};
[h,mu,sigma,q,notch] = al_goodplot(leftHem,0.1,.2,[ 0    0.4470    0.7410], 'left');
[h,mu,sigma,q,notch] = al_goodplot(rightHem,0.11,.2,[0.8500    0.3250
0.0980], 'right');
xticks([0.105])
xticklabels('Average Total Variant Territory')
ylabel('Surface Area in mm^2')
axis([-0.2, 0.5, min([leftHem; rightHem])-std([leftHem; rightHem]), max([leftHem;
rightHem])+std([leftHem; rightHem])])
ax = gca;
ax.FontSize = 24;
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.3, 0.9]); %first and second
control position on screen, third controls width, and fourth controls height
m = findobj(gca, 'Type', 'Scatter');
hleg1 = legend(m(1:2), 'Right Hemisphere', 'Left Hemisphere', 'Location',
'SouthOutside');
print(gcf, [output_dir cell2mat(out_str(g)) '_avgVarTerritory_mm2.jpg'], '-dpng', '-r
300')
leftHem = variants_info.left_hem.group_avg{:,5};
rightHem = variants_info.right_hem.group_avg{:,5};
[h,mu,sigma,q,notch] = al_goodplot(leftHem,0.1,.2,[ 0    0.4470    0.7410], 'left');
[h,mu,sigma,q,notch] = al_goodplot(rightHem,0.11,.2,[0.8500    0.3250
0.0980], 'right');
xticks([0.105])
xticklabels('Average Variant Size')
ylabel('Surface area (mm^2)')
axis([-0.2, 0.5, min([leftHem; rightHem])-std([leftHem; rightHem]), max([leftHem;
rightHem])+std([leftHem; rightHem])])
ax = gca;
ax.FontSize = 24;
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.3, 0.9]); %first and second
control position on screen, third controls width, and fourth controls height
m = findobj(gca, 'Type', 'Scatter');
hleg1 = legend(m(1:2), 'Right Hemisphere', 'Left Hemisphere', 'Location',
'SouthOutside');
print(gcf, [output_dir cell2mat(out_str(g)) '_avgSizeSurfArea.jpg'], '-dpng', '-r300')
network_names = {'DMN' 'Vis' 'FP' 'DAN' 'Lang' 'Sal' 'CO' 'SMd' 'SMl'
'Aud' 'Tpole' 'MTL' 'PMN' 'PON'};
good_nets_LH = [networksxHem.clustersLH(:,1:3) networksxHem.verticesLH(:,5) networksxHem.

```



```

verticesLH(:,7:end)];
good_nets_RH = [networksxHem.clustersRH(:,1:3) networksxHem.verticesRH(:,5) networksxHem.
verticesRH(:,7:end)];
good_nets_LH = mean(good_nets_LH);
good_nets_RH = mean(good_nets_RH);
means = [good_nets_LH' good_nets_RH'];
bar(1:14, means)
legend('Left Hem', 'Right Hem')
xticks(1:14)
xticklabels(network_names)
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.9, 0.7]);
ylabel('Average Number of Variant vertices');
xlabel('Network');
title('Variant vertices Assigned to Each Network Across Hemispheres');
ax = gca;
ax.FontSize = 24;
network_names = {'DMN' 'Vis' 'FP' 'DAN' 'Lang' 'Sal' 'CO' 'SMd' 'SML'
'Aud' 'Tpole' 'MTL' 'PMN' 'PON'};
good_nets_LH = [networksxHem.clustersLH(:,1:3) networksxHem.clustersLH(:,5) networksxHem.
clustersLH(:,7:end)];
good_nets_RH = [networksxHem.clustersRH(:,1:3) networksxHem.clustersRH(:,5) networksxHem.
clustersRH(:,7:end)];
good_nets_LH = mean(good_nets_LH);
good_nets_RH = mean(good_nets_RH);
means = [good_nets_LH' good_nets_RH'];
bar(1:14, means)
legend('Left Hem', 'Right Hem')
xticks(1:14)
xticklabels(network_names)
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.9, 0.7]);
ylabel('Average Number of Variant vertices');
xlabel('Network');
title('Variant vertices Assigned to Each Network Across Hemispheres');
ax = gca;
ax.FontSize = 24;
title('Variant vertices Assigned to Each Network Across Hemispheres');
title('Variants Assigned to Each Network Across Hemispheres');
print(gcf,[output_dir cell2mat(out_str(g)) '_netAssignment_clusters.jpg'],'-dpng','-
r300');
network_names = {'DMN' 'Vis' 'FP' 'DAN' 'Lang' 'Sal' 'CO' 'SMd' 'SML'
'Aud' 'Tpole' 'MTL' 'PMN' 'PON'};
good_nets_LH = [networksxHem.surfaceLH(:,1:3) networksxHem.surfaceLH(:,5) networksxHem.
surfaceLH(:,7:end)];
good_nets_RH = [networksxHem.surfaceRH(:,1:3) networksxHem.surfaceRH(:,5) networksxHem.
surfaceRH(:,7:end)];
good_nets_LH = mean(good_nets_LH);
good_nets_RH = mean(good_nets_RH);
means = [good_nets_LH' good_nets_RH'];
bar(1:14, means)
legend('Left Hem', 'Right Hem')
xticks(1:14)
xticklabels(network_names)
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.9, 0.7]);
ylabel('Surface Area (mm^2)');
xlabel('Network');
title('Variant Territory Assigned to Each Network Across Hemispheres');
ax = gca;
ax.FontSize = 24;
print(gcf,[output_dir cell2mat(out_str(g)) '_netAssignment_territorymm2.jpg'],'-dpng','-
r300');

```

```

network_names = {'DMN' 'Vis' 'FP' 'DAN' 'Lang' 'Sal' 'CO' 'SMd' 'SMl'
'Aud' 'Tpole' 'MTL' 'PMN' 'PON'};
good_nets_LH = [networksxHem.avgSizeLH(:,1:3) networksxHem.avgSizeLH(:,5) networksxHem.
avgSizeLH(:,7:end)];
good_nets_RH = [networksxHem.avgSizeRH(:,1:3) networksxHem.avgSizeRH(:,5) networksxHem.
avgSizeRH(:,7:end)];
good_nets_LH = mean(good_nets_LH);
good_nets_RH = mean(good_nets_RH);
means = [good_nets_LH' good_nets_RH'];
bar(1:14, means)
legend('Left Hem', 'Right Hem')
xticks(1:14)
xticklabels(network_names)
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.9, 0.7]);
ylabel('Surface Area (mm^2)');
xlabel('Network');
title('Average Size of Variants Assigned to Each Network Across Hemispheres');
ax = gca;
ax.FontSize = 24;
print(gcf,[output_dir cell2mat(out_str(g)) '_netAssignment_varSizemm2.jpg'],'-dpng','-
r300');
perms_var_freq_size
title_str = 'Average Variant Territory in Surface Area Across Hemispheres';
y_label = 'Difference in Average Variant Territory in Surface Area (left hem - right
hem)';
outfile = [output_dir out_str '_PermutationTesting_VariantTerritorySurfArea.jpg'];
[true_diff perm_diffs] = permute_values(allSubs_info.variants_info.left_hem.group_avg{:,
5}, allSubs_info.variants_info.right_hem.group_avg{:,5}, flip_switch, numperms, 0);
p = min([length(find(perm_diffs<true_diff))/numperms, length(find(perm_diffs>true_diff))
/numperms]);
p_vals = [p_vals; p];
disp(['Average Variant Size in Surface Area (permute hemispheres) - ' num2str(p)])
output_mat = [true_diff; perm_diffs'];
save([output_dir out_str 'var_size_surf_area.mat'], 'output_mat');
if plot
title_str = 'Average Variant Size in Surface Area Across Hemispheres';
y_label = 'Difference in Average Variant Size in Surface Area (left hem - right hem)';
outfile = [output_dir out_str '_PermutationTesting_VariantSizeSurfArea.jpg'];
plot_permtesting(output_mat, title_str, y_label, outfile);
end
permute_net_assignment
save([output_dir 'network_perms_varVertices.mat'], 'output_mat');
[p_fdr p_masked] = FDR(p_vals, .025);
permute_net_assignment
scatter([1:14],good_nets(1,:), 'MarkerEdgeColor', 'k', 'MarkerFaceColor', 'k',
'SizeData', 50)
load
('/Users/dianaperez/Desktop/Research/Lateralization/lateralization_code/testing_output/ne
twork_perms_varVertices.mat')
network_names =
{'DMN','Vis','FP','DAN','Lang','Sal','CO','SMd','SMl','Aud','Tpole','MTL','PMN','PON'};
rgb_colors = [1 0 0; %DMN
0 0 .6; %Vis
1 1 0; %FP
0 .8 0; %DAN
0 .6 .6;%VAN
0 0 0; % Sal
.3 0 .6; %CON
.2 1 1; %SMd
1 .5 0; % SMl

```

```

.6 .2 1; %Aud
0 .2 .4;%Tpole
.2 1 .2; %MTL
0 0 1; %PMN
.8 .8 .6]; %PON
good_nets = [output_mat(:,1:3) output_mat(:,5) output_mat(:,7:end)];
rgb_for_plot = {};
for n = 1:numel(network_names)
    rgb_for_plot{n} = rgb_colors(n,:);
end
ind = zeros(numperms+1,1);
ind(1,1) = 1;
net_inds = [ind ind ind ind ind ind ind ind ind ind ind ind ind];
handles = plotSpread(good_nets, 'categoryMarkers', {'x', '.'}, 'categoryLabels', \
{'Permuted Differences', 'True Difference'}, 'distributionColors', rgb_for_plot', \
'categoryIdx', net_inds, 'xNames', network_names)
ax = gca;
ax.FontSize = 24;
scatter([1:14],good_nets(1,:), 'MarkerEdgeColor', 'k', 'MarkerFaceColor', 'k', \
'SizeData', 50)
scatter(6,good_nets(1,6), 'MarkerEdgeColor', 'w', 'MarkerFaceColor', 'w', 'SizeData', 50)
ylabel('Difference in Number of Variant Vertices')
%ylabel('Difference in Variant Size')
xlabel('Assigned Network')
min_ax = min(min(good_nets(:,:)))-(min(std(good_nets)/2));
max_ax = max(max(good_nets(:,:)))+(min(std(good_nets)/2));
axis([0, 15, min_ax, max_ax])
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.9, 0.7]);
print(gcf,[output_dir out_str '_PermutationTesting_networkAssignment_numVerts.jpg'],'- \
dpng', '-r300');
close gcf
mean(allSubs_info.networksxHem.clustersLH,1)
mean(allSubs_info.networksxHem.clustersLH,1)'
mean(allSubs_info.networksxHem.clustersRH,1)'
mean(allSubs_info.networksxHem.verticesRH,1)'
mean(allSubs_info.networksxHem.verticesLH,1)'
mean(allSubs_info.networksxHem.surfaceLH,1)'
mean(allSubs_info.networksxHem.surfaceRH,1)'
mean(allSubs_info.networksxHem.avgSizeRH,1)'
mean(allSubs_info.networksxHem.avgSizeLH,1)'
var_territory_scatterplot
clear all
var_territory_scatterplot
axis([100 400 100 400])
print(gcf, \
'/Users/dianaperez/Desktop/Research/Lateralization/lateralization_code/HCP384_variant_clu \
sters_by_net.jpg', '-dpng', '-r300')
var_territory_scatterplot
axis([1 3 1 3])
print(gcf, \
'/Users/dianaperez/Desktop/Research/Lateralization/lateralization_code/HCP384_variant_clu \
sters_by_net.jpg', '-dpng', '-r300')
var_territory_scatterplot
axis([100 400 100 400])
print(gcf, \
'/Users/dianaperez/Desktop/Research/Lateralization/lateralization_code/HCP384_variant_ver \
tices_by_net.jpg', '-dpng', '-r300')
var_territory_scatterplot
axis([200 800 200 800])
print(gcf, \

```

```

'/Users/dianaperez/Desktop/Research/Lateralization/lateralization_code/HCP384_variant_ter
ritory_by_net.jpg', '-dpng', '-r300')
var_territory_scatterplot
axis([200 800 200 800])
var_territory_scatterplot
axis([50 600 50 600])
axis([20 600 20 600])
print(gcf,
'/Users/dianaperez/Desktop/Research/Lateralization/lateralization_code/HCP384_variant_ter
ritory_by_net.jpg', '-dpng', '-r300')
var_territory_scatterplot
axis([50 300 50 300])
axis([10 300 10 300])
axis([20 300 20 300])
print(gcf,
'/Users/dianaperez/Desktop/Research/Lateralization/lateralization_code/HCP384_variant_siz
e_by_net.jpg', '-dpng', '-r300')
close gcf
var_territory_scatterplot
axis([0 3 0 3])
line = reffline(1,0);
line.Color = 'black';
print(gcf,
'/Users/dianaperez/Desktop/Research/Lateralization/lateralization_code/HCP384_variant_clu
sters_by_net.jpg', '-dpng', '-r300')
var_territory_scatterplot
print(gcf,
'/Users/dianaperez/Desktop/Research/Lateralization/lateralization_code/HCP384_variant_ver
tices_by_net.jpg', '-dpng', '-r300')
help ttest
load
(''/Users/dianaperez/Desktop/Research/Lateralization/lateralization_code/testing_output/HC
P384_networksxHem.mat')
root_dir = '/Users/dianaperez/Desktop/Research/Lateralization/lateralization_code/'; %
location of code directory
surf_areas = ft_read_cifti_mod([root_dir '/needed_files/surf_areas_verts.dtseries.nii']);
surf_areas_LHem = surf_areas.data(1:29696);
surf_areas_RHem = surf_areas.data(29697:end);
clear all\
clear all
load
(''/Users/dianaperez/Desktop/Research/Lateralization/lateralization_code/testing_output/HC
P384_networksxHem.mat')
root_dir = '/Users/dianaperez/Desktop/Research/Lateralization/lateralization_code/'; %
location of code directory
surf_areas = ft_read_cifti_mod([root_dir '/needed_files/surf_areas_verts.dtseries.nii']);
surf_areas_LHem = surf_areas.data(1:29696);
surf_areas_RHem = surf_areas.data(29697:end);
sum(surf_areas_LHem)
sum(surf_areas_RHem)
var_territory_scatterplot
axis([0 .02 0 .02])
axis([0 .01 0 .01])
print(gcf,
'/Users/dianaperez/Desktop/Research/Lateralization/lateralization_code/HCP384_variant_ter
ritory_proportion_scatterplot.jpg', '-dpng', '-r300')
%-- 7/16/22, 8:13 PM --%
cd /Volumes
%-- 8/18/22, 4:20 PM --%
perms_var_freq_size

```

```

%-- 8/19/22, 1:37 PM --%
var_freq_size_net
permute_net_assignment
order = [7,11,2,14,10,12,15,8,5,16,3,9,1];
network_names = ␣
{'DMN','Vis','FP','DAN','Lang','Sal','CO','SMd','SML','Aud','Tpole','MTL','PMN','PON'};
rgb_colors = [1 0 0; %DMN
0 0 .6; %Vis
1 1 0; %FP
0 .8 0; %DAN
0 .6 .6;%VAN
0 0 0; % Sal
.3 0 .6; %CON
.2 1 1; %SMd
1 .5 0; % SML
.6 .2 1; %Aud
0 .2 .4;%Tpole
.2 1 .2; %MTL
0 0 1; %PMN
.8 .8 .6]; %PON
good_nets = [output_mat(:,1:3) output_mat(:,5) output_mat(:,7:end)];
rgb_for_plot = {};
for n = 1:numel(network_names)
    rgb_for_plot{n} = rgb_colors(n,:);
end
ind = zeros(numperms+1,1);
ind(1,1) = 1;
nets_sorted = output_mat(:,order);
net_inds = [ind ind ind ind ind ind ind ind ind ind ind ind ind];
rgb_sorted = rgb_for_plot{order};
rgb_sorted = rgb_for_plot{:,order};
for rgb = 1:length(order)
    rgb_sorted{rgb} = rgb_for_plot{order(rgb)};
end
rgb_sorted=[];
for rgb = 1:length(order)
    rgb_sorted{rgb} = rgb_for_plot{order(rgb)};
end
order = [5,9,2,12,8,10,13,6,4,14,3,7,1];
nets_sorted = output_mat(:,order);
rgb_sorted=[];
for rgb = 1:length(order)
    rgb_sorted{rgb} = rgb_for_plot{order(rgb)};
end
names_sorted = [];
for name = 1:length(order)
    names_sorted{name} = network_names{order(name)};
end
handles = plotSpread(nets_sorted, 'categoryMarkers', {'x', '.'}, 'categoryLabels', ␣
{'Permuted Differences','True Difference'}, 'distributionColors', rgb_sorted, ␣
'categoryIdx', net_inds, 'xNames', names_sorted)
ax = gca;
ax.FontSize = 24;
scatter([1:13],nets_sorted(1,:), 'MarkerEdgeColor', 'k', 'MarkerFaceColor', 'k', ␣
'SizeData', 50)
scatter(8,nets_sorted(1,8), 'MarkerEdgeColor', 'w', 'MarkerFaceColor', 'w', 'SizeData', ␣
50)
clear all
permute_net_assignment
ylabel('Difference in Total Variant Territory')

```

```
print(gcf,[output_dir out_str '_PermutationTesting_networkAssignment_VariantTerritory.↵
jpg'],'-dpng','-r300');
close gcf
clear all
make_overlapmap
overlap_nonbrain = insert_nonbrain(overlap_map, 'both', out_cifti);
length(template.brainstructure)
length(template.brainstructure)/2
overlap_nonbrain = insert_nonbrain(overlap_map.data, 'both', template);
left = overlap_nonbrain(1:32492);
right = overlap_nonbrain(32493:end);
diff_map = left - right;
diff_map = [diff_map; diff_map];
diff_map = diff_map(template.brainstructure>0);
template.data = diff_map;
ft_write_cifti_mod(strrep(out_cifti, 'overlap', 'diff'), template)
make_overlapmap
out_cifti
make_overlapmap
clear all
make_overlapmap
max(overlap_map.data
max(overlap_map.data)
overlap_map.data*1000
max(overlap_map.data)
test = overlap_map.data*1000;
max(test)
test = overlap_map.data*100;
max(test)
overlap_map.data = test;
ft_write_cifti_mod(out_cifti, overlap_map);
if create_diff
disp('Creating difference map...')
overlap_nonbrain = insert_nonbrain(overlap_map.data, 'both', template);
left = overlap_nonbrain(1:32492);
right = overlap_nonbrain(32493:end);
diff_map = left - right;
diff_map = [diff_map; diff_map];
diff_map = diff_map(template.brainstructure>0);
template.data = diff_map;
ft_write_cifti_mod(strrep(out_cifti, 'overlap', 'diff'), template)
end
out_cifti
make_overlapmap
test = overlap_map.data*100;
max(test)
overlap_map.data = test;
make_overlapmap
max(overlap_map)
max(overlap_map.data)
make_overlapmap
wb_dir = '/Applications/workbench/bin_macosx64/';
group_avg =↵
'/Users/dianaperez/Desktop/Research/Lateralization/lateralization_code/group_avgs/120_LR_↵
minsize400_recolored_manualconsensus_LR.dlabel.nii';
borderlabel_file = '/Users/dianaperez/Downloads/wbBorderLabelFile.txt';
output_dlabel = '/Users/dianaperez/Desktop/WashU120_labelfile_fornetoutlines.dlabel.nii';
output_lefthem = '/Users/dianaperez/Desktop/WashU120_labelfile_fornetoutlines_left.label.↵
gii';
output_righthem = '/Users/dianaperez/Desktop/WashU120_labelfile_fornetoutlines_right.↵
```

```

label.gii';
system([wb_dir 'wb_command -cifti-label-import ' group_avg ' ' borderlabel_file ' '
output_dlabel])
system([wb_dir 'wb_command -cifti-separate ' output_dlabel ' COLUMN -label CORTEX_LEFT '
output_lefthem ' -label CORTEX_RIGHT ' output_righthem])
system([wb_dir 'wb_command -label-to-border
/Users/dianaperez/Documents/Dependencies/Resources/Conte69_atlas-v2.LR.32k_fs_LR.
wb/Conte69.L.midthickness.32k_fs_LR.surf.gii ' output_lefthem ' WashU120_netborders_L.
border'])
system([wb_dir 'wb_command -label-to-border
/Users/dianaperez/Documents/Dependencies/Resources/Conte69_atlas-v2.LR.32k_fs_LR.
wb/Conte69.R.midthickness.32k_fs_LR.surf.gii ' output_righthem ' WashU120_netborders_R.
border'])
make_overlapmap
clear all
load
('Users/dianaperez/Desktop/Research/Lateralization/lateralization_code/testing_output/HCP
P752_RH_networksxHem.mat')
good_nets_LH = [networksxHem.surfaceLH(:,1:3) networksxHem.surfaceLH(:,5) networksxHem.
surfaceLH(:,7:end)];
good_nets_RH = [networksxHem.surfaceRH(:,1:3) networksxHem.surfaceRH(:,5) networksxHem.
surfaceRH(:,7:end)];
good_nets_LH = mean(good_nets_LH);
good_nets_RH = mean(good_nets_RH);
means = [good_nets_LH' good_nets_RH'];
bar(1:14, means)
legend('Left Hem', 'Right Hem')
xticks(1:14)
xticklabels(network_names)
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.9, 0.7]);
ylabel('Average Total Variant Territory');
network_names = {'DMN' 'Vis' 'FP' 'DAN' 'Lang' 'Sal' 'CO' 'SMd' 'SMl'
'Aud' 'Tpole' 'MTL' 'PMN' 'PON'};
xticklabels(network_names)
title('Variant Territory Assigned to Each Network Across Hemispheres - Right Handers');
ax = gca;
ax.FontSize = 24;
print(gcf, '/Users/dianaperez/Desktop/HCP752_RightHanders_netAssignment_VariantTerritory.
jpg', '-dpng', '-r300');
print(gcf, '/Users/dianaperez/Desktop/HCP752_RightHanders_netAssignment_VariantTerritory.
jpg', '-dpng', '-r300');
title('Variant Territory Across Networks - Right Handers');
ylabel('Average Variant Territory (mm^2)');
print(gcf, '/Users/dianaperez/Desktop/HCP752_RightHanders_netAssignment_VariantTerritory.
jpg', '-dpng', '-r300');
close gcf
load
('Users/dianaperez/Desktop/Research/Lateralization/lateralization_code/testing_output/HCP
P752_LH_networksxHem.mat')
good_nets_LH = [networksxHem.surfaceLH(:,1:3) networksxHem.surfaceLH(:,5) networksxHem.
surfaceLH(:,7:end)];
good_nets_RH = [networksxHem.surfaceRH(:,1:3) networksxHem.surfaceRH(:,5) networksxHem.
surfaceRH(:,7:end)];
good_nets_LH = mean(good_nets_LH);
good_nets_RH = mean(good_nets_RH);
means = [good_nets_LH' good_nets_RH'];
bar(1:14, means)
legend('Left Hem', 'Right Hem')
xticks(1:14)
xticklabels(network_names)

```



```

set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.9, 0.7]);
title('Variant Territory Assigned to Each Network Across Hemispheres - Left Handers');
ylabel('Average Variant Territory (mm^2)');
ax = gca;
ax.FontSize = 24;
print(gcf, '/Users/dianaperez/Desktop/HCP752_LeftHanders_netAssignment_VariantTerritory.
jpg', '-dpng', '-r300');
load
('/Users/dianaperez/Desktop/Research/Lateralization/lateralization_code/testing_output/HCP
P752_RH_networksxHem.mat')
good_nets_LH = [networksxHem.surfaceLH(:,1:3) networksxHem.surfaceLH(:,5) networksxHem.
surfaceLH(:,7:end)];
good_nets_RH = [networksxHem.surfaceRH(:,1:3) networksxHem.surfaceRH(:,5) networksxHem.
surfaceRH(:,7:end)];
good_nets_LH = mean(good_nets_LH);
good_nets_RH = mean(good_nets_RH);
means = [good_nets_LH' good_nets_RH'];
bar(1:14, means)
legend('Left Hem', 'Right Hem')
xticks(1:14)
xticklabels(network_names)
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.9, 0.7]);
ax = gca;
ax.FontSize = 24;
title('Variant Territory Across Networks - Right Handers');
ylabel('Average Variant Territory (mm^2)');
print(gcf, '/Users/dianaperez/Desktop/HCP752_RightHanders_netAssignment_VariantTerritory.
jpg', '-dpng', '-r300');
close gcf
load
('/Users/dianaperez/Desktop/Research/Lateralization/lateralization_code/testing_output/HCP
P752_new_split_LH_networksxHem.mat')
network_names = {'DMN' 'Vis' 'FP' 'DAN' 'Lang' 'Sal' 'CO' 'SMd' 'SML'
'Aud' 'Tpole' 'MTL' 'PMN' 'PON'};
%good_nets_LH = [networksxHem.verticesLH(:,1:3) networksxHem.verticesLH(:,5)
networksxHem.verticesLH(:,7:end)];
good_nets_LH = [networksxHem.surfaceLH(:,1:3) networksxHem.surfaceLH(:,5) networksxHem.
surfaceLH(:,7:end)];
good_nets_RH = [networksxHem.surfaceRH(:,1:3) networksxHem.surfaceRH(:,5) networksxHem.
surfaceRH(:,7:end)];
good_nets_LH = mean(good_nets_LH);
good_nets_RH = mean(good_nets_RH);
means = [good_nets_LH' good_nets_RH'];
for net = 1:16
se_lh(net) = std(networksxHem.surfaceLH(:,net))/sqrt(40);
se_rh(net) = std(networksxHem.surfaceRH(:,net))/sqrt(40);
end
se_lh(4) = [];
se_lh(5) = [];
se_rh(4) = [];
se_rh(5) = [];
bar(1:14, means)
legend('Left Hem', 'Right Hem')
xticks(1:14)
xticklabels(network_names)
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.9, 0.7]);
%ylabel('Average Number of Variant Regions')
ylabel('Average Variant Territory (mm^2)');
xlabel('Network');
%title('Variant Regions Assigned to Each Network Across Hemispheres')

```



```

title('Variant Territory Across Networks – Right Handers');
ax = gca;
ax.FontSize = 24;
hold on
h = errorbar(.85:1:13.85, means(:,1), se_lh, se_lh)
h.Color = [0,0,0];
g = errorbar(1.15:1:14.15, means(:,2), se_rh, se_rh)
g.Color = [0,0,0];
h.LineStyle = 'none';
g.LineStyle = 'none';
h.LineWidth = 2;
g.LineWidth = 2;
legend('Left Hem', 'Right Hem')
clear all
load
('Users/dianaperez/Desktop/Research/Lateralization/lateralization_code/testing_output/HCP752_new_split_LH_networksxHem.mat')
network_names = {'DMN' 'Vis' 'FP' 'DAN' 'Lang' 'Sal' 'CO' 'SMd' 'SMl'
'Aud' 'Tpole' 'MTL' 'PMN' 'PON'};
%good_nets_LH = [networksxHem.verticesLH(:,1:3) networksxHem.verticesLH(:,5)
networksxHem.verticesLH(:,7:end)];
good_nets_LH = [networksxHem.surfaceLH(:,1:3) networksxHem.surfaceLH(:,5) networksxHem.
surfaceLH(:,7:end)];
good_nets_RH = [networksxHem.surfaceRH(:,1:3) networksxHem.surfaceRH(:,5) networksxHem.
surfaceRH(:,7:end)];
good_nets_LH = mean(good_nets_LH);
good_nets_RH = mean(good_nets_RH);
means = [good_nets_LH' good_nets_RH'];
load
('Users/dianaperez/Desktop/Research/Lateralization/lateralization_code/testing_output/HCP752_LH_networksxHem.mat')
load
('Users/dianaperez/Desktop/Research/Lateralization/lateralization_code/testing_output/HCP752_LH_networksxHem.mat')
%load
('Users/dianaperez/Desktop/Research/Lateralization/lateralization_code/testing_output/HCP752_new_split_LH_networksxHem.mat')
network_names = {'DMN' 'Vis' 'FP' 'DAN' 'Lang' 'Sal' 'CO' 'SMd' 'SMl'
'Aud' 'Tpole' 'MTL' 'PMN' 'PON'};
%good_nets_LH = [networksxHem.verticesLH(:,1:3) networksxHem.verticesLH(:,5)
networksxHem.verticesLH(:,7:end)];
good_nets_LH = [networksxHem.surfaceLH(:,1:3) networksxHem.surfaceLH(:,5) networksxHem.
surfaceLH(:,7:end)];
good_nets_RH = [networksxHem.surfaceRH(:,1:3) networksxHem.surfaceRH(:,5) networksxHem.
surfaceRH(:,7:end)];
good_nets_LH = mean(good_nets_LH);
good_nets_RH = mean(good_nets_RH);
means = [good_nets_LH' good_nets_RH'];
for net = 1:16
se_lh(net) = std(networksxHem.surfaceLH(:,net))/sqrt(40);
se_rh(net) = std(networksxHem.surfaceRH(:,net))/sqrt(40);
end
se_lh(4) = [];
se_lh(5) = [];
se_rh(4) = [];
se_rh(5) = [];
bar(1:14, means)
legend('Left Hem', 'Right Hem')
xticks(1:14)
xticklabels(network_names)

```

```

set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.9, 0.7]);
ylabel('Average Number of Variant Regions')
ylabel('Average Variant Territory (mm^2)');
xlabel('Network');
%title('Variant Regions Assigned to Each Network Across Hemispheres')
title('Variant Territory Across Networks – Right Handers');
ax = gca;
ax.FontSize = 24;
hold on
h = errorbar(.85:1:13.85, means(:,1), se_lh, se_lh)
h.Color = [0,0,0];
g = errorbar(1.15:1:14.15, means(:,2), se_rh, se_rh)
g.Color = [0,0,0];
h.LineStyle = 'none';
g.LineStyle = 'none';
h.LineWidth = 2;
g.LineWidth = 2;
legend('Left Hem', 'Right Hem')
print(gcf, '/Users/dianaperez/Desktop/HCP752_numvars_Lhand_leftvrighthem.jpg', '-dpng',
'-r300')
close gcf
load
('/Users/dianaperez/Desktop/Research/Lateralization/lateralization_code/testing_output/HCP
P752_RH_networksxHem.mat')
network_names = {'DMN' 'Vis' 'FP' 'DAN' 'Lang' 'Sal' 'CO' 'SMD' 'SML'
'Aud' 'Tpole' 'MTL' 'PMN' 'PON'};
%good_nets_LH = [networksxHem.verticesLH(:,1:3) networksxHem.verticesLH(:,5)
networksxHem.verticesLH(:,7:end)];
good_nets_LH = [networksxHem.surfaceLH(:,1:3) networksxHem.surfaceLH(:,5) networksxHem.
surfaceLH(:,7:end)];
good_nets_RH = [networksxHem.surfaceRH(:,1:3) networksxHem.surfaceRH(:,5) networksxHem.
surfaceRH(:,7:end)];
good_nets_LH = mean(good_nets_LH);
good_nets_RH = mean(good_nets_RH);
means = [good_nets_LH' good_nets_RH'];
for net = 1:16
se_lh(net) = std(networksxHem.surfaceLH(:,net))/sqrt(670);
se_rh(net) = std(networksxHem.surfaceRH(:,net))/sqrt(670);
end
se_lh(4) = [];
se_lh(5) = [];
se_rh(4) = [];
se_rh(5) = [];
bar(1:14, means)
legend('Left Hem', 'Right Hem')
xticks(1:14)
xticklabels(network_names)
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.9, 0.7]);
ylabel('Average Variant Territory (mm^2)');
xlabel('Network');
title('Variant Territory Across Networks – Right Handers');
%title('Variant Regions Assigned to Each Network Across Hemispheres')
ax = gca;
ax.FontSize = 24;
hold on
h = errorbar(.85:1:13.85, means(:,1), se_lh, se_lh)
h.Color = [0,0,0];
g = errorbar(1.15:1:14.15, means(:,2), se_rh, se_rh)
g.Color = [0,0,0];
h.LineStyle = 'none';

```

```

g.LineStyle = 'none';
h.LineWidth = 2;
g.LineWidth = 2;
legend('Left Hem', 'Right Hem')
axis([0,15,0,2.5])
close gcf
load
('/Users/dianaperez/Desktop/Research/Lateralization/lateralization_code/testing_output/HCP752_RH_networksxHem.mat')
network_names = {'DMN' 'Vis' 'FP' 'DAN' 'Lang' 'Sal' 'CO' 'SMd' 'SMl'
'Aud' 'Tpole' 'MTL' 'PMN' 'PON'};
%good_nets_LH = [networksxHem.verticesLH(:,1:3) networksxHem.verticesLH(:,5)
networksxHem.verticesLH(:,7:end)];
good_nets_LH = [networksxHem.surfaceLH(:,1:3) networksxHem.surfaceLH(:,5) networksxHem.
surfaceLH(:,7:end)];
good_nets_RH = [networksxHem.surfaceRH(:,1:3) networksxHem.surfaceRH(:,5) networksxHem.
surfaceRH(:,7:end)];
good_nets_LH = mean(good_nets_LH);
good_nets_RH = mean(good_nets_RH);
means = [good_nets_LH good_nets_RH];
for net = 1:16
se_lh(net) = std(networksxHem.surfaceLH(:,net))/sqrt(670);
se_rh(net) = std(networksxHem.surfaceRH(:,net))/sqrt(670);
end
se_lh(4) = [];
se_lh(5) = [];
se_rh(4) = [];
se_rh(5) = [];
bar(1:14, means)
legend('Left Hem', 'Right Hem')
xticks(1:14)
xticklabels(network_names)
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.9, 0.7]);
ylabel('Average Number of Variant Regions')
ylabel('Average Variant Territory (mm^2)');
xlabel('Network');
title('Variant Territory Across Networks - Right Handers');
%title('Variant Regions Assigned to Each Network Across Hemispheres')
ax = gca;
ax.FontSize = 24;
hold on
h = errorbar(.85:1:13.85, means(:,1), se_lh, se_lh)
h.Color = [0,0,0];
g = errorbar(1.15:1:14.15, means(:,2), se_rh, se_rh)
g.Color = [0,0,0];
h.LineStyle = 'none';
g.LineStyle = 'none';
h.LineWidth = 2;
g.LineWidth = 2;
legend('Left Hem', 'Right Hem')
print(gcf, '/Users/dianaperez/Desktop/HCP752_numvars_Rhand_leftvrighthem.jpg', '-dpng',
'-r300')
close gcf
load
('/Users/dianaperez/Desktop/Research/Lateralization/lateralization_code/testing_output/HCP752_LH_networksxHem.mat')
%load
('/Users/dianaperez/Desktop/Research/Lateralization/lateralization_code/testing_output/HCP752_new_split_LH_networksxHem.mat')
network_names = {'DMN' 'Vis' 'FP' 'DAN' 'Lang' 'Sal' 'CO' 'SMd' 'SMl'

```

```

'Aud' 'Tpole' 'MTL' 'PMN' 'PON'};
%good_nets_LH = [networksxHem.verticesLH(:,1:3) networksxHem.verticesLH(:,5)
networksxHem.verticesLH(:,7:end)];
good_nets_LH = [networksxHem.surfaceLH(:,1:3) networksxHem.surfaceLH(:,5) networksxHem.
surfaceLH(:,7:end)];
good_nets_RH = [networksxHem.surfaceRH(:,1:3) networksxHem.surfaceRH(:,5) networksxHem.
surfaceRH(:,7:end)];
good_nets_LH = mean(good_nets_LH);
good_nets_RH = mean(good_nets_RH);
means = [good_nets_LH' good_nets_RH'];
for net = 1:16
se_lh(net) = std(networksxHem.surfaceLH(:,net))/sqrt(40);
se_rh(net) = std(networksxHem.surfaceRH(:,net))/sqrt(40);
end
se_lh(4) = [];
se_lh(5) = [];
se_rh(4) = [];
se_rh(5) = [];
bar(1:14, means)
legend('Left Hem', 'Right Hem')
xticks(1:14)
xticklabels(network_names)
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.9, 0.7]);
ylabel('Average Number of Variant Regions')
ylabel('Average Variant Territory (mm^2)');
xlabel('Network');
%title('Variant Regions Assigned to Each Network Across Hemispheres')
title('Variant Territory Across Networks – Left Handers');
ax = gca;
ax.FontSize = 24;
hold on
h = errorbar(.85:1:13.85, means(:,1), se_lh, se_lh)
h.Color = [0,0,0];
g = errorbar(1.15:1:14.15, means(:,2), se_rh, se_rh)
g.Color = [0,0,0];
h.LineStyle = 'none';
g.LineStyle = 'none';
h.LineWidth = 2;
g.LineWidth = 2;
legend('Left Hem', 'Right Hem')
%print(gcf,
'/Users/dianaperez/Desktop/Research/lateralization_code/HCP752_numvars_Lhand_leftvrighthe
m.jpg', '-dpng', '-r300')
print(gcf, '/Users/dianaperez/Desktop/HCP752_numvars_Lhand_leftvrighthem.jpg', '-dpng',
'-r300')
close gcf
left_handers = networksxHem;
%% right handers
load
('/Users/dianaperez/Desktop/Research/Lateralization/lateralization_code/testing_output/HCP
P752_RH_networksxHem.mat')
network_names = {'DMN' 'Vis' 'FP' 'DAN' 'Lang' 'Sal' 'CO' 'SMd' 'SMl'
'Aud' 'Tpole' 'MTL' 'PMN' 'PON'};
%good_nets_LH = [networksxHem.verticesLH(:,1:3) networksxHem.verticesLH(:,5)
networksxHem.verticesLH(:,7:end)];
good_nets_LH = [networksxHem.surfaceLH(:,1:3) networksxHem.surfaceLH(:,5) networksxHem.
surfaceLH(:,7:end)];
good_nets_RH = [networksxHem.surfaceRH(:,1:3) networksxHem.surfaceRH(:,5) networksxHem.
surfaceRH(:,7:end)];
good_nets_LH = mean(good_nets_LH);

```

```

good_nets_RH = mean(good_nets_RH);
means = [good_nets_LH' good_nets_RH'];
for net = 1:16
se_lh(net) = std(networksxHem.surfaceLH(:,net))/sqrt(670);
se_rh(net) = std(networksxHem.surfaceRH(:,net))/sqrt(670);
end
se_lh(4) = [];
se_lh(5) = [];
se_rh(4) = [];
se_rh(5) = [];
bar(1:14, means)
legend('Left Hem', 'Right Hem')
xticks(1:14)
xticklabels(network_names)
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.9, 0.7]);
ylabel('Average Number of Variant Regions')
ylabel('Average Variant Territory (mm^2)');
xlabel('Network');
title('Variant Territory Across Networks – Right Handers');
%title('Variant Regions Assigned to Each Network Across Hemispheres')
ax = gca;
ax.FontSize = 24;
hold on
h = errorbar(.85:1:13.85, means(:,1), se_lh, se_lh)
h.Color = [0,0,0];
g = errorbar(1.15:1:14.15, means(:,2), se_rh, se_rh)
g.Color = [0,0,0];
h.LineStyle = 'none';
g.LineStyle = 'none';
h.LineWidth = 2;
g.LineWidth = 2;
legend('Left Hem', 'Right Hem')
axis([0,15,0,2.5])
%print(gcf,
'/Users/dianaperez/Desktop/Research/lateralization_code/HCP752_numvars_Rhand_leftvrighthe
m.jpg', '-dpng', '-r300')
print(gcf, '/Users/dianaperez/Desktop/HCP752_numvars_Rhand_leftvrighthem.jpg', '-dpng',
'-r300')
close(gcf)
right_handers = networksxHem;
bar(1:14, means)
legend('Left Hem', 'Right Hem')
xticks(1:14)
xticklabels(network_names)
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.9, 0.7]);
ylabel('Average Number of Variant Regions')
ylabel('Average Variant Territory (mm^2)');
xlabel('Network');
title('Variant Territory Across Networks – Right Handers');
%title('Variant Regions Assigned to Each Network Across Hemispheres')
ax = gca;
ax.FontSize = 24;
hold on
h = errorbar(.85:1:13.85, means(:,1), se_lh, se_lh)
h.Color = [0,0,0];
g = errorbar(1.15:1:14.15, means(:,2), se_rh, se_rh)
g.Color = [0,0,0];
h.LineStyle = 'none';
g.LineStyle = 'none';
h.LineWidth = 2;

```

```
g.LineWidth = 2;
legend('Left Hem', 'Right Hem')
%print(gcf,
'/Users/dianaperez/Desktop/Research/lateralization_code/HCP752_numvars_Rhand_leftvrighthe
m.jpg', '-dpng', '-r300')
print(gcf, '/Users/dianaperez/Desktop/HCP752_numvars_Rhand_leftvrighthem.jpg', '-dpng',
'-r300')
close(gcf)
right_handlers = networksxHem;
diff_lefthanders = left_handlers.surfaceLH - left_handlers.surfaceRH;
mean_diff_lefthanders = mean(diff_lefthanders, 1);
diff_righthanders = right_handlers.surfaceLH - right_handlers.surfaceRH;
mean_diff_righthanders = mean(diff_righthanders, 1);
for net = 1:16
se_lh(net) = std(diff_lefthanders(:,net))/sqrt(40);
se_rh(net) = std(diff_righthanders(:,net))/sqrt(670);
end
se_lh(4) = [];
se_lh(5) = [];
se_rh(4) = [];
se_rh(5) = [];
diffs = [mean_diff_lefthanders; mean_diff_righthanders];
diffs(:,4) = []; diffs(:,5) = [];
b = bar(1:14, diffs)
b(1).FaceColor = [0,0,0];%[.2 .6 .5];
b(2).FaceColor = [.5,.5,.5];%[0.4940 0.1840 0.5560];
legend('Left Handlers', 'Right Handlers')
xticks(1:14)
xticklabels(network_names)
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.9, 0.7]);
ylabel(' ')
xlabel(' ');
ax = gca;
ax.FontSize = 24;
hold on
h = errorbar(.85:1:13.85, diffs(1,:), se_lh, se_lh)
h.Color = [0.5,0.5,0.5];
g = errorbar(1.15:1:14.15, diffs(2,:), se_rh, se_rh)
g.Color = [0,0,0];
h.LineStyle = 'none';
g.LineStyle = 'none';
h.LineWidth = 2;
g.LineWidth = 2;
legend('Left Handlers', 'Right Handlers')
close(gcf)
diff_lefthanders = left_handlers.surfaceLH - left_handlers.surfaceRH;
mean_diff_lefthanders = mean(diff_lefthanders, 1);
diff_righthanders = right_handlers.surfaceLH - right_handlers.surfaceRH;
mean_diff_righthanders = mean(diff_righthanders, 1);
for net = 1:16
se_lh(net) = std(diff_lefthanders(:,net))/sqrt(40);
se_rh(net) = std(diff_righthanders(:,net))/sqrt(670);
end
se_lh(4) = [];
se_lh(5) = [];
se_rh(4) = [];
se_rh(5) = [];
diffs = [mean_diff_lefthanders; mean_diff_righthanders];
diffs(:,4) = []; diffs(:,5) = [];
b = bar(1:14, diffs)
```

```

b(1).FaceColor = [.2 .6 .5];%[0,0,0];
b(2).FaceColor = [0.4940 0.1840 0.5560];%[.5,.5,.5]
legend('Left Handers', 'Right Handers')
xticks(1:14)
xticklabels(network_names)
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.5, 0.7, 0.9, 0.7]);
ylabel(' ')
xlabel(' ');
ax = gca;
ax.FontSize = 24;
hold on
h = errorbar(.85:1:13.85, diffs(1,:), se_lh, se_lh)
%h.Color = [0.5,0.5,0.5];
g = errorbar(1.15:1:14.15, diffs(2,:), se_rh, se_rh)
%g.Color = [0,0,0];
h.LineStyle = 'none';
g.LineStyle = 'none';
h.LineWidth = 2;
g.LineWidth = 2;
legend('Left Handers', 'Right Handers')
g.Color = [0,0,0];
h.Color = [0,0,0];
ylabel('Average Difference in Variant Territory (left-right hem)');
xlabel('Network');
ylabel('Average Difference in Variant Territory');
print(gcf, '/Users/dianaperez/Desktop/HCP752_numvars_LvRhand_L-Rhem.jpg', '-dpng', '-r300')
permute_net_assignment
RandStream
permute_net_assignment
allSubs_info = load([data_location in_str '_LH_networksxHem.mat']);
allSubs_info = load
('/Users/dianaperez/Desktop/Research/Lateralization/lateralization_code/testing_output/HCP752_LH_networksxHem.mat')
output_dir = [output_dir 'Left_handers_'];
permute_net_assignment
allSubs_info = load
('/Users/dianaperez/Desktop/Research/Lateralization/lateralization_code/testing_output/HCP752_RH_networksxHem.mat')
numsubs = height(allSubs_info.networksxHem.clustersLH);
permute_net_assignment
allSubs_info = load
('/Users/dianaperez/Desktop/Research/Lateralization/lateralization_code/testing_output/HCP752_RH_networksxHem.mat')
permute_net_assignment
allSubs_info = load
('/Users/dianaperez/Desktop/Research/Lateralization/lateralization_code/testing_output/HCP752_RH_networksxHem.mat')
output_dir = [output_dir 'Right_handers_'];
permute_net_assignment
clear all
%-- 8/31/22, 9:38 AM --%
load
('/Users/dianaperez/Desktop/Research/Lateralization/lateralization_code/testing_output/HCP384_variants_info.mat')
load
('/Users/dianaperez/Desktop/Research/Lateralization/lateralization_code/testing_output/HCP384_networksxHem.mat')
numVars = variants_info.left_hem.group_avg{:,1};
[h,p,ci,stats] = ttest(variants_info.left_hem.group_avg{:,1},variants_info.right_hem.

```

```

group_avg{:,1});
[h,p,ci,stats] = ttest(variants_info.left_hem.group_avg{:,2},variants_info.right_hem.
group_avg{:,2});
[h,p,ci,stats] = ttest(variants_info.left_hem.group_avg{:,3},variants_info.right_hem.
group_avg{:,3});
DMN_LH = networksxHem.clustersLH(:,1);
[h,p,ci,stats] = ttest(networksxHem.clustersLH(:,1),networksxHem.clustersRH(:,1));
[h,p,ci,stats] = ttest(networksxHem.clustersLH(:,2),networksxHem.clustersRH(:,2));
[h,p,ci,stats] = ttest(networksxHem.clustersLH(:,7),networksxHem.clustersRH(:,7));
[h,p,ci,stats] = ttest(networksxHem.clustersLH(:,9),networksxHem.clustersRH(:,9));
[h,p,ci,stats] = ttest(networksxHem.clustersLH(:,11),networksxHem.clustersRH(:,11));
clear all
permute_net_assignment
addpath '/Users/dianaperez/Documents/GitHub/PerezEtAl_HemAsymmetries/needed_files'
permute_net_assignment
addpath
'/Users/dianaperez/Documents/GitHub/PerezEtAl_HemAsymmetries/needed_files/plotSpread/plot
Spread'
permute_net_assignment
clear all
library(tidyverse) #load tidyverse
data <- read.csv("C:/Users/dianaperez/Downloads/DIC_Data2022.csv")
data = readtable('/Users/dianaperez/Downloads/DIC_Data2022.csv');
DOI = data(:,18:90);
enc = DOI(~isnan(DOI(:,1)), 1:11);
help isnan
DOI(:,1)
DOI{:,1}
enc = DOI(~isnan(DOI{:,1}), 1:11);
mean_enc = mean(enc, 2);
mean_enc = mean(table2array(enc), 2);
mean_enc = mean(table2array(enc), 1);
bar(mean_enc)
groups = {'Race/Ethnicity', 'Gender', 'Sexual Orientation', 'Economic Status', 'Ability
Status', 'Age', 'Religion', 'Familial Obligations', 'Political Ideology',
'Nationality/Immigration', 'Mental Health'};
xlabel(groups)
bar(mean_enc, 1:11)
bar(1:11, mean_enc)
xlabel(groups)
help xtick
bar(1:11,mean_enc)
xticks(groups)
xticklabels(groups)
xtickangle(90)
xtickangle(70)
xtickangle(60)
yaxis([1:6])
axis([1:6])
clos(gcf)
close(gcf)
bar(1:11,mean_enc)
xticklabels(groups)
xtickangle(60)
yticks([1:6])
yticks([1:6])
help axis
axis([1 11 1 6])
axis([0 12 1 6])
answers = {'Not at all', 'A little', 'A moderate amount', 'Quite a bit', 'A great deal'};

```



```

axis([0 12 1 5])
ylabel('Average Response')
title('How much do you think the department acts to encourage diversity and inclusion as related to:')
title(' ')
answers = {'Never', 'Once a year', 'Once/twice a quarter', 'Several times', 'Weekly', 'Daily'};
imp_bias = DOI(~isnan(DOI(:,1)), 12:22);
mean_IB = mean(table2array(imp_bias), 2);
bar(1:11, mean_IB)
xticklabels(groups)
xtickangle(60)
axis([0 12 1 6])
ylabel('Average Response')
print(gcf, '/Users/dianaperez/Desktop/observe_imp_bias.jpg', '-dpng', '-r300');
answers = {'Never', 'Once a year', 'Once/twice a quarter', 'Several times', 'Weekly', 'Daily'};
imp_bias = DOI(~isnan(DOI(:,1)), 12:22);
mean_IB = mean(table2array(imp_bias), 2);
bar(1:11, mean_IB)
xticklabels(groups)
xtickangle(60)
axis([0 12 1 6])
ylabel('Average Response')
print(gcf, '/Users/dianaperez/Desktop/observe_imp_bias.jpg', '-dpng', '-r300');
imp_bias2 = DOI(~isnan(DOI(:,1)), 25:35);
mean_IB2 = mean(table2array(imp_bias2), 2);
bar(1:11, mean_IB2)
xticklabels(groups)
xtickangle(60)
axis([0 12 1 6])
ylabel('Average Response')
print(gcf, '/Users/dianaperez/Desktop/hear_imp_bias.jpg', '-dpng', '-r300');
clear mean*
mean_IB = mean(table2array(imp_bias), 2);
mean_IB = mean(table2array(imp_bias), 1);
bar(1:11, mean_IB)
xticklabels(groups)
xtickangle(60)
axis([0 12 1 6])
ylabel('Average Response')
print(gcf, '/Users/dianaperez/Desktop/observe_imp_bias.jpg', '-dpng', '-r300');
imp_bias = DOI(~isnan(DOI(:,1)), 12:22);
imp_bias(isnan(imp_bias)) = 0;
imp_bias(isnan(imp_bias{:,:})) = 0;
length(imp_bias)
width(imp_bias)
for q = 1:width(imp_bias)
col = table2array(imp_bias(~isnan(imp_bias(:,q)), q));
mean_IB(q) = mean(col);
end
bar(1:11, mean_IB)
xticklabels(groups)

```

```
xtickangle(60)
axis([0 12 1 6])
yticklabels(answers)
yticks([1:6])
yticklabels(answers)
print(gcf, '/Users/dianaperez/Desktop/observe_imp_bias.jpg', '-dpng', '-r300');
imp_bias2 = DOI(:, 12:22);
for q = 1:width(imp_bias2)
col = table2array(imp_bias2(~isnan(imp_bias2(:,q)),q));
mean_IB2(q) = mean(col);
end
bar(1:11,mean_IB2)
xticklabels(groups)
xtickangle(60)
axis([0 12 1 6])
yticklabels(answers)
ylabel('Average Response')
print(gcf, '/Users/dianaperez/Desktop/hear_imp_bias.jpg', '-dpng', '-r300');
yticks([1:6])
imp_bias2 = DOI(:, 12:22);
for q = 1:width(imp_bias2)
col = table2array(imp_bias2(~isnan(imp_bias2(:,q)),q));
mean_IB2(q) = mean(col);
end
bar(1:11,mean_IB2)
xticklabels(groups)
xtickangle(60)
axis([0 12 1 6])
yticks([1:6])
yticklabels(answers)
ylabel('Average Response')
print(gcf, '/Users/dianaperez/Desktop/hear_imp_bias.jpg', '-dpng', '-r300');
sources = DOI(:,38);
height(sources)
cell2mat(sources{1})
cell2mat(sources{4})
sources{4}
str2num(sources{4})
sources = DOI(:,38);
sources_col = [];
for s = 1:height(sources)
resp = str2num(sources{s});
sources_col = [sources_col; resp'];
end
sources = DOI(:,38);
sources_col = [];
count = 0;
for s = 1:height(sources)
resp = str2num(sources{s});
sources_col = [sources_col; resp'];
if ~isempty(resp)
count = count + 1;
end
end
max(sources_col)
for a = 1:max(sources_col)
perc = length(find(sources_col==a))/count;
end
for a = 1:max(sources_col)
perc(a) = length(find(sources_col==a))/count;
```

```
end
bar(1:11,perc)
ylabel('% responses')
print(gcf, '/Users/dianaperez/Desktop/sources_imp_bias.jpg', '-dpng', '-r300');
answers = {'Never', 'Once a year', 'Once/twice a quarter', 'Severla times', 'Weekly', 'Daily'};
exp_bias = DOI(:, 41:51);
for q = 1:width(exp_bias)
col = table2array(exp_bias(~isnan(exp_bias{:,q}),q));
mean_EB(q) = mean(col);
end
bar(1:11,mean_EB)
xticklabels(groups)
xtickangle(60)
axis([0 12 1 6])
yticks([1:6])
yticklabels(answers)
ylabel('Average Response')
print(gcf, '/Users/dianaperez/Desktop/observe_exp_bias.jpg', '-dpng', '-r300');
exp_bias2 = DOI(:, 54:64);
for q = 1:width(exp_bias2)
col = table2array(exp_bias2(~isnan(exp_bias2{:,q}),q));
mean_EB2(q) = mean(col);
end
bar(1:11,mean_EB2)
xticklabels(groups)
xtickangle(60)
axis([0 12 1 6])
yticks([1:6])
yticklabels(answers)
ylabel('Average Response')
print(gcf, '/Users/dianaperez/Desktop/hear_exp_bias.jpg', '-dpng', '-r300');
sources = DOI{:,67};
sources_col = [];
count = 0;
for s = 1:height(sources)
resp = str2num(sources{s});
sources_col = [sources_col; resp'];
if ~isempty(resp)
count = count + 1;
end
end
for a = 1:max(sources_col)
perc(a) = length(find(sources_col==a))/count;
end
bar(1:11,perc)
ylabel('% responses')
print(gcf, '/Users/dianaperez/Desktop/sources_exp_bias.jpg', '-dpng', '-r300');
sources = DOI{:,67};
sources_col = [];
count = 0;
for s = 1:height(sources)
resp = str2num(sources{s});
sources_col = [sources_col; resp'];
if ~isempty(resp)
count = count + 1;
end
end
perc = [];
for a = 1:max(sources_col)
```

```
perc(a) = length(find(sources_col==a))/count;
end
bar(1:11,perc)
ylabel('% responses')
contxt = DOI{:,70};
contxt_col = [];
count = 0;
for s = 1:height(contxt)
    resp = str2num(contxt{s});
    sources_col = [contxts_col; resp'];
    if ~isempty(resp)
        count = count + 1;
    end
end
perc = [];
for a = 1:max(contxt_col)
    perc(a) = length(find(contxt_col==a))/count;
end
bar(1:11,perc)
ylabel('% responses')
contxt = DOI{:,70};
contxt_col = [];
count = 0;
for s = 1:height(contxt)
    resp = str2num(contxt{s});
    sources_col = [contxt_col; resp'];
    if ~isempty(resp)
        count = count + 1;
    end
end
perc = [];
for a = 1:max(contxt_col)
    perc(a) = length(find(contxt_col==a))/count;
end
bar(1:11,perc)
ylabel('% responses')
bar(1:a,perc)
resp = str2num(contxt{1});
resp = str2num(contxt{4});
contxt = DOI{:,70};
contxt_col = [];
count = 0;
for s = 1:height(contxt)
    resp = str2num(contxt{s});
    contxt_col = [contxt_col; resp'];
    if ~isempty(resp)
        count = count + 1;
    end
end
perc = [];
for a = 1:max(contxt_col)
    perc(a) = length(find(contxt_col==a))/count;
end
bar(1:a,perc)
axis([0 12 0 .6])
ylabel('% responses')
print(gcf, '/Users/dianaperez/Desktop/context_bias.jpg', '-dpng', '-r300');
rec = DOI{:,73};
rec_col = [];
count = 0;
```

```
for s = 1:height(rec)
    resp = str2num(rec{s});
    rec_col = [rec_col; resp'];
    if ~isempty(resp)
        count = count + 1;
    end
end
perc = [];
for a = 1:max(rec_col)
    perc(a) = length(find(rec_col==a))/count;
end
bar(1:a,perc)
axis([0 12 0 .6])
ylabel('% responses')
axis([0 12 0 1])
print(gcf, '/Users/dianaperez/Desktop/what_can_dpt_do.jpg', '-dpng', '-r300');
%-- 9/1/22, 11:29 AM --%
DIC_data
imp_bias2 = DOI(:, 25:35);
for q = 1:width(imp_bias2)
    col = table2array(imp_bias2(~isnan(imp_bias2{:,q}),q));
    mean_IB2(q) = mean(col);
end
bar(1:11,mean_IB2)
xticklabels(groups)
xtickangle(60)
axis([0 12 1 6])
yticks([1:6])
yticklabels(answers)
ylabel('Average Response')
print(gcf, '/Users/dianaperez/Desktop/hear_imp_bias.jpg', '-dpng', '-r300');
sources = DOI{:,67};
sources_col = [];
count = 0;
for s = 1:height(sources)
    resp = str2num(sources{s});
    sources_col = [sources_col; resp'];
    if ~isempty(resp)
        count = count + 1;
    end
end
perc = [];
for a = 1:max(sources_col)
    perc(a) = length(find(sources_col==a))/count;
end
bar(1:11,perc)
ylabel('% responses')
xticklabels(sources_leg)
xtickangle(60)
sources_leg = {'grad students', 'post-docs', 'undergrads', 'faculty', 'staff', 'area
members', 'advisor', 'dept policies', 'dept/area culture', 'other', 'none'};
ylabel('% responses')
xticklabels(sources_leg)
xtickangle(60)
sources_leg = {'grad students', 'post-docs', 'undergrads', 'faculty', 'staff', 'area
members', 'advisor', 'dept policies', 'dept/area culture', 'other', 'none'};
sources = DOI{:,38};
sources_col = [];
count = 0;
for s = 1:height(sources)
```

```

resp = str2num(sources{s});
sources_col = [sources_col; resp'];
if ~isempty(resp)
count = count + 1;
end
end
for a = 1:max(sources_col)
perc(a) = length(find(sources_col==a))/count;
end
bar(1:11,perc)
ylabel('% responses')
xticklabels(sources_leg)
xtickangle(60)
print(gcf, '/Users/dianaperez/Desktop/sources_imp_bias.jpg', '-dpng', '-r300');
sources = DOI{:,67};
sources_col = [];
count = 0;
for s = 1:height(sources)
resp = str2num(sources{s});
sources_col = [sources_col; resp'];
if ~isempty(resp)
count = count + 1;
end
end
perc = [];
for a = 1:max(sources_col)
perc(a) = length(find(sources_col==a))/count;
end
bar(1:11,perc)
ylabel('% responses')
xticklabels(sources_leg)
xtickangle(60)
print(gcf, '/Users/dianaperez/Desktop/sources_exp_bias.jpg', '-dpng', '-r300');
[h, mu, sigma, q, notch] = al_goodplot(enc, [], [ 0    0.4470    0.7410], 'left');
addpath '/Users/dianaperez/Documents/GitHub/PerezEtAl_HemAsymmetries/needed_files/upload'
[h, mu, sigma, q, notch] = al_goodplot(enc, [], [ 0    0.4470    0.7410], 'left');
[h, mu, sigma, q, notch] = al_goodplot(enc{:, :}, [], [ 0    0.4470    0.7410], 'left');
[h, mu, sigma, q, notch] = al_goodplot(enc{:, :}, [], [ 0    0.4470    0.7410], 'left');
enc = table2array(DOI(~isnan(DOI{:,1}), 1:11));
[h, mu, sigma, q, notch] = al_goodplot(enc, [], [ 0    0.4470    0.7410], 'left');
enc1 = enc(:,1);
[h, mu, sigma, q, notch] = al_goodplot(enc1, [], [ 0    0.4470    0.7410], 'left');
[h, mu, sigma, q, notch] = al_goodplot(enc1, 0.1, .2, [ 0    0.4470    0.7410], 'left');
[h, mu, sigma, q, notch] = al_goodplot(enc, 0.1, .2, [ 0    0.4470    0.7410], 'left');
[h, mu, sigma, q, notch] = al_goodplot(enc, [0.1:1:11.1], .2, [ 0    0.4470    0.7410], 'left');
[h, mu, sigma, q, notch] = al_goodplot(enc, [0.1:1:11.1], .5, [ 0    0.4470    0.7410], 'left');
axis([-1 12 1 6])
axis([-1 11 1 5])
groups = {'Race/Ethnicity', 'Gender', 'Sexual Orientation', 'Economic Status', 'Ability Status', 'Age', 'Religion', 'Familial Obligations', 'Political Ideology', 'Nationality/Immigration', 'Mental Health'};
answers = {'Not at all', 'A little', 'A moderate amount', 'Quite a bit', 'A great deal'};
enc = table2array(DOI(~isnan(DOI{:,1}), 1:11));
%mean_enc = mean(table2array(enc), 1);
%bar(1:11,mean_enc)
[h, mu, sigma, q, notch] = al_goodplot(enc, [0.1:1:11.1], .5, [ 0    0.4470    0.7410], 'bilateral');
xticklabels(groups)

```

```

[h, mu, sigma, q, notch] = al_goodplot(enc,[1:1:11],.5,[ 0    0.4470
0.7410], 'bilateral');
xticks([1:11])
yticklabels(answers)
yticks([1:5])
xticklabels(groups)
xtickangle(60)
print(gcf, '/Users/dianaperez/Desktop/encouragement.jpg', '-dpng', '-r300');
al_goodplot([], 1, .5, [], 'man')
[h, mu, sigma, q, notch] = al_goodplot(enc,[1:1:11],.5,[ 0    0.4470
0.7410], 'bilateral');
xticklabels(groups)
xtickangle(60)
axis([0 12 1 5])
yticklabels(answers)
ylabel('Average Response')
xticks([1:11])
[h, mu, sigma, q, notch] = al_goodplot(enc,[1:1:11],.5,[ 0    0.4470
0.7410], 'bilateral');
xticks([1:11])
xticklabels(groups)
xtickangle(60)
axis([0 12 1 5])
yticks([1:5])
yticklabels(answers)
ylabel('Average Response')
axis([0 12 .5 5.5])
print(gcf, '/Users/dianaperez/Desktop/encouragement.jpg', '-dpng', '-r300');
answers = {'Never', 'Once a year', 'Once/twice a quarter', 'Several times', 'Weekly',
'Daily'};
imp_bias = DOI(:, 12:22);
for q = 1:width(imp_bias)
col(:,q) = table2array(imp_bias(~isnan(imp_bias(:,q)),q));
mean_IB(q) = mean(col(:,q));
end
for q = 1:width(imp_bias)
col = table2array(imp_bias(~isnan(imp_bias(:,q)),q));
mean_IB(q) = mean(col);
[h, mu, sigma, q, notch] = al_goodplot(col,q,.5,[ 0    0.4470    0.7410], 'left');
hold on
end
xticks([1:11])
xticklabels(groups)
xtickangle(60)
axis([0 12 .5 6.5])
yticks([1:6])
DIC_data
dist(col)
figure
hist(col)
DIC_data
answers = {'Never', 'Once a year', 'Once/twice a quarter', 'Several times', 'Weekly',
'Daily'};
imp_bias = DOI(:, 12:22);
for q = 1:width(imp_bias)
col = table2array(imp_bias(~isnan(imp_bias(:,q)),q));
mean_IB(q) = mean(col);
[h, mu, sigma, q, notch] = al_goodplot(col,q,.2,[ 0    0.4470    0.7410], 'left', .5,.
0011);
hold on

```

```

end
%bar(1:11,mean_IB)
xticks([1:11])
xticklabels(groups)
xtickangle(60)
axis([0 12 .5 6.5])
yticks([1:6])
yticklabels(answers)
% xticklabels(groups)
% xtickangle(60)
% axis([0 12 1 6])
% yticks([1:6])
% yticklabels(answers)
% ylabel('Average Response')
% observe explicit bias
answers = {'Never', 'Once a year', 'Once/twice a quarter', 'Severla times', 'Weekly',↵
'Daily'};
exp_bias = DOI(:, 41:51);
for q = 1:width(exp_bias)
col = table2array(exp_bias(~isnan(exp_bias{:,q}),q));
mean_EB(q) = mean(col);
[h, mu, sigma, q, notch] = al_goodplot(col,q,.2,[0.8500    0.3250    0.0980], 'right', .↵
5,.0011);
end
figure
imp_bias = DOI(:, 12:22);
for q = 1:width(imp_bias)
col = table2array(imp_bias(~isnan(imp_bias{:,q}),q));
mean_IB(q) = mean(col);
[h, mu, sigma, q, notch] = al_goodplot(col,q,.3,[ 0    0.4470    0.7410], 'left', .5,↵
0011);
hold on
end
%bar(1:11,mean_IB)
xticks([1:11])
xticklabels(groups)
xtickangle(60)
axis([0 12 .5 6.5])
yticks([1:6])
yticklabels(answers)
% xticklabels(groups)
% xtickangle(60)
% axis([0 12 1 6])
% yticks([1:6])
% yticklabels(answers)
% ylabel('Average Response')
% observe explicit bias
answers = {'Never', 'Once a year', 'Once/twice a quarter', 'Severla times', 'Weekly',↵
'Daily'};
exp_bias = DOI(:, 41:51);
for q = 1:width(exp_bias)
col = table2array(exp_bias(~isnan(exp_bias{:,q}),q));
mean_EB(q) = mean(col);
[h, mu, sigma, q, notch] = al_goodplot(col,q,.3,[0.8500    0.3250    0.0980], 'right', .↵
5,.0011);
end
figure
imp_bias = DOI(:, 12:22);
for q = 1:width(imp_bias)
col = table2array(imp_bias(~isnan(imp_bias{:,q}),q));

```



```

mean_IB(q) = mean(col);
[h, mu, sigma, q, notch] = al_goodplot(col,(q-.1),.3,[ 0    0.4470    0.7410],'left', .5,.0011);
hold on
end
%bar(1:11,mean_IB)
xticks([1:11])
xticklabels(groups)
xtickangle(60)
axis([0 12 .5 6.5])
yticks([1:6])
yticklabels(answers)
% xticklabels(groups)
% xtickangle(60)
% axis([0 12 1 6])
% yticks([1:6])
% yticklabels(answers)
% ylabel('Average Response')
% observe explicit bias
answers = {'Never', 'Once a year', 'Once/twice a quarter', 'Severla times', 'Weekly',
'Daily'};
exp_bias = DOI(:, 41:51);
for q = 1:width(exp_bias)
col = table2array(exp_bias(~isnan(exp_bias{:,q}),q));
mean_EB(q) = mean(col);
[h, mu, sigma, q, notch] = al_goodplot(col,(q+.1),.3,[0.8500    0.3250
0.0980],'right', .5,.0011);
end
figure; imp_bias = DOI(:, 12:22);
for q = 1:width(imp_bias)
col = table2array(imp_bias(~isnan(imp_bias{:,q}),q));
mean_IB(q) = mean(col);
[h, mu, sigma, q, notch] = al_goodplot(col,(q-.01),.3,[ 0    0.4470    0.7410],'left', .5,.0011);
hold on
end
%bar(1:11,mean_IB)
xticks([1:11])
xticklabels(groups)
xtickangle(60)
axis([0 12 .5 6.5])
yticks([1:6])
yticklabels(answers)
% xticklabels(groups)
% xtickangle(60)
% axis([0 12 1 6])
% yticks([1:6])
% yticklabels(answers)
% ylabel('Average Response')
% observe explicit bias
answers = {'Never', 'Once a year', 'Once/twice a quarter', 'Severla times', 'Weekly',
'Daily'};
exp_bias = DOI(:, 41:51);
for q = 1:width(exp_bias)
col = table2array(exp_bias(~isnan(exp_bias{:,q}),q));
mean_EB(q) = mean(col);
[h, mu, sigma, q, notch] = al_goodplot(col,(q+.01),.3,[0.8500    0.3250
0.0980],'right', .5,.0011);
end
figure; imp_bias = DOI(:, 12:22);

```

```

for q = 1:width(imp_bias)
col = table2array(imp_bias(~isnan(imp_bias(:,q)),q));
mean_IB(q) = mean(col);
[h, mu, sigma, q, notch] = al_goodplot(col,(q-.02),.3,[ 0    0.4470    0.7410], 'left', .5, .0011);
hold on
end
%bar(1:11,mean_IB)
xticks([1:11])
xticklabels(groups)
xtickangle(60)
axis([0 12 .5 6.5])
yticks([1:6])
yticklabels(answers)
% xticklabels(groups)
% xtickangle(60)
% axis([0 12 1 6])
% yticks([1:6])
% yticklabels(answers)
% ylabel('Average Response')
% observe explicit bias
answers = {'Never', 'Once a year', 'Once/twice a quarter', 'Severla times', 'Weekly', 'Daily'};
exp_bias = DOI(:, 41:51);
for q = 1:width(exp_bias)
col = table2array(exp_bias(~isnan(exp_bias(:,q)),q));
mean_EB(q) = mean(col);
[h, mu, sigma, q, notch] = al_goodplot(col,(q+.02),.3,[0.8500    0.3250    0.0980], 'right', .5, .0011);
end
print(gcf, '/Users/dianaperez/Desktop/observe_imp_exp_bias.jpg', '-dpng', '-r300');
imp_bias2 = DOI(:, 25:35);
for q = 1:width(imp_bias2)
col = table2array(imp_bias2(~isnan(imp_bias2(:,q)),q));
mean_IB2(q) = mean(col);
[h, mu, sigma, q, notch] = al_goodplot(col,(q-.02),.3,[ 0    0.4470    0.7410], 'left', .5, .0011);
hold on
end
%bar(1:11,mean_IB2)
xticks([1:11])
xticklabels(groups)
xtickangle(60)
axis([0 12 .5 6.5])
yticks([1:6])
yticklabels(answers)
%print(gcf, '/Users/dianaperez/Desktop/hear_imp_bias.jpg', '-dpng', '-r300');
% hear of explicit bias
exp_bias2 = DOI(:, 54:64);
for q = 1:width(exp_bias2)
col = table2array(exp_bias2(~isnan(exp_bias2(:,q)),q));
mean_EB2(q) = mean(col);
[h, mu, sigma, q, notch] = al_goodplot(col,(q+.02),.3,[0.8500    0.3250    0.0980], 'right', .5, .0011);
end
print(gcf, '/Users/dianaperez/Desktop/hear_imp_exp_bias.jpg', '-dpng', '-r300');
sources_leg = {'grad students', 'post-docs', 'undergrads', 'faculty', 'staff', 'area members', 'advisor', 'dept policies', 'dept/area culture', 'other', 'none'};
sources = DOI(:,38);
sources_col = [];

```

```

count = 0;
for s = 1:height(sources)
    resp = str2num(sources{s});
    sources_col = [sources_col; resp'];
    if ~isempty(resp)
        count = count + 1;
    end
end
perc1=[];
for a = 1:max(sources_col)
    perc1(a) = length(find(sources_col==a))/count;
end
% bar(1:11,perc)
% ylabel('% responses')
% xticklabels(sources_leg)
% xtickangle(60)
% print(gcf,'/Users/dianaperez/Desktop/sources_imp_bias.jpg','-dpng','-r300');
% sources of explicit bias
sources = DOI{:,67};
sources_col = [];
count = 0;
for s = 1:height(sources)
    resp = str2num(sources{s});
    sources_col = [sources_col; resp'];
    if ~isempty(resp)
        count = count + 1;
    end
end
perc2 = [];
for a = 1:max(sources_col)
    perc2(a) = length(find(sources_col==a))/count;
end
bar(1:11, [perc1 perc2])
bar(1:11, [perc1; perc2])
width(50)
bar(1:11, [perc1; perc2], 'width', 50)
bar(1:11, [perc1; perc2])
h = bar(1:11, [perc1; perc2])
h.width = 1
h = bar(1:11, [perc1; perc2])
ylabel('% responses')
xticklabels(sources_leg)
xtickangle(60)
print(gcf,'/Users/dianaperez/Desktop/sources_exp_bias.jpg','-dpng','-r300');
print(gcf,'/Users/dianaperez/Desktop/sources_imp_exp_bias.jpg','-dpng','-r300');
m = findobj(gca, 'Type', 'Bar');
hleg1 = legend(m(1:2), 'Explicit Bias', 'Implicit Bias', 'Location', 'NorthEast');
print(gcf,'/Users/dianaperez/Desktop/sources_imp_exp_bias.jpg','-dpng','-r300');
%-- 9/18/22, 10:25 PM --%
fcimage_corrmat_volume
('/Users/dianaperez/Desktop/Research/Lifespan/datalists/lifespan_datalist_LS03.xlsx',
'/Volumes/fsmresfiles/PBS/Gratton_Lab/Lifespan/Post-
COVID/BIDS/derivatives/preproc_FCProc/', 'Seitzman300');
addpath '/Users/dianaperez/Documents/GitHub/GrattonLab-General-Repo/FCProcess'
fcimage_corrmat_volume
('/Users/dianaperez/Desktop/Research/Lifespan/datalists/lifespan_datalist_LS03.xlsx',
'/Volumes/fsmresfiles/PBS/Gratton_Lab/Lifespan/Post-
COVID/BIDS/derivatives/preproc_FCProc/', 'Seitzman300');
FCPROCESS_GrattonLab
('/Users/dianaperez/Desktop/Research/Lifespan/datalists/lifespan_datalist_LS03.xlsx',

```

```

'/Users/dianaperez/Desktop/', 'defaults2')
fcimage_corrmat_volume
fcimage_corrmat_volume↵
('/Users/dianaperez/Desktop/Research/Lifespan/datalists/lifespan_datalist_LS03.xlsx',↵
'/Volumes/fsmresfiles/PBS/Gratton_Lab/Lifespan/Post-↵
COVID/BIDS/derivatives/preproc_FCProc/', 'Seitzman300');
cb_vox = find(roi_data==7);
cb_vox_inds = find(roi_data==7);
cb_vox = sess_data(cb_vox_inds,:);
%-- 9/20/22, 9:01 AM --%
help nnz
%-- 10/6/22, 5:39 PM --%
data = readtable('/Users/dianaperez/Downloads/DIC_Data2022.csv');
DOI = data(:,18:90);
imp_bias = [DOI(:,12) DOI(:,13) DOI(:,15) DOI(:,16) DOI(:,21)];
groups = {'Race/Ethnicity', 'Gender', 'SES', 'Ability Status', 'Mental Health'};
answers = {'Not at all', 'A little', 'A moderate amount', 'Quite a bit', 'A great deal'};
answers = {'Never', 'Once a year', 'Once/twice a quarter', 'Several times', 'Weekly',↵
'Daily'};
imp_bias = [DOI(:,12) DOI(:,13) DOI(:,15) DOI(:,16) DOI(:,21)];
for q = 1:width(imp_bias)
col = table2array(imp_bias(~isnan(imp_bias{:,q}),q));
mean_IB(q) = mean(col);
[h, mu, sigma, q, notch] = al_goodplot(col,(q-.02),.3,[ 0    0.4470    0.7410], 'left', .↵
5,.0011);
hold on
end
addpath '/Users/dianaperez/Documents/GitHub/PerezEtAl_HemAsymmetries/needed_files/upload'
imp_bias = [DOI(:,12) DOI(:,13) DOI(:,15) DOI(:,16) DOI(:,21)];
for q = 1:width(imp_bias)
col = table2array(imp_bias(~isnan(imp_bias{:,q}),q));
mean_IB(q) = mean(col);
[h, mu, sigma, q, notch] = al_goodplot(col,(q-.02),.3,[ 0    0.4470    0.7410], 'left', .↵
5,.0011);
hold on
end
exp_bias = [DOI(:,25:26) DOI(:,28:29) DOI(:,35)];
for q = 1:width(exp_bias)
col = table2array(exp_bias(~isnan(exp_bias{:,q}),q));
mean_EB(q) = mean(col);
[h, mu, sigma, q, notch] = al_goodplot(col,(q+.02),.3,[0.8500    0.3250↵
0.0980], 'right', .5,.0011);
end
xticklabels(groups)
xtickangle(60)
xticks([1:2:10])
xticks([1:5])
yticklabels(answers)
yticks([1:6])
ylabel('Average Response')
print(gcf, '/Users/dianaperez/Desktop/observe_imp_exp_bias_CONDENSED.jpg', '-dpng', '-↵
r300');
title('Experienced implicit and explicit bias')
print(gcf, '/Users/dianaperez/Desktop/observe_imp_exp_bias_CONDENSED.jpg', '-dpng', '-↵
r300');
for q = 1:width(imp_bias)
col = table2array(imp_bias(~isnan(imp_bias{:,q}),q));
mean_IB(q) = mean(col);
[h, mu, sigma, q, notch] = al_goodplot(col,(q-.02),.3,[ 0    0.4470    0.7410], 'left', .↵
5,.0011);

```

```

hold on
end
for q = 1:width(exp_bias)
col = table2array(exp_bias(~isnan(exp_bias(:,q)),q));
mean_EB(q) = mean(col);
[h, mu, sigma, q, notch] = al_goodplot(col,(q+.02),.3,[0.8500    0.3250
0.0980], 'right', .5,.0011);
end
xticklabels(groups)
xtickangle(60)
axis([0 12 1 6])
yticks([1:6])
yticklabels(answers)
axis([0 6 1 6])
axis([0 5 1 6])
axis([0 6 1 6])
axis([0 5.5 1 6])
axis([0.5 5.5 1 6])
xticks([1:5])
yticks([1:6])
title('How often do you hear of implicit/explicit bias?')
print(gcf, '/Users/dianaperez/Desktop/observe_imp_exp_bias_CONDENSED.jpg', '-dpng', '-r300');
close gcf
imp_bias2 = [DOI(:,41:42) DOI(:,44:45) DOI(:,51)];
for q = 1:width(imp_bias2)
col = table2array(imp_bias2(~isnan(imp_bias2(:,q)),q));
mean_IB2(q) = mean(col);
[h, mu, sigma, q, notch] = al_goodplot(col,(q-.02),.3,[ 0    0.4470    0.7410], 'left', .5,.0011);
hold on
end
exp_bias2 = [DOI(:,54:55) DOI(:,57:58) DOI(:,64)];
for q = 1:width(exp_bias2)
col = table2array(exp_bias2(~isnan(exp_bias2(:,q)),q));
mean_EB2(q) = mean(col);
[h, mu, sigma, q, notch] = al_goodplot(col,(q+.02),.3,[0.8500    0.3250
0.0980], 'right', .5,.0011);
end
for q = 1:width(imp_bias2)
col = table2array(imp_bias2(~isnan(imp_bias2(:,q)),q));
mean_IB2(q) = mean(col);
[h, mu, sigma, q, notch] = al_goodplot(col,(q-.02),.3,[ 0    0.4470    0.7410], 'left', .5,.0011);
hold on
end
for q = 1:width(exp_bias2)
col = table2array(exp_bias2(~isnan(exp_bias2(:,q)),q));
mean_EB2(q) = mean(col);
[h, mu, sigma, q, notch] = al_goodplot(col,(q+.02),.3,[0.8500    0.3250
0.0980], 'right', .5,.0011);
end
xticks([1:5])
yticks([1:6])
xticklabels(groups)
xtickangle(60)
yticklabels(answers)
title('How often do you observe/experience implicit/explicit bias?')
print(gcf, '/Users/dianaperez/Desktop/observe_imp_exp_bias_CONDENSED2.jpg', '-dpng', '-r300');

```

```

imp_bias = [DOI(:,12) DOI(:,13) DOI(:,15) DOI(:,16) DOI(:,21)];
for q = 1:width(imp_bias)
col = table2array(imp_bias(~isnan(imp_bias(:,q)),q));
mean_IB(q) = mean(col);
[h, mu, sigma, q, notch] = al_goodplot(col,(q-.02),.3,[ 0      0.4470      0.7410],'left', .5, .0011);
hold on
end
exp_bias = [DOI(:,41:42) DOI(:,44:45) DOI(:,51)];%
for q = 1:width(exp_bias)
col = table2array(exp_bias(~isnan(exp_bias(:,q)),q));
mean_EB(q) = mean(col);
[h, mu, sigma, q, notch] = al_goodplot(col,(q+.02),.3,[0.8500      0.3250
0.0980],'right', .5, .0011);
end
imp_bias = [DOI(:,12) DOI(:,13) DOI(:,15) DOI(:,16) DOI(:,21)];
for q = 1:width(imp_bias)
col = table2array(imp_bias(~isnan(imp_bias(:,q)),q));
mean_IB(q) = mean(col);
[h, mu, sigma, q, notch] = al_goodplot(col,(q-.02),.3,[ 0      0.4470      0.7410],'left', .5, .0011);
hold on
end
exp_bias = [DOI(:,41:42) DOI(:,44:45) DOI(:,51)];%
for q = 1:width(exp_bias)
col = table2array(exp_bias(~isnan(exp_bias(:,q)),q));
mean_EB(q) = mean(col);
[h, mu, sigma, q, notch] = al_goodplot(col,(q+.02),.3,[0.8500      0.3250
0.0980],'right', .5, .0011);
end
xticks([1:5])
xticklabels(groups)
xtickangle(60)
yticks([1:6])
yticklabels(answers)
ax = gca;
ax.FontSize = 20;
ax.FontSize = 15;
print(gcf, '/Users/dianaperez/Desktop/observe_imp_exp_bias_CONDENSED.jpg', '-dpng', '-r300')
imp_bias2 = [DOI(:,25:26) DOI(:,28:29) DOI(:,35)];
for q = 1:width(imp_bias2)
col = table2array(imp_bias2(~isnan(imp_bias2(:,q)),q));
mean_IB2(q) = mean(col);
[h, mu, sigma, q, notch] = al_goodplot(col,(q-.02),.3,[ 0      0.4470      0.7410],'left', .5, .0011);
hold on
end
exp_bias2 = [DOI(:,54:55) DOI(:,57:58) DOI(:,64)];
for q = 1:width(exp_bias2)
col = table2array(exp_bias2(~isnan(exp_bias2(:,q)),q));
mean_EB2(q) = mean(col);
[h, mu, sigma, q, notch] = al_goodplot(col,(q+.02),.3,[0.8500      0.3250
0.0980],'right', .5, .0011);
end
xticks([1:5])
yticks([1:6])
xticklabels(groups)
xtickangle(60)
yticklabels(answers)
ax = gca;

```

```
ax.FontSize = 15;
print(gcf, '/Users/dianaperez/Desktop/hear_imp_exp_bias_CONDENSED.jpg', '-dpng', '-r300')
sources_leg = {'grad students', 'post-docs', 'undergrads', 'faculty', 'staff', 'area  
members', 'advisor', 'dept policies', 'dept/area culture', 'other', 'none'};
sources = DOI{:,38};
sources_col = [];
count = 0;
for s = 1:height(sources)
    resp = str2num(sources{s});
    sources_col = [sources_col; resp'];
    if ~isempty(resp)
        count = count + 1;
    end
end
perc1=[];
for a = 1:max(sources_col)
    perc1(a) = length(find(sources_col==a))/count;
end
sources = DOI{:,67};
sources_col = [];
count = 0;
for s = 1:height(sources)
    resp = str2num(sources{s});
    sources_col = [sources_col; resp'];
    if ~isempty(resp)
        count = count + 1;
    end
end
perc2 = [];
for a = 1:max(sources_col)
    perc2(a) = length(find(sources_col==a))/count;
end
perc = [perc1; perc2];
avg = mean(perc,2)
avg = mean(perc,1)
sortrows(avg')
sortrows(avg', 'descend')
ind = sortrows(avg', 'descend');
[avg_sorted, ind] = sortrows(avg', 'descend');
perc1(ind)
perc1=perc1(ind);
perc2=perc2(ind);
perc = [perc1; perc2];
sources_leg{ind}
sources_leg{end} = [];
sources_leg(end) = [];
sources = DOI{:,38};
sources_col = [];
count = 0;
for s = 1:height(sources)
    resp = str2num(sources{s});
    sources_col = [sources_col; resp'];
    if ~isempty(resp)
        count = count + 1;
    end
end
perc1=[];
for a = 1:max(sources_col)
    perc1(a) = length(find(sources_col==a))/count;
end
```

```

sources = DOI{:,67};
sources_col = [];
count = 0;
for s = 1:height(sources)
    resp = str2num(sources{s});
    sources_col = [sources_col; resp'];
    if ~isempty(resp)
        count = count + 1;
    end
end
perc2 = [];
for a = 1:max(sources_col)
    perc2(a) = length(find(sources_col==a))/count;
end
perc1(end) = [];
perc2(end) = [];
avg = mean(perc,1)
[avg_sorted, ind] = sortrows(avg, 'descend');
perc1=perc1(ind);
perc = [perc1; perc2];
avg = mean(perc,1)
[avg_sorted, ind] = sortrows(avg, 'descend');
perc1=perc1(ind);
perc2=perc2(ind);
perc = [perc1; perc2];
barh(1:10, perc)
flip(perc)
flip(perc,1)
flip(perc,2)
barh(1:10, flip(perc,2))
xlabel('% responses')
yticklabels(sources_leg{ind})
sources_leg_sorted = sources_leg{ind};
yticklabels(sources_leg_sorted)
yticks(1:10)
yticklabels(sources_leg_sorted)
sources_leg{ind}
sources_leg_sorted = {'faculty', 'other', 'grad students', 'advisor', 'dept policies',
'undergrads', 'area members', 'post-docs', 'dept/area culture', 'staff'};
yticklabels(sources_leg_sorted)
sources_leg_sorted = {'staff', 'dept/area culture', 'post-docs', 'area members',
'undergrads', 'dept policies', 'advisor', 'grad students', 'other', 'faculty'};
yticklabels(sources_leg_sorted)
ax = gca;
ax.FontSize = 14;
barh(1:10, perc, .7)
barh(1:10, perc, 1.2)
barh(1:10, perc, 1)
barh(1:10, flip(perc,2), 1)
xlabel('% responses')
yticklabels(sources_leg_sorted)
ax.FontSize = 14;
print(gcf, '/Users/dianaperez/Desktop/sources_exp_imp_bias_horizontal.jpg', '-dpng', '-r300')
bar(1:10, perc)
bar(1:10, perc, 1)
ylabel('% responses')
xticklabels(sources_leg_sorted)
xticklabels(flip(sources_leg_sorted))
xtickangle(60)

```



```
ax.FontSize = 14;
print(gcf, '/Users/dianaperez/Desktop/sources_exp_imp_bias.jpg', '-dpng', '-r300')
rec = DOI{:,73};
rec_col = [];
count = 0;
for s = 1:height(rec)
    resp = str2num(rec{s});
    rec_col = [rec_col; resp'];
    if ~isempty(resp)
        count = count + 1;
    end
end
perc = [];
for a = 1:max(rec_col)
    perc(a) = length(find(rec_col==a))/count;
end
[recs_sorted, ind] = sortrows(perc, 'descend');
%-- 10/11/22, 5:15 PM --%
load_nii_wrapper
addpath '/Users/dianaperez/Documents/GitHub/GrattonLab-General-Repo/FCProcess'
uiopen('/Users/dianaperez/Desktop/tpl-MNI152Nlin2009cAsym_res-01_desc-  
BrainCerebellumExtraction_mask.nii.gz',1)
load_nii_wrapper('/Users/dianaperez/Desktop/tpl-MNI152Nlin2009cAsym_res-01_desc-  
BrainCerebellumExtraction_mask.nii.gz')
cd '/Users/dianaperez/Documents/GitHub/GrattonLab-General-Repo/FCProcess'
load_nii_wrapper('/Users/dianaperez/Desktop/tpl-MNI152Nlin2009cAsym_res-01_desc-  
BrainCerebellumExtraction_mask.nii.gz')
load_nii_wrapper('/Users/dianaperez/Desktop/tpl-MNI152Nlin2009cAsym_res-01_desc-  
BrainCerebellumExtraction_mask.nii.gz');
load_nii_wrapper('/Users/dianaperez/Downloads/tpl-MNI152Nlin2009cAsym_res-01_desc-  
carpet_dseg.nii.gz');
nifti = ans;
unique(nifti)
uiopen('/Volumes/fsmresfiles/PBS/Gratton_Lab/Lifespan/Post-  
COVID/BIDS/derivatives/preproc_fmriprep-20.2.0/fmriprep/sub-LS03/ses-1/func/sub-LS03_ses-  
1_task-rest_run-1_space-MNI152Nlin6Asym_res-2_desc-brain_mask.nii.gz',1)
load_nii_wrapper('/Users/dianaperez/Desktop/Buckner_CBmask_resampled.nii.gz')
load_nii_wrapper('/Users/dianaperez/Desktop/Buckner_CBmask_resampled.nii.gz')
load_nii_wrapper('/Users/dianaperez/Desktop/Buckner_CBmask_resampled.nii.gz');
buck_mask = ans;
unique(buck_mask)
new_mask = zeros(buck_mask);
new_mask = logical(buck_mask);
unique(new_mask)
help save_nii
save_nii(new_mask, '/Users/dianaperez/Desktop/Buckner_CBmask_resampled_binary.nii.gz')
new_mask = single(logical(buck_mask));
save_nii(new_mask, '/Users/dianaperez/Desktop/Buckner_CBmask_resampled_binary.nii.gz')
%-- 10/12/22, 12:57 PM --%
atlas = 'Seitzman300';
FCdir = '/Users/dianaperez/Desktop/preprofc_FCProc';
outDir_top = [FCdir '/cormats_' atlas '/'];
if ~exist(outDir_top)
    mkdir(outDir_top);
end
clear all
fcimage_cormat_volume('/Users/dianaperez/Desktop/preprofc_FCProc/lifespan_datalist_LS03.  
xlsx', '/Users/dianaperez/Desktop/preprofc_FCProc/', 'Seitzman300')
addpath '/Users/dianaperez/Documents/GitHub/GrattonLab-General-Repo/FCProcess'
fcimage_cormat_volume('/Users/dianaperez/Desktop/preprofc_FCProc/lifespan_datalist_LS03.  
xlsx', '/Users/dianaperez/Desktop/preprofc_FCProc/', 'Seitzman300')
```

```

xlsx', '/Users/dianaperez/Desktop/preprofc_FCProc/', 'Seitzman300')
addpath '/Users/dianaperez/Documents/GitHub/Lifespan-Analysis'
fcimage_corrmat_volume('/Users/dianaperez/Desktop/preprofc_FCProc/lifespan_datalist_LS03.
xlsx', '/Users/dianaperez/Desktop/preprofc_FCProc/', 'Seitzman300')
length(find(sess_roi_timeseries_concat(:,1)~=0))
unique(roi_data)
length(find(atlas_params.mods_array==7))
length(find(roi_data==7))
fcimage_corrmat_volume('/Users/dianaperez/Desktop/preprofc_FCProc/lifespan_datalist_LS03.
xlsx', '/Users/dianaperez/Desktop/preprofc_FCProc/', 'Seitzman300')
atlas_params = atlas_parameters_GrattonLab(atlas,atlas_dir);
atlas = 'Seitzman300';
atlas_params = atlas_parameters_GrattonLab(atlas,atlas_dir);
atlas_dir = '/Volumes/fsmresfiles/PBS/Gratton_Lab/Atlases/';
atlas_params = atlas_parameters_GrattonLab(atlas,atlas_dir);
unique(atlas_params.mods_array)
fcimage_corrmat_volume
fcimage_corrmat_volume('/Users/dianaperez/Desktop/preprofc_FCProc/lifespan_datalist_LS03.
xlsx', '/Users/dianaperez/Desktop/preprofc_FCProc/', 'Seitzman300')
only_cb = sess_data(sess_data(:,1)~=0,:);
tmaskFile = sprintf('%s/sub-%s/ses-%d/func/FD_outputs/sub-%s_ses-%d_task-%s_run-%d_desc-
tmask_%s.txt',...
FCdir,subInfo(i).subjectID,subInfo(i).session,...
subInfo(i).subjectID,subInfo(i).session,subInfo(i).condition,subInfo(i).runs(j),subInfo
(i).FDtype);
tmask{j} = table2array(readtable(tmaskFile));
tmask_concat = [tmask_concat; tmask{j}];
fcimage_corrmat_volume('/Users/dianaperez/Desktop/preprofc_FCProc/lifespan_datalist_LS03.
xlsx', '/Users/dianaperez/Desktop/preprofc_FCProc/', 'Seitzman300')
fcimage_corrmat_volume_only_cbvox
('/Users/dianaperez/Desktop/preprofc_FCProc/lifespan_datalist_LS03.xlsx',
'/Users/dianaperez/Desktop/preprofc_FCProc/', 'Seitzman300')
h = figure('Color',[0.8275 0.8275 0.8275],'Position',[56 143 1295 807]);
imagesc(corrmat)
load better_jet_colormap.mat; % assume this is in the same folder
colormap(better_jet_colormap_diff);
colorbar
fcimage_corrmat_volume_only_cbvox
fcimage_corrmat_volume_only_cbvox
('/Users/dianaperez/Desktop/preprofc_FCProc/lifespan_datalist_LS03.xlsx',
'/Users/dianaperez/Desktop/preprofc_FCProc/', 'Seitzman300')
nrois = [274:300];
nrois(nr)
fcimage_corrmat_volume_only_cbvox
fcimage_corrmat_volume_only_cbvox
('/Users/dianaperez/Desktop/preprofc_FCProc/lifespan_datalist_LS03.xlsx',
'/Users/dianaperez/Desktop/preprofc_FCProc/', 'Seitzman300')
h = figure('Color',[0.8275 0.8275 0.8275],'Position',[56 143 1295 807]); %[56 143 1095
807]
imagesc(matrix_orig)
load better_jet_colormap.mat; % assume this is in the same folder
colormap(better_jet_colormap_diff);
colorbar;
axis square;
fcimage_corrmat_volume_only_cbvox
('/Users/dianaperez/Desktop/preprofc_FCProc/lifespan_datalist_LS03.xlsx',
'/Users/dianaperez/Desktop/preprofc_FCProc/', 'Seitzman300')
fcimage_corrmat_volume_only_cbvox
('/Users/dianaperez/Desktop/preprofc_FCProc/lifespan_datalist_LS03.xlsx',
'/Volumes/fsmresfiles/PBS/Gratton_Lab/Lifespan/Post-

```

```
COVID/BIDS/derivatives/preproc_FCProc/', 'Seitzman300')
%-- 10/18/22, 5:09 PM --%
fcimage_corrmat_volume_only_cbvox✓
('/Users/dianaperez/Desktop/preprofc_FCProc/lifespan_datalist_LS03.xlsx', ✓
'/Users/dianaperez/Desktop/preprofc_FCProc/norm_preproc/', 'Seitzman300')
addpath '/Users/dianaperez/Documents/GitHub/GrattonLab-General-Repo/FCProcess'
fcimage_corrmat_volume_only_cbvox✓
('/Users/dianaperez/Desktop/preprofc_FCProc/lifespan_datalist_LS03.xlsx', ✓
'/Users/dianaperez/Desktop/preprofc_FCProc/norm_preproc/', 'Seitzman300')
addpath '/Users/dianaperez/Documents/GitHub/GrattonLab-General-Repo/FCProcess'
fcimage_corrmat_volume_only_cbvox✓
('/Users/dianaperez/Desktop/preprofc_FCProc/lifespan_datalist_LS03.xlsx', ✓
'/Users/dianaperez/Desktop/preprofc_FCProc/norm_preproc/', 'Seitzman300')
addpath '/Users/dianaperez/Documents/GitHub/Lifespan-Analysis'
fcimage_corrmat_volume_only_cbvox✓
('/Users/dianaperez/Desktop/preprofc_FCProc/lifespan_datalist_LS03.xlsx', ✓
'/Users/dianaperez/Desktop/preprofc_FCProc/norm_preproc/', 'Seitzman300')
reliability_cbdata
plot(times_all, corr(1,:), 'Color', [1,0.5,0], 'LineWidth', 3)
hold on
dataDir = '/Users/dianaperez/Desktop/preprofc_FCProc/norm_preproc/corrmats_Seitzman300/';
for s = 1:numel(subject)
    catData = [];
    catTmask = [];
    for i = 1:sessions
        %load mat file
        load([dataDir 'sub-' subject{s} '/sub-' subject{s} '_sess-' num2str(i) '_task-'
rest_corrmat_Seitzman300.mat'])
        masked_data = sess_roi_timeseries_concat(:,logical(tmask_concat'));
        % catData = [catData masked_data];
        catData = [catData masked_data];
        catTmask = [catTmask tmask_concat'];
    end
    %I think this should be 10,816
    disp(sprintf('Total number of sample points for subject %s is %d by %d...', subject{s}, ✓
size(catData,1), size(catData,2)))
    %true-half is 150min=8181 samp points (TR = 1.1; (8181*1.1)/60=149.99)
    truehalf = catData(:,1:pts2sample);
    corrmatrix_truehalf = paircorr_mod(truehalf);
    fout_truehalf = sprintf('%s/sub-%s_truehalf_100min_corrmatrix_reliabilityanalysis', ✓
output_dir, subject{s});
    %figure_corrmatrix_GrattonLab(corrmatrix_truehalf,atlas_params,-1,1);
    %saveas(gcf,[fout_truehalf '.tiff'],'tiff');
    %close(gcf);
    maskmat = ones(size(corrmatrix_truehalf, 1));
    maskmat = logical(triu(maskmat, 1));
    truehalf_corrlin(1,:) = corrmatrix_truehalf(maskmat);
    times = floor((size(catData,2)-pts2sample)/sampstep);
    times = [5:5:(times*5)];
    %sample data
    for t = 1:numel(times)
        sampledData{t} = catData(:, (pts2sample+1):(pts2sample+(sampstep*t)));
        corrmatrix = paircorr_mod(sampledData{t});
        fout = sprintf('%s/sub-%s_sampled_%dmin_corrmatrix_reliabilityanalysis', output_dir, subject{
{s}}, times(t));
        %figure_corrmatrix_GrattonLab(corrmatrix,atlas_params,-1,1);
        %saveas(gcf,[fout '.tiff'],'tiff');
        %close(gcf);
        corrs{t} = paircorr_mod(triu(corrmatrix_truehalf), triu(corrmatrix));
        corrlins(t,:) = corrmatrix(maskmat);
```

```

end
sampledDatas{t+1} = catData(:, (pts2sample+1):end);
corrmat = paircorr_mod(sampledDatas{t+1}');
corrs{t+1} = paircorr_mod(triu(corrmat_truehalf), triu(corrmat));
corrlins((t+1),:) = corrmat(maskmat);
times = [5:5:((numel(times)+1)*5)];
for j = 1:size(corrlins,1)
    tmpcorr = corrcoef(truehalf_corrlin', corrlins(j,:)', 'rows', 'complete');
    corr(s,j) = tmpcorr(2,1);
clear tmpcorr
end
times_all(s,1:size(times,2)) = times;
clear times
clear catData
clear catTmask
clear corrlins
clear corrmat
clear corrmat_truehalf
clear corrs
clear masked_data
clear sampledDatas
clear tmask
clear tmask_concat
clear truehalf
clear truehalf_corrlin
clear sess_roi_timeseries
clear sess_roi_timeseries_concat
end
plot(times_all ,corr(1,:), 'Color',[0, 0, 1], 'LineWidth', 3)
m = findobj(gca,'Type','line');
hleg1 = legend(m(1:2), 'normal gsr', 'cb gsr', 'Location', 'SouthEast');
print(gcf,[output_dir '/Reliability_cb-gsr.jpg'],'-dpng','-r300');
print(gcf, '/Users/dianaperez/Desktop/preprofc_FCPProc/Reliability_cb-gsr.jpg', '-dpng', '-r300');
%-- 10/31/22, 4:44 PM --%
system('pwsh')
system('pwd')
system('~/')
system('cd ~/')
system('pwd')
system('cd ~/Users/dianaperez')
system('cd ..')
system('pwd')
cd /Users/dianaperez
system('pwsh')
system('pwd')
[a, b] = system('pwsh')
[a, b] = system('/usr/local/bin/pwsh')
%-- 10/31/22, 5:07 PM --%
[a, b] = system('/usr/local/bin/pwsh help')
[a, b] = system('/usr/local/bin/pwsh -Login')
%-- 11/28/22, 7:13 AM --%
similarity_analysis_wLongitudinal
%-- 2/1/23, 12:20 AM --%
cdmscale_lifespan
similarity_analysis_forced_same_amt_data
addpath '/Users/dianaperez/Documents/GitHub/General_Scripts'
cdmscale_lifespan
addpath '/Users/dianaperez/Documents/GitHub/Lifespan-Analysis'
cdmscale_lifespan

```

```
axes([-15 30 -15 15])
axis([-15 30 -15 15])
axis([-15 30 -15 25])
cdmscale_lifespan
%-- 2/3/23, 10:57 AM --%
cdmscale_lifespan
addpath '/Users/dianaperez/Documents/GitHub/Lifespan-Analysis'
cdmscale_lifespan
vline_new(atlas_params.transitions,'k',3);
hline_new(atlas_params.transitions,'k',3);
tickpos = atlas_params.centers;
ax = axis;
%-- 2/3/23, 3:01 PM --%
cdmscale_lifespan
addpath '/Users/dianaperez/Documents/GitHub/GrattonLab-General-Repo/FCProcess'
cdmscale_lifespan
addpath '/Users/dianaperez/Documents/GitHub/Lifespan-Analysis'
cdmscale_lifespan
[Y E] = cmdscale_mat_MSC(input,groups,'euclidean',colors,subs)
figure
[Y E] = cmdscale_mat_MSC(input,groups,'correlation',colors,subs)
help pdist
[Y E] = cmdscale_mat_MSC(input,groups,'correlation',colors,subs)
figure
imagesc(mat_col')
clear all
cdmscale_lifespan
figure
imagesc(mat_col')
axis([-10 30 -20 25])
figure;
[Y E] = cmdscale_mat_MSC(input,groups,'correlation',colors,subs)
D = pdist(double(mat_col'),dist_type);
axis([-0.4 0.6 -0.4 0.4])
cdmscale_lifespan
axis([-0.4 0.6 -0.4 0.4])
axis([-10 30 -20 25])
axis([-5 30 -20 25])
axis([-20 30 -20 25])
axis([-15 30 -20 25])
axis([-15 40 -20 25])
axis([-15 45 -20 25])
cdmscale_lifespan
sum(allSubs_amtData,1)
sum(allSubs_amtData,2)
min(total_data)
total_data = sum(allSubs_amtData,2);
smallest_amt = min(total_data);
split_half = floor(smallest_amt/2);
cdmscale_lifespan_splitses
size(masked_data,2)
cdmscale_lifespan_splitses
total_data = sum(allSubs_amtData,1);
smallest_amt = min(total_data);
amt_data = floor(smallest_amt/2);
size(masked_data,2)
size(masked_data,2)-amt_data
matched_data_2 = masked_data(:,(size(masked_data,2)-amt_data):end);
matched_data_2 = masked_data(:,(size(masked_data,2)-amt_data+1):end);
cdmscale_lifespan_splitses
```

```
cdmscale_lifespan
%-- 2/13/23, 5:44 PM --%
load('/Users/dianaperez/Desktop/FC_Parcel_333/sub-LS02_rest_ses-1_parcel_timecourse.
mat')
reliability_analysis_G333
maskmat = ones(333);
maskmat = logical(triu(maskmat, 1));
truehalf_corrln(1,:) = corrmatrix_truehalf(maskmat);
reliability_analysis_G333
axis([0 1 0 100])
axis([0 100 0 1])
plot(times_all(1,1:48),mean(1:48), ':', 'Color', [0,0,0], 'LineWidth',3) %average
mean(means{1:8})
reliability_analysis_G333
means = [0.558299 0.655694765 0.707371651 0.73979493 0.762946177 0.779178405
0.790522018 0.802113188 0.812064613 0.820295988 0.827574146 0.833868939 0.839342416
0.843669992 0.84802989 0.852440158 0.855737711 0.8592886 0.861987349 0.864990617
0.867766204 0.870012403 0.871921675 0.874183608 0.876164527 0.871685358 0.873252072
0.875060784 0.876514917 0.877787231 0.905295217 0.906622921 0.90778087 0.908984651
0.910107888 0.910929481 0.912091361 0.912735012 0.913586008 0.914280476 0.910357217
0.911032658 0.911704121 0.91238943 0.913110063 0.913701016 0.914340857 0.914802561
0.915357006 0.915758258 0.918267278];
reliability_analysis_G333
%-- 2/15/23, 12:36 PM --%
load('/Volumes/fsmresfiles/PBS/Gratton_Lab/Lifespan/Post-
COVID/BIDS/derivatives/postFCproc_CIFTI/FC_Parcel_333/sub-INET001_rest_ses-
1_parcel_timecourse.mat')
similarity_analysis_forced_same_amt_data
parcel_time(:,logical(tmask_concat))
parcel_time(:,logical(tmask_concat));
parcel_time(logical(tmask_concat),:);
similarity_analysis_forced_same_amt_data
amt_data = min(min(allSubs_amtData));
similarity_analysis_forced_same_amt_data
amt_data = min(allSubs_amtData);
similarity_analysis_forced_same_amt_data
addpath '/Users/dianaperez/Documents/GitHub/General_Scripts'
similarity_analysis_forced_same_amt_data
figure('Position',[1 1 1000 800]);
imagesc(simmat,[0 1]); colormap('jet');
hline_new([0,4,8,12,16,20,24,28,32,36,40]+0.5,'k',2);
vline_new([0,4,8,12,16,20,24,28,32,36,40]+0.5,'k',2);
set(gca,'XTick',[2,6,10,14,18,22,26,30,34,38], 'YTick', [2,6,10,14,18,22,26,30,34,38],
'XTickLabel',...
{'INET001','INET002',
'INET003','INET005','INET006','INET010','INET016','INET018','INET019','INET030'},
'YTickLabel', {'LS02','LS03','LS05','LS08','LS11','LS14','LS16','LS17'});
set(gca,'XTick',[2,6,10,14,18,22,26,30,34,38]+0.5, 'YTick', [2,6,10,14,18,22,26,30,34,38]
+0.5, 'XTickLabel',...
{'INET001','INET002',
'INET003','INET005','INET006','INET010','INET016','INET018','INET019','INET030'},
'YTickLabel', {'LS02','LS03','LS05','LS08','LS11','LS14','LS16','LS17'});
hline_new([0,4,8,12,16,20,24,28,32,36,40]+0.5,'k',2);
vline_new([0,4,8,12,16,20,24,28,32,36,40]+0.5,'k',2);
set(gca,'XTick',[2,6,10,14,18,22,26,30,34,38]+0.5, 'YTick', [2,6,10,14,18,22,26,30,34,38]
+0.5, 'XTickLabel',...
{'INET001','INET002',
'INET003','INET005','INET006','INET010','INET016','INET018','INET019','INET030'},
'YTickLabel', {'INET001','INET002',
'INET003','INET005','INET006','INET010','INET016','INET018','INET019','INET030'});
```

```
axis square;
colorbar;
title('Correlation Matrix Similarity');
if match_data
saveas(gcf,[outdir data_set '_' data_type '_similarityMat_matchedData_' num2str(amt_data)
'.tiff'],'tiff');
else
saveas(gcf,[outdir data_set '_' data_type '_similarityMat.tiff'],'tiff');
end
similarity_analysis_forced_same_amt_data
hline_new([0,4,8,12,16,20,24,28,32,36,40]+0.5,'k',2);
vline_new([0,4,8,12,16,20,24,28,32,36,40]+0.5,'k',2);
set(gca,'XTick',[2,6,10,14,18,22,26,30,34,38]+0.5,'YTick',[2,6,10,14,18,22,26,30,34,38]
+0.5,'XTickLabel',...
{'INET001','INET002'},
'INET003','INET005','INET006','INET010','INET016','INET018','INET019','INET030'},
'YTickLabel',{ 'INET001','INET002'},
'INET003','INET005','INET006','INET010','INET016','INET018','INET019','INET030'});
axis square;
colorbar;
title('Correlation Matrix Similarity');
if match_data
saveas(gcf,[outdir data_set '_' data_type '_similarityMat_matchedData_' num2str(amt_data)
'.tiff'],'tiff');
else
saveas(gcf,[outdir data_set '_' data_type '_similarityMat.tiff'],'tiff');
end
close('all');
count = 1;
within = [];
between = [];
for s = 1:numel(subject)
lines = [count:(count+sessions-1)];
sub_vals = simmat(lines,:);
maskmat = ones(sessions,sessions);
maskmat = logical(triu(maskmat, 1));
within_sub = sub_vals(:,lines);
within = [within; within_sub(maskmat)];
maskmat = ones(size(sub_vals));
maskmat(:,lines) = 0;
between = [between; sub_vals(maskmat==1)];
count = count+sessions;
end
mean(between)
mean(within)
similarity_analysis_forced_same_amt_data
dataDir = '/Volumes/fsmresfiles/PBS/Gratton_Lab/Lifespan/Post-
COVID/derivatives/preproc_FCProc/corrmats_Seitzman300/';
cd dataDir
cd /Volumes/fsmresfiles/PBS/Gratton_Lab/Lifespan/Post-COVID/
dataDir = '/Volumes/fsmresfiles/PBS/Gratton_Lab/Lifespan/Post-
COVID/BIDS/derivatives/preproc_FCProc/corrmats_Seitzman300/';
cd dataDir
pwd
dataDir = '/Volumes/fsmresfiles/PBS/Gratton_Lab/Lifespan/Post-
COVID/BIDS/derivatives/preproc_FCProc/corrmats_Seitzman300';
cd dataDir = '/Volumes/fsmresfiles/PBS/Gratton_Lab/Lifespan/Post-
COVID/BIDS/derivatives/preproc_FCProc/corrmats_Seitzman300';
cd dataDir
dataDir = '/Volumes/fsmresfiles/PBS/Gratton_Lab/Lifespan/Post-
```



```
COVID/BIDS/derivatives/preproc_FCPProc/corrmats_Seitzman300/';
similarity_analysis_forced_same_amt_data
tick_marks = [0:5:(5*numel(LS_subject))]+0.5;
tick_marks+(sessions/2)
tick_marks(1:numel(subject))+(sessions/2)
figure('Position',[1 1 1000 800]);
imagesc(simmat,[0 1]); colormap('jet');
tick_marks = [0:sessions:(5*numel(subject))]+0.5;
hline_new(tick_marks,'k',2);
vline_new(tick_marks,'k',2);
set(gca,'XTick',tick_marks(1:numel(subject))+(sessions/2), 'YTick', tick_marks(1:numel(
(subject))+(sessions/2), 'XTickLabel',...
subject, 'YTickLabel', subject);
similarity_analysis_forced_same_amt_data
cd /Users/dianaperez/Documents/GitHub/Lifespan-Analysis/
load('Lifespan_reliability_data_for_plot.mat')
clear all
cd /Users/dianaperez/Documents/GitHub/Lifespan-Analysis/
load('Lifespan_reliability_data_for_plot.mat')
addpath '/Users/dianaperez/Desktop'
load('Lifespan_reliability_data_for_plot.mat')
for s = 1:numel(subject)
plot(times_all(s,1:size(means{1,s},2)),means{1,s},'Color',rgb_colors(s,:), 'LineWidth', 3)
hold on
end
subject = {'LS02', 'LS03', 'LS05', 'LS08', 'LS11', 'LS14', 'LS16', 'LS17'};
sessions = 5;
for s = 1:numel(subject)
plot(times_all(s,1:size(means{1,s},2)),means{1,s},'Color',rgb_colors(s,:), 'LineWidth', 3)
hold on
end
for t = 1:48
tmp = [];
for s = 1:numel(subject)
if t <= size(means{1,s},2)
tmp = [tmp;means{1,s}(t)]
else
continue;
end
end
mean_of_means(t) = mean(tmp);
end
plot(times_all(1,1:48),mean_of_means(1:48), ':', 'Color', [0,0,0], 'LineWidth',3) %<
average
ylabel('Pearson Correlation (r)');
xlabel('Time (Minutes)');
m = findobj(gca,'Type','line');
hleg1 = legend(m(1:9), 'Mean', 'LS17', 'LS16', 'LS14', 'LS11', 'LS08', 'LS05', 'LS03',<
'LS02', 'Location', 'SouthEast');
hleg1.FontSize = 20;
ax = gca;
ax.FontSize = 24;
output_dir = '/Users/dianaperez/Desktop/';
print(gcf,[output_dir 'Lifespan_Reliability_truehalf_' num2str(pts2sample) '.jpg'],'-<
dpng','-r300');
pts2sample = 3808;
print(gcf,[output_dir 'Lifespan_Reliability_truehalf_' num2str(pts2sample) '.jpg'],'-<
dpng','-r300');
dmat_file = '/Users/dianaperez/Documents/Dependencies/Resources/Conte69_atlas-v2.LR.<
32k_fs_LR.wb/Cifti_geo_distances_xhemisphere_large.mat'
```



```
dmat = smartload(dmat_file);
clear all
dmat = smartload(dmat_file);
dmat_file = '/Users/dianaperez/Documents/Dependencies/Resources/Conte69_atlas-v2.LR.
32k_fs_LR.wb/Cifti_geo_distances_xhemisphere_large.mat'
dmat = smartload(dmat_file);
%-- 2/17/23, 2:19 PM --%
gifti_read('/Volumes/fsmresfiles/PBS/GrattonLab/Lifespan/Pre-
COVID/Nifti/derivatives/freesurfer-6.0.1/FREESURFER_fs_LR/sub-
LS02/NativeVol/fsaverage_LR32k/sub-LS02.R.midthickness.32k_fs_LR.surf.gii')
gifti_read('/Volumes/fsmresfiles/PBS/Gratton_Lab/Lifespan/Pre-
COVID/Nifti/derivatives/freesurfer-6.0.1/FREESURFER_fs_LR/sub-
LS02/NativeVol/fsaverage_LR32k/sub-LS02.R.midthickness.32k_fs_LR.surf.gii')
gifti_read('/Volumes/fsmresfiles/PBS/Gratton_Lab/Lifespan/Pre-
COVID/BIDS/Nifti/derivatives/freesurfer-6.0.1/FREESURFER_fs_LR/sub-
LS02/NativeVol/fsaverage_LR32k/sub-LS02.R.midthickness.32k_fs_LR.surf.gii')
gifti_read('/Volumes/fsmresfiles/PBS/Gratton_Lab/Lifespan/Pre-
COVID/BIDS/Nifti/derivatives/freesurfer-6.0.1/FREESURFER_fs_LR/sub-
LS02/NativeVol/fsaverage_LR32k/sub-LS02.R.midthickness.32k_fs_LR.surf.gii', this)
this.data = [];
this.metaData = [];
this.label = [];
gifti_read('/Volumes/fsmresfiles/PBS/Gratton_Lab/Lifespan/Pre-
COVID/BIDS/Nifti/derivatives/freesurfer-6.0.1/FREESURFER_fs_LR/sub-
LS02/NativeVol/fsaverage_LR32k/sub-LS02.R.midthickness.32k_fs_LR.surf.gii', this)
cd /Users/dianaperez/Applications/
cd ~/Applications/workbench
cd /Applications/workbench
wb_path = '/Applications/workbench/bin_macosx64/';
right_surface = '/Volumes/fsmresfiles/PBS/GrattonLab/Lifespan/Pre-
COVID/Nifti/derivatives/freesurfer-6.0.1/FREESURFER_fs_LR/sub-
LS02/NativeVol/fsaverage_LR32k/sub-LS02.R.midthickness.32k_fs_LR.surf.gii';
vertex = 1;
system([wb_path ' wb_command -surface-geodesic-distance ' surface ' ' num2str(vertex) ' '
output_fname])
output = [];
system([wb_path ' wb_command -surface-geodesic-distance ' surface ' ' num2str(vertex) ' '
output_fname])
system([wb_path ' wb_command -surface-geodesic-distance ' surface ' ' num2str(vertex) ' '
output])
output = '/Users/dianaperez/Desktop/testing_output_dist';
system([wb_path ' wb_command -surface-geodesic-distance ' surface ' ' num2str(vertex) ' '
output])
[x y] = system([wb_path ' wb_command -surface-geodesic-distance ' surface ' ' num2str
(vertex) ' ' output])
[x y] = system([wb_path ' wb_command -surface-geodesic-distance ' surface ' ' num2str
(vertex) ' ' output_f])
[x y] = system([wb_path ' wb_command -surface-geodesic-distance ' surface ' ' num2str
(vertex) ' ' output_fname])
output_fname = {};
[x y] = system([wb_path ' wb_command -surface-geodesic-distance ' surface ' ' num2str
(vertex) ' ' output_fname])
[x y] = system([wb_path ' wb_command -surface-geodesic-distance-all-to-all ' surface ' '
output '.dtseries.nii'])
[x y] = system([wb_path ' wb_command -surface-geodesic-distance-all-to-all ' surface ' '
output ])
system([wb_path ' wb_command -surface-geodesic-distance-all-to-all ' surface ' '
output_fname])
system([wb_path ' wb_command -surface-geodesic-distance-all-to-all ' surface ' '
output_fname]);
```

```
system([wb_path 'wb_command -surface-geodesic-distance-all-to-all ' sub-  
LS02_distances_right.dconn.nii'])  
system('/Applications/workbench/bin_macosx64/wb_command -surface-geodesic-distance-all-  
to-all /Volumes/fsmresfiles/PBS/GrattonLab/Lifespan/Pre-  
COVID/Nifti/derivatives/freesurfer-6.0.1/FREESURFER_fs_LR/sub-  
LS02/NativeVol/fsaverage_LR32k/sub-LS02.R.midthickness.32k_fs_LR.surf.gii sub-  
LS02_distances_right.dconn.nii')  
right_surface = '/Volumes/fsmresfiles/PBS/Gratton_Lab/Lifespan/Pre-  
COVID/BIDS/Nifti/derivatives/freesurfer-6.0.1/FREESURFER_fs_LR/sub-  
LS02/NativeVol/fsaverage_LR32k/sub-LS02.R.midthickness.32k_fs_LR.surf.gii';  
command = [wb_path 'wb_command -surface-geodesic-distance-all-to-all ' right_surface '  
sub-LS02_distances_right.dconn.nii'];  
command  
system(command)  
command = [wb_path 'wb_command -surface-geodesic-distance-all-to-all ' right_surface '  
/Users/dianaperez/Desktop/sub-LS02_distances_right.dconn.nii'];  
system(command)  
test = ft_read_cifti_mod('/Users/dianaperez/Desktop/sub-LS02_distances_right.dconn.nii')  
cd /Users/dianaperez/Documents/  
test = ft_read_cifti_mod('/Users/dianaperez/Desktop/sub-LS02_distances_right.dconn.nii')  
right_surface = '/Volumes/fsmresfiles/PBS/Gratton_Lab/Lifespan/Pre-  
COVID/BIDS/Nifti/derivatives/freesurfer-6.0.1/FREESURFER_fs_LR/sub-  
LS02/NativeVol/fsaverage_LR32k/sub-LS02.L.midthickness.32k_fs_LR.surf.gii';  
command = [wb_path 'wb_command -surface-geodesic-distance-all-to-all ' right_surface '  
/Users/dianaperez/Desktop/sub-LS02_distances_left.dconn.nii'];  
system(command)  
%-- 2/22/23, 12:36 PM --%  
test = ft_read_cifti_mod('/Users/dianaperez/Desktop/sub-LS02_distances_right.dconn.nii')  
network_seg_new  
cd(atlasdir)  
network_seg_new  
rois = atlas_params.mods{net};  
network_seg_new  
cd /Users/dianaperez/Documents/GitHub/Lifespan-Analysis/  
within_sub_reliability_v2  
data = parcel_time';  
masked_data = data(:,logical(tmask_concat));  
within_sub_reliability_v2  
size(atlas_params.networks)  
length(atlas_params.networks)  
within_sub_reliability_v2  
figure;  
for s = 1:numel(subject)  
plot(times_all(s,1:size(allsubs_seg_ind{s},2)),mean(allsubs_abs_diffs{1,s}(:,1:size_  
(allsubs_seg_ind{s},2)),1),'LineWidth', 3)  
hold on  
means_diff{s} = mean(allsubs_abs_diffs{1,s}(:,1:size(allsubs_seg_ind{s},2)),1);  
end  
within_sub_reliability_v2  
for t = 1:length(times)  
tmp = [];  
for s = 1:numel(subject)  
if t <= size(means_diff{1,s},2)  
tmp = [tmp;means_diff{1,s}(t)]  
else  
continue;  
end  
end  
mean_of_means(t) = mean(tmp);  
end
```

```
plot(times_all,mean_of_means, ':', 'Color', [0,0,0], 'LineWidth',3)
plot(times_all(1:40),mean_of_means(1:40), ':', 'Color', [0,0,0], 'LineWidth',3)
plot(times_all(1,1:40),mean_of_means(1:40), ':', 'Color', [0,0,0], 'LineWidth',3)
within_sub_reliability_v2
save('/Users/dianaperez/Desktop/segregation_index_data_for_plots.mat',↵
'allsubs_abs_diffs', 'allsubs_seg_ind', 'times_all', 'mean_of_means', 'rgb_colors_iNet',↵
'rgb_colors_LS', 'subject')
within_sub_reliability_v2
figure;
for s = 1:numel(subject)
plot(times_all(s,1:size(allsubs_seg_ind{s},2)),mean(allsubs_seg_ind{1,s}(:,1:size↵
(allsubs_seg_ind{s},2)),1), 'Color', rgb_colors_LS(s,:), 'LineWidth', 3)
hold on
end
figure;
for s = 1:numel(subject)
plot(times_all(s,1:size(allsubs_seg_ind{s},2)),mean(allsubs_seg_ind{1,s}(:,1:size↵
(allsubs_seg_ind{s},2)),1), 'Color', rgb_colors_LS(s,:), 'LineWidth', 3)
hold on
end
for t = 1:length(times)
tmp = [];
for s = 1:numel(subject)
if t <= size(allsubs_seg_ind{1,s},2)
tmp = [tmp;allsubs_seg_ind{1,s}(t)];
else
continue;
end
end
mean_of_means_SI(t) = mean(tmp);
end
%means = [0.228121374    0.282854956 0.311825923 0.330394855 0.344075468 0.3540615↵
0.362085299 0.368580906 0.374353669 0.378886496 0.38303392 0.386997298 0.39018919↵
0.392779018 0.395671349 0.3921087 0.394652817 0.396349383 0.398223 0.39998822↵
0.401678269 0.402921549 0.404130596 0.405287927 0.406779144 0.407857626 0.408819803↵
0.40779906 0.4087539 0.409494202 0.41042813 0.411371703 0.4121878 0.412706817↵
0.413360333 0.414032785 0.41465492 0.419794232 0.420634614 0.420983464 0.421506702↵
0.421954376 0.422355276 0.42278862 0.423175282 0.42350195 0.423740618 0.42402081↵
0.434813233 0.42669529 0.42691267 0.42712188 0.42740375];
plot(times_all(1,1:40),mean_of_means_SI(1:40), ':', 'Color', [0,0,0], 'LineWidth',3)
figure;
for s = 1:numel(subject)
plot(times_all(s,1:size(allsubs_between_FC{s},2)),mean(allsubs_between_FC{1,s}(:,1:size↵
(allsubs_between_FC{s},2)),1), 'Color', rgb_colors_LS(s,:), 'LineWidth', 3)
hold on
end
for t = 1:length(times)
tmp = [];
for s = 1:numel(subject)
if t <= size(allsubs_between_FC{1,s},2)
tmp = [tmp;allsubs_between_FC{1,s}(t)];
else
continue;
end
end
mean_of_means_bFC(t) = mean(tmp);
end
%means = [0.228121374    0.282854956 0.311825923 0.330394855 0.344075468 0.3540615↵
0.362085299 0.368580906 0.374353669 0.378886496 0.38303392 0.386997298 0.39018919↵
0.392779018 0.395671349 0.3921087 0.394652817 0.396349383 0.398223 0.39998822↵
```

```
0.401678269 0.402921549 0.404130596 0.405287927 0.406779144 0.407857626 0.408819803✓
0.40779906 0.4087539 0.409494202 0.41042813 0.411371703 0.4121878 0.412706817✓
0.413360333 0.414032785 0.41465492 0.419794232 0.420634614 0.420983464 0.421506702✓
0.421954376 0.422355276 0.42278862 0.423175282 0.42350195 0.423740618 0.42402081✓
0.434813233 0.42669529 0.42691267 0.42712188 0.42740375];
plot(times_all(1,1:40),mean_of_means_bFC(1:40), ':', 'Color', [0,0,0], 'LineWidth',3)
%axis([0 120 0.1 .5])
ylabel('Between FC');
xlabel('Time (Minutes)');
figure;
for s = 1:numel(subject)
plot(times_all(s,1:size(allsubs_within_FC{s},2)),mean(allsubs_within_FC{1,s}(:,1:size(
allsubs_within_FC{s},2)),1), 'Color', rgb_colors_LS(s,:), 'LineWidth', 3)
hold on
end
for t = 1:length(times)
tmp = [];
for s = 1:numel(subject)
if t <= size(allsubs_within_FC{1,s},2)
tmp = [tmp;allsubs_within_FC{1,s}(t)];
else
continue;
end
end
mean_of_means_wFC(t) = mean(tmp);
end
%means = [0.228121374 0.282854956 0.311825923 0.330394855 0.344075468 0.3540615✓
0.362085299 0.368580906 0.374353669 0.378886496 0.38303392 0.386997298 0.39018919✓
0.392779018 0.395671349 0.3921087 0.394652817 0.396349383 0.398223 0.39998822✓
0.401678269 0.402921549 0.404130596 0.405287927 0.406779144 0.407857626 0.408819803✓
0.40779906 0.4087539 0.409494202 0.41042813 0.411371703 0.4121878 0.412706817✓
0.413360333 0.414032785 0.41465492 0.419794232 0.420634614 0.420983464 0.421506702✓
0.421954376 0.422355276 0.42278862 0.423175282 0.42350195 0.423740618 0.42402081✓
0.434813233 0.42669529 0.42691267 0.42712188 0.42740375];
plot(times_all(1,1:40),mean_of_means_wFC(1:40), ':', 'Color', [0,0,0], 'LineWidth',3)
%axis([0 120 0.1 .5])
ylabel('Within FC');
xlabel('Time (Minutes)');
figure;
for s = 1:numel(subject)
plot(times_all(s,1:size(allsubs_within_FC{s},2)),mean(allsubs_within_FC{1,s}(:,1:size(
allsubs_within_FC{s},2)),1), 'Color', rgb_colors_LS(s,:), 'LineWidth', 3)
hold on
end
for t = 1:length(times)
tmp = [];
for s = 1:numel(subject)
if t <= size(allsubs_within_FC{1,s},2)
tmp = [tmp;allsubs_within_FC{1,s}(t)];
else
continue;
end
end
mean_of_means_wFC(t) = mean(tmp);
end
%means = [0.228121374 0.282854956 0.311825923 0.330394855 0.344075468 0.3540615✓
0.362085299 0.368580906 0.374353669 0.378886496 0.38303392 0.386997298 0.39018919✓
0.392779018 0.395671349 0.3921087 0.394652817 0.396349383 0.398223 0.39998822✓
0.401678269 0.402921549 0.404130596 0.405287927 0.406779144 0.407857626 0.408819803✓
0.40779906 0.4087539 0.409494202 0.41042813 0.411371703 0.4121878 0.412706817✓
```

```
0.413360333 0.414032785 0.41465492 0.419794232 0.420634614 0.420983464 0.421506702✓
0.421954376 0.422355276 0.42278862 0.423175282 0.42350195 0.423740618 0.42402081✓
0.434813233 0.42669529 0.42691267 0.42712188 0.42740375];
plot(times_all(1,1:40),mean_of_means_wFC(1:40), ':', 'Color', [0,0,0], 'LineWidth',3)
%axis([0 120 0.1 .5])
ylabel('Within FC');
xlabel('Time (Minutes)');
leg_names = ['Mean' subject]
leg_names = ['Mean' flip(subject)]
within_sub_reliability_v2
figure;
for s = 1:numel(subject)
plot(times_all(s,1:size(allsubs_between_FC{s},2)),mean(allsubs_between_FC{1,s}(:,1:size(
allsubs_between_FC{s},2)),1), 'Color', rgb_colors_LS(s,:), 'LineWidth', 3)
hold on
end
for t = 1:length(times)
tmp = [];
for s = 1:numel(subject)
if t <= size(allsubs_between_FC{1,s},2)
tmp = [tmp;allsubs_between_FC{1,s}(t)];
else
continue;
end
end
mean_of_means_bFC(t) = mean(tmp);
end
%means = [0.228121374 0.282854956 0.311825923 0.330394855 0.344075468 0.3540615✓
0.362085299 0.368580906 0.374353669 0.378886496 0.38303392 0.386997298 0.39018919✓
0.392779018 0.395671349 0.3921087 0.394652817 0.396349383 0.398223 0.39998822✓
0.401678269 0.402921549 0.404130596 0.405287927 0.406779144 0.407857626 0.408819803✓
0.40779906 0.4087539 0.409494202 0.41042813 0.411371703 0.4121878 0.412706817✓
0.413360333 0.414032785 0.41465492 0.419794232 0.420634614 0.420983464 0.421506702✓
0.421954376 0.422355276 0.42278862 0.423175282 0.42350195 0.423740618 0.42402081✓
0.434813233 0.42669529 0.42691267 0.42712188 0.42740375];
plot(times_all(1,1:40),mean_of_means_bFC(1:40), ':', 'Color', [0,0,0], 'LineWidth',3)
%axis([0 120 0.1 .5])
ylabel('Between FC');
xlabel('Time (Minutes)');
figure;
for s = 1:numel(subject)
plot(times_all(s,1:size(allsubs_within_FC{s},2)),mean(allsubs_within_FC{1,s}(:,1:size(
allsubs_within_FC{s},2)),1), 'Color', rgb_colors_LS(s,:), 'LineWidth', 3)
hold on
end
for t = 1:length(times)
tmp = [];
for s = 1:numel(subject)
if t <= size(allsubs_within_FC{1,s},2)
tmp = [tmp;allsubs_within_FC{1,s}(t)];
else
continue;
end
end
mean_of_means_wFC(t) = mean(tmp);
end
%means = [0.228121374 0.282854956 0.311825923 0.330394855 0.344075468 0.3540615✓
0.362085299 0.368580906 0.374353669 0.378886496 0.38303392 0.386997298 0.39018919✓
0.392779018 0.395671349 0.3921087 0.394652817 0.396349383 0.398223 0.39998822✓
0.401678269 0.402921549 0.404130596 0.405287927 0.406779144 0.407857626 0.408819803✓
```

```

0.40779906 0.4087539 0.409494202 0.41042813 0.411371703 0.4121878 0.412706817✓
0.413360333 0.414032785 0.41465492 0.419794232 0.420634614 0.420983464 0.421506702✓
0.421954376 0.422355276 0.42278862 0.423175282 0.42350195 0.423740618 0.42402081✓
0.434813233 0.42669529 0.42691267 0.42712188 0.42740375];
plot(times_all(1,1:40),mean_of_means_wFC(1:40), ':', 'Color', [0,0,0], 'LineWidth',3)
%axis([0 120 0.1 .5])
ylabel('Within FC');
xlabel('Time (Minutes)');
within_sub_reliability_v2
figure;
for s = 1:numel(subject)
plot(times_all(s,1:size(allsubs_seg_ind{s},2)),mean(allsubs_abs_diffs{1,s}(:,1:size(
allsubs_seg_ind{s},2)),1), 'Color', rgb_colors_LS, 'LineWidth', 3)
hold on
means_diff{s} = mean(allsubs_abs_diffs{1,s}(:,1:size(allsubs_seg_ind{s},2)),1);
end
rgb_colors_LS{s}
rgb_colors_LS{s,:}
rgb_colors_LS(s,:)
for s = 1:numel(subject)
plot(times_all(s,1:size(allsubs_seg_ind{s},2)),mean(allsubs_abs_diffs{1,s}(:,1:size(
allsubs_seg_ind{s},2)),1), 'Color', rgb_colors_LS(s,:), 'LineWidth', 3)
hold on
means_diff{s} = mean(allsubs_abs_diffs{1,s}(:,1:size(allsubs_seg_ind{s},2)),1);
end
for t = 1:length(times)
tmp = [];
for s = 1:numel(subject)
if t <= size(means_diff{1,s},2)
tmp = [tmp;means_diff{1,s}(t)];
else
continue;
end
end
mean_of_means(t) = mean(tmp);
end
%mean_diff_all = [0.451054915 0.31958749 0.250078328 0.205589374 0.173270193✓
0.149834654 0.130898036 0.116096011 0.103579279 0.09357542 0.085252215 0.077510628✓
0.071893195 0.067765274 0.063014982 0.0609474 0.057319998 0.055589204 0.053587631✓
0.051688755 0.049855826 0.049189074 0.047716066 0.047485058 0.046371319 0.046092851✓
0.045026644 0.049051036 0.048147125 0.04798667 0.047803287 0.048079218 0.047927852✓
0.047976362 0.048676489 0.048785449 0.048605224 0.048932758 0.049221036 0.049052662✓
0.049501161 0.049540971 0.049734258 0.04993298 0.050447245 0.050401381 0.0504621✓
0.050506165 0.048969311 0.042753616 0.042789566 0.043138017 0.043480223 0.026972895];
plot(times_all(1,1:40),mean_of_means(1:40), ':', 'Color', [0,0,0], 'LineWidth',3)
axis([0 100 0 .6])
ylabel('% Difference');
xlabel('Time (Minutes)');
m = findobj(gca,'Type','line');
hleg1 = legend(m(1:(numel(subject)+1)), ['Mean' flip(subject)], 'Location', 'NorthEast');
hleg1.FontSize = 20;
ax = gca;
ax.FontSize = 24;
print(gcf, '/Users/dianaperez/Desktop/ReliabilityLifespanSegIndex_Diff.jpg', '-dpng', '-r300');
print(gcf, '/Users/dianaperez/Desktop/ReliabilityLifespanSegIndex_Diff.jpg', '-dpng', '-r300');
figure;
for s = 1:numel(subject)
plot(times_all(s,1:size(allsubs_seg_ind{s},2)),mean(allsubs_seg_ind{1,s}(:,1:size(

```

```
(allsubs_seg_ind{s},2)),1),'Color',rgb_colors_LS(s,:),'LineWidth', 3)
hold on
end
for t = 1:length(times)
tmp = [];
for s = 1:numel(subject)
if t <= size(allsubs_seg_ind{1,s},2)
tmp = [tmp;allsubs_seg_ind{1,s}(t)];
else
continue;
end
end
mean_of_means_SI(t) = mean(tmp);
end
size(allsubs_seg_ind{1,s},2)
for t = 1:length(times)
tmp = [];
for s = 1:numel(subject)
if t <= size(allsubs_seg_ind{1,s},2)
tmp = [tmp;mean(allsubs_seg_ind{1,s}(t))];
else
continue;
end
end
mean_of_means_SI(t) = mean(tmp);
end
mean(allsubs_seg_ind{1,1}(1))
mean(allsubs_seg_ind{1,2}(1))
mean(allsubs_seg_ind{1,3}(1))
mean(allsubs_seg_ind{1,1}(2))
mean(allsubs_seg_ind{1,1}(3))
t = 1;
tmp = [];
for s = 1:numel(subject)
if t <= size(allsubs_seg_ind{1,s},2)
tmp = [tmp;mean(allsubs_seg_ind{1,s}(t))];
else
continue;
end
end
mean_of_means_SI(t) = mean(tmp);
mean_of_means_SI(t) = mean(tmp)
t = 2;
tmp = [];
for s = 1:numel(subject)
if t <= size(allsubs_seg_ind{1,s},2)
tmp = [tmp;mean(allsubs_seg_ind{1,s}(t))];
else
continue;
end
end
mean_of_means_SI(t) = mean(tmp)
t = 3;
tmp = [];
for s = 1:numel(subject)
if t <= size(allsubs_seg_ind{1,s},2)
tmp = [tmp;mean(allsubs_seg_ind{1,s}(t))];
else
continue;
end
end
```



```

end
mean_of_means_SI(t) = mean(tmp)
mean(tmp)
length(times)
allsubs_seg_ind{1,t}
allsubs_seg_ind{1,1}(1)
allsubs_seg_ind{1,1}(:,1)
for t = 1:length(times)
    tmp = [];
    for s = 1:numel(subject)
        if t <= size(allsubs_seg_ind{1,s},2)
            tmp = [tmp;mean(allsubs_seg_ind{1,s}(:,t))];
        else
            continue;
        end
    end
    mean_of_means_SI(t) = mean(tmp);
end
plot(times_all(1,1:40),mean_of_means_SI(1:40), ':', 'Color', [0,0,0], 'LineWidth',3)
plot(times_all(1,1:40),smooth(mean_of_means_SI(1:40)), ':', 'Color', [0,0,0], '
'LineWidth',3)
help smooth
axis([0 100 0.1 .5])
ylabel('Segregation Index');
xlabel('Time (Minutes)');
m = findobj(gca,'Type','line');
hleg1 = legend(m(1:(numel(subject)+1)), ['Mean' flip(subject)], 'Location', 'NorthEast');
hleg1.FontSize = 20;
ax = gca;
ax.FontSize = 20;
axis([0 100 0 .4])
axis([0 100 -0.1 .4])
hleg1 = legend(m(1:(numel(subject)+1)), ['Mean' flip(subject)], 'Location', 'SouthEast');
axis([0 100 -0.2 .5])
axis([0 100 -0.2 .4])
print(gcf,[output_dir '/ReliabilityLifespanSegIndex.jpg'],'-dpng','-r300');
print(gcf,['/Users/dianaperez/Desktop/ReliabilityLifespanSegIndex.jpg'],'-dpng','-r300');
size(allsubs_between_FC{1},2)
size(allsubs_between_FC{2},2)
allsubs_between_FC{1,s}(:,1:size(allsubs_between_FC{s},2))
mean(allsubs_between_FC{1,s}(:,1:size(allsubs_between_FC{s},2)),1)
figure;
s = 1;
plot(times_all(s,1:size(allsubs_between_FC{s},2)),mean(allsubs_between_FC{1,s}(:,1:size
(allsubs_between_FC{s},2)),1),'Color',rgb_colors_LS(s,:), 'LineWidth', 3)
hold on
s = 2;
plot(times_all(s,1:size(allsubs_between_FC{s},2)),mean(allsubs_between_FC{1,s}(:,1:size
(allsubs_between_FC{s},2)),1),'Color',rgb_colors_LS(s,:), 'LineWidth', 3)
hold on
s = 3;
plot(times_all(s,1:size(allsubs_between_FC{s},2)),mean(allsubs_between_FC{1,s}(:,1:size
(allsubs_between_FC{s},2)),1),'Color',rgb_colors_LS(s,:), 'LineWidth', 3)
hold on
times_all(s,1:size(allsubs_between_FC{s},2)
times_all(s,1:size(allsubs_between_FC{s},2))
figure;
for s = 1:numel(subject)
    plot(times_all(2,1:size(allsubs_between_FC{s},2)),mean(allsubs_between_FC{1,s}(:,1:size
(allsubs_between_FC{s},2)),1),'Color',rgb_colors_LS(s,:), 'LineWidth', 3)

```



```

hold on
end
for s = 1:numel(subject)
x = mean(allsubs_between_FC{1,s}(:,1:size(allsubs_between_FC{s},2)),1);
end
for s = 1:numel(subject)
x(s,:) = mean(allsubs_between_FC{1,s}(:,1:size(allsubs_between_FC{s},2)),1);
end
x = mean(allsubs_between_FC{1,s}(:,1:size(allsubs_between_FC{s},2)),1);
figure;
plot(times_all(2,1:size(allsubs_between_FC{s},2)),mean(allsubs_between_FC{1,s}(:,1:size(
allsubs_between_FC{s},2)),1), 'Color', rgb_colors_LS(s,:), 'LineWidth', 3)
hold on
s = 2;
plot(times_all(2,1:size(allsubs_between_FC{s},2)),mean(allsubs_between_FC{1,s}(:,1:size(
allsubs_between_FC{s},2)),1), 'Color', rgb_colors_LS(s,:), 'LineWidth', 3)
hold on
s = 3;
plot(times_all(2,1:size(allsubs_between_FC{s},2)),mean(allsubs_between_FC{1,s}(:,1:size(
allsubs_between_FC{s},2)),1), 'Color', rgb_colors_LS(s,:), 'LineWidth', 3)
hold on
mean(allsubs_between_FC{1,s}(:,1:size(allsubs_between_FC{s},2)),1)
s = 4;
plot(times_all(2,1:size(allsubs_between_FC{s},2)),mean(allsubs_between_FC{1,s}(:,1:size(
allsubs_between_FC{s},2)),1), 'Color', rgb_colors_LS(s,:), 'LineWidth', 3)
s = 5;
hold on
plot(times_all(2,1:size(allsubs_between_FC{s},2)),mean(allsubs_between_FC{1,s}(:,1:size(
allsubs_between_FC{s},2)),1), 'Color', rgb_colors_LS(s,:), 'LineWidth', 3)
s = 6;
hold on
plot(times_all(2,1:size(allsubs_between_FC{s},2)),mean(allsubs_between_FC{1,s}(:,1:size(
allsubs_between_FC{s},2)),1), 'Color', rgb_colors_LS(s,:), 'LineWidth', 3)
hold on
s = 7;
plot(times_all(2,1:size(allsubs_between_FC{s},2)),mean(allsubs_between_FC{1,s}(:,1:size(
allsubs_between_FC{s},2)),1), 'Color', rgb_colors_LS(s,:), 'LineWidth', 3)
figure;
for s = 1:numel(subject)
plot(times_all(2,1:size(allsubs_seg_ind{s},2)),mean(allsubs_between_FC{1,s}(:,1:size(
allsubs_seg_ind{s},2)),1), 'Color', rgb_colors_LS(s,:), 'LineWidth', 3)
hold on
end
for t = 1:length(times)
tmp = [];
for s = 1:numel(subject)
if t <= size(allsubs_seg_ind{1,s},2)
tmp = [tmp;allsubs_between_FC{1,s}(t)];
else
continue;
end
end
mean_of_means_bFC(t) = mean(tmp);
end
plot(times_all(1,1:40),mean_of_means_bFC(1:40), ':', 'Color', [0,0,0], 'LineWidth',3)
for t = 1:length(times)
tmp = [];
for s = 1:numel(subject)
if t <= size(allsubs_seg_ind{1,s},2)
tmp = [tmp;mean(allsubs_between_FC{1,s}(t))];

```

```

else
continue;
end
end
mean_of_means_bFC(t) = mean(tmp);
end
for t = 1:length(times)
tmp = [];
for s = 1:numel(subject)
if t <= size(allsubs_seg_ind{1,s},2)
tmp = [tmp;mean(allsubs_between_FC{1,s}(:,t))];
else
continue;
end
end
mean_of_means_bFC(t) = mean(tmp);
end
%means = [0.228121374    0.282854956  0.311825923  0.330394855  0.344075468  0.3540615
0.362085299  0.368580906  0.374353669  0.378886496  0.38303392  0.386997298  0.39018919
0.392779018  0.395671349  0.3921087    0.394652817  0.396349383  0.398223    0.39998822
0.401678269  0.402921549  0.404130596  0.405287927  0.406779144  0.407857626  0.408819803
0.40779906   0.4087539    0.409494202  0.41042813   0.411371703  0.4121878   0.412706817
0.413360333  0.414032785  0.41465492   0.419794232  0.420634614  0.420983464  0.421506702
0.421954376  0.422355276  0.42278862   0.423175282  0.42350195   0.423740618  0.42402081
0.434813233  0.42669529   0.42691267   0.42712188   0.42740375];
plot(times_all(1,1:40),mean_of_means_bFC(1:40), ':', 'Color', [0,0,0], 'LineWidth',3)
figure;
for s = 1:numel(subject)
plot(times_all(2,1:size(allsubs_seg_ind{s},2)),mean(allsubs_between_FC{1,s}(:,1:size
(allsubs_seg_ind{s},2)),1), 'Color',rgb_colors_LS(s,:), 'LineWidth', 3)
hold on
end
for t = 1:length(times)
tmp = [];
for s = 1:numel(subject)
if t <= size(allsubs_seg_ind{1,s},2)
tmp = [tmp;mean(allsubs_between_FC{1,s}(:,t))];
else
continue;
end
end
mean_of_means_bFC(t) = mean(tmp);
end
%means = [0.228121374    0.282854956  0.311825923  0.330394855  0.344075468  0.3540615
0.362085299  0.368580906  0.374353669  0.378886496  0.38303392  0.386997298  0.39018919
0.392779018  0.395671349  0.3921087    0.394652817  0.396349383  0.398223    0.39998822
0.401678269  0.402921549  0.404130596  0.405287927  0.406779144  0.407857626  0.408819803
0.40779906   0.4087539    0.409494202  0.41042813   0.411371703  0.4121878   0.412706817
0.413360333  0.414032785  0.41465492   0.419794232  0.420634614  0.420983464  0.421506702
0.421954376  0.422355276  0.42278862   0.423175282  0.42350195   0.423740618  0.42402081
0.434813233  0.42669529   0.42691267   0.42712188   0.42740375];
plot(times_all(1,1:40),mean_of_means_bFC(1:40), ':', 'Color', [0,0,0], 'LineWidth',3)
axis([0 100 0 .2])
axis([0 100 0.06 .2])
ylabel('Between-network FC');
xlabel('Time (Minutes)');
m = findobj(gca,'Type','line');
hleg1 = legend(m(1:(numel(subject)+1)), ['Mean' flip(subject)], 'Location', 'NorthEast');
hleg1.FontSize = 20;
ax = gca;

```

```

ax.FontSize = 20;
axis([0 100 0.06 .25])
print(gcf, '/Users/dianaperez/Desktop/ReliabilityBetweenFC.jpg', '-dpng', '-r300')
figure;
for s = 1:numel(subject)
plot(times_all(s,1:size(allsubs_seg_ind{s},2)),mean(allsubs_within_FC{1,s}(:,1:size(
(allsubs_seg_ind{s},2)),1), 'Color', rgb_colors_LS(s,:), 'LineWidth', 3)
hold on
end
for t = 1:length(times)
tmp = [];
for s = 1:numel(subject)
if t <= size(allsubs_seg_ind{1,s},2)
tmp = [tmp;allsubs_within_FC{1,s}(:,t)];
else
continue;
end
end
mean_of_means_wFC(t) = mean(tmp);
end
%means = [0.228121374    0.282854956 0.311825923 0.330394855 0.344075468 0.3540615
0.362085299 0.368580906 0.374353669 0.378886496 0.38303392 0.386997298 0.39018919
0.392779018 0.395671349 0.3921087    0.394652817 0.396349383 0.398223    0.39998822
0.401678269 0.402921549 0.404130596 0.405287927 0.406779144 0.407857626 0.408819803
0.40779906 0.4087539    0.409494202 0.41042813 0.411371703 0.4121878    0.412706817
0.413360333 0.414032785 0.41465492 0.419794232 0.420634614 0.420983464 0.421506702
0.421954376 0.422355276 0.42278862 0.423175282 0.42350195 0.423740618 0.42402081
0.434813233 0.42669529 0.42691267 0.42712188 0.42740375];
plot(times_all(1,1:40),mean_of_means_wFC(1:40), ':', 'Color', [0,0,0], 'LineWidth',3)
axis([0 100 0.09 .3])
ylabel('Within FC');
xlabel('Time (Minutes)');
output_dir = '/Users/dianaperez/Desktop/';
m = findobj(gca,'Type','line');
hleg1 = legend(m(1:(numel(subject)+1)), ['Mean' flip(subject)], 'Location', 'NorthEast');
hleg1.FontSize = 20;
ax = gca;
ax.FontSize = 20;
print(gcf,[output_dir '/ReliabilityWithinFC.jpg'],'-dpng','-r300');
xlabel('Time (Minutes)');
ylabel('Pearson correlation (r)');
ax = gca;
ax.FontSize = 20;
within_sub_reliability_v2
length(atlas_params.networks)
size(atlas_params.networks,ind)
within = tmp_within(maskmat);
within(within<0) = [];
system_divisions(sys)
similarity_analysis_by_system
maskmat = ones(size(corrmat_matched_data,1));
maskmat = logical(triu(maskmat, 1));
matcheddata_corrlin(count,:) = single(FisherTransform(corrmat_matched_data(maskmat)));
similarity_analysis_by_system
title('Correlation Matrix Similarity - sensorimotor systems');
output_str = {'sensorimotor_systems', 'control_systems'};
title(['Correlation Matrix Similarity - ']);
title(['Correlation Matrix Similarity - ' output_str(1)]);
title(['Correlation Matrix Similarity - ' output_str{1}]);
similarity_analysis_by_system

```

```

title('Correlation Matrix Similarity - sensorimotor');
similarity_analysis_by_system
disp(['The average similarity between subjects for ' output_str{sys} ' is ' num2str(mean(
(between)))]
disp(['The average similarity within subjects for ' output_str{sys} ' is ' num2str(mean(
(within)))]
disp(['The average similarity within subjects for ' output_str{sys} ' is ' num2str(mean(
(within)))]
disp(['The average similarity within subjects for ' output_str{sys} ' is ' num2str(mean(
(within)))]
disp(['The average similarity between subjects for ' output_str{sys} ' is ' num2str(mean(
(between)))]
disp(['The average similarity within subjects for ' output_str{sys} ' is ' num2str(mean(
(within)))]
similarity_analysis_by_system
help gifti
clear all
gii_surf = gifti('/Volumes/fsmresfiles/PBS/Gratton_Lab/Lifespan/Pre-
COVID/BIDS/Nifti/derivatives/freesurfer-6.0.1/FREESURFER_fs_LR/sub-
LS02/NativeVol/fsaverage_LR32k/sub-LS02.L.midthickness.32k_fs_LR.surf.gii');
gii_surf = gifti('/Volumes/fsmresfiles/PBS/Gratton_Lab/Lifespan/Pre-
COVID/BIDS/Nifti/derivatives/freesurfer-6.0.1/FREESURFER_fs_LR/sub-
LS02/NativeVol/fsaverage_LR32k/sub-LS02.L.midthickness.32k_fs_LR.surf.gii');
gii_coord = gifti('/Volumes/fsmresfiles/PBS/Gratton_Lab/Lifespan/Pre-
COVID/BIDS/Nifti/derivatives/freesurfer-6.0.1/FREESURFER_fs_LR/sub-
LS02/NativeVol/fsaverage_LR32k/sub-LS02.L.midthickness.32k_fs_LR.coord.gii');
gii_surf(:,1)
gii_surf.vertices(:,1)
gii_surf.vertices(1,:)
gii_coord.vertices(1,:)
gii_surf = load('/Volumes/fsmresfiles/PBS/Gratton_Lab/Lifespan/Pre-
COVID/BIDS/Nifti/derivatives/freesurfer-6.0.1/FREESURFER_fs_LR/sub-
LS02/NativeVol/fsaverage_LR32k/sub-LS02.L.midthickness.32k_fs_LR.surf.gii');
gii_surf = load('/Volumes/fsmresfiles/PBS/Gratton_Lab/Lifespan/Pre-
COVID/BIDS/Nifti/derivatives/freesurfer-6.0.1/FREESURFER_fs_LR/sub-
LS02/NativeVol/fsaverage_LR32k/sub-LS02.L.midthickness.32k_fs_LR.surf.gii');
gii_coord.mat;
indparcels_dismats_gw
clear gii*
clear ans
%-- 2/27/23, 4:50 PM --%
make_geo_dist_matrix
addpath '/Users/dianaperez/Documents/GitHub/General_Scripts/gifti-master/@gifti'
make_geo_dist_matrix
addpath '/Users/dianaperez/Documents/GitHub/General_Scripts/gifti-master/@gifti'
make_geo_dist_matrix
cd '/Users/dianaperez/Documents/GitHub/General_Scripts/gifti-master/@gifti'
make_geo_dist_matrix
L_surf_name
L_surf_fname
make_geo_dist_matrix
%-- 2/28/23, 3:06 PM --%
similarity_analysis_by_system
addpath '/Users/dianaperez/Documents/GitHub/GrattonLab-General-Repo/FCProcess'
similarity_analysis_by_system
addpath '/Users/dianaperez/Documents/GitHub/Lifespan-Analysis/utilities'
similarity_analysis_by_system
addpath '/Users/dianaperez/Documents/GitHub/General_Scripts'
similarity_analysis_by_system
output_str{sys}

```

```

sys
output_str = {'sensorimotor', 'control'};
title(['Correlation Matrix Similarity - ' output_str{sys}]);
if match_data
saveas(gcf,[output_dir data_set '_' atlas '_' output_str{sys}
'_similarityMat_matchedData_' num2str(amt_data) '.tiff'],'tiff');
else
saveas(gcf,[output_dir data_set '_' atlas '_' output_str{sys}
'_similarityMat_unMatchedData.tiff'],'tiff');
end
close('all');
%% CALCULATE average within- and between-subject correlations
count = 1;
within = [];
between = [];
for s = 1:numel(subject)
lines = [count:(count+sessions-1)];
sub_vals = simmat(lines,:);
maskmat = ones(sessions,sessions);
maskmat = logical(triu(maskmat, 1));
within_sub = sub_vals(:,lines);
within = [within; within_sub(maskmat)];
maskmat = ones(size(sub_vals));
maskmat(:,lines) = 0;
between = [between; sub_vals(maskmat==1)];
count = count+sessions;
end
disp(['The average similarity between subjects for ' output_str{sys} ' is ' num2str(mean(
(between))])
disp(['The average similarity within subjects for ' output_str{sys} ' is ' num2str(mean(
(within))])
similarity_analysis_by_system
%-- 3/5/23, 2:16 PM --%
segregation_ind_by_ses
addpath '/Users/dianaperez/Documents/GitHub/GrattonLab-General-Repo/FCProcess'
segregation_ind_by_ses
addpath '/Users/dianaperez/Documents/GitHub/Lifespan-Analysis/utilities'
segregation_ind_by_ses
addpath '/Users/dianaperez/Documents/GitHub/General_Scripts'
segregation_ind_by_ses
matrix_sorted = matrix(atlas_params.sorti);
matched_data_sorted = matched_data(atlas_params.sorti);
matched_data_sorted = matched_data(atlas_params.sorti,:);
matched_data_sorted = matched_data(atlas_params.sorti,:);
matrix = paircorr_mod(matched_data_sorted');
matrix = single(FisherTransform(matrix));
imagesc(matrix)
imagesc(matrix)
load better_jet_colormap.mat; % assume this is in the same folder
colormap(better_jet_colormap_diff);
imagesc(matrix, -1, 1)
imagesc(matrix, [-1 1])
net == 1
segregation_ind_by_ses
rois(end)
segregation_ind_by_ses
strcmpr(neg_corrs, 'nan')
segregation_ind_by_ses
[output_dir data_set '_allsubs_seg_index_ses_negcorrs' neg_corrs '_' atlas '.mat']
segregation_ind_by_ses

```

```

mean(all_within)
segregation_ind_by_ses
mean(all_within)
mean(all_between)
mean(all_within)
segregation_ind_by_ses
size(atlas_params.networks)
length(atlas_params.networks)
segregation_ind_by_ses
clear all
load
('Users/dianaperez/Desktop/Research/Segregation_Analyses/Lifespan_allsubs_seg_index_ses_
negcorrnsnan_Seitzman300.mat')
all_OA_subs = load
('Users/dianaperez/Desktop/Research/Segregation_Analyses/Lifespan_allsubs_seg_index_ses_
negcorrnsnan_Seitzman300.mat')
all_OA_subs = all_OA_subs.ses_SI;
make_figures
pre_first_ses = [0.40465483 0.47502443 0.42045888];
pre_first_ses = [0.40465483 0.47502443 0.42045888];
post_first_ses = [0.35866937 0.35903123 0.39794636 0.39622128 0.37635228 0.48386648
0.45387810 0.50429422 0.49092054 0.35866398 0.49722296 0.45307758 0.44976237 0.40819031
0.39566854];
[pre_first_ses post_first_ses]
make_figures
handles = plotSpread(first_ses_long_OA', 'distributionMarkers', {'+'},
'distributionColors', rgb_for_long, 'xNames', subs)
make_figures
close all
make_figures
axis([0 19 .5 1.5])
axis([0 19 .7 1.1])
axis([0 19 .8 1.1])
axis([0 19 .8 1.05])
ax = gca;
ax.FontSize = 20;
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.3, 0.7]); %first and second
control position on screen, third controls width, and fourth controls height
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.7, 0.3]); %first and second
control position on screen, third controls width, and fourth controls height
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 0.7, 0.5]); %first and second
control position on screen, third controls width, and fourth controls height
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 1, 0.5]); %first and second
control position on screen, third controls width, and fourth controls height
set(gcf, 'Units', 'Normalized', 'OuterPosition', [0.3, 0.3, 1, 0.7]); %first and second
control position on screen, third controls width, and fourth controls height
print(gcf, [output_dir 'SegInd_stability_negcorrs' neg_corrs '_' atlas '.jpg'], '-dpng',
'-r300')
make_figures
addpath
'Users/dianaperez/Documents/GitHub/PerezEtAl_HemAsymmetries/needed_files/plotSpread/plot
Spread'
make_figures
axis([0 19 0.6 0.8])
make_figures
axis([0 19 0.4 0.6])
make_figures
close all
clear all
make_figures

```

```
axis([0 19 .25 .6])
make_figures
close all
make_figures
axis([0 19 .4 .8])
axis([0 19 .5 .8])
make_figures
axis([0 19 .8 1.05]) % for the as is neg corrs parcels333
axis([0 19 .8 1.1]) % for the as is neg corrs parcels333
axis([0 19 .8 1.2]) % for the as is neg corrs parcels333
axis([0 19 .8 1.15]) % for the as is neg corrs parcels333
axis([0 19 .9 1.15]) % for the as is neg corrs parcels333
axis([0 19 .85 1.15]) % for the as is neg corrs parcels333
axis([0 19 .85 1.15])
clear all
%% some paths
data_dir = '/Users/dianaperez/Desktop/Research/Segregation_Analyses/';
output_dir = '/Users/dianaperez/Desktop/Research/Segregation_Analyses/';
%% some variables
% I don't think I'll need these, but just in case
subs = {'LS02', 'LS03', 'LS05', 'LS08', 'LS11', 'LS14', 'LS16', 'LS17', 'INET001', 'INET002', 'INET003', 'INET005', 'INET006', 'INET010', 'INET016', 'INET018', 'INET019', 'INET030'};
%% first load the first batch
neg_corrs = 'asis'; % choose: 'nan', 'zero', 'asis'
atlas = 'Seitzman300'; % 'Seitzman300' or 'Parcels333'
%% load all the files
% these are the TimeB segregation indices for older adults
OA_vol_asis = load([data_dir 'Lifespan_allsubs_seg_index_ses_negcorrs' neg_corrs '_' atlas '.mat']);
OA_vol_asis = OA_vol_asis.ses_SI;
% these are the segregation indices for young adults
YA_vol_asis = load([data_dir 'iNetworks_allsubs_seg_index_ses_negcorrs' neg_corrs '_' atlas '.mat']);
YA_vol_asis = YA_vol_asis.ses_SI;
%% load the next batch
neg_corrs = 'zero'; % choose: 'nan', 'zero', 'asis'
atlas = 'Seitzman300'; % 'Seitzman300' or 'Parcels333'
%% load all the files
% these are the TimeB segregation indices for older adults
OA_vol_zero = load([data_dir 'Lifespan_allsubs_seg_index_ses_negcorrs' neg_corrs '_' atlas '.mat']);
OA_vol_zero = OA_vol_zero.ses_SI;
% these are the segregation indices for young adults
YA_vol_zero = load([data_dir 'iNetworks_allsubs_seg_index_ses_negcorrs' neg_corrs '_' atlas '.mat']);
YA_vol_zero = YA_vol_zero.ses_SI;
%% load the next batch
neg_corrs = 'nan'; % choose: 'nan', 'zero', 'asis'
atlas = 'Seitzman300'; % 'Seitzman300' or 'Parcels333'
%% load all the files
% these are the TimeB segregation indices for older adults
OA_vol_nan = load([data_dir 'Lifespan_allsubs_seg_index_ses_negcorrs' neg_corrs '_' atlas '.mat']);
OA_vol_nan = OA_vol_nan.ses_SI;
% these are the segregation indices for young adults
YA_vol_nan = load([data_dir 'iNetworks_allsubs_seg_index_ses_negcorrs' neg_corrs '_' atlas '.mat']);
YA_vol_nan = YA_vol_nan.ses_SI;
%% load the first batch
```



```
neg_corrs = 'asis'; % choose: 'nan', 'zero', 'asis'
atlas = 'Parcels333'; % 'Seitzman300' or 'Parcels333'
%% load all the files
% these are the TimeB segregation indices for older adults
OA_surf_asis = load([data_dir 'Lifespan_allsubs_seg_index_ses_negcorrs' neg_corrs '_'
atlas '.mat']);
OA_surf_asis = OA_surf_asis.ses_SI;
% these are the segregation indices for young adults
YA_surf_asis = load([data_dir 'iNetworks_allsubs_seg_index_ses_negcorrs' neg_corrs '_'
atlas '.mat']);
YA_surf_asis = YA_surf_asis.ses_SI;
%% load the next batch
neg_corrs = 'zero'; % choose: 'nan', 'zero', 'asis'
atlas = 'Parcels333'; % 'Seitzman300' or 'Parcels333'
%% load all the files
% these are the TimeB segregation indices for older adults
OA_surf_zero = load([data_dir 'Lifespan_allsubs_seg_index_ses_negcorrs' neg_corrs '_'
atlas '.mat']);
OA_surf_zero = OA_surf_zero.ses_SI;
% these are the segregation indices for young adults
YA_surf_zero = load([data_dir 'iNetworks_allsubs_seg_index_ses_negcorrs' neg_corrs '_'
atlas '.mat']);
YA_surf_zero = YA_surf_zero.ses_SI;
%% load the next batch
neg_corrs = 'nan'; % choose: 'nan', 'zero', 'asis'
atlas = 'Parcels'; % 'Seitzman300' or 'Parcels333'
%% load all the files
% these are the TimeB segregation indices for older adults
OA_surf_nan = load([data_dir 'Lifespan_allsubs_seg_index_ses_negcorrs' neg_corrs '_'
atlas '.mat']);
OA_surf_nan = OA_surf_nan.ses_SI;
% these are the segregation indices for young adults
YA_surf_nan = load([data_dir 'iNetworks_allsubs_seg_index_ses_negcorrs' neg_corrs '_'
atlas '.mat']);
YA_surf_nan = YA_surf_nan.ses_SI;
neg_corrs = 'nan'; % choose: 'nan', 'zero', 'asis'
atlas = 'Parcels333'; % 'Seitzman300' or 'Parcels333'
%% load all the files
% these are the TimeB segregation indices for older adults
OA_surf_nan = load([data_dir 'Lifespan_allsubs_seg_index_ses_negcorrs' neg_corrs '_'
atlas '.mat']);
OA_surf_nan = OA_surf_nan.ses_SI;
% these are the segregation indices for young adults
YA_surf_nan = load([data_dir 'iNetworks_allsubs_seg_index_ses_negcorrs' neg_corrs '_'
atlas '.mat']);
YA_surf_nan = YA_surf_nan.ses_SI;
test = OA_surf_asis(:);
YA_surf_nan = YA_surf_nan.ses_SI(:);
YA_surf_nan(:) = YA_surf_nan.ses_SI(:);
YA_surf_nan(:) = YA_surf_nan.ses_SI;
YA_vol_zero = YA_vol_zero(:)
load('sub-LS03_infomap_conBensus_weighted_minsize400.mat')
clear all
load('sub-LS03_infomap_conBensus_weighted_minsize400.mat')
load('sub-LS02_infomap_conBensus_weighted_minsize400.mat')
tmp = ft_read_cifti_mod('/corrofcrr_allgrad_LR_subcort_smooth2.55.dtseries.nii');
tmp = ft_read_cifti_mod('corrofcrr_allgrad_LR_subcort_smooth2.55.dtseries.nii');
data = consen(brainstructure>0);
data = consen(tmp.brainstructure>0);
tmp.data = data;
```



```
ft_write_cifti_mod('/sub-LS02_infomap_conBensus_weighted_minsize400.dtseries.nii', tmp);
ft_write_cifti_mod('sub-LS02_infomap_conBensus_weighted_minsize400.dtseries.nii', tmp);
unique(tmp.brainstructure)
load('sub-LS03_infomap_conBensus_weighted_minsize400.mat')
data = consen(tmp.brainstructure>0);
tmp.data = data;
ft_write_cifti_mod('sub-LS03_infomap_conBensus_weighted_minsize400.dtseries.nii', tmp);
dlmread('rawassn.txt')
rawassn = ans;
data = rawassn(:,1);
data = data(tmp.brainstructure>0);
tmp.data = data;
ft_write_cifti_mod('test_raw_assn_col1.dtseries.nii', tmp);
data = rawassn(:,2);
data = data(tmp.brainstructure>0);
tmp.data = data;
ft_write_cifti_mod('test_raw_assn_col2.dtseries.nii', tmp);
data = rawassn(:,3);
data = data(tmp.brainstructure>0);
tmp.data = data;
ft_write_cifti_mod('test_raw_assn_col3.dtseries.nii', tmp);
data = rawassn(:,4);
data = data(tmp.brainstructure>0);
tmp.data = data;
ft_write_cifti_mod('test_raw_assn_col4.dtseries.nii', tmp);
data = rawassn(:,5);
data = data(tmp.brainstructure>0);
tmp.data = data;
ft_write_cifti_mod('test_raw_assn_col5.dtseries.nii', tmp);
data = rawassn(:,6);
data = data(tmp.brainstructure>0);
tmp.data = data;
ft_write_cifti_mod('test_raw_assn_col6.dtseries.nii', tmp);
data = rawassn(:,7);
data = data(tmp.brainstructure>0);
tmp.data = data;
ft_write_cifti_mod('test_raw_assn_col7.dtseries.nii', tmp);
data = rawassn(:,8);
data = data(tmp.brainstructure>0);
tmp.data = data;
ft_write_cifti_mod('test_raw_assn_col8.dtseries.nii', tmp);
data = rawassn(:,9);
data = data(tmp.brainstructure>0);
tmp.data = data;
ft_write_cifti_mod('test_raw_assn_col9.dtseries.nii', tmp);
data = rawassn(:,10);
data = data(tmp.brainstructure>0);
tmp.data = data;
ft_write_cifti_mod('test_raw_assn_col10.dtseries.nii', tmp);
data = rawassn(:,11);
data = data(tmp.brainstructure>0);
tmp.data = data;
ft_write_cifti_mod('test_raw_assn_col11.dtseries.nii', tmp);
data = rawassn(:,12);
data = data(tmp.brainstructure>0);
tmp.data = data;
ft_write_cifti_mod('test_raw_assn_col12.dtseries.nii', tmp);
clear all
%% some paths
data_dir = '/Users/dianaperez/Desktop/Research/Segregation_Analyses/';
```

```
output_dir = '/Users/dianaperez/Desktop/Research/Segregation_Analyses/';
%% some variables
% I don't think I'll need these, but just in case
subs = {'LS02', 'LS03', 'LS05', 'LS08', 'LS11', 'LS14', 'LS16', 'LS17', 'INET001', 'INET002', 'INET003', 'INET005', 'INET006', 'INET010', 'INET016', 'INET018', 'INET019', 'INET030'};
rgb_colors = {[1 0 0], [0, 0, 1]}; % red and blue
%% first load the first batch
neg_corrs = 'asis'; % choose: 'nan', 'zero', 'asis'
atlas = 'Seitzman300'; % 'Seitzman300' or 'Parcels333'
%% load all the files
% these are the TimeB segregation indices for older adults
OA_vol_asis = load([data_dir 'Lifespan_allsubs_seg_index_ses_negcorrs' neg_corrs '_' atlas '.mat']);
OA_vol_asis = OA_vol_asis.ses_SI;
OA_vol_asis = OA_vol_asis(:);
% these are the segregation indices for young adults
YA_vol_asis = load([data_dir 'iNetworks_allsubs_seg_index_ses_negcorrs' neg_corrs '_' atlas '.mat']);
YA_vol_asis = YA_vol_asis.ses_SI;
YA_vol_asis = YA_vol_asis(:);
%% load the next batch
neg_corrs = 'zero'; % choose: 'nan', 'zero', 'asis'
atlas = 'Seitzman300'; % 'Seitzman300' or 'Parcels333'
%% load all the files
% these are the TimeB segregation indices for older adults
OA_vol_zero = load([data_dir 'Lifespan_allsubs_seg_index_ses_negcorrs' neg_corrs '_' atlas '.mat']);
OA_vol_zero = OA_vol_zero.ses_SI;
OA_vol_zero = OA_vol_zero(:);
% these are the segregation indices for young adults
YA_vol_zero = load([data_dir 'iNetworks_allsubs_seg_index_ses_negcorrs' neg_corrs '_' atlas '.mat']);
YA_vol_zero = YA_vol_zero.ses_SI;
YA_vol_zero = YA_vol_zero(:);
%% load the next batch
neg_corrs = 'nan'; % choose: 'nan', 'zero', 'asis'
atlas = 'Seitzman300'; % 'Seitzman300' or 'Parcels333'
%% load all the files
% these are the TimeB segregation indices for older adults
OA_vol_nan = load([data_dir 'Lifespan_allsubs_seg_index_ses_negcorrs' neg_corrs '_' atlas '.mat']);
OA_vol_nan = OA_vol_nan.ses_SI;
OA_vol_nan = OA_vol_nan(:);
% these are the segregation indices for young adults
YA_vol_nan = load([data_dir 'iNetworks_allsubs_seg_index_ses_negcorrs' neg_corrs '_' atlas '.mat']);
YA_vol_nan = YA_vol_nan.ses_SI;
YA_vol_nan = YA_vol_nan(:);
%% load the first batch
neg_corrs = 'asis'; % choose: 'nan', 'zero', 'asis'
atlas = 'Parcels333'; % 'Seitzman300' or 'Parcels333'
%% load all the files
% these are the TimeB segregation indices for older adults
OA_surf_asis = load([data_dir 'Lifespan_allsubs_seg_index_ses_negcorrs' neg_corrs '_' atlas '.mat']);
OA_surf_asis = OA_surf_asis.ses_SI;
OA_surf_asis = OA_surf_asis(:);
% these are the segregation indices for young adults
YA_surf_asis = load([data_dir 'iNetworks_allsubs_seg_index_ses_negcorrs' neg_corrs '_' atlas '.mat']);
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atlas '.mat']));
YA_surf_asis = YA_surf_asis.ses_SI;
YA_surf_asis = YA_surf_asis(:);
%% load the next batch
neg_corrs = 'zero'; % choose: 'nan', 'zero', 'asis'
atlas = 'Parcels333'; % 'Seitzman300' or 'Parcels333'
%% load all the files
% these are the TimeB segregation indices for older adults
OA_surf_zero = load([data_dir 'Lifespan_allsubs_seg_index_ses_negcorrs' neg_corrs '_'
atlas '.mat']));
OA_surf_zero = OA_surf_zero.ses_SI;
OA_surf_zero = OA_surf_zero(:);
% these are the segregation indices for young adults
YA_surf_zero = load([data_dir 'iNetworks_allsubs_seg_index_ses_negcorrs' neg_corrs '_'
atlas '.mat']));
YA_surf_zero = YA_surf_zero.ses_SI;
YA_surf_zero = YA_surf_zero(:);
%% load the next batch
neg_corrs = 'nan'; % choose: 'nan', 'zero', 'asis'
atlas = 'Parcels333'; % 'Seitzman300' or 'Parcels333'
%% load all the files
% these are the TimeB segregation indices for older adults
OA_surf_nan = load([data_dir 'Lifespan_allsubs_seg_index_ses_negcorrs' neg_corrs '_'
atlas '.mat']));
OA_surf_nan = OA_surf_nan.ses_SI;
OA_surf_nan = OA_surf_nan(:);
% these are the segregation indices for young adults
YA_surf_nan = load([data_dir 'iNetworks_allsubs_seg_index_ses_negcorrs' neg_corrs '_'
atlas '.mat']));
YA_surf_nan = YA_surf_nan.ses_SI(:);
YA_surf_nan = YA_surf_nan(:);
%% Now how do I plot these?
OA_surf = [OA_surf_asis; OA_surf_zero; OA_surf_nan];
OA_vol = [OA_vol_asis; OA_vol_zero; OA_vol_nan];
YA_surf = [YA_surf_asis; YA_surf_zero; YA_surf_nan];
YA_vol = [YA_vol_asis; YA_vol_zero; YA_vol_nan];
handles = plotSpread([OA_surf_asis OA_vol_asis], 'distributionMarkers', {'o'},
'distributionColors', [0,0,1], 'binWidth', .5);
close all
handles = plotSpread_v2([OA_surf_asis OA_vol_asis], 'distributionMarkers', {'o'},
'distributionColors', [0,0,1], 'binWidth', .5);
hold on
close all
handles = plotSpread([OA_surf_asis OA_vol_asis], 'distributionMarkers', {'o'},
'distributionColors', [0,0,1], 'binWidth', .5);
hold on
handles = plotSpread([YA_surf_asis YA_vol_asis], 'distributionMarkers', {'o'},
'distributionColors', [1,0,0], 'binWidth', .5);
hold on
clear all
handles = plotSpread([OA_surf_asis OA_vol_asis], 'distributionMarkers', {'+'},
'distributionColors', [0,0,1], 'binWidth', .5);
hold on
handles = plotSpread([YA_surf_asis YA_vol_asis], 'distributionMarkers', {'+'},
'distributionColors', [1,0,0], 'binWidth', .5);
hold on
handles = plotSpread([OA_surf_zero OA_vol_zero], 'distributionMarkers', {'o'},
'distributionColors', [0,0,1], 'binWidth', .5);
hold on
handles = plotSpread([YA_surf_zero YA_vol_zero], 'distributionMarkers', {'o'},
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'distributionColors', [1,0,0], 'binWidth', .5);
hold on
handles = plotSpread([OA_surf_nan OA_vol_nan], 'distributionMarkers', {'*'}, ↵
'distributionColors', [0,0,1], 'binWidth', .5);
hold on
handles = plotSpread([YA_surf_nan YA_vol_nan], 'distributionMarkers', {'*'}, ↵
'distributionColors', [1,0,0], 'binWidth', .5);
hold on
%% some variables
% I don't think I'll need these, but just in case
subs = {'LS02', 'LS03', 'LS05', 'LS08', 'LS11', 'LS14', ↵
'LS16', 'LS17', 'INET001', 'INET002', ↵
'INET003', 'INET005', 'INET006', 'INET010', 'INET016', 'INET018', 'INET019', 'INET030'};
rgb_colors = {[1 0 0], [0, 0, 1]}; % red and blue
%% first load the first batch
neg_corrs = 'asis'; % choose: 'nan', 'zero', 'asis'
atlas = 'Seitzman300'; % 'Seitzman300' or 'Parcels333'
%% load all the files
% these are the TimeB segregation indices for older adults
OA_vol_asis = load([data_dir 'Lifespan_allsubs_seg_index_ses_negcorrs' neg_corrs '_' ↵
atlas '.mat']);
OA_vol_asis = OA_vol_asis.ses_SI;
OA_vol_asis = OA_vol_asis(:);
% these are the segregation indices for young adults
YA_vol_asis = load([data_dir 'iNetworks_allsubs_seg_index_ses_negcorrs' neg_corrs '_' ↵
atlas '.mat']);
YA_vol_asis = YA_vol_asis.ses_SI;
YA_vol_asis = YA_vol_asis(:);
%% load the next batch
neg_corrs = 'zero'; % choose: 'nan', 'zero', 'asis'
atlas = 'Seitzman300'; % 'Seitzman300' or 'Parcels333'
%% load all the files
% these are the TimeB segregation indices for older adults
OA_vol_zero = load([data_dir 'Lifespan_allsubs_seg_index_ses_negcorrs' neg_corrs '_' ↵
atlas '.mat']);
OA_vol_zero = OA_vol_zero.ses_SI;
OA_vol_zero = OA_vol_zero(:);
% these are the segregation indices for young adults
YA_vol_zero = load([data_dir 'iNetworks_allsubs_seg_index_ses_negcorrs' neg_corrs '_' ↵
atlas '.mat']);
YA_vol_zero = YA_vol_zero.ses_SI;
YA_vol_zero = YA_vol_zero(:);
%% load the next batch
neg_corrs = 'nan'; % choose: 'nan', 'zero', 'asis'
atlas = 'Seitzman300'; % 'Seitzman300' or 'Parcels333'
%% load all the files
% these are the TimeB segregation indices for older adults
OA_vol_nan = load([data_dir 'Lifespan_allsubs_seg_index_ses_negcorrs' neg_corrs '_' atlas ↵
'.mat']);
OA_vol_nan = OA_vol_nan.ses_SI;
OA_vol_nan = OA_vol_nan(:);
% these are the segregation indices for young adults
YA_vol_nan = load([data_dir 'iNetworks_allsubs_seg_index_ses_negcorrs' neg_corrs '_' ↵
atlas '.mat']);
YA_vol_nan = YA_vol_nan.ses_SI;
YA_vol_nan = YA_vol_nan(:);
%% load the first batch
neg_corrs = 'asis'; % choose: 'nan', 'zero', 'asis'
atlas = 'Parcels333'; % 'Seitzman300' or 'Parcels333'
%% load all the files
```

```
% these are the TimeB segregation indices for older adults
OA_surf_asis = load([data_dir 'Lifespan_allsubs_seg_index_ses_negcorrs' neg_corrs '_'
atlas '.mat']);
OA_surf_asis = OA_surf_asis.ses_SI;
OA_surf_asis = OA_surf_asis(:);
% these are the segregation indices for young adults
YA_surf_asis = load([data_dir 'iNetworks_allsubs_seg_index_ses_negcorrs' neg_corrs '_'
atlas '.mat']);
YA_surf_asis = YA_surf_asis.ses_SI;
YA_surf_asis = YA_surf_asis(:);
%% load the next batch
neg_corrs = 'zero'; % choose: 'nan', 'zero', 'asis'
atlas = 'Parcels333'; % 'Seitzman300' or 'Parcels333'
%% load all the files
% these are the TimeB segregation indices for older adults
OA_surf_zero = load([data_dir 'Lifespan_allsubs_seg_index_ses_negcorrs' neg_corrs '_'
atlas '.mat']);
OA_surf_zero = OA_surf_zero.ses_SI;
OA_surf_zero = OA_surf_zero(:);
% these are the segregation indices for young adults
YA_surf_zero = load([data_dir 'iNetworks_allsubs_seg_index_ses_negcorrs' neg_corrs '_'
atlas '.mat']);
YA_surf_zero = YA_surf_zero.ses_SI;
YA_surf_zero = YA_surf_zero(:);
%% load the next batch
neg_corrs = 'nan'; % choose: 'nan', 'zero', 'asis'
atlas = 'Parcels333'; % 'Seitzman300' or 'Parcels333'
%% load all the files
% these are the TimeB segregation indices for older adults
OA_surf_nan = load([data_dir 'Lifespan_allsubs_seg_index_ses_negcorrs' neg_corrs '_'
atlas '.mat']);
OA_surf_nan = OA_surf_nan.ses_SI;
OA_surf_nan = OA_surf_nan(:);
% these are the segregation indices for young adults
YA_surf_nan = load([data_dir 'iNetworks_allsubs_seg_index_ses_negcorrs' neg_corrs '_'
atlas '.mat']);
YA_surf_nan = YA_surf_nan.ses_SI(:);
YA_surf_nan = YA_surf_nan(:);
%% Now how do I plot these?
handles = plotSpread([OA_surf_asis OA_vol_asis], 'distributionMarkers', {'+'},
'distributionColors', [0,0,1], 'binWidth', .5);
hold on
handles = plotSpread([YA_surf_asis YA_vol_asis], 'distributionMarkers', {'+'},
'distributionColors', [1,0,0], 'binWidth', .5);
hold on
handles = plotSpread([OA_surf_zero OA_vol_zero], 'distributionMarkers', {'o'},
'distributionColors', [0,0,1], 'binWidth', .5);
hold on
handles = plotSpread([YA_surf_zero YA_vol_zero], 'distributionMarkers', {'o'},
'distributionColors', [1,0,0], 'binWidth', .5);
hold on
handles = plotSpread([OA_surf_nan OA_vol_nan], 'distributionMarkers', {'*'},
'distributionColors', [0,0,1], 'binWidth', .5);
hold on
handles = plotSpread([YA_surf_nan YA_vol_nan], 'distributionMarkers', {'*'},
'distributionColors', [1,0,0], 'binWidth', .5);
hold on
data_dir = '/Users/dianaperez/Desktop/Research/Segregation_Analyses/';
output_dir = '/Users/dianaperez/Desktop/Research/Segregation_Analyses/';
%% some variables
```

```
% I don't think I'll need these, but just in case
subs = {'LS02', 'LS03', 'LS05', 'LS08', 'LS11', 'LS14', '
'LS16', 'LS17', 'INET001', 'INET002', '
'INET003', 'INET005', 'INET006', 'INET010', 'INET016', 'INET018', 'INET019', 'INET030'};
rgb_colors = {[1 0 0], [0, 0, 1]};% red and blue
%% first load the first batch
neg_corrs = 'asis'; % choose: 'nan', 'zero', 'asis'
atlas = 'Seitzman300'; % 'Seitzman300' or 'Parcels333'
%% load all the files
% these are the TimeB segregation indices for older adults
OA_vol_asis = load([data_dir 'Lifespan_allsubs_seg_index_ses_negcorrs' neg_corrs '_' atlas
'.mat']);
OA_vol_asis = OA_vol_asis.ses_SI;
OA_vol_asis = OA_vol_asis(:);
% these are the segregation indices for young adults
YA_vol_asis = load([data_dir 'iNetworks_allsubs_seg_index_ses_negcorrs' neg_corrs '_' atlas
'.mat']);
YA_vol_asis = YA_vol_asis.ses_SI;
YA_vol_asis = YA_vol_asis(:);
%% load the next batch
neg_corrs = 'zero'; % choose: 'nan', 'zero', 'asis'
atlas = 'Seitzman300'; % 'Seitzman300' or 'Parcels333'
%% load all the files
% these are the TimeB segregation indices for older adults
OA_vol_zero = load([data_dir 'Lifespan_allsubs_seg_index_ses_negcorrs' neg_corrs '_' atlas
'.mat']);
OA_vol_zero = OA_vol_zero.ses_SI;
OA_vol_zero = OA_vol_zero(:);
% these are the segregation indices for young adults
YA_vol_zero = load([data_dir 'iNetworks_allsubs_seg_index_ses_negcorrs' neg_corrs '_' atlas
'.mat']);
YA_vol_zero = YA_vol_zero.ses_SI;
YA_vol_zero = YA_vol_zero(:);
%% load the next batch
neg_corrs = 'nan'; % choose: 'nan', 'zero', 'asis'
atlas = 'Seitzman300'; % 'Seitzman300' or 'Parcels333'
%% load all the files
% these are the TimeB segregation indices for older adults
OA_vol_nan = load([data_dir 'Lifespan_allsubs_seg_index_ses_negcorrs' neg_corrs '_' atlas
'.mat']);
OA_vol_nan = OA_vol_nan.ses_SI;
OA_vol_nan = OA_vol_nan(:);
% these are the segregation indices for young adults
YA_vol_nan = load([data_dir 'iNetworks_allsubs_seg_index_ses_negcorrs' neg_corrs '_' atlas
'.mat']);
YA_vol_nan = YA_vol_nan.ses_SI;
YA_vol_nan = YA_vol_nan(:);
%% load the first batch
neg_corrs = 'asis'; % choose: 'nan', 'zero', 'asis'
atlas = 'Parcels333'; % 'Seitzman300' or 'Parcels333'
%% load all the files
% these are the TimeB segregation indices for older adults
OA_surf_asis = load([data_dir 'Lifespan_allsubs_seg_index_ses_negcorrs' neg_corrs '_' atlas
'.mat']);
OA_surf_asis = OA_surf_asis.ses_SI;
OA_surf_asis = OA_surf_asis(:);
% these are the segregation indices for young adults
YA_surf_asis = load([data_dir 'iNetworks_allsubs_seg_index_ses_negcorrs' neg_corrs '_' atlas
'.mat']);
YA_surf_asis = YA_surf_asis.ses_SI;
```



```
YA_surf_asis = YA_surf_asis(:);
%% load the next batch
neg_corrs = 'zero'; % choose: 'nan', 'zero', 'asis'
atlas = 'Parcels333'; % 'Seitzman300' or 'Parcels333'
%% load all the files
% these are the TimeB segregation indices for older adults
OA_surf_zero = load([data_dir 'Lifespan_allsubs_seg_index_ses_negcorrs' neg_corrs '_'
atlas '.mat']);
OA_surf_zero = OA_surf_zero.ses_SI;
OA_surf_zero = OA_surf_zero(:);
% these are the segregation indices for young adults
YA_surf_zero = load([data_dir 'iNetworks_allsubs_seg_index_ses_negcorrs' neg_corrs '_'
atlas '.mat']);
YA_surf_zero = YA_surf_zero.ses_SI;
YA_surf_zero = YA_surf_zero(:);
%% load the next batch
neg_corrs = 'nan'; % choose: 'nan', 'zero', 'asis'
atlas = 'Parcels333'; % 'Seitzman300' or 'Parcels333'
%% load all the files
% these are the TimeB segregation indices for older adults
OA_surf_nan = load([data_dir 'Lifespan_allsubs_seg_index_ses_negcorrs' neg_corrs '_'
atlas '.mat']);
OA_surf_nan = OA_surf_nan.ses_SI;
OA_surf_nan = OA_surf_nan(:);
% these are the segregation indices for young adults
YA_surf_nan = load([data_dir 'iNetworks_allsubs_seg_index_ses_negcorrs' neg_corrs '_'
atlas '.mat']);
YA_surf_nan = YA_surf_nan.ses_SI(:);
YA_surf_nan = YA_surf_nan(:);
%% Now how do I plot these?
handles = plotSpread([OA_surf_asis OA_vol_asis], 'distributionMarkers', {'+'},
'distributionColors', [0,0,1], 'binWidth', .5);
hold on
handles = plotSpread([YA_surf_asis YA_vol_asis], 'distributionMarkers', {'+'},
'distributionColors', [1,0,0], 'binWidth', .5);
hold on
handles = plotSpread([OA_surf_zero OA_vol_zero], 'distributionMarkers', {'o'},
'distributionColors', [0,0,1], 'binWidth', .5);
hold on
handles = plotSpread([YA_surf_zero YA_vol_zero], 'distributionMarkers', {'o'},
'distributionColors', [1,0,0], 'binWidth', .5);
hold on
handles = plotSpread([OA_surf_nan OA_vol_nan], 'distributionMarkers', {'*'},
'distributionColors', [0,0,1], 'binWidth', .5);
hold on
handles = plotSpread([YA_surf_nan YA_vol_nan], 'distributionMarkers', {'*'},
'distributionColors', [1,0,0], 'binWidth', .5);
hold on
close all
data_dir = '/Users/dianaperez/Desktop/Research/Segregation_Analyses/';
output_dir = '/Users/dianaperez/Desktop/Research/Segregation_Analyses/';
%% some variables
% I don't think I'll need these, but just in case
subs = {'LS02', 'LS03', 'LS05', 'LS08', 'LS11', 'LS14',
'LS16', 'LS17', 'INET001', 'INET002',
'INET003', 'INET005', 'INET006', 'INET010', 'INET016', 'INET018', 'INET019', 'INET030'};
rgb_colors = {[1 0 0], [0, 0, 1]}; % red and blue
%% first load the first batch
neg_corrs = 'asis'; % choose: 'nan', 'zero', 'asis'
atlas = 'Seitzman300'; % 'Seitzman300' or 'Parcels333'
```

```
%% load all the files
% these are the TimeB segregation indices for older adults
OA_vol_asis = load([data_dir 'Lifespan_allsubs_seg_index_ses_negcorrs' neg_corrs '_' atlas '.mat']);
OA_vol_asis = OA_vol_asis.ses_SI;
OA_vol_asis = OA_vol_asis(:);
% these are the segregation indices for young adults
YA_vol_asis = load([data_dir 'iNetworks_allsubs_seg_index_ses_negcorrs' neg_corrs '_' atlas '.mat']);
YA_vol_asis = YA_vol_asis.ses_SI;
YA_vol_asis = YA_vol_asis(:);
%% load the next batch
neg_corrs = 'zero'; % choose: 'nan', 'zero', 'asis'
atlas = 'Seitzman300'; % 'Seitzman300' or 'Parcels333'
%% load all the files
% these are the TimeB segregation indices for older adults
OA_vol_zero = load([data_dir 'Lifespan_allsubs_seg_index_ses_negcorrs' neg_corrs '_' atlas '.mat']);
OA_vol_zero = OA_vol_zero.ses_SI;
OA_vol_zero = OA_vol_zero(:);
% these are the segregation indices for young adults
YA_vol_zero = load([data_dir 'iNetworks_allsubs_seg_index_ses_negcorrs' neg_corrs '_' atlas '.mat']);
YA_vol_zero = YA_vol_zero.ses_SI;
YA_vol_zero = YA_vol_zero(:);
%% load the next batch
neg_corrs = 'nan'; % choose: 'nan', 'zero', 'asis'
atlas = 'Seitzman300'; % 'Seitzman300' or 'Parcels333'
%% load all the files
% these are the TimeB segregation indices for older adults
OA_vol_nan = load([data_dir 'Lifespan_allsubs_seg_index_ses_negcorrs' neg_corrs '_' atlas '.mat']);
OA_vol_nan = OA_vol_nan.ses_SI;
OA_vol_nan = OA_vol_nan(:);
% these are the segregation indices for young adults
YA_vol_nan = load([data_dir 'iNetworks_allsubs_seg_index_ses_negcorrs' neg_corrs '_' atlas '.mat']);
YA_vol_nan = YA_vol_nan.ses_SI;
YA_vol_nan = YA_vol_nan(:);
%% load the first batch
neg_corrs = 'asis'; % choose: 'nan', 'zero', 'asis'
atlas = 'Parcels333'; % 'Seitzman300' or 'Parcels333'
%% load all the files
% these are the TimeB segregation indices for older adults
OA_surf_asis = load([data_dir 'Lifespan_allsubs_seg_index_ses_negcorrs' neg_corrs '_' atlas '.mat']);
OA_surf_asis = OA_surf_asis.ses_SI;
OA_surf_asis = OA_surf_asis(:);
% these are the segregation indices for young adults
YA_surf_asis = load([data_dir 'iNetworks_allsubs_seg_index_ses_negcorrs' neg_corrs '_' atlas '.mat']);
YA_surf_asis = YA_surf_asis.ses_SI;
YA_surf_asis = YA_surf_asis(:);
%% load the next batch
neg_corrs = 'zero'; % choose: 'nan', 'zero', 'asis'
atlas = 'Parcels333'; % 'Seitzman300' or 'Parcels333'
%% load all the files
% these are the TimeB segregation indices for older adults
OA_surf_zero = load([data_dir 'Lifespan_allsubs_seg_index_ses_negcorrs' neg_corrs '_' atlas '.mat']);
```



```

OA_surf_zero = OA_surf_zero.ses_SI;
OA_surf_zero = OA_surf_zero(:);
% these are the segregation indices for young adults
YA_surf_zero = load([data_dir 'iNetworks_allsubs_seg_index_ses_negcorrs' neg_corrs '_' atlas '.mat']);
YA_surf_zero = YA_surf_zero.ses_SI;
YA_surf_zero = YA_surf_zero(:);
%% load the next batch
neg_corrs = 'nan'; % choose: 'nan', 'zero', 'asis'
atlas = 'Parcels333'; % 'Seitzman300' or 'Parcels333'
%% load all the files
% these are the TimeB segregation indices for older adults
OA_surf_nan = load([data_dir 'Lifespan_allsubs_seg_index_ses_negcorrs' neg_corrs '_' atlas '.mat']);
OA_surf_nan = OA_surf_nan.ses_SI;
OA_surf_nan = OA_surf_nan(:);
% these are the segregation indices for young adults
YA_surf_nan = load([data_dir 'iNetworks_allsubs_seg_index_ses_negcorrs' neg_corrs '_' atlas '.mat']);
YA_surf_nan = YA_surf_nan.ses_SI(:);
YA_surf_nan = YA_surf_nan(:);
%% Now how do I plot these?
handles = plotSpread([OA_surf_asis OA_vol_asis], 'distributionMarkers', {'+'}, 'distributionColors', [0,0,1], 'binWidth', .5);
hold on
handles = plotSpread([YA_surf_asis YA_vol_asis], 'distributionMarkers', {'+'}, 'distributionColors', [1,0,0], 'binWidth', .5);
hold on
handles = plotSpread([OA_surf_zero OA_vol_zero], 'distributionMarkers', {'o'}, 'distributionColors', [0,0,1], 'binWidth', .5);
hold on
handles = plotSpread([YA_surf_zero YA_vol_zero], 'distributionMarkers', {'o'}, 'distributionColors', [1,0,0], 'binWidth', .5);
hold on
handles = plotSpread([OA_surf_nan OA_vol_nan], 'distributionMarkers', {'*'}, 'distributionColors', [0,0,1], 'binWidth', .5);
hold on
handles = plotSpread([YA_surf_nan YA_vol_nan], 'distributionMarkers', {'*'}, 'distributionColors', [1,0,0], 'binWidth', .5);
hold on
segregation_ind_split_fist_ses
matched_data_1 = matched_data_sorted(:,1:amt_data);
matched_data_2 = matched_data_sorted(:,end-amt_data:end);
matched_data_2 = matched_data_sorted(:,end-amt_data-1:end);
matched_data_2 = matched_data_sorted(:,end-amt_data+1:end);
segregation_ind_split_fist_ses
load([output_dir '_allsubs_seg_index_splitfirstses_negcorrs' neg_corrs '_' atlas '.mat'])
load([output_dir data_set '_allsubs_seg_index_splitfirstses_negcorrs' neg_corrs '_' atlas '.mat'])
rgb_colors = [1 0 0;%LS02
0, 1, 0;%LS03
0, 0, 1;%LS05
0, 1, 1;%LS08
1, 0, 1;%LS11
0.4660 0.6740 0.188;%LS14
0.9290 0.6940 0.1250;%LS16
0.4940 0.1840 0.5560;%LS17];
subs = {'LS02', 'LS03', 'LS05', 'LS08', 'LS11', 'LS14', 'LS16', 'LS17'};
for n = 1:numel(subs)
    rgb_for_plot{n} = rgb_colors(n,:);

```

```

end
handles = plotSpread_v2(ses_SI', 'distributionMarkers', {'o'}, 'distributionColors', ↵
rgb_for_plot', 'xNames', subs, 'binWidth', 1);
hold on
close all
handles = plotSpread_v2(ses_SI', 'distributionMarkers', {'o'}, 'distributionColors', ↵
rgb_for_plot', 'xNames', subs, 'binWidth', 1);
rgb_colors = [0 0 0;
0 0 0;
1 1 1;
0 0 0;
0 0 0;
0 0 0;
0 0 0;
0 0 0];
for n = 1:numel(subs)
rgb_for_plot{n} = rgb_colors(n,:);
end
handles = plotSpread(ses_SI', 'distributionMarkers', {'o'}, 'distributionColors', ↵
rgb_for_plot', 'xNames', subs, 'binWidth', 1);
clear all
parcels = ft_read_cifti_mod('/Users/dianaperez/Desktop/parcellations/sub-↵
LS02/lifespan_individual_parcels_edgethresh_0.5.dtseries.nii')
ft_read_cifti_mod('lifespan_individual_parcels_edgethresh_0.5.dtseries.nii')
load('label.mat')
template = ft_read_cifti_mod('/Users/dianaperez/Desktop/sub-↵
LS03_infomap_conBensus_weighted_minsize400.dtseries.nii')
template.data = label;
ft_write_cifti_mod('/Users/dianaperez/Desktop/sub-LS02_individual_parcels.dtseries.nii', ↵
template);
raw_assn = dlmread('/Users/dianaperez/Desktop/rawassn.txt');
data = raw_assn(:,1);
template.data = data;
ft_write_cifti_mod('/Users/dianaperez/Desktop/test_LS02_thresh1.dtseries.nii', template);
template.data = raw_assn(:,2);
ft_write_cifti_mod('/Users/dianaperez/Desktop/test_LS02_thresh2.dtseries.nii', template);
template.data = raw_assn(:,12);
ft_write_cifti_mod('/Users/dianaperez/Desktop/test_LS02_thresh12.dtseries.nii', ↵
template);
template.data = raw_assn(:,11);
ft_write_cifti_mod('/Users/dianaperez/Desktop/test_LS02_thresh11.dtseries.nii', ↵
template);
load('sub-LS02_dice_to_templates.mat')
clear all
load('sub-LS02_dice_to_templates.mat')
help max
for vert = 1:59412
[M,I] = max(dice_to_templates(:,vert)); template_match(vert,1) = I;
end
template = ft_read_cifti_mod('/Users/dianaperez/Desktop/sub-↵
LS03_infomap_conBensus_weighted_minsize400.dtseries.nii')
template.data = template_match;
ft_write_cifti_mod('/Users/dianaperez/Desktop/sub_LS02_template_match_nets.dtseries.nii', ↵
template);
colors = [1,1;2,2; 3,3; 4,5; 5,7; 6,8;7,9;8,10;9,11;10,12;;11,13;12,14;13,15;14,16];
outmat = raw2colors_mat(template_match, colors);
template.data = outmat;
ft_write_cifti_mod('/Users/dianaperez/Desktop/sub_LS02_template_match_nets.dtseries.nii', ↵
template);
load('sub-LS03_dice_to_templates.mat')

```

```
for vert = 1:59412
[M,I] = max(dice_to_templates(:,vert));
template_match(vert,1) = I;
end
colors = [1,1;2,2; 3,3; 4,5; 5,7; 6,8;7,9;8,10;9,11;10,12;;11,13;12,14;13,15;14,16];
outmat = raw2colors_mat(template_match, colors);
template.data = outmat;
ft_write_cifti_mod('/Users/dianaperez/Desktop/sub_LS03_template_match_nets.dtseries.nii',
template);
clear all
uiopen('/Users/dianaperez/Desktop/Lifespan_parcellations/subsample_10/sub-LS02/sub-
LS02_individual_parcel_edgethresh_0.5.dtseries.nii',1)
ind_parcel = ft_read_cifti_mod
('/Users/dianaperez/Desktop/Lifespan_parcellations/subsample_10/sub-LS02/sub-
LS02_individual_parcel_edgethresh_0.5.dtseries.nii');
unique_ind_parcel = unique(ind_parcel.data);
ind_parcel = ind_parcel; clear ind_parcel
ind_parcel = ft_read_cifti_mod('sub-
LS02_infomap_conBensus_weighted_minsize20_consecutive_parcelopt.dtseries.nii');
post_infomap_proc
open conBensus.m
post_infomap_proc
%-- 4/12/23, 8:25 PM --%
help evalc
load('/Users/dianaperez/Desktop/Lifespan_parcellations/GordonParcellation/sub-LS05/sub-
LS05_corr_parcel_by_template_binarized.mat')
load('/Volumes/fsmresfiles/PBS/Gratton_Lab/Lifespan/seg_index/sub-
LS02_segregation_indices_by_net_indiv_parcs.mat')
clear all
load('/Volumes/fsmresfiles/PBS/Gratton_Lab/Lifespan/seg_index/sub-
LS02_segregation_indices_by_net_indiv_parcs.mat')
load('/Volumes/fsmresfiles/PBS/Gratton_Lab/Lifespan/seg_index/Lifespan_seg_index_by_sub.
mat')
clear all
similarity_analysis_main
similarity_analysis_by_system
addpath '/Users/dianaperez/Documents/GitHub/GrattonLab-General-Repo/FCProcess'
similarity_analysis_by_system
similarity_analysis_longitudinal
clear all
similarity_analysis_longitudinal
clear simmat
simmat = corr(corrln);
simmat = corr(corrln');
simmat = single(corr(corrln'));
simmat_single = single(simmat)
similarity_analysis_longitudinal
%-- 4/20/23, 3:31 PM --%
load('/Users/dianaperez/Downloads/DartmouthMIND_tutorial-master/data/Parcel_params.mat')
ROI_params = {};
ROI_params.atlas = 'Seitzman300';
atlas_params = atlas_parameters_GrattonLab
('Seitzman300','/Volumes/fsmresfiles/PBS/Gratton_Lab/Atlases/')
ROI_params.networks = atlas_params.networks
ROI_params.networks = atlas_params.networks'
ROI_params.mods = atlas_params.mods'
ROI_params.mods = atlas_params.mods
ROI_params.mods_array = atlas_params.mods_array';
ROI_params.dist_thresh = atlas_params.dist_thresh;
ROI_params.num_rois = atlas_params.num_rois;
```

```
ROI_params.colors = atlas_params.colors;
ROI_params.sorti = atlas_params.sorti';
ROI_params.transitions = atlas_params.transitions';
ROI_params.centers = atlas_params.centers;
clear all
load('/Volumes/fsmresfiles/PBS/Gratton_Lab/Lifespan/Post-
COVID/BIDS/derivatives/preproc_FCProc/corrmats_Seitzman300/sub-LS02/sub-LS02_sess-1_task-
rest_corrmats_Seitzman300.mat')
corrmats_1 = corrmats;
load('/Volumes/fsmresfiles/PBS/Gratton_Lab/Lifespan/Post-
COVID/BIDS/derivatives/preproc_FCProc/corrmats_Seitzman300/sub-LS02/sub-LS02_sess-2_task-
rest_corrmats_Seitzman300.mat')
corrmats_2 = corrmats;
all_corrmats(:,:,1,2) = corrmats_1;
all_corrmats(:,:,1,1) = corrmats_2;
all_corrmats_2(:,:,1,1) = corrmats_1;
avg_corrmats = mean(all_corrmats);
avg_corrmats = squeeze(mean(all_corrmats));
avg_corrmats = squeeze(mean(all_corrmats(:,:,,:)));
clear all
subs = {'LS02', 'LS03', 'LS05', 'LS08', 'LS11', 'LS14', 'LS16', 'LS17'};
sessions = 5;
corrmats_dir = '/Volumes/fsmresfiles/PBS/Gratton_Lab/Lifespan/Post-
COVID/BIDS/derivatives/preproc_FCProc/corrmats_Seitzman300/';
count = 1;
for sub = 1:numel(subs)
for ses = 1:sessions
% load the matrix
corrmats_fname = sprintf('%s/sub-%s/sub-%s_sess-%d_task-rest_corrmats_Seitzman300.mat',
corrmats_dir, subs{sub}, subs{ses}, ses);
load(corrmats_fname)
all_corrmats(count,(:,:,,:)) = corrmats;
count = count + 1;
end
end
subs = {'LS02', 'LS03', 'LS05', 'LS08', 'LS11', 'LS14', 'LS16', 'LS17'};
sessions = 5;
corrmats_dir = '/Volumes/fsmresfiles/PBS/Gratton_Lab/Lifespan/Post-
COVID/BIDS/derivatives/preproc_FCProc/corrmats_Seitzman300/';
count = 1;
for sub = 1:numel(subs)
for ses = 1:sessions
% load the matrix
corrmats_fname = sprintf('%s/sub-%s/sub-%s_sess-%d_task-rest_corrmats_Seitzman300.mat',
corrmats_dir, subs{sub}, subs{sub}, ses);
load(corrmats_fname)
all_corrmats(count,(:,:,,:)) = corrmats;
count = count + 1;
end
end
mean_corrmats = squeeze(mean(all_corrmats));
imagesc(mean_corrmats,[-1 1])
atlas_params = atlas_parameters_GrattonLab
('Seitzman300','/Volumes/fsmresfiles/PBS/Gratton_Lab/Atlases/');
clear all_corrmats count corrmats corrmats_dir sess_roi*
clear tmask*
figure_corrmats_GrattonLab(mean_corrmats,atlas_params,-1,1);
close('all')
figure_corrmats_GrattonLab(mean_corrmats,atlas_params,-1,1);
addpath '/Users/dianaperez/Documents/GitHub/GrattonLab-General-Repo/FCProcess'
```

```
figure_corrmat_GrattonLab(mean_corrmat,atlas_params,-1,1);
clear all
dataset = 'Lifespan';
subs = {'LS02', 'LS03', 'LS05', 'LS08', 'LS11', 'LS14', 'LS16', 'LS17'};
sessions = 5;
corrmat_dir = '/Volumes/fsmresfiles/PBS/Gratton_Lab/Lifespan/Post-✓
COVID/BIDS/derivatives/preproc_FCPProc/corrmats_Seitzman300/';
out_dir = '/Users/dianaperez/Desktop/';
fout_str = sprintf('%s/%s_average_correlation_matrix', out_dir, dataset);
count = 1;
for sub = 1:numel(subs)
for ses = 1:sessions
% load the matrix
corrmat_fname = sprintf('%s/sub-%s/sub-%s_sess-%d_task-rest_corrmat_Seitzman300.mat',✓
corrmat_dir, subs{sub}, subs{sub}, ses);
load(corrmat_fname)
all_corrmats(count,.,:,:) = corrmat;
count = count + 1;
end
end
%calculate mean
mean_corrmat = squeeze(mean(all_corrmats));
%clear some memory
clear all_corrmats count corrmat corrmat_dir sess_roi* tmask*
%load atlas parameters for plotting
atlas_params = atlas_parameters_GrattonLab✓
('Seitzman300','/Volumes/fsmresfiles/PBS/Gratton_Lab/Atlases/');
%plot mean correlation matrix
figure_corrmat_GrattonLab(mean_corrmat,atlas_params,-1,1);
%save mean correlation matrix image
saveas(gcf,[fout_str '.tiff'],'tiff');
close(gcf);
%save the corrmat data
save([fout_str '.mat'], 'mean_corrmat');
clear all
dataset = 'iNetworks';
if strcmpi(dataset, 'Lifespan')
subs = {'LS02', 'LS03', 'LS05', 'LS08', 'LS11', 'LS14', 'LS16', 'LS17'};
sessions = 5;
elseif strcmpi(dataset, 'iNetworks')
subs = {'INET001','INET002',✓
'INET003','INET005','INET006','INET010','INET016','INET018','INET019','INET030'};
sessions = 4;
end
corrmat_dir = '/Volumes/fsmresfiles/PBS/Gratton_Lab/Lifespan/Post-✓
COVID/BIDS/derivatives/preproc_FCPProc/corrmats_Seitzman300/';
out_dir = '/Users/dianaperez/Desktop/';
fout_str = sprintf('%s/%s_average_correlation_matrix', out_dir, dataset);
count = 1;
for sub = 1:numel(subs)
for ses = 1:sessions
% load the matrix
corrmat_fname = sprintf('%s/sub-%s/sub-%s_sess-%d_task-rest_corrmat_Seitzman300.mat',✓
corrmat_dir, subs{sub}, subs{sub}, ses);
load(corrmat_fname)
all_corrmats(count,.,:,:) = corrmat;
count = count + 1;
end
end
%calculate mean
```

```

mean_corrmat = squeeze(mean(all_corrmat));
%clear some memory
clear all_corrmat count corrmat corrmat_dir sess_roi* tmask*
%load atlas parameters for plotting
atlas_params = atlas_parameters_GrattonLab(
('Seitzman300','/Volumes/fsmresfiles/PBS/Gratton_Lab/Atlases/'));
%plot mean correlation matrix
figure_corrmat_GrattonLab(mean_corrmat,atlas_params,-1,1);
%save mean correlation matrix image
saveas(gcf,[fout_str '.tiff'],'tiff');
close(gcf);
%save the corrmat data
save([fout_str '.mat'], 'mean_corrmat');
clear all
dataset = 'iNetworks';
if strcmpi(dataset, 'Lifespan')
subs = {'LS02', 'LS03', 'LS05', 'LS08', 'LS11', 'LS14', 'LS16', 'LS17'};
sessions = 5;
corrmat_dir = '/Volumes/fsmresfiles/PBS/Gratton_Lab/Lifespan/Post-
COVID/BIDS/derivatives/preproc_FCPProc/corrmats_Seitzman300/';
elseif strcmpi(dataset, 'iNetworks')
subs = {'INET001','INET002',
'INET003','INET005','INET006','INET010','INET016','INET018','INET019','INET030'};
sessions = 4;
corrmat_dir =
'/Volumes/fsmresfiles/PBS/Gratton_Lab/Lifespan/Segregation_analyses/iNetworks/Nifti/deriv
atives/preproc_FCPProc/corrmats_Seitzman300';
end
out_dir = '/Users/dianaperez/Desktop/';
fout_str = sprintf('%s/%s_average_correlation_matrix', out_dir, dataset);
count = 1;
for sub = 1:numel(subs)
for ses = 1:sessions
% load the matrix
corrmat_fname = sprintf('%s/sub-%s/sub-%s_sess-%d_task-rest_corrmat_Seitzman300.mat',
corrmat_dir, subs{sub}, subs{sub}, ses);
load(corrmat_fname)
all_corrmat(count,:,:,:) = corrmat;
count = count + 1;
end
end
%calculate mean
mean_corrmat = squeeze(mean(all_corrmat));
%clear some memory
clear all_corrmat count corrmat corrmat_dir sess_roi* tmask*
%load atlas parameters for plotting
atlas_params = atlas_parameters_GrattonLab(
('Seitzman300','/Volumes/fsmresfiles/PBS/Gratton_Lab/Atlases/'));
%plot mean correlation matrix
figure_corrmat_GrattonLab(mean_corrmat,atlas_params,-1,1);
%save mean correlation matrix image
saveas(gcf,[fout_str '.tiff'],'tiff');
close(gcf);
%save the corrmat data
save([fout_str '.mat'], 'mean_corrmat');
clear all
load('/Users/dianaperez/Desktop/Neurohackademy_Tutorial-
master/data/MS02_333parcels_infomapassn.mat')
clear all
clear all

```

```
dataset = 'iNetworks';
if strcmpi(dataset, 'Lifespan')
subs = {'LS02', 'LS03', 'LS05', 'LS08', 'LS11', 'LS14', 'LS16', 'LS17'};
sessions = 5;
corrmat_dir = '/Volumes/fsmresfiles/PBS/Gratton_Lab/Lifespan/Post-
COVID/BIDS/derivatives/preproc_FCProc/corrmats_Seitzman300/';
elseif strcmpi(dataset, 'iNetworks')
subs = {'INET001', 'INET002',
'INET003', 'INET005', 'INET006', 'INET010', 'INET016', 'INET018', 'INET019', 'INET030'};
sessions = 4;
corrmat_dir =
'/Volumes/fsmresfiles/PBS/Gratton_Lab/Lifespan/Segregation_analyses/iNetworks/Nifti/deriv
atives/preproc_FCProc/corrmats_Seitzman300';
end
out_dir = '/Users/dianaperez/Desktop/';
fout_str = sprintf('%s/%s_average_correlation_matrix', out_dir, dataset);
count = 1;
for sub = 1:numel(subs)
for ses = 1:sessions
% load the matrix
corrmat_fname = sprintf('%s/sub-%s/sub-%s_sess-%d_task-rest_corrmat_Seitzman300.mat',
corrmat_dir, subs{sub}, subs{sub}, ses);
load(corrmat_fname)
all_corrmats(count, :, :, :) = corrmat;
sub_corrmat(ses, :, :, :) = corrmat;
count = count + 1;
end
all_subs_corrmats(sub, :, :, :) = squeeze(mean(sub_corrmat));
end
mean_corrmat = squeeze(mean(all_subs_corrmats));
atlas_params = atlas_parameters_GrattonLab
('Seitzman300', '/Volumes/fsmresfiles/PBS/Gratton_Lab/Atlases/');
%plot mean correlation matrix
figure_corrmat_GrattonLab(mean_corrmat, atlas_params, -1, 1);
load('/Users/dianaperez/Desktop/Neurohackademy_Tutorial-
master/data/Parcel_info/ROI_params.mat')
clear all
load('/Users/dianaperez/Desktop/Neurohackademy_Tutorial-
master/data/Parcel_info/ROI_params.mat')
load
('/Volumes/fsmresfiles/PBS/Gratton_Lab/Atlases/Seitzman300/old_vers/BigBrain300/BigBrain3
00_dmat.mat')
ROI_params.dmat = dmat;
mods_array = ROI_params.mods_array;
load('/Users/dianaperez/Desktop/Neurohackademy_Tutorial-
master/data/Parcel_info/Gordon333_net_colors.mat')
net_colors = ROI_params.colors
load('/Users/dianaperez/Desktop/Neurohackademy_Tutorial-
master/data/Parcel_info/Gordon333_roi_sort.mat')
roi_sort = ROI_params.sorti;
load('/Users/dianaperez/Desktop/Neurohackademy_Tutorial-
master/data/Lifespan_average_correlation_matrix.mat')
%-- 4/27/23, 5:07 PM --%
make_group_avg_corrmat
addpath '/Users/dianaperez/Documents/GitHub/GrattonLab-General-Repo/FCProcess'
make_group_avg_corrmat
clear all
load('/Users/dianaperez/Desktop/Neurohackademy_Tutorial-
master/data/iNetworks_average_correlation_matrix.mat')
avg_corrmat = load('/Users/dianaperez/Desktop/Neurohackademy_Tutorial-
```



```

master/data/iNetworks_average_correlation_matrix.mat')
clear all
%% Spring-embedding plots
load('/Users/dianaperez/Desktop/Neurohackademy_Tutorial-master/data/iNetworks_average_correlation_matrix.mat')
corrmat = mean_corrmat;
clear mean_corrmat
colors = zeros([size(corrmat,1),3]);
atlas_params = atlas_parameters_GrattonLab('Seitzman300','/Volumes/fsmresfiles/PBS/Gratton_Lab/Atlases/');
for m = 1:length(atlas_params.mods)
    colors(atlas_params.mods{m},:) = repmat(atlas_params.colors(m,:),[length(atlas_params.mods{m}),1]);
end
addpath(genpath('/Users/dianaperez/Downloads/DartmouthMIND_tutorial-master/'))
make_spring_fig_MIND(groupmat,0.02,colors,atlas_params.networks,atlas_params.colors);
make_spring_fig_MIND(corrmat,0.02,colors,atlas_params.networks,atlas_params.colors);
thresholds = [0.01:0.01:0.04]; %[0.01:0.01:0.1,0.2,0.3]; %these take a while, esp for higher thresholds. Only do a few.
outdir = '/Users/dianaperez/Desktop/';
for t = 1:length(thresholds)
    %tic;
    make_spring_fig_MIND(corrmat,thresholds(t),colors,atlas_params.networks,atlas_params.colors);
    saveas(gcf,sprintf('%sSpring_group_t%.02f.tiff',outdir,thresholds(t)),'tiff');
    %toc
end
thresholds = [0.01:0.01:0.04]; %[0.01:0.01:0.1,0.2,0.3]; %these take a while, esp for higher thresholds. Only do a few.
outdir = '/Users/dianaperez/Desktop/';
for t = 1:length(thresholds)
    %tic;
    make_spring_fig_MIND(corrmat,thresholds(t),colors,atlas_params.networks,atlas_params.colors);
    saveas(gcf,sprintf('%sSpring_group_t%.02f.tiff',outdir,thresholds(t)),'tiff');
    %toc
end
load('/Users/dianaperez/Desktop/Neurohackademy_Tutorial-master/data/Lifespan_average_correlation_matrix.mat')
corrmat = mean_corrmat;
clear mean_corrmat
% Start with a single correlation matrix per subject and network assignment
% [For the group and for the individual separately?]
%submats = squeeze(mean(corrmat,2));
% and colormap variable needed for function:
% information about prespecified networks
colors = zeros([size(corrmat,1),3]);
atlas_params = atlas_parameters_GrattonLab('Seitzman300','/Volumes/fsmresfiles/PBS/Gratton_Lab/Atlases/');
for m = 1:length(atlas_params.mods)
    colors(atlas_params.mods{m},:) = repmat(atlas_params.colors(m,:),[length(atlas_params.mods{m}),1]);
end
% Now create across a range of thresholds (1% - 10%, 20%, 30%) for the
% group -- IF TIME
% 1. Discuss consequences of different thresholds
thresholds = [0.01:0.01:0.04]; %[0.01:0.01:0.1,0.2,0.3]; %these take a while, esp for higher thresholds. Only do a few.
outdir = '/Users/dianaperez/Desktop/';
for t = 1:length(thresholds)

```



```
%tic;
make_spring_fig_MIND(corrmat,thresholds(t),colors,atlas_params.networks,atlas_params.
colors);
saveas(gcf,sprintf('%sSpring_group_t%.02f.tiff',outdir,thresholds(t)),'tiff');
%toc
end
thresholds = 0.5;%[0.01:0.01:0.04]; %[0.01:0.01:0.1,0.2,0.3]; %these take a while, esp
for higher thresholds. Only do a few.
outdir = '/Users/dianaperez/Desktop/';
for t = 1:length(thresholds)
%tic;
make_spring_fig_MIND(corrmat,thresholds(t),colors,atlas_params.networks,atlas_params.
colors);
saveas(gcf,sprintf('%sSpring_group_t%.02f.tiff',outdir,thresholds(t)),'tiff');
%toc
end
thresholds = 0.05;%[0.01:0.01:0.04]; %[0.01:0.01:0.1,0.2,0.3]; %these take a while, esp
for higher thresholds. Only do a few.
outdir = '/Users/dianaperez/Desktop/';
for t = 1:length(thresholds)
%tic;
make_spring_fig_MIND(corrmat,thresholds(t),colors,atlas_params.networks,atlas_params.
colors);
saveas(gcf,sprintf('%sSpring_group_t%.02f.tiff',outdir,thresholds(t)),'tiff');
%toc
end
close all;
load('/Users/dianaperez/Desktop/Neurohackademy_Tutorial-
master/data/iNetworks_average_correlation_matrix.mat')
corrmat = mean_corrmat;
clear mean_corrmat
% Start with a single correlation matrix per subject and network assignment
% [For the group and for the individual separately?]
%submats = squeeze(mean(corrmat,2));
% and colormap variable needed for function:
% information about prespecified networks
colors = zeros([size(corrmat,1),3]);
atlas_params = atlas_parameters_GrattonLab
('Seitzman300','/Volumes/fsmresfiles/PBS/Gratton_Lab/Atlases/');
for m = 1:length(atlas_params.mods)
colors(atlas_params.mods{m},:) = repmat(atlas_params.colors(m,:),[length(atlas_params.
mods{m}),1]);
end
% Now create across a range of thresholds (1% - 10%, 20%, 30%) for the
% group -- IF TIME
% 1. Discuss consequences of different thresholds
thresholds = 0.05;%[0.01:0.01:0.04]; %[0.01:0.01:0.1,0.2,0.3]; %these take a while, esp
for higher thresholds. Only do a few.
outdir = '/Users/dianaperez/Desktop/';
for t = 1:length(thresholds)
%tic;
make_spring_fig_MIND(corrmat,thresholds(t),colors,atlas_params.networks,atlas_params.
colors);
saveas(gcf,sprintf('%sSpring_group_t%.02f.tiff',outdir,thresholds(t)),'tiff');
%toc
end
close all;
%-- 4/27/23, 6:18 PM --%
load('/Users/dianaperez/Desktop/Neurohackademy_Tutorial-
master/data/Lifespan_average_correlation_matrix.mat')
```

```

corrmat = mean_corrmat;
clear mean_corrmat
% Start with a single correlation matrix per subject and network assignment
% [For the group and for the individual separately?]
%submats = squeeze(mean(corrmat,2));
% and colormap variable needed for function:
% information about prespecified networks
colors = zeros([size(corrmat,1),3]);
atlas_params = atlas_parameters_GrattonLab
('Seitzman300','/Volumes/fsmresfiles/PBS/Gratton_Lab/Atlases/');
for m = 1:length(atlas_params.mods)
colors(atlas_params.mods{m},:) = repmat(atlas_params.colors(m,:),[length(atlas_params.
mods{m}),1]);
end
% Now create across a range of thresholds (1% - 10%, 20%, 30%) for the
% group -- IF TIME
% 1. Discuss consequences of different thresholds
thresholds = 0.05;:[0.01:0.01:0.04]; %[0.01:0.01:0.1,0.2,0.3]; %these take a while, esp
for higher thresholds. Only do a few.
outdir = '/Users/dianaperez/Desktop/';
for t = 1:length(thresholds)
%tic;
make_spring_fig_MIND(corrmat,thresholds(t),colors,atlas_params.networks,atlas_params.
colors);
saveas(gcf,sprintf('%sSpring_group_t%.02f.tiff',outdir,thresholds(t)),'tiff');
%toc
end
close all;
addpath '/Users/dianaperez/Documents/GitHub/GrattonLab-General-Repo/FCProcess'
addpath(genpath('/Users/dianaperez/Downloads/DartmouthMIND_tutorial-master/'))
load('/Users/dianaperez/Desktop/Neurohackademy_Tutorial-
master/data/Lifespan_average_correlation_matrix.mat')
corrmat = mean_corrmat;
clear mean_corrmat
% Start with a single correlation matrix per subject and network assignment
% [For the group and for the individual separately?]
%submats = squeeze(mean(corrmat,2));
% and colormap variable needed for function:
% information about prespecified networks
colors = zeros([size(corrmat,1),3]);
atlas_params = atlas_parameters_GrattonLab
('Seitzman300','/Volumes/fsmresfiles/PBS/Gratton_Lab/Atlases/');
for m = 1:length(atlas_params.mods)
colors(atlas_params.mods{m},:) = repmat(atlas_params.colors(m,:),[length(atlas_params.
mods{m}),1]);
end
% Now create across a range of thresholds (1% - 10%, 20%, 30%) for the
% group -- IF TIME
% 1. Discuss consequences of different thresholds
thresholds = 0.05;:[0.01:0.01:0.04]; %[0.01:0.01:0.1,0.2,0.3]; %these take a while, esp
for higher thresholds. Only do a few.
outdir = '/Users/dianaperez/Desktop/';
for t = 1:length(thresholds)
%tic;
make_spring_fig_MIND(corrmat,thresholds(t),colors,atlas_params.networks,atlas_params.
colors);
saveas(gcf,sprintf('%sSpring_group_t%.02f.tiff',outdir,thresholds(t)),'tiff');
%toc
end
close all;

```

```
%-- 4/29/23, 1:34 PM --%
simple = modify_clrfile('simplify','rawassn.txt',minNetSize); %makes a file called
rawassn_minsizeX.txt
minNetSize = 20; %400 for vertices, 20 for parcels?
simple = modify_clrfile('simplify','rawassn.txt',minNetSize); %makes a file called
rawassn_minsizeX.txt
addpath(genpath('/Users/dianaperez/Documents/Dependencies'))
simple = modify_clrfile('simplify','rawassn.txt',minNetSize); %makes a file called
rawassn_minsizeX.txt
regularized = rawoutput2clr(simple);
regularized(regularized < 2) = 0;
regularized = regularized - 1;
dlmwrite(['rawassn_minsize' num2str(minNetSize) '_regularized.txt'],
regularized,'delimiter',' ') %makes a file called rawassn_minizeX_regularized.txt
post_infomap_proc
open modify_clrfile
open modify_clrfile
type = 'simplify'
clrfile = 'rawassn.txt';
varargin = 20;
modify_clrfile
modify_clrfile('simplify','rawassn.txt',minNetSize);
help nnz
modify_clrfile('simplify','rawassn.txt',minNetSize);
clrs_orig = clrs;
clrs = clrs_orig(:,8:18);
clrs = clrs_orig(:,9:18);
%-- 5/2/23, 9:37 AM --%
timeseries = ft_read_cifti_mod
('/Volumes/fsmresfiles/PBS/Gratton_Lab/iNetworks/BIDS/Nifti/derivatives/postFCproc_CIFTI_
20.2.0/sub-INET001/ses-1/cifti_timeseries_normalwall/sub-INET001_ses-1_task-rest_run-
1_LR_surf_subcort_222_32k_fsLR_smooth2.55.dtseries.nii')
help chol
thresh = 0.05;
bins = [1, 0.9, 0.8, 0.7, 0.6, 0.5, 0.4, 0.3, 0.2, 0.1, thresh];
mod_threshold = 0.05;
fake_signals = {};
i = 1;
count = 0;
bin = bins(i);
real_signal = timeseries;
real_fc = corr(real_signal');
rand_signal = randn(size(real_signal));
real_fc = paircorr_mod(real_signal');
real_fc = paircorr_mod(real_signal);
find(real_signal==NaN)
find(real_signal=NaN)
real_signal = timeseries.data;
real_fc = corr(real_signal');
real_fc = paircorr_mod(real_signal');
load('/Volumes/fsmresfiles/PBS/Gratton_Lab/Lifespan/Post-
COVID/BIDS/derivatives/postFCproc_CIFTI/FC_Parcel_333/sub-INET001_rest_ses-
1_parcel_timecourse.mat')
real_signal = parcel_time;
real_fc = paircorr_mod(real_signal');
real_fc = paircorr_mod(real_signal);
imagesc(real_fc,[-1 1])
rand_signal = randn(size(real_signal));
try
R = chol(real_fc);
```

```
catch
fc_sim(i) = NaN;
mod_sim(i) = NaN;
continue;
end
try
R = chol(real_fc);
catch
fc_sim(i) = NaN;
mod_sim(i) = NaN;
%continue;
end
[V D] = eig(real_fc);
V_use = V*sqrt(D);
fake_signal = V_use*rand_signal;
fake_signals{j,k} = fake_signal;
fake_fc = corr(fake_signal);
fake_signal = V_use*rand_signal;
fake_signal = V_use*rand_signal';
fake_fc = corr(fake_signal);
fake_fc = corr(fake_signal');
imagesc(fake_fc, [-1 1])
good_corrs = find(fake_fc>.8);
good_corrs = find(fake_fc>.8 && fake_fc<1);
good_corrs = find(1>fake_fc>.8);
good_corrs = find(fake_fc(1>fake_fc>.8));
upper_tri = upper(fake_fc);
upper_tri = triu(fake_fc);
upper_tri = triu(fake_fc, 1);
max = upper_tri
max(upper_tri)
clear max
max(upper_tri)
sig = rand_signal';
plot(sig(78,:))
plot(sig(78,1:100))
smooth_sig = smoothdata(sig(78,1:100));
plot(smooth_sig)
plot(smooth_sig, 'LineWidth', 5)
plot(smooth_sig, 'LineWidth', 3)
hold on
plot(sig(78,1:100))
smooth_sig = smoothdata(smooth_sig);
hold on
plot(smooth_sig)
close all
plot(smooth_sig)
hold on
smooth_sig = smoothdata(smooth_sig);
plot(smooth_sig)
close all
second_sig = fake_sig(79,1:100);
second_sig = fake_signal(79,1:100);
smooth_sig1 = smoothdata(second_sig);
smooth_sig1 = smoothdata(smooth_sig2);
smooth_sig1 = smoothdata(smooth_sig1);
plot(smooth_sig1)
plot(smooth_sig1, 'LineWidth', 3)
hold on
plot(smooth_sig, 'LineWidth', 3)
```

```
close all
smooth_signal_1 = smoothdata(fake_signal(78,:));
smooth_signal_2 = smoothdata(fake_signal(79,:));
plot(smooth_signal_1, 'LineWidth', 2)
close all
plot(smooth_signal_1)
hold on
plot(smooth_signal_2)
figure
smooth_signal_1 = smoothdata(smooth_signal_1);
smooth_signal_2 = smoothdata(smooth_signal_2);
plot(smooth_signal_1)
hold on
plot(smooth_signal_2)
smooth_signal_1 = smoothdata(smooth_signal_1);
smooth_signal_2 = smoothdata(smooth_signal_2);
figure
plot(smooth_signal_1)
hold on
plot(smooth_signal_2)
smooth_signal_1 = smoothdata(smooth_signal_1);
smooth_signal_2 = smoothdata(smooth_signal_2);
figure
plot(smooth_signal_1)
hold on
plot(smooth_signal_2)
smooth_signal_1 = smoothdata(smooth_signal_1);
smooth_signal_2 = smoothdata(smooth_signal_2);
figure
plot(smooth_signal_2)
hold on
plot(smooth_signal_1)
sub_smooth_signal_1 = smooth_signal_1(600:1000);
sub_smooth_signal_2 = smooth_signal_2(600:1000);
figure
plot(sub_smooth_signal_1, 'LineWidth', 3)
hold on
plot(sub_smooth_signal_2, 'LineWidth', 3)
xlabel('BOLD signal')
ylabel('Time')
axis([-1 1 0 400])
axis([0 400 -1 1])
set(gca, 'fontsize', 18)
[height, width] = size(gca)
smooth_signal_3 = smoothdata(fake_signal(300,:));
sub_smooth_signal_3 = smooth_signal_3(600:1000);
smooth_signal_3 = smoothdata(smooth_signal_3);
plot(smooth_signal_3)
figure
plot(smooth_signal_3, 'LineWidth', 3, 'Color', 'Green')
plot(smooth_signal_3(1:400), 'LineWidth', 3, 'Color', [0.4660 0.6740 0.1880])
hold on
plot(smooth_signal_2(1:400), 'LineWidth', 3, 'Color', [0.4940 0.1840 0.5560])
plot(smooth_signal_2(1:400), 'LineWidth', 3, 'Color', [0.4940 0.1840 0.5560])
plot(smooth_signal_1(1:400), 'LineWidth', 3, 'Color', [0.4940 0.1840 0.5560])
plot(smooth_signal_1(1:400), 'LineWidth', 3, 'Color', [0.4940 0.1840 0.5560])
figure
plot(smooth_signal_2(5:405), 'LineWidth', 3, 'Color', [0.4940 0.1840 0.5560])
plot(smooth_signal_2(5:405), 'LineWidth', 3, 'Color', [0.4940 0.1840 0.5560])
hold on
```

```

plot(smooth_signal_3(1:400), 'LineWidth', 3, 'Color', [0.4660 0.6740 0.1880])
axis([0 400 -1 1])
xlabel('BOLD signal')
ylabel('Time')
set(gca, 'fontsize', 18)
corr(sub_smooth_signal_1, sub_smooth_signal_2)
corrcoef(sub_smooth_signal_1, sub_smooth_signal_2)
corrcoef(smooth_signal_2(5:405), smooth_signal_3(1:400))
corrcoef(smooth_signal_2(6:405), smooth_signal_3(1:400))
figure
plot(sub_smooth_signal_2, 'LineWidth', 3, 'Color', [0.8500 0.3250 0.0980])
hold on
plot(smooth_signal_3(1:400), 'LineWidth', 3, 'Color', [0.4660 0.6740 0.1880])
corrcoef(sub_smooth_signal_2, smooth_signal_3(1:400))
corrcoef(sub_smooth_signal_2, smooth_signal_3(1:401))
axis([0 400 -1 1])
%-- 5/8/23, 5:28 PM --%
post_infomap_proc
tmp.data = consen;
ft_write_cifti_mod([str '_conBensus_weighted_minsize' num2str(minNetSize) '_vertices.'
dtseries.nii'], tmp);
colorChange = [];
inds = unique(consen);
colorChange(:,1) = inds;
colorChange(:,2) = 0:length(inds)-1;
outmat = raw2colors_mat(consen, colorChange);
tmp.data = zeros(size(tmp.data));
tmp.data = outmat;
ft_write_cifti_mod([str '_conBensus_weighted_minsize' num2str(minNetSize)
'_consecutive_individual_vertices.dtseries.nii'], tmp);
post_infomap_proc
conBensus
post_infomap_proc
%-- 5/11/23, 2:47 PM --%
load('HCP384_variants_info.mat')
load_nii('/Users/dianaperez/Desktop/thalamus_wta_output_111_mask_MNI.nii')
load_untouch_nii('/Users/dianaperez/Desktop/thalamus_wta_output_111_mask_MNI.nii')
thal_data = ans.img;
thal_data = ans.img;
unique_IDs = unique(thal_data);
unique_IDs(1) = [];
relabel = [1, 10, 2, 3, 7, 17, 9, 5, 11, 8, 12];
input = [unique_IDs, relabel'];
raw2colors_mat(thal_data, input);
load_nii('/Users/dianaperez/Desktop/thalamus_wta_output_111_mask_MNI.nii')
nifti_file = ans;
raw2colors_mat(thal_data, input);
addpath '/Users/dianaperez/Documents/GitHub/Lifespan-Analysis'
raw2colors_mat(thal_data, input);
nifti_file.img = ans;
save_untouch_nii(nifti_file, '/Users/dianaperez/Desktop/thalamus_relabeled.nii')
save_nii(nifti_file, '/Users/dianaperez/Desktop/thalamus_relabeled.nii')
input(6,2) = 4;
unique_relabeled = unique(ans);
thal_relabeled = raw2colors_mat(thal_data, input);
unique_relabeled = unique(thal_relabeled);
thal_relabeled = raw2colors_mat(thal_data, input);
size(regularNodes, 2)
thal_relabeled(thal_data==25) = 12
thal_relabeled(thal_data==25) = 12;

```

```

unique_relabeled = unique(thal_relabeled)
thal_relabeled(thal_data==21) = 8;
thal_relabeled(thal_data==17) = 11;
thal_relabeled(thal_data==16) = 5;
thal_relabeled(thal_data==15) = 9;
thal_relabeled(thal_data==12) = 4;
thal_relabeled(thal_data==8) = 7;
thal_relabeled(thal_data==7) = 3;
thal_relabeled(thal_data==6) = 2;
thal_relabeled(thal_data==5) = 10;
thal_relabeled(thal_data==4) = 1;
unique_relabeled = unique(thal_relabeled)
save_nii(nifti_file, '/Users/dianaperez/Desktop/thalamus_relabeled.nii')
nifti_file.img = thal_relabeled;
save_nii(nifti_file, '/Users/dianaperez/Desktop/thalamus_relabeled.nii')
load_nii('/Users/dianaperez/Desktop/basalGanglia_wta_output_111_mask_MNI.nii')
load_nii('/Users/dianaperez/Desktop/basalGanglia_wta_output_111_mask_MNI_wbcolors.nii')
bg_nifti = load_nii(
('/Users/dianaperez/Desktop/basalGanglia_wta_output_111_mask_MNI_wbcolors.nii');
bg_data = bg_nifti.img;
bg_data_relabeled = bg_data;
bg_data_relabeled(bg_data==16) = 4;
nifti_file.img = bg_data_relabeled;
save_nii(nifti_file, '/Users/dianaperez/Desktop/basalGanglia_relabeled.nii')
cb_nifti = load_nii(
('/Users/dianaperez/Desktop/cerebellum_wta_output_111_mask_MNI_wbcolors.nii')
cb_data = cb_nifti.img;
cb_relabeled = cb_data;
cb_data_relabeled(cb_data==16) = 4;
cb_nifti.img = cb_data_relabeled;
save_nii(cb_nifti, '/Users/dianaperez/Desktop/cerebellum_relabeled.nii')
cb_nifti = load_nii(
('/Users/dianaperez/Desktop/cerebellum_wta_output_111_mask_MNI_wbcolors.nii');
cb_data = cb_nifti.img;
cb_relabeled = cb_data;
cb_relabeled(cb_data==16) = 4;
cb_nifti.img = cb_relabeled;
save_nii(cb_nifti, '/Users/dianaperez/Desktop/cerebellum_relabeled.nii')
%-- 5/24/23, 10:57 AM --%
timeseries = load('/Volumes/fsmresfiles/PBS/Gratton_Lab/Lifespan/Post-
COVID/BIDS/derivatives/postFCproc_CIFTI/FC_Parcel333/sub-INET001_rest_ses-
1_parcel_timecourse.mat');
thresh = 0.05;
bins = [1, 0.9, 0.8, 0.7, 0.6, 0.5, 0.4, 0.3, 0.2, 0.1, thresh];
mod_threshold = 0.05;
fake_signals = {};
count = 0;
real_signal = timeseries.parcel_time;
real_fc = paircorr_mod(real_signal);
rand_signal = randn(size(real_signal));
addpath '/Users/dianaperez/Documents/GitHub/Lifespan-Analysis/utilities'
real_signal = timeseries.parcel_time;
real_fc = paircorr_mod(real_signal);
rand_signal = randn(size(real_signal));
R = chol(real_fc);
help chol
[V D] = eig(real_fc);
V_use = V*sqrt(D);
fake_signal = V_use*rand_signal;
fake_signals{j,k} = fake_signal;

```

```
plot(rand_signal(:,1))  
plot(rand_signal(1:400,1))  
smooth_signal_1 = smooth(rand_signal(1:400,1));  
fake_signal = V_use*rand_signal;
```