COMP0118: Coursework 2

Mingzhou Hu

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Part 1

Q1 (abc)

```
% set random seed
rng(3)
% size of sample 1
s1 size = 25;
% size of sample 2
s2 size = 25;
% set the means to g1 and g2
mean s1 = 1;
mean s2 = 1.5;
% stochastic error
e1 = 0.25*randn(s1 size, 1);
e2 = 0.25*randn(s2 size, 1);
% generate datasets
sample1 = mean_s1+e1;
sample2 = mean s2+e2;
%% 1 (a)
% computer the mean for new datasets
new_mean_s1 = mean(sample1);
new mean s2 = mean(sample2);
\mbox{\%} computer the std for new datasets
new std s1 = std(sample1);
new std s2 = std(sample2);
%% 1 (b)
% perform two-sample t-test
[hyp,p,ci,stats] = ttest2(sample1, sample2);
%% 1 (c) (i)
% generate design matrix
oness = ones(s1_size,1);
zeross = zeros(s2 size,1);
design_mat_X = [oness, zeross; zeross, oness];
```

```
% dimension of col space of design matrix
dim X = rank(design mat X);
%% 1 (c) (ii)
% perpendicular projection operator Px corresponding to any C(X)
projection X =
design mat X*inv(design mat X'*design mat X)*design mat X';
% check idempotence
proj id = projection X^2;
% check symmetric
proj sy = projection X';
%% 1 (c) (iii)
% both groups
bothgroups = [sample1; sample2];
% use projection X to determine projection Y
projection Yhat = projection X*bothgroups;
%% 1 (c) (iv)
% compute Rx = I - Px
Rx = eye(size(bothgroups,1)) - projection X;
% check idempotence
Rxsq = Rx^2;
% check symmetric
Rxt = Rx';
%% 1 (c) (v)
% determine ehat, the projection of Y into the error space
ehat = Rx*bothgroups;
%% 1 (c) (vi)
% determine the angle between ehat and Yhat.
angle = acos(dot(ehat,projection Yhat));
%% 1 (c) (vii)
% use derived formula to determine betahat
betahat = inv(design mat X'*design mat X)*design mat X'*bothgroups;
%% 1 (c) (vii)
% estimate the variance of the stochastic component ehat
variance = (ehat'*ehat)/48;
%% 1_(c)_(ix)
% covariance matrix of estimated model parameters
```

```
covariance mat = variance*inv(design mat X'*design mat X);
% use covariance mat to determine std of model parameters
std parameters = sqrt(covariance mat(1,1));
%% 1 (c) (x)
% derive contrast vector for comparing group differences in means
lambda = [1 -1]';
% reduced model XO corresponding to the null hypothesis
X0 = ones(length(bothgroups),1);
%% 1 (c) (xi)
% recalculate new betahat
new betahat = inv(X0'*X0)*X0'*bothgroups;
% determine error via e = Y-X0B0
new errorhat = bothgroups-X0*new betahat;
% additional error as a result of placing the constraint
additional error = new errorhat - ehat;
% projection matrix for new design mat X0
projection X0 = X0*inv(X0'*X0)*X0';
% calculate m and n-k for f-statistic
m = trace(projection X-projection X0);
n k = trace(eye(length(bothgroups))-projection X);
% estimate f-statistic of comparing X0 to X
SSE1 = new_errorhat'*new_errorhat;
SSE2 = ehat'*ehat;
F = ((SSE1 - SSE2)/m)/(SSE2/n_k);
%% 1 (c) (xii)
% determine t-statistic to test whether one group has higher mean
t= (lambda'*betahat)/(sqrt(lambda'*covariance mat*lambda));
%% 1 (c) (xiii)
\ensuremath{\$} ground truth beta composed of ground truth means
beta gt = [1 \ 1.5]';
%% 1 (c) (xiv)
% compute the projection of the ground truth deviation e into C(X)
egtd = bothgroups - design_mat_X*beta_gt;
```

Q1 (def)

```
% set random seed
rng(3)
```

```
% size of sample 1
s1 \text{ size} = 25;
% size of sample 2
s2 size = 25;
% set the means to g1 and g2
mean s1 = 1;
mean s2 = 1.5;
% stochastic error
e1 = 0.25*randn(s1 size, 1);
e2 = 0.25*randn(s2 size, 1);
% generate datasets
sample1 = mean s1+e1;
sample2 = mean s2+e2;
% %% 1 (d)
% % generate design matrix
% oness = ones(s1 size,1);
% zeross = zeros(s2 size,1);
% design mat X = [oness, oness, zeross; oness, zeross, oness];
% % dimension of col space of design matrix
% dim X = rank(design mat X);
% % perpendicular projection operator Px corresponding to any C(X)
% projection X =
design mat X*pinv(design mat X'*design mat X) *design mat X';
% % both groups
% bothgroups = [sample1; sample2];
% % use projection X to determine projection Y
% projection Yhat = projection X*bothgroups;
% % compute Rx = I - Px
% Rx = eye(size(bothgroups,1)) - projection X;
% % determine ehat, the projection of Y into the error space
% ehat = Rx*bothgroups;
% % use derived formula to determine betahat
% betahat =
pinv(design mat X'*design mat X)*design mat X'*bothgroups;
% % estimate the variance of the stochastic component ehat
% variance = (ehat'*ehat)/48;
```

```
% % covariance matrix of estimated model parameters
% covariance_mat = variance*pinv(design_mat_X'*design_mat X);
% % use covariance mat to determine std of model parameters
% std parameters = sqrt(covariance mat(1,1));
% derive contrast vector for comparing group differences in means
% lambda = [0 1 -1]';
% % reduced model XO corresponding to the null hypothesis
% X0 = ones(length(bothgroups),1);
% % determine t-statistic to test whether one group has higher mean
% t= (lambda'*betahat)/(sqrt(lambda'*covariance mat*lambda));
%% 1 (e)
% generate design matrix
oness = ones(s1 size,1);
zeross = zeros(s2 size,1);
design mat X = [oness, oness; oness, zeross];
% dimension of col space of design matrix
dim_X = rank(design_mat_X);
% perpendicular projection operator Px corresponding to any C(X)
projection X =
design mat X*pinv(design mat X'*design mat X) *design mat X';
응응
% both groups
bothgroups = [sample1; sample2];
% use projection X to determine projection Y
projection Yhat = projection X*bothgroups;
% compute Rx = I - Px
Rx = eye(size(bothgroups,1)) - projection X;
% determine ehat, the projection of Y into the error space
ehat = Rx*bothgroups;
% use derived formula to determine betahat
betahat = pinv(design mat_X'*design_mat_X) *design_mat_X'*bothgroups;
% estimate the variance of the stochastic component ehat
variance = (ehat'*ehat)/48;
```

```
% covariance matrix of estimated model parameters
covariance_mat = variance*pinv(design_mat_X'*design_mat_X);
% use covariance_mat to determine std of model parameters
std_parameters = sqrt(covariance_mat(1,1));
%% 1_(d)_(iii)
% derive contrast vector for comparing group differences in means
lambda = [0 1]';
% reduced model X0 corresponding to the null hypothesis
X0 = ones(length(bothgroups),1);
%% 1_(d)_(iv)
% determine t-statistic to test whether one group has higher mean
t= (lambda'*betahat)/(sqrt(lambda'*covariance_mat*lambda));
```

Q2(a)

```
% set random seed
rng(3)
% size of sample 1
s1 size = 25;
% size of sample 2
s2 size = 25;
% set the means to g1 and g2
mean s1 = 1;
mean s2 = 1.5;
% stochastic error
e1 = 0.25*randn(s1 size, 1);
e2 = 0.25*randn(s2 size, 1);
% generate datasets
sample1 = mean s1+e1;
sample2 = mean s2+e2;
% perform two-sample t-test
[hyp,p,ci,stats] = ttest(sample1, sample2);
```

Q2(b)

```
% set random seed
rng(3);
% size of sample 1
```

```
s1 \text{ size} = 25;
% size of sample 2
s2 size = 25;
% set the means to g1 and g2
mean s1 = 1;
mean s2 = 1.5;
% stochastic error
e1 = 0.25*randn(s1 size, 1);
e2 = 0.25*randn(s2\_size, 1);
% generate datasets
sample1 = mean s1+e1;
sample2 = mean s2+e2;
% perform two-sample t-test
[hyp,p,ci,stats] = ttest(sample1, sample2);
% generate design matrix
oness = ones(s1 size,1);
zeross = zeros(s2 size,1);
identity = eye(s1_size);
design mat X = [oness,oness,identity;oness,zeross,identity];
% dimension of design matrix
dim X = rank(design mat X);
% perpendicular projection operator Px corresponding to any C(X)
projection X =
design mat X*pinv(design mat X'*design mat X) *design mat X';
% both groups
bothgroups = [sample1; sample2];
\mbox{\ensuremath{\$}} use projection_X to determine projection_Y
projection Yhat = projection X*bothgroups;
% compute Rx = I - Px
Rx = eye(size(bothgroups,1)) - projection_X;
% determine ehat, the projection of Y into the error space
ehat = Rx*bothgroups;
% use derived formula to determine betahat
betahat = pinv(design mat X'*design mat X) *design mat X'*bothgroups;
% estimate the variance of the stochastic component ehat
```

Part 2

Q1 (abc)

```
% set random seed
rng(10)
% size of sample 1
s1 size = 6;
% size of sample 2
s2 size = 8;
% set the means to g1 and g2
mean s1 = 1;
mean s2 = 1.5;
% stochastic error
e1 = 0.25*randn(s1 size, 1);
e2 = 0.25*randn(s2 size, 1);
% generate datasets
sample1 = mean s1+e1;
sample2 = mean s2+e2;
%% 1 (a)
% perform two-sample t-test
[hyp,p1,ci,stats] = ttest2(sample1, sample2);
original tstatistic = stats.tstat;
original meandiff = mean(sample1) -mean(sample2);
```

```
%% 1 (b) (i)
% both groups
bothgroups = [sample1; sample2];
% construct an one-dimensional array D
D=bothgroups';
%% 1 (b) (ii)&(iii)
% construct all the valid permutations of D
C1 = combnk(D, s1 size);
% initialise C2
C2 = zeros(length(C1),s2 size);
% initialise array to store all t-statistics
tstatistic array = zeros(1, length(C1)); % initialise array to store
all t-statistics
for i=1:length(C1)
   % make sure there is no repeat for valid permutations
   C2(i,:) = setdiff(D, C1(i,:));
   % compute the t-statistics for all the membership permutations
   [hyp,p2,ci,stats] = ttest2(C1(i,:), C2(i,:));
   tstatistic array(i) = stats.tstat;
end
% construct empirical distribution of t-statistic
figure,
hist(tstatistic_array, 50);
title('The empirical distribution of t-statistic');
%% 1 (b) (iv)
% determine the p-value by finding the percentage of the permutations
with a t-statistic greater
% than and equal to that of the original labeling.
p value b = nnz(tstatistic array <=</pre>
original tstatistic)/numel(tstatistic array);
%% 1 (c)
% repeat (b) but rather than using the t-statistic, use the
difference between
% the means as the test statistic
mean diff array = zeros(1, length(C1));
for i=1:length(C1)
  mean diff = mean(C1(i,:)) - mean(C2(i,:));
  mean diff array(i) = mean diff;
end
figure,
```

```
hist(mean_diff_array, 50);
title('The empirical distribution of the difference between means');
% p-value computed with mean difference
p_value_c = nnz(mean_diff_array <=
original meandiff)/numel(mean diff array);</pre>
```

Q1 (d)

```
% set random seed
rng(10);
% size of sample 1
s1 size = 6;
% size of sample 2
s2 size = 8;
% set the means to g1 and g2
mean s1 = 1;
mean s2 = 1.5;
% stochastic error
e1 = 0.25*randn(s1 size, 1);
e2 = 0.25*randn(s2 size, 1);
% generate datasets
sample1 = mean s1+e1;
sample2 = mean s2+e2;
% perform two-sample t-test
[hyp,p1,ci,stats] = ttest2(sample1, sample2);
original tstatistic = stats.tstat;
original_meandiff = mean(sample1) -mean(sample2);
% both groups
bothgroups = [sample1; sample2];
% construct an one-dimensional array D
D=bothgroups';
%% d(i)
total permutations = 1000;
tstatistic_array_d = zeros(1,total_permutations);
all permutations = zeros(total permutations, 6);
for i=1:total permutations
   % use randperm to generate a random set of permutations of the
   % intergers
   random_permutations = randperm(14);
   all_permutations(i,:) = random_permutations(1:6);
```

```
bothgroups = D(random permutations);
   group1 = bothgroups(1:6);
   group2 = bothgroups(7:14);
   [hyp,p2,ci,stats] = ttest2(group1, group2);
   tstatistic array d(i) = stats.tstat;
end
figure,
hist(tstatistic array d,50);
title('The empirical distribution of t-statistic');
p value d = nnz(tstatistic array d
<=original tstatistic)/numel(tstatistic array d);</pre>
%% (diii)
% sort the elements of each row of permutations
sort permutations = sort(all permutations(:,1:6),2);
% get the position of elments without repetition
[~, id1, ~]=unique(sort permutations, 'rows');
extended permutations = 1500;
tstatistic array diii = zeros(1, total permutations);
all permutations iii= zeros(extended permutations, 14);
for i = 1:extended permutations
   % randperm generates random set of integer permutations
   % integers are indices for D
   random permutations iii = randperm(14);
   all permutations iii(i,:) = random permutations iii;
   bothgroups iii = D(random permutations iii);
   group1 iii = bothgroups iii(1:6);
   group2 iii = bothgroups iii(7:14);
   [hyp,p3,ci,stats] = ttest2(group1_iii, group2_iii);
   tstatistic_array_diii(i) = stats.tstat;
end
% sort the elements of each row of permutations
sort permutations iii = sort(all permutations iii(:,1:6),2);
% get the position of elments without repetition
[~, id2, ~]=unique(sort permutations iii, 'rows');
new permutations = sort permutations iii(id2,:);
new permutations = new permutations(1:total permutations, :);
tstatistic array diii = tstatistic array diii(id2);
tstatistic_array_diii = tstatistic_array_diii(10:1010);
% p-value without repetition
p value diii = nnz(tstatistic array diii <=</pre>
```

```
original tstatistic)/numel(tstatistic array diii);
```

Q2

```
% load files
CPA I = [4,5,6,7,8,9,10,11]; % identifier for CPA file names
PPA I = [3,6,9,10,13,14,15,16]; % identifier for PPA file names
number subjects = 8;
% group1 had 8 subjects, 64000=40*40*40
CPA = zeros(64000, number subjects);
\dot{j} = 0;
for i=drange(CPA I)
   filename = sprintf('CPA%d diffeo fa.img',i);
   fid = fopen(filename, 'r', 'l'); % little-endian
   CPA sub = fread(fid, 'float'); % 16-bit floating point
   j=j+1;
   CPA(:,j) = CPA sub;
% group2 had 8 subjects, 64000=40*40*40
PPA = zeros(64000, number subjects);
\dot{j} = 0;
for i=drange(PPA I)
   filename = sprintf('PPA%d diffeo fa.img',i);
   fid = fopen(filename, 'r', 'l'); % little-endian
   PPA sub = fread(fid, 'float'); % 16-bit floating point
   j=j+1;
   PPA(:,j) = PPA sub;
end
% wm mask.img is an additional binary volume defining the ROI for
% statistical analysis
fid = fopen('wm mask.img', 'r', 'l');
mask = fread(fid, 'float');
voxels = [CPA PPA];
응응 2 (a)
% we use the GLM in part1 1c Y = X1b1 + X2b2 + e.
% generate design matrix
oness = ones(number subjects,1);
zeross = zeros(number subjects,1);
design mat X = [oness, zeross; zeross, oness];
% contrast vector
lambda = [1 -1]';
% determine betahat
```

```
betahat = voxels*(inv(design mat X'*design mat X) *design mat X')';
% determine ehat
ehat = voxels-(betahat*design mat X');
% covariance matrix of estimated model parameters
variance = sum((ehat.*ehat), 2)./ 14;
t statiatic =
(betahat*lambda)./sqrt(variance.*(lambda'*(inv(design mat X'*design m
at X)) *lambda));
% compute t-statisti for ROI
ROItstatistic = t statiatic.*mask;
% compute maximum t-statistic.
maxtstatistic = max(ROItstatistic);
%% 2 (b)
s1 size = 8;
s2 size = 8;
indices = 1:(s1 size+s2 size);
% construct all the valid permutations of indices
C1 = combnk(indices, s1 size);
% initialise C2
C2 = zeros(length(C1),s2 size);
for i=1:length(C1)
   % make sure there is no repeat for valid permutations
   C2(i,:) = setdiff(indices, C1(i,:));
end
bothgroups = [C1 C2];
% initialise array to store all max t-statistic
max tstatistic array = zeros((length(C1)),1);
% just keep the values in ROI
voxels = voxels.*mask;
for i=1:length(C1)
   voxels now = voxels(:,bothgroups(i,:));
   betahat now =
voxels now*(inv(design mat X'*design mat X) *design mat X')';
   ehat now = voxels now-betahat now*design mat X';
   % find std of error
   variance = sum((ehat now.*ehat now), 2)./ 14;
   tstatistic =
(betahat now*lambda)./sqrt(variance.*(lambda'*(inv(design mat X'*desi
gn mat X))*lambda));
   max_tstatistic_array(i,1) = max(tstatistic);
end
hist(max tstatistic array,50);
```

```
title('The empirical distribution of max t-statistic');

%% (c)
p_value_d =
nnz(max_tstatistic_array >=maxtstatistic)/numel(max_tstatistic_array);

%% (d)
% determine maximum t-statistic threshold corresponding to p-value of
5%
threeshold=prctile(max tstatistic array, 95);
```