Coefficient covariance for rotated data with identical weights

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This document contains R-code for analysing the coefficient covariance for two features with sample weights equal 1. We compare coefficient covariance of non-rotated data and rotated data. Therefore we assume a sample design with age and group (case, control) as covariates. group is the hypothesis coefficient.

age	group
7	case
11	case
17	case
27	case
6	case
26	case

The correlation coefficient of the error terms between feature 1 and feature 2 is assumed as 0.8. However, as it is a constant factor in the covariance of the coefficients, the value of the correlation coefficient solely changes the scales of the plots below, but not the pattern.

```
X <- model.matrix(~ 1 + age + group, samp.inf)</pre>
rho12 <- 0.8
# assume sigma1 = sigma2 = 1
sigma12 <- rho12
# define random uniformly distributed weights
W <- matrix(runif(10), nrow = 2, ncol = n, byrow = TRUE)
          [,1]
                  [,2]
                         [,3]
                                 [,4]
                                        [,5]
                                                [,6]
                                                        [,7]
                                                                [,8]
                                                                        [,9]
                                                                             [,10]
```

```
## [,1] [,2] [,3] [,4] [,5] [,6] [,7] [,8] [,9] [,10]
## [1,] 0.9347 0.2121 0.6517 0.1256 0.2672 0.3861 0.01339 0.3824 0.8697 0.3403
```

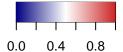
```
## [2,] 0.9347 0.2121 0.6517 0.1256 0.2672 0.3861 0.01339 0.3824 0.8697 0.3403 ## [,11] [,12] [,13] [,14] [,15] [,16] [,17] [,18] [,19] [,20] ## [1,] 0.9347 0.2121 0.6517 0.1256 0.2672 0.3861 0.01339 0.3824 0.8697 0.3403 ## [2,] 0.9347 0.2121 0.6517 0.1256 0.2672 0.3861 0.01339 0.3824 0.8697 0.3403
```

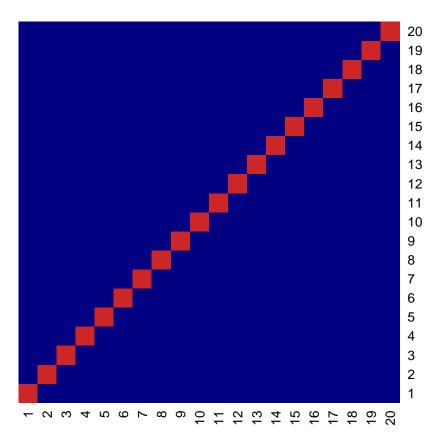
The following steps of whitening, QR-decomposition and generation of random (restricted) rotation matrices are outlined in the main manuscript.

```
# whitening of X
X1 <- sqrt(W[1,]) * X</pre>
X2 \leftarrow sqrt(W[2,]) * X
# qr decomposition
# group as "hypothesis coefficient"
# intercept and age as "determined coefficients"
coef.d <- 1:2
Q1 <- qr.Q(qr(X1), complete = TRUE)
Xd1 <- Q1[,coef.d]</pre>
Xhe1 <- Q1[,-coef.d]</pre>
Q2 <- qr.Q(qr(X2), complete = TRUE)
Xd2 \leftarrow Q2[,coef.d]
Xhe2 \leftarrow Q2[,-coef.d]
r <- ncol(Xhe1)
E.R1R2 <- Xd1 %*% t(Xd1) %*% Xd2 %*% t(Xd2) +
    1/r * Xhe1 %*% t(Xhe2) * sum(diag(t(Xhe1)%*%Xhe2))
```

The heatmap shows the expected value E.R1R2

$$\mathbb{E}_{R}\left[ilde{oldsymbol{R}}_{r1}^{*}\, ilde{oldsymbol{R}}_{r2}^{*T}
ight]$$





The diagonal elements of E.R1R2 are

diag(E.R1R2)

In the following, we calculate

$$\mathbb{E}_{R}\left[ilde{oldsymbol{R}}_{r1}^{*}\, ilde{oldsymbol{R}}_{r2}^{*T}
ight]$$

and estimate the element wise standard deviation

$$\operatorname{sd}_{R}\left[\tilde{\boldsymbol{R}}_{r1}^{*}\,\tilde{\boldsymbol{R}}_{r2}^{*T}\right]$$

```
E.cov.beta.r <- sigma12 *
    solve(t(X1)%*%X1) %*% t(X1) %*% E.R1R2 %*% X2 %*% solve(t(X2)%*%X2)

covs <- vapply(1:100, function(i){
    R <- randorth(ncol(Xhe1))
    R1 <- Xd1 %*% t(Xd1) + Xhe1 %*% R %*% t(Xhe1)
    R2 <- Xd2 %*% t(Xd2) + Xhe2 %*% R %*% t(Xhe2)

sigma12 * solve(t(X1)%*%X1) %*% t(X1) %*% R1 %*%
    t(R2) %*% X2 %*% solve(t(X2)%*%X2) - E.cov.beta.r</pre>
```

```
}, matrix(1.2, 3,3))
sd.cov.beta.r <- apply(covs, 1:2, function(i)sqrt(mean(i^2)))
cov.beta <- sigma12 * solve(t(X1)%*%X1) %*% t(X1) %*% X2 %*% solve(t(X2)%*%X2)</pre>
```

Coefficient covariance for non-rotated data:

kable(cov.beta)

	(Intercept)	age	groupcontrol
(Intercept)	0.5223	-0.0236	-0.1756
age	-0.0236	0.0017	-0.0011
groupcontrol	-0.1756	-0.0011	0.3832

Expected coefficient covariance for rotated data:

kable(E.cov.beta.r)

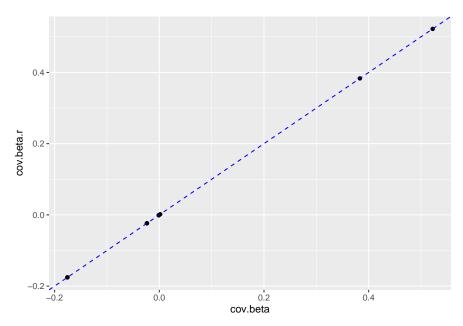
	(Intercept)	age	groupcontrol
(Intercept)	0.5223	-0.0236	-0.1756
age	-0.0236	0.0017	-0.0011
groupcontrol	-0.1756	-0.0011	0.3832

Standard deviation of coefficient covariance for rotated data:

kable(sd.cov.beta.r)

	(Intercept)	age	groupcontrol
(Intercept)	0	0	0
age	0	0	0
groupcontrol	0	0	0

The graphical representation of these tables is:



The given example shows, that for identical weights, the dependence structure of coefficient estimates is exactly retained with random rotation for the assumed experimental design.

Session Info

sessionInfo()

```
## R Under development (unstable) (2020-11-14 r79432)
## Platform: x86_64-w64-mingw32/x64 (64-bit)
## Running under: Windows 10 x64 (build 19041)
##
## Matrix products: default
##
## locale:
## [1] LC_COLLATE=German_Austria.1252 LC_CTYPE=German_Austria.1252
## [3] LC_MONETARY=German_Austria.1252 LC_NUMERIC=C
## [5] LC_TIME=German_Austria.1252
##
## attached base packages:
  [1] stats
                 graphics grDevices utils
                                                datasets methods
                                                                    base
## other attached packages:
## [1] heatmap3_1.1.7
                          knitr_1.30
                                              ggplot2_3.3.2
                                                                 randRotation_1.3.4
##
## loaded via a namespace (and not attached):
##
   [1] xm12_1.3.2
                           magrittr_1.5
                                               munsell_0.5.0
                                                                  colorspace_2.0-0
##
   [5] R6_2.5.0
                           rlang_0.4.8
                                               highr_0.8
                                                                  fastcluster_1.1.25
   [9] stringr_1.4.0
                           tools_4.1.0
                                               rbibutils_1.4
                                                                  grid_4.1.0
## [13] gtable_0.3.0
                           xfun_0.19
                                               withr_2.3.0
                                                                  ellipsis_0.3.1
  [17] htmltools 0.5.0
                           yaml 2.2.1
                                               digest 0.6.27
                                                                  tibble 3.0.4
  [21] lifecycle_0.2.0
                           crayon_1.3.4
                                               farver_2.0.3
                                                                  vctrs_0.3.4
## [25] Rdpack_2.1
                           gbRd_0.4-11
                                               glue_1.4.2
                                                                  evaluate_0.14
  [29] rmarkdown_2.5
                           labeling_0.4.2
                                               stringi_1.5.3
                                                                  pillar_1.4.6
## [33] compiler_4.1.0
                           scales_1.1.1
                                               pkgconfig_2.0.3
```