

Investigation: does motion influence the correlation between PE and FA?

Loading packages we'll need

```
% Adding packages we need
% For Plotting
addpath('C:\Users\dk00549\OneDrive - University of Surrey\Documents\Surrey\MATLAB\PluginsPrograms\Plotting');
% For emmeans
addpath('C:\Users\dk00549\OneDrive - University of Surrey\Documents\Surrey\MATLAB\PluginsPrograms\emmeans');
```

Loading tables containing, PE, FA, and MO

PE = parameter estimates. In this case, these are parameter estimates for regions where white matter structure influences the tDCS-induced changes in brain activity.

FA = our metric for white matter structure. We are evaluating three tracts- whole skeleton (WS), the rAI-dACC/preSMA, or Salience Network tract (SN), and the mPFC-PCC/PRE, or Default Mode Network (DMN) tract.

MO = motion outliers. Our metric are DVARS, though several others may be used as well.

```
PEandFA = readtable("PEandFA.xlsx");
MO = table2dataset(readtable("PTandHCMO.csv"));

% The below lines average the DVARS values per participant so they can be
% added to the PE and FA table. Two TBI participants had one fewer session
% than the others, and two healthy control participants had three more DVARS values
% than the others, so their means were calculated seperately.

df = [];
for k=1:width(MO)
    if k==5
        df(:,k) = mean(table2array(dataset2table(MO(1:784,k)))));
    elseif k==30
        df(:,k) = mean(table2array(dataset2table(MO(1:784,k)))));
    elseif k==43
        df(:,k) = mean(table2array(dataset2table(MO(1:1179,k)))));
    elseif k==51
        df(:,k) = mean(table2array(dataset2table(MO(1:1179,k)))));
    else
        df(:,k) = mean(table2array(dataset2table(MO(1:1176,k)))));
    end
end

df = array2table(df');
PEandFA(:,12) = df ;
```

Influence of motion on the relationship between WS FA and tDCS-induced changes in brain activity

```
% Four WS analysis:
%Columns 1:4 = WS PE
```

```
%Column 5 = WS FA
```

```
T = table2dataset(PEandFA);
```

```
T.Properties.VarNames{1} = 'PE_Wholeskel_cope3';
T.Properties.VarNames{2} = 'PE_Wholeskel_cope5';
T.Properties.VarNames{3} = 'PE_Wholeskel_cope10';
T.Properties.VarNames{4} = 'PE_Wholeskel_cope11';
T.Properties.VarNames{5} = 'FA_WHOLESKEL';
```

```
T.Properties.VarNames{6} = 'PE_SN_cope3';
T.Properties.VarNames{7} = 'PE_SN_cope5';
T.Properties.VarNames{8} = 'FA_SN';
```

```
T.Properties.VarNames{9} = 'PE_DMN_cope3';
T.Properties.VarNames{10} = 'PE_DMN_cope10';
T.Properties.VarNames{11} = 'FA_DMN';
```

```
T.Properties.VarNames{12} = 'MO';
```

```
% Column 1
```

```
lme_light_clean1= fitglme(T,'PE_Wholeskel_cope3 ~ MO*FA_WHOLESKEL');
disp(lme_light_clean1)
```

Generalized linear mixed-effects model fit by PL

Model information:

Number of observations	55
Fixed effects coefficients	4
Random effects coefficients	0
Covariance parameters	1
Distribution	Normal
Link	Identity
FitMethod	MPL

Formula:

PE_Wholeskel_cope3 ~ 1 + FA_WHOLESKEL*MO

Model fit statistics:

AIC	BIC	LogLikelihood	Deviance
354.49	364.52	-172.24	344.49

Fixed effects coefficients (95% CIs):

Name	Estimate	SE	tStat
{ '(Intercept)' }	-71.255	39.839	-1.7886
{ 'FA_WHOLESKEL' }	229.18	127.14	1.8026
{ 'MO' }	1.0303	5.8485	0.17617
{ 'FA_WHOLESKEL:MO' }	-3.6669	18.706	-0.19603

DF	pValue	Lower	Upper
51	0.079624	-151.24	8.7252
51	0.07736	-26.06	484.43
51	0.86086	-10.711	12.772
51	0.84536	-41.22	33.886

Random effects covariance parameters:

Group: Error

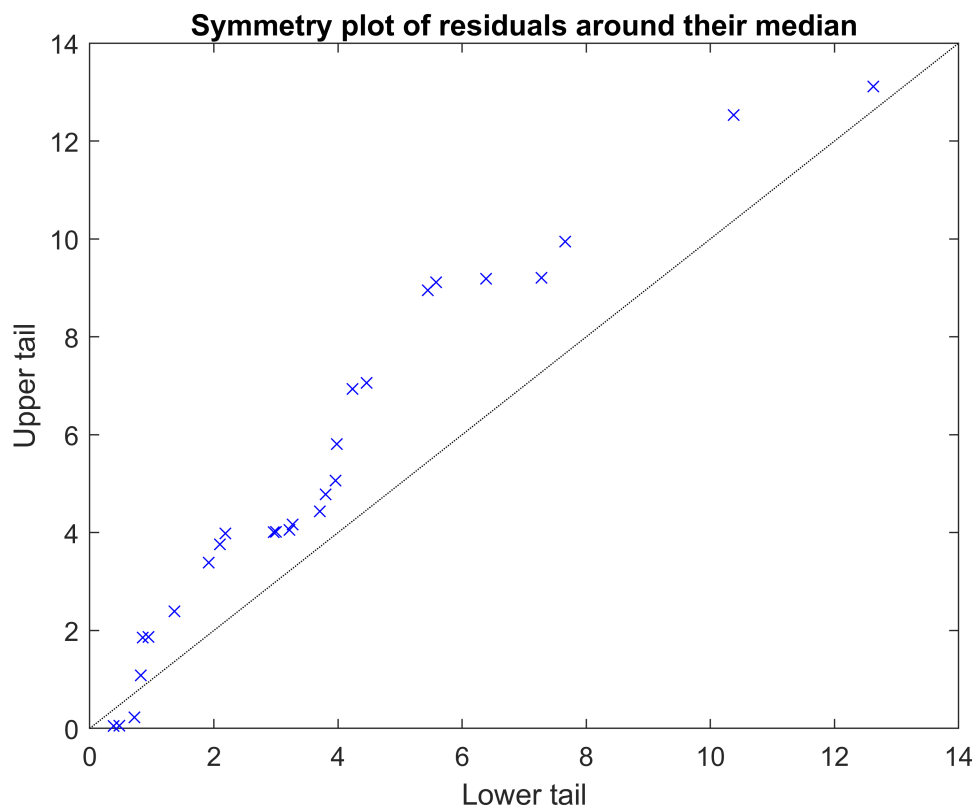
Name	Estimate
{ 'sqrt(Dispersion)' }	5.5442

```
anova(lme_light_clean1)
```

```
ans =
ANOVA marginal tests: DFMethod = 'residual'
```

Term	FStat	DF1	DF2	pValue
{'(Intercept)'} }	3.199	1	51	0.079624
{'FA_WHOLESKEL' } }	3.2494	1	51	0.07736
{'MO' } }	0.031036	1	51	0.86086
{'FA_WHOLESKEL:MO' }	0.038429	1	51	0.84536

```
figure
plotResiduals(lme_light_clean1, 'symmetry')
```



```
% Interaction p val = 0.8
```

```
% Column 2
```

```
lme_light_clean1= fitglme(T,'PE_Wholeskel_cope5 ~ MO*FA_WHOLESKEL');
disp(lme_light_clean1)
```

Generalized linear mixed-effects model fit by PL

Model information:

Number of observations	55
Fixed effects coefficients	4
Random effects coefficients	0
Covariance parameters	1
Distribution	Normal
Link	Identity
FitMethod	MPL

Formula:

```
PE_Wholeskel_cope5 ~ 1 + FA_WHOLESKEL*MO
```

Model fit statistics:

AIC	BIC	LogLikelihood	Deviance
393.53	403.57	-191.77	383.53

Fixed effects coefficients (95% CIs):

Name	Estimate	SE	tStat
{'(Intercept)'} }	-61.556	56.815	-1.0834
{'FA_WHOLESKEL' }	175.29	181.32	0.96677
{'MO' }	-1.0709	8.3407	-0.12839
{'FA_WHOLESKEL:MO'}	5.9746	26.676	0.22396

DF	pValue	Lower	Upper
51	0.28371	-175.62	52.505
51	0.33822	-188.72	539.3
51	0.89834	-17.815	15.674
51	0.82368	-47.58	59.53

Random effects covariance parameters:

Group: Error

Name	Estimate
{'sqrt(Dispersion)'} }	7.9066

```
anova(lme_light_clean1)
```

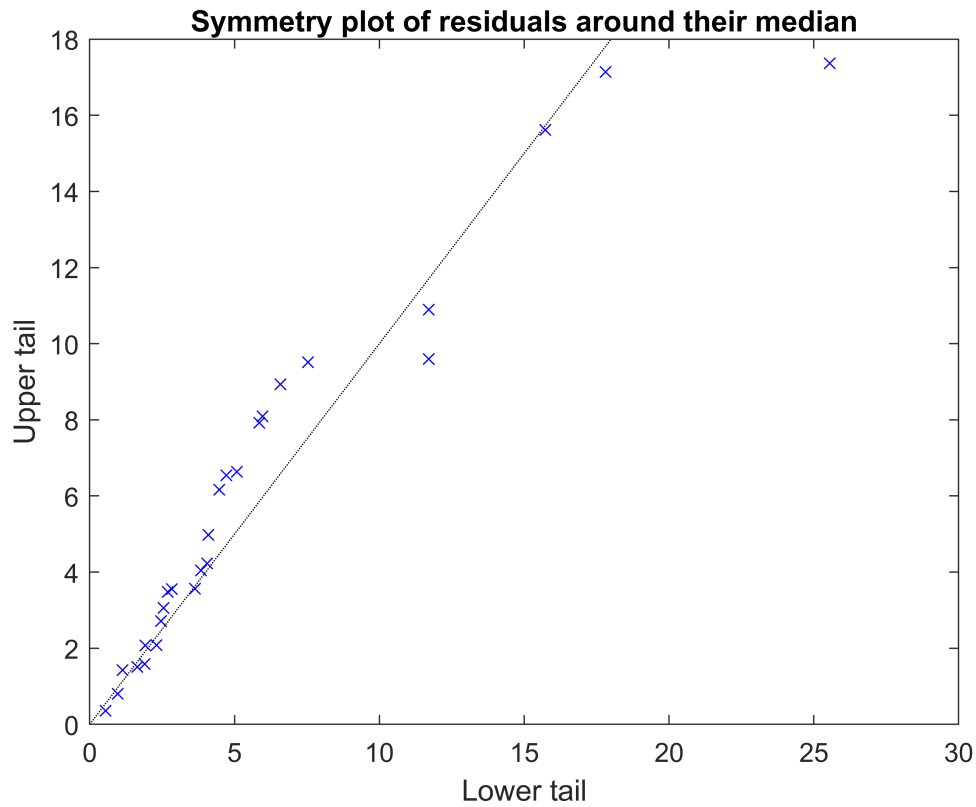
```
ans =
```

```
ANOVA marginal tests: DFMethod = 'residual'
```

Term	FStat	DF1	DF2	pValue
{'(Intercept)'} }	1.1739	1	51	0.28371
{'FA_WHOLESKEL' }	0.93464	1	51	0.33822
{'MO' }	0.016485	1	51	0.89834
{'FA_WHOLESKEL:MO'}	0.05016	1	51	0.82368

```
figure
```

```
plotResiduals(lme_light_clean1, 'symmetry')
```



```
% Interaction p val = 0.8
```

```
% Column 3
```

```
lme_light_clean1= fitglme(T,'PE_Wholeskel_cope10 ~ MO*FA_WHOLESKEL');
disp(lme_light_clean1)
```

Generalized linear mixed-effects model fit by PL

Model information:

Number of observations	55
Fixed effects coefficients	4
Random effects coefficients	0
Covariance parameters	1
Distribution	Normal
Link	Identity
FitMethod	MPL

Formula:

```
PE_Wholeskel_cope10 ~ 1 + FA_WHOLESKEL*MO
```

Model fit statistics:

AIC	BIC	LogLikelihood	Deviance
415.26	425.3	-202.63	405.26

Fixed effects coefficients (95% CIs):

Name	Estimate	SE	tStat
{ '(Intercept)' }	-46.637	69.225	-0.6737
{ 'FA_WHOLESKEL' }	132.9	220.92	0.60159
{ 'MO' }	-7.4665	10.163	-0.73471
{ 'FA_WHOLESKEL:MO' }	25.7	32.503	0.7907

DF	pValue	Lower	Upper
51	0.50354	-185.61	92.339
51	0.55011	-310.61	576.42
51	0.46588	-27.869	12.936
51	0.43278	-39.553	90.954

Random effects covariance parameters:

Group: Error

Name	Estimate
{'sqrt(Dispersion)'} }	9.6337

```
anova(lme_light_clean1)
```

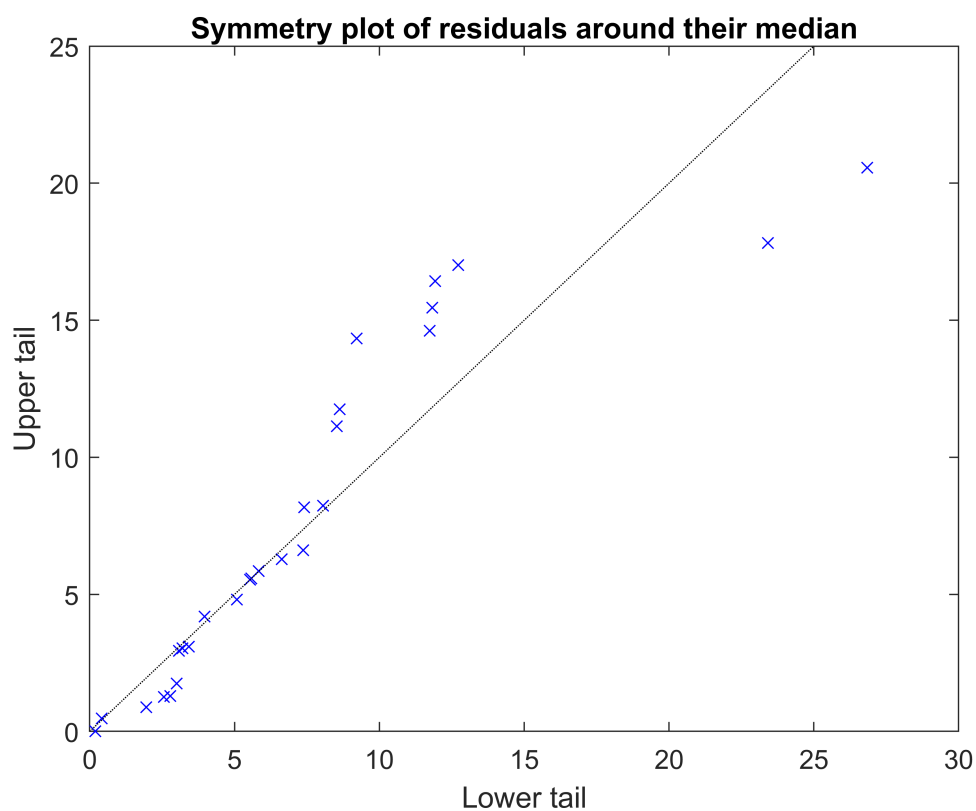
ans =

ANOVA marginal tests: DFMethod = 'residual'

Term		FStat	DF1	DF2	pValue
{' (Intercept) ' }		0.45387	1	51	0.50354
{'FA_WHOLESKEL ' }		0.36191	1	51	0.55011
{'MO ' }		0.5398	1	51	0.46588
{'FA_WHOLESKEL:MO ' }		0.6252	1	51	0.43278

figure

```
plotResiduals(lme_light_clean1, 'symmetry')
```



```
% Interaction p val = 0.4
```

```
% Column 4
```

```
lme_light_clean1= fitglme(T,'PE_Wholeskel_cope11 ~ MO*FA_WHOLESKEL');  
disp(lme_light_clean1)
```

Generalized linear mixed-effects model fit by PL

Model information:

Number of observations	55
Fixed effects coefficients	4
Random effects coefficients	0
Covariance parameters	1
Distribution	Normal
Link	Identity
FitMethod	MPL

Formula:

```
PE_Wholeskel_cope11 ~ 1 + FA_WHOLESKEL*MO
```

Model fit statistics:

AIC	BIC	LogLikelihood	Deviance
422.07	432.11	-206.04	412.07

Fixed effects coefficients (95% CIs):

Name	Estimate	SE	tStat
{'(Intercept)'} }	-170.92	73.646	-2.3208
{'FA_WHOLESKEL' }	526.28	235.03	2.2392
{'MO' }	13.954	10.812	1.2907
{'FA_WHOLESKEL:MO' }	-41.931	34.579	-1.2126

DF	pValue	Lower	Upper
51	0.024334	-318.77	-23.066
51	0.02953	54.443	998.13
51	0.20263	-7.7508	35.659
51	0.23087	-111.35	27.49

Random effects covariance parameters:

Group: Error

Name	Estimate
{'sqrt(Dispersion)'} }	10.249

```
anova(lme_light_clean1)
```

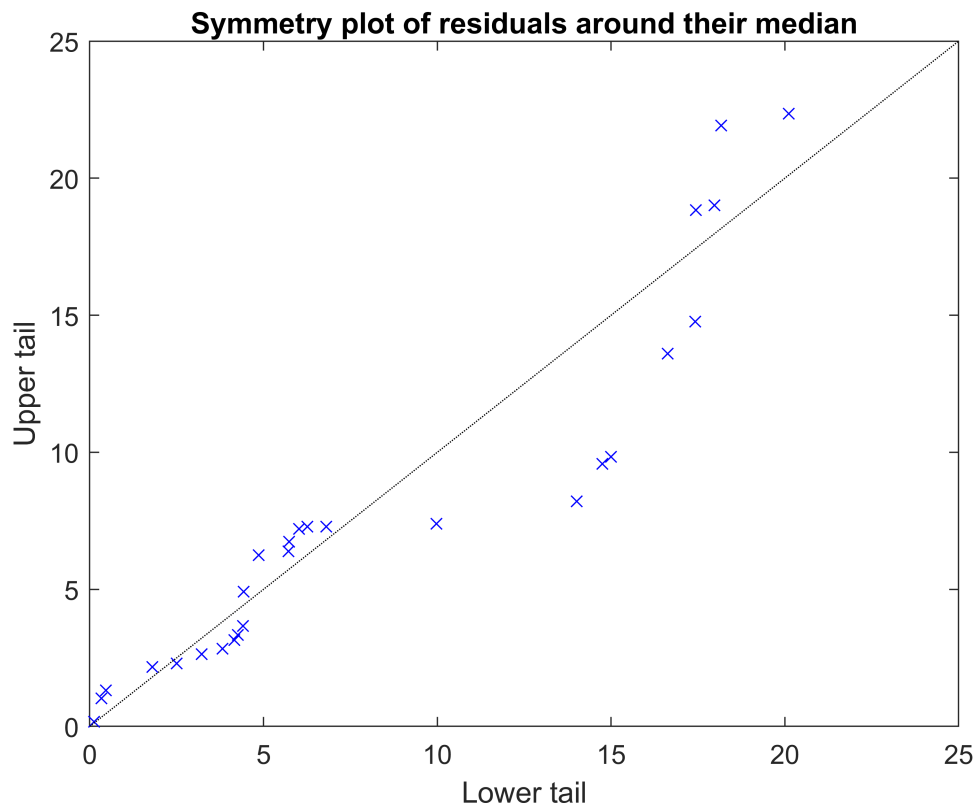
```
ans =
```

```
ANOVA marginal tests: DFMethod = 'residual'
```

Term	FStat	DF1	DF2	pValue
{'(Intercept)'} }	5.3861	1	51	0.024334
{'FA_WHOLESKEL' }	5.0141	1	51	0.02953
{'MO' }	1.6659	1	51	0.20263
{'FA_WHOLESKEL:MO' }	1.4704	1	51	0.23087

```
figure
```

```
plotResiduals(lme_light_clean1, 'symmetry')
```



```
% Interaction p val = 0.2
```

Influence of motion on the relationship between SN FA and tDCS-induced changes in brain activity

```
% Two SN analysis:
%Column 6:7 = SN PE
%Column 8 = SN FA

% Column 6
lme_light_clean1= fitglme(T, 'PE_SN_cope3 ~ MO*FA_WHOLESKEL');
disp(lme_light_clean1)
```

Generalized linear mixed-effects model fit by PL

Model information:

Number of observations	55
Fixed effects coefficients	4
Random effects coefficients	0
Covariance parameters	1
Distribution	Normal
Link	Identity
FitMethod	MPL

Formula:

```
PE_SN_cope3 ~ 1 + FA_WHOLESKEL*MO
```

Model fit statistics:

AIC	BIC	LogLikelihood	Deviance
379.01	389.05	-184.51	369.01

Fixed effects coefficients (95% CIs):

Name	Estimate	SE	tStat
{'(Intercept)'} }	-21.865	49.79	-0.43915
{'FA_WHOLESKEL' }	87.299	158.89	0.54941
{'MO' }	-4.8009	7.3093	-0.65682
{'FA_WHOLESKEL:MO' }	13.064	23.378	0.55883

DF	pValue	Lower	Upper
51	0.66241	-121.82	78.091
51	0.58512	-231.7	406.29
51	0.51425	-19.475	9.8731
51	0.57872	-33.869	59.997

Random effects covariance parameters:

Group: Error

Name	Estimate
{'sqrt(Dispersion)'} }	6.9289

```
anova(lme_light_clean1)
```

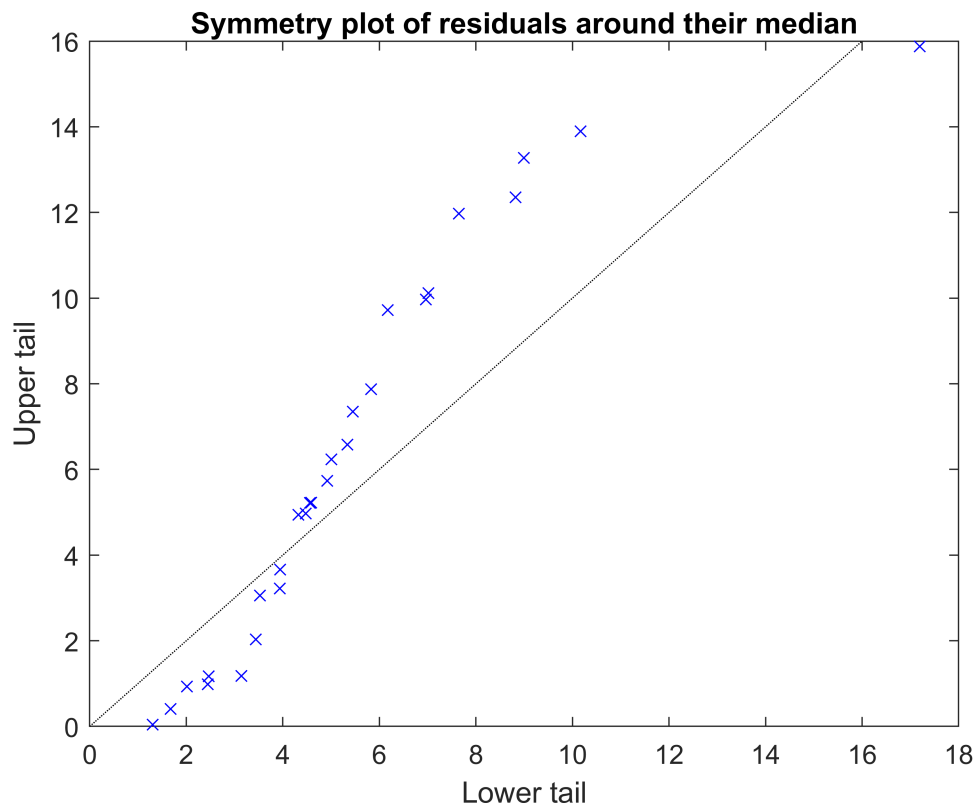
ans =

ANOVA marginal tests: DFMethod = 'residual'

Term	FStat	DF1	DF2	pValue
{'(Intercept)'} }	0.19285	1	51	0.66241
{'FA_WHOLESKEL' }	0.30185	1	51	0.58512
{'MO' }	0.43142	1	51	0.51425
{'FA_WHOLESKEL:MO' }	0.31229	1	51	0.57872

figure

```
plotResiduals(lme_light_clean1, 'symmetry')
```



```
% Interaction p val = 0.6
```

```
% Column 7
```

```
lme_light_clean1= fitglme(T,'PE_SN_cope5 ~ MO*FA_WHOLESKEL');  
disp(lme_light_clean1)
```

Generalized linear mixed-effects model fit by PL

Model information:

Number of observations	55
Fixed effects coefficients	4
Random effects coefficients	0
Covariance parameters	1
Distribution	Normal
Link	Identity
FitMethod	MPL

Formula:

```
PE_SN_cope5 ~ 1 + FA_WHOLESKEL*MO
```

Model fit statistics:

AIC	BIC	LogLikelihood	Deviance
387.97	398.01	-188.98	377.97

Fixed effects coefficients (95% CIs):

Name		Estimate	SE	tStat
{'(Intercept)'	}	46.842	54.014	0.86722
{'FA_WHOLESKEL'	}	-151.04	172.38	-0.87626
{'MO'	}	-10.23	7.9294	-1.2902
{'FA_WHOLESKEL:MO'}		32.065	25.361	1.2643

DF	pValue	Lower	Upper
51	0.38988	-61.595	155.28
51	0.385	-497.1	195.01
51	0.2028	-26.149	5.6885
51	0.21186	-18.85	82.979

Random effects covariance parameters:

Group: Error

Name	Estimate
{'sqrt(Dispersion)'}	7.5168

```
anova(lme_light_clean1)
```

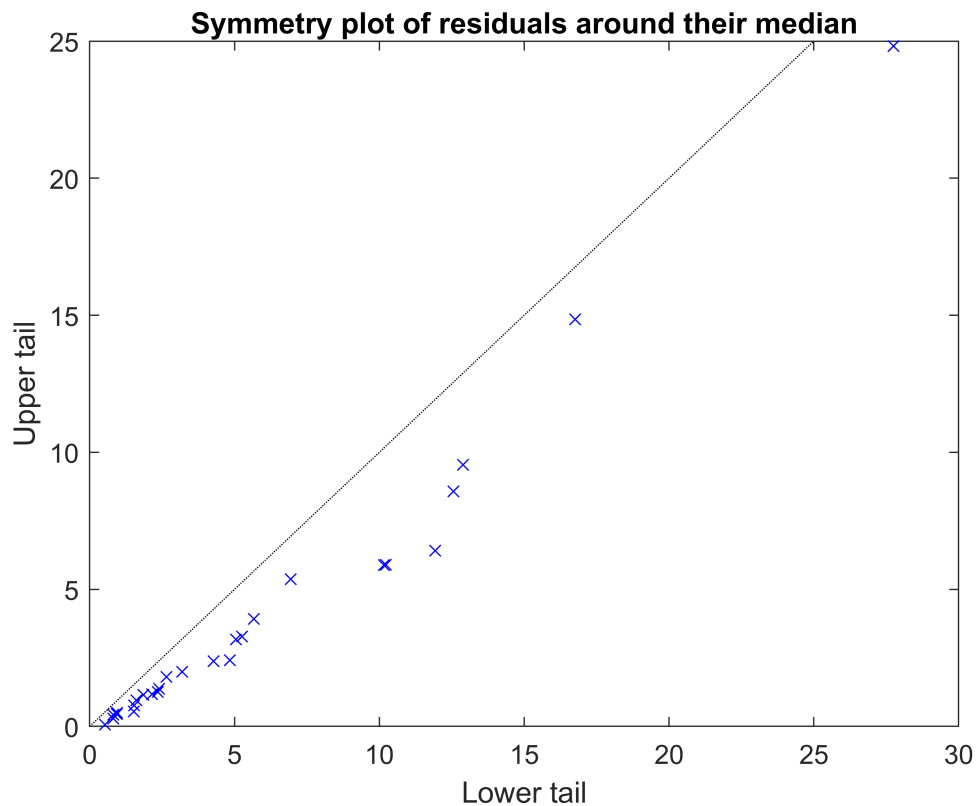
```
ans =
```

```
ANOVA marginal tests: DFMethod = 'residual'
```

Term		FStat	DF1	DF2	pValue
{'(Intercept)'	}	0.75208	1	51	0.38988
{'FA_WHOLESKEL'	}	0.76782	1	51	0.385
{'MO'	}	1.6646	1	51	0.2028
{'FA_WHOLESKEL:MO'}		1.5985	1	51	0.21186

figure

```
plotResiduals(lme_light_clean1, 'symmetry')
```



```
% Interaction p val = 0.2
```

Influence of motion on the relationship between DMN FA and tDCS-induced changes in brain activity

```
% Two DMN analysis:
%Column 9:10 = DMN PE
%Column 11 = DMN FA

% Column 9
lme_light_clean1= fitglm(T, 'PE_DMN_cope3 ~ MO*FA_WHOLESKEL');
disp(lme_light_clean1)
```

Generalized linear mixed-effects model fit by PL

Model information:

Number of observations	55
Fixed effects coefficients	4
Random effects coefficients	0
Covariance parameters	1
Distribution	Normal
Link	Identity
FitMethod	MPL

Formula:

```
PE_DMN_cope3 ~ 1 + FA_WHOLESKEL*MO
```

Model fit statistics:

AIC	BIC	LogLikelihood	Deviance
383.1	393.14	-186.55	373.1

Fixed effects coefficients (95% CIs):

Name	Estimate	SE	tStat
{'(Intercept)'} }	42.416	51.676	0.82081
{'FA_WHOLESKEL' }	-114.3	164.91	-0.69307
{'MO' }	-12.606	7.5862	-1.6617
{'FA_WHOLESKEL:MO' }	37.089	24.263	1.5286

DF	pValue	Lower	Upper
51	0.41557	-61.328	146.16
51	0.49141	-445.38	216.78
51	0.10271	-27.836	2.624
51	0.13255	-11.622	85.8

Random effects covariance parameters:

Group: Error

Name	Estimate
{'sqrt(Dispersion)'} }	7.1915

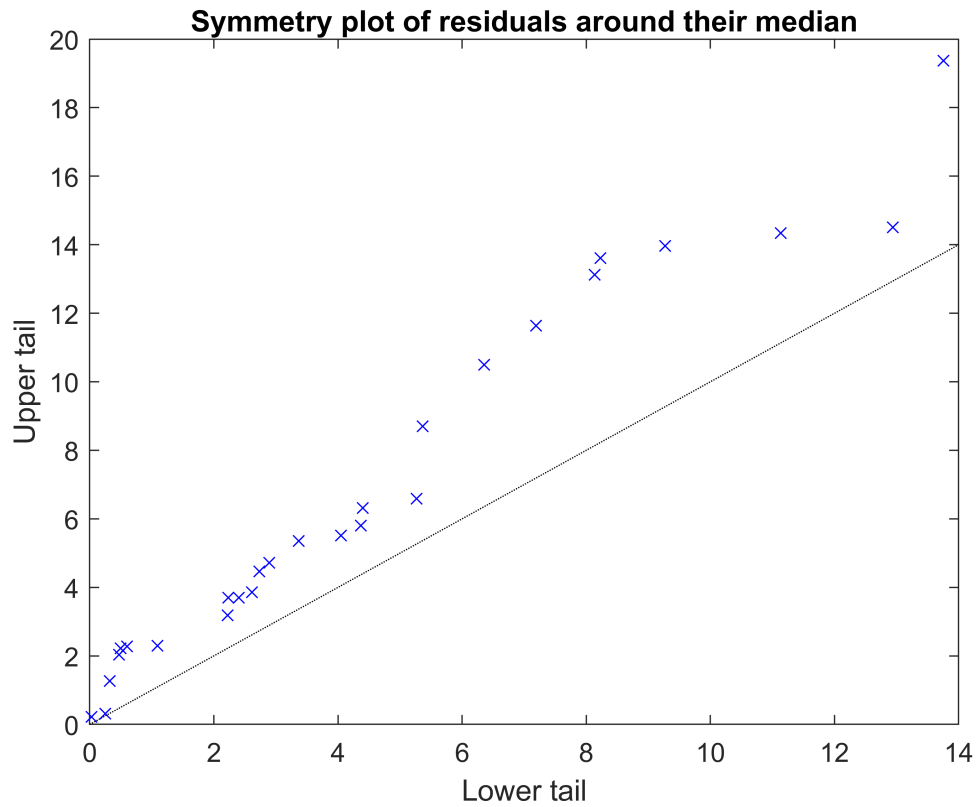
```
anova(lme_light_clean1)
```

ans =

ANOVA marginal tests: DFMethod = 'residual'

Term	FStat	DF1	DF2	pValue
{'(Intercept)'} }	0.67373	1	51	0.41557
{'FA_WHOLESKEL' }	0.48034	1	51	0.49141
{'MO' }	2.7612	1	51	0.10271
{'FA_WHOLESKEL:MO' }	2.3366	1	51	0.13255

```
figure
plotResiduals(lme_light_clean1, 'symmetry')
```



```
% Interaction p val = 0.1
```

```
% Column 10
```

```
lme_light_clean1= fitglme(T, 'PE_DMN_cope10 ~ MO*FA_WHOLESKEL');  
disp(lme_light_clean1)
```

Generalized linear mixed-effects model fit by PL

Model information:

Number of observations	55
Fixed effects coefficients	4
Random effects coefficients	0
Covariance parameters	1
Distribution	Normal
Link	Identity
FitMethod	MPL

Formula:

```
PE_DMN_cope10 ~ 1 + FA_WHOLESKEL*MO
```

Model fit statistics:

AIC	BIC	LogLikelihood	Deviance
385.35	395.38	-187.67	375.35

Fixed effects coefficients (95% CIs):

Name	Estimate	SE	tStat
{ '(Intercept)' }	18.506	52.742	0.35089
{ 'FA_WHOLESKEL' }	-62.451	168.32	-0.37103
{ 'MO' }	-8.4656	7.7427	-1.0934
{ 'FA_WHOLESKEL:MO' }	27.124	24.764	1.0953

DF	pValue	Lower	Upper
51	0.72712	-87.377	124.39
51	0.71215	-400.36	275.46
51	0.27937	-24.01	7.0786
51	0.27853	-22.592	76.84

Random effects covariance parameters:

Group: Error	
Name	Estimate
{'sqrt(Dispersion)'} }	7.3398

anova(lme_light_clean1)

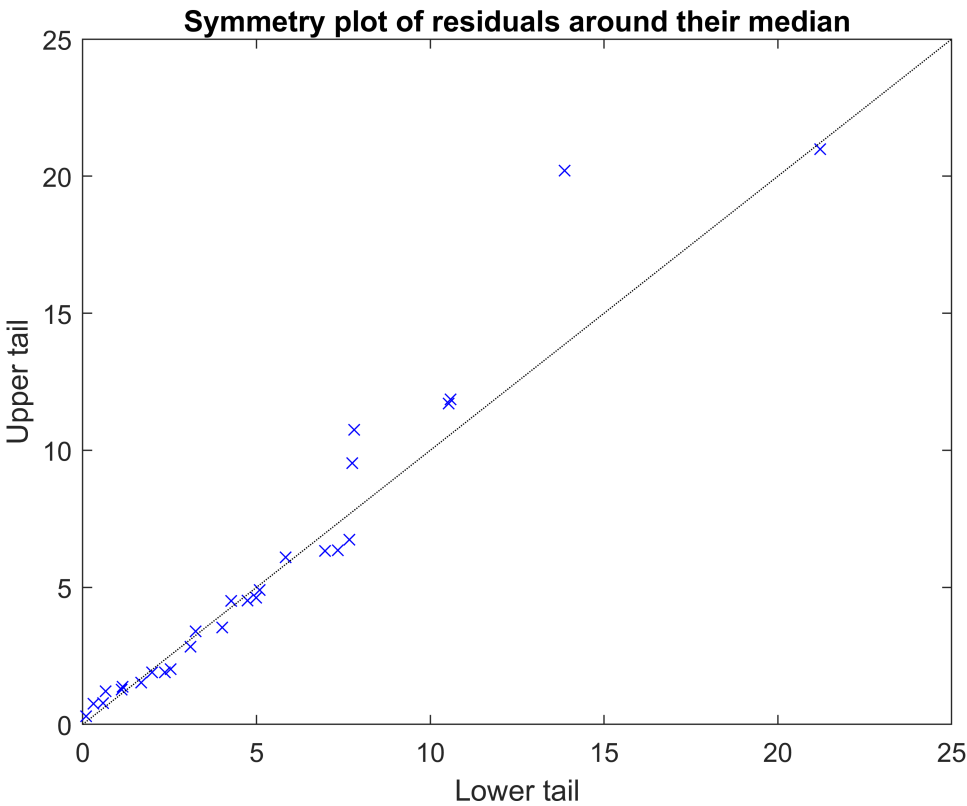
ans =

ANOVA marginal tests: DFMethod = 'residual'

Term		FStat	DF1	DF2	pValue
{' (Intercept) ' }		0.12312	1	51	0.72712
{' FA_WHOLESKEL ' }		0.13767	1	51	0.71215
{' MO ' }		1.1954	1	51	0.27937
{' FA_WHOLESKEL:MO ' }		1.1997	1	51	0.27853

figure

plotResiduals(lme_light_clean1, 'symmetry')



% Interaction p val = 0.3