

main

August 12, 2021

1 Boxplots for permutations analysis (male downsampling to female level)

```
[1]: library(tidyverse)
library(ggpubr)
```

```
Attaching packages: tidyverse
1.3.1
```

```
ggplot2 3.3.5    purrr  0.3.4
tibble  3.1.2    dplyr  1.0.7
tidyr   1.1.3    stringr 1.4.0
readr   1.4.0    forcats 0.5.1
```

Conflicts

```
tidyverse_conflicts()
dplyr::filter() masks stats::filter()
dplyr::lag()     masks stats::lag()
```

1.1 Function and configuration

```
[2]: save_ggplots <- function(p, fn, w=6, h=6){
  for(ext in c('.svg', '.png', '.pdf')){
    ggsave(p, filename=paste0(fn, ext), width=w, height=h)
  }
}
```

```
[3]: config = list(
  "Caudate" = "../.../caudate/subsampling_male/deg_summary/_m/permutations.
↪ csv",
  "DLPFC" = "../.../dlpfc/subsampling_male/deg_summary/_m/permutations.csv",
  "Hippocampus" = "../.../hippocampus/subsampling_male/deg_summary/_m/
↪ permutations.csv"
)
```

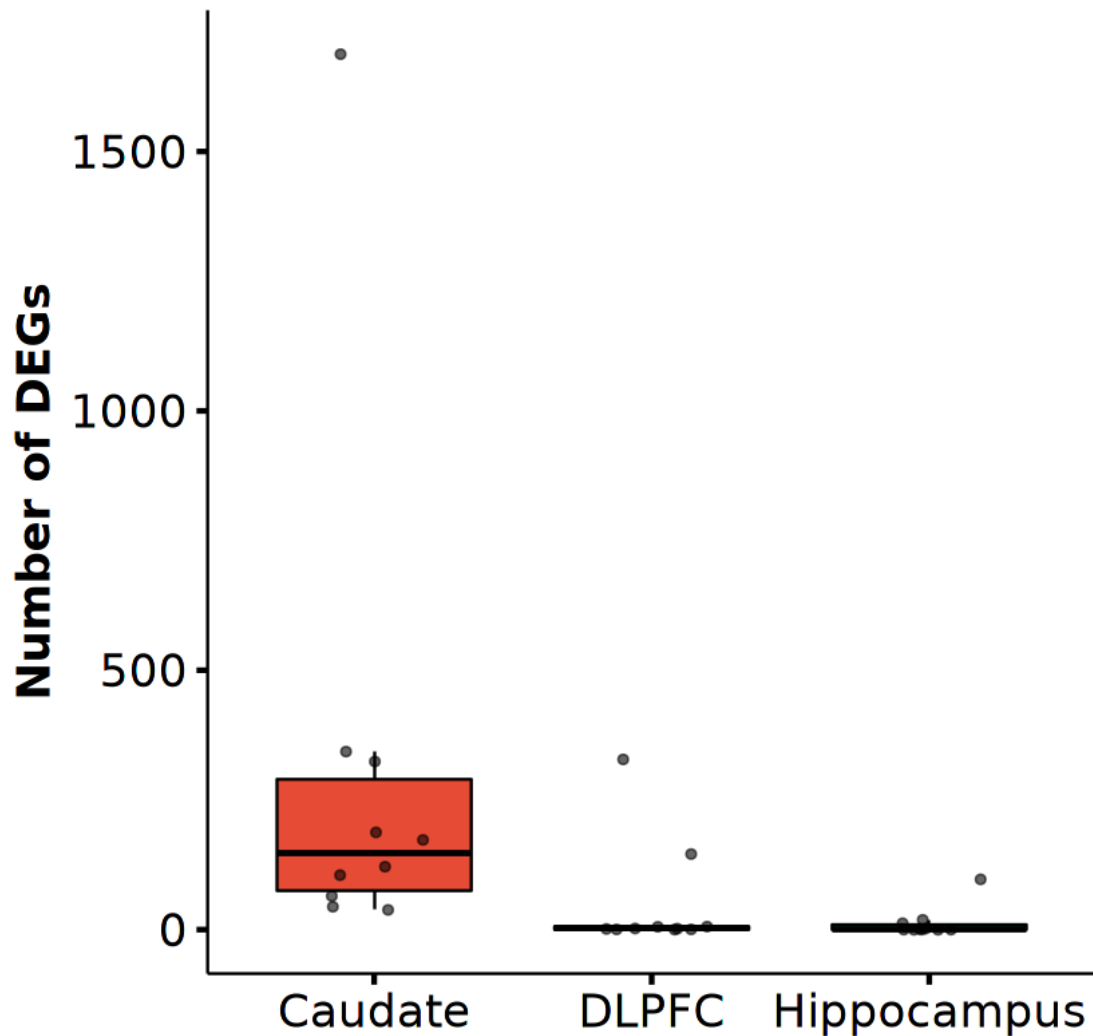
1.2 Merge dataframes

```
[4]: datalist = list()
for(tissue in c("Caudate", "DLPFC", "Hippocampus")){
  df = data.table::fread(config[[tissue]])
  df$tissue <- tissue # maybe you want to keep track of which iteration
  # produced it?
  datalist[[tissue]] <- df
}
big_df <- bind_rows(datalist) %>%
  select(Symbol, gencodeID, logFC, t, adj.P.Val, Permutation, tissue)
big_df %>% head()
```

	Symbol <chr>	gencodeID <chr>	logFC <dbl>	t <dbl>	adj.P.Val <dbl>	Permutation <int>
A data.table: 6 × 7	GDNF-AS1	ENSG00000248587.7	1.0159219	8.553727	1.566672e-09	2
	GDNF	ENSG00000168621.14	0.8188297	6.045744	1.541499e-04	2
	NSUN5P2	ENSG00000106133.17	-0.3648580	-6.031454	1.541499e-04	2
	KCNH5	ENSG00000140015.19	0.5558007	5.980020	1.541499e-04	2
	PAPPA	ENSG00000182752.9	0.7535862	5.862024	2.130430e-04	2
	RMDN2	ENSG00000115841.19	-0.3193313	-5.796196	2.403024e-04	2

```
[5]: bxp = big_df %>% group_by(Permutation, tissue) %>% summarize(Size = n()) %>%
  as.data.frame %>% pivot_wider(names_from = tissue, values_from = Size) %>%
  replace_na(list(DLPFC = 0, Hippocampus = 0)) %>%
  pivot_longer(-Permutation, names_to="Tissue", values_to="DEGs") %>%
  mutate_if(is.character, as.factor) %>%
  ggboxplot(x="Tissue", y="DEGs", fill="Tissue", add="jitter",
    xlab='', palette="npg", ylab="Number of DEGs",
    add.params=list(alpha=0.6), outlier.shape=NA,
    panel.labs.font=list(face='bold'), legend="",
    ggtheme=theme_pubr(base_size=20))+
    font("xy.title", face="bold")
save_ggplots(bxp, "permutation_boxplots_3regions", 6, 5)
bxp
```

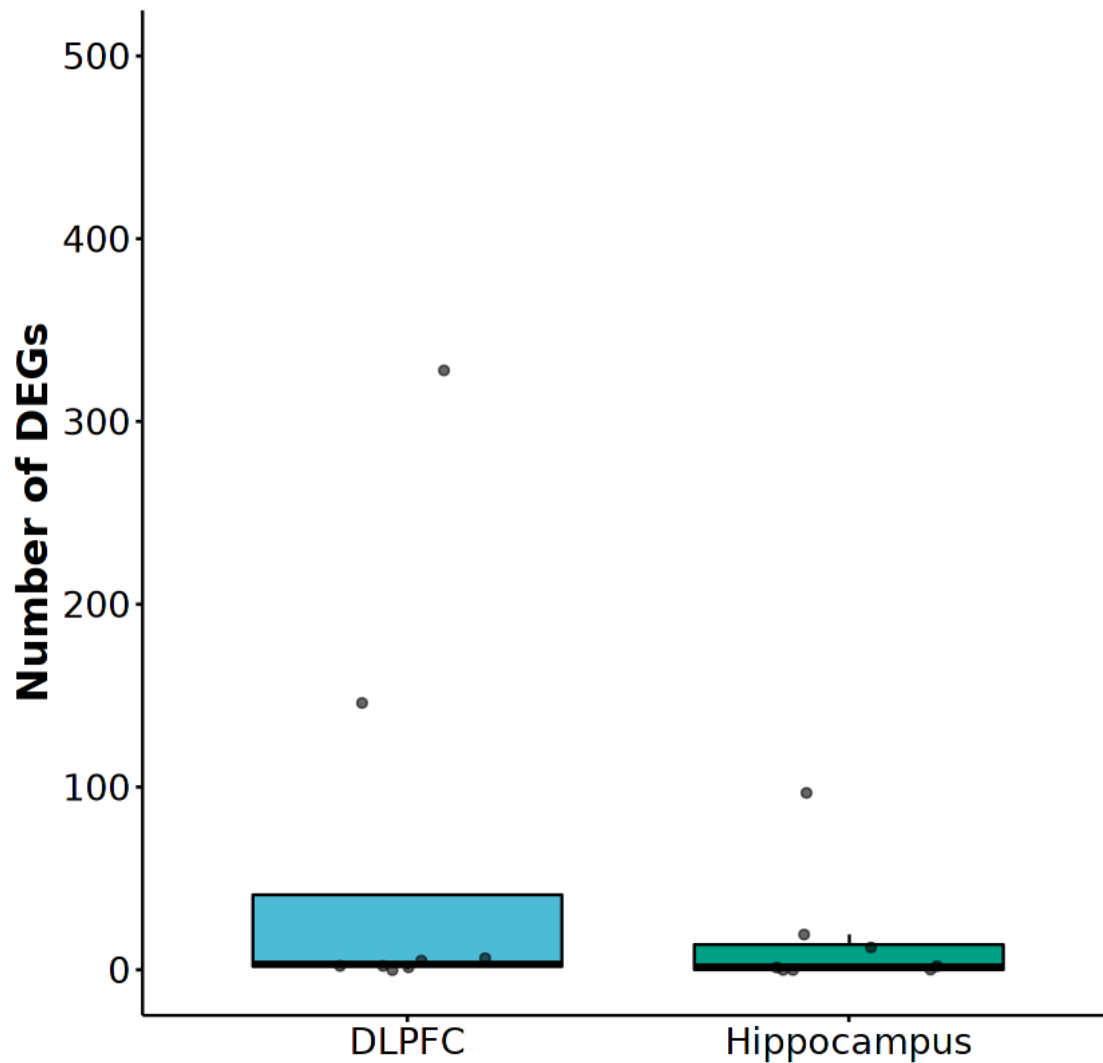
`summarise()` has grouped output by 'Permutation'. You can override using the
`.groups` argument.



```
[6]: bxp = big_df %>% filter(tissue != "Caudate") %>%
  group_by(Permutation, tissue) %>% summarize(Size = n()) %>%
  as.data.frame %>% pivot_wider(names_from = tissue, values_from = Size) %>%
  replace_na(list(DLPFC = 0, Hippocampus = 0)) %>%
  pivot_longer(-Permutation, names_to="Tissue", values_to="DEGs") %>%
  mutate_if(is.character, as.factor) %>%
  ggboxplot(x="Tissue", y="DEGs", fill="Tissue", add="jitter",
    xlab='', palette=get_palette("npg", 3)[2:3], ylab="Number of DEGs",
    add.params=list(alpha=0.6), outlier.shape=NA,
    panel.labs.font=list(face='bold', size = 18),
    legend="", ylim=c(0, 500))+
  font("xy.title", size=18, face="bold") +
```

```
font("xy.text", size=16)
save_ggplots(bxp, "permutation_boxplots_2regions", 6, 5)
bxp
```

`summarise()` has grouped output by 'Permutation'. You can override using the `.groups` argument.



```
[7]: big_df %>% group_by(Permutation, tissue) %>% summarize(Size = n()) %>%
  as.data.frame %>% pivot_wider(names_from = tissue, values_from = Size) %>%
  replace_na(list(DLPFC = 0, Hippocampus = 0)) %>%
  pivot_longer(-Permutation, names_to="Tissue", values_to="DEGs") %>%
  mutate_if(is.character, as.factor) %>%
```

```
group_by(Tissue) %>% summarize(Mean = mean(DEGs), Median = median(DEGs),  
  Std = sd(DEGs))
```

`summarise()` has grouped output by 'Permutation'. You can override using the
`.groups` argument.

	Tissue <fct>	Mean <dbl>	Median <dbl>	Std <dbl>
A tibble: 3 × 4	Caudate	308.8	147.0	496.27094
	DLPFC	49.0	2.0	107.99177
	Hippocampus	13.1	0.5	30.18995

1.3 Reproducibility Information

```
[8]: Sys.time()  
proc.time()  
options(width = 120)  
sessioninfo::session_info()
```

```
[1] "2021-08-12 19:05:34 EDT"
```

```
  user system elapsed  
4.731  0.222   5.261
```

Session info

```
setting value  
version R version 4.0.3 (2020-10-10)  
os Arch Linux  
system x86_64, linux-gnu  
ui X11  
language (EN)  
collate en_US.UTF-8  
ctype en_US.UTF-8  
tz America/New_York  
date 2021-08-12
```

Packages

package	* version	date	lib	source
abind	1.4-5	2016-07-21	[1]	CRAN (R 4.0.2)
assertthat	0.2.1	2019-03-21	[1]	CRAN (R 4.0.2)
backports	1.2.1	2020-12-09	[1]	CRAN (R 4.0.2)
base64enc	0.1-3	2015-07-28	[1]	CRAN (R 4.0.2)
broom	0.7.8	2021-06-24	[1]	CRAN (R 4.0.3)
Cairo	1.5-12.2	2020-07-07	[1]	CRAN (R 4.0.2)
car	3.0-11	2021-06-27	[1]	CRAN (R 4.0.3)
carData	3.0-4	2020-05-22	[1]	CRAN (R 4.0.2)
cellranger	1.1.0	2016-07-27	[1]	CRAN (R 4.0.2)
cli	3.0.0	2021-06-30	[1]	CRAN (R 4.0.3)

colorspace	2.0-2	2021-06-24	[1]	CRAN	(R 4.0.3)
crayon	1.4.1	2021-02-08	[1]	CRAN	(R 4.0.3)
curl	4.3.2	2021-06-23	[1]	CRAN	(R 4.0.3)
data.table	1.14.0	2021-02-21	[1]	CRAN	(R 4.0.3)
DBI	1.1.1	2021-01-15	[1]	CRAN	(R 4.0.2)
dbplyr	2.1.1	2021-04-06	[1]	CRAN	(R 4.0.3)
digest	0.6.27	2020-10-24	[1]	CRAN	(R 4.0.2)
dplyr	* 1.0.7	2021-06-18	[1]	CRAN	(R 4.0.3)
ellipsis	0.3.2	2021-04-29	[1]	CRAN	(R 4.0.3)
evaluate	0.14	2019-05-28	[1]	CRAN	(R 4.0.2)
fansi	0.5.0	2021-05-25	[1]	CRAN	(R 4.0.3)
farver	2.1.0	2021-02-28	[1]	CRAN	(R 4.0.3)
forcats	* 0.5.1	2021-01-27	[1]	CRAN	(R 4.0.2)
foreign	0.8-80	2020-05-24	[2]	CRAN	(R 4.0.3)
fs	1.5.0	2020-07-31	[1]	CRAN	(R 4.0.2)
generics	0.1.0	2020-10-31	[1]	CRAN	(R 4.0.2)
ggplot2	* 3.3.5	2021-06-25	[1]	CRAN	(R 4.0.3)
ggpubr	* 0.4.0	2020-06-27	[1]	CRAN	(R 4.0.2)
ggsci	2.9	2018-05-14	[1]	CRAN	(R 4.0.2)
ggsignif	0.6.2	2021-06-14	[1]	CRAN	(R 4.0.3)
glue	1.4.2	2020-08-27	[1]	CRAN	(R 4.0.2)
gtable	0.3.0	2019-03-25	[1]	CRAN	(R 4.0.2)
haven	2.4.1	2021-04-23	[1]	CRAN	(R 4.0.3)
hms	1.1.0	2021-05-17	[1]	CRAN	(R 4.0.3)
htmltools	0.5.1.1	2021-01-22	[1]	CRAN	(R 4.0.2)
httr	1.4.2	2020-07-20	[1]	CRAN	(R 4.0.2)
IRdisplay	1.0	2021-01-20	[1]	CRAN	(R 4.0.2)
IRkernel	1.2	2021-05-11	[1]	CRAN	(R 4.0.3)
jsonlite	1.7.2	2020-12-09	[1]	CRAN	(R 4.0.2)
labeling	0.4.2	2020-10-20	[1]	CRAN	(R 4.0.2)
lifecycle	1.0.0	2021-02-15	[1]	CRAN	(R 4.0.3)
lubridate	1.7.10	2021-02-26	[1]	CRAN	(R 4.0.3)
magrittr	2.0.1	2020-11-17	[1]	CRAN	(R 4.0.2)
modelr	0.1.8	2020-05-19	[1]	CRAN	(R 4.0.2)
munsell	0.5.0	2018-06-12	[1]	CRAN	(R 4.0.2)
openxlsx	4.2.4	2021-06-16	[1]	CRAN	(R 4.0.3)
pbdZMQ	0.3-5	2021-02-10	[1]	CRAN	(R 4.0.3)
pillar	1.6.1	2021-05-16	[1]	CRAN	(R 4.0.3)
pkgconfig	2.0.3	2019-09-22	[1]	CRAN	(R 4.0.2)
purrr	* 0.3.4	2020-04-17	[1]	CRAN	(R 4.0.2)
R6	2.5.0	2020-10-28	[1]	CRAN	(R 4.0.2)
Rcpp	1.0.7	2021-07-07	[1]	CRAN	(R 4.0.3)
readr	* 1.4.0	2020-10-05	[1]	CRAN	(R 4.0.2)
readxl	1.3.1	2019-03-13	[1]	CRAN	(R 4.0.2)
repr	1.1.3	2021-01-21	[1]	CRAN	(R 4.0.2)
reprex	2.0.0	2021-04-02	[1]	CRAN	(R 4.0.3)
rio	0.5.27	2021-06-21	[1]	CRAN	(R 4.0.3)
rlang	0.4.11	2021-04-30	[1]	CRAN	(R 4.0.3)

rstatix	0.7.0	2021-02-13	[1]	CRAN	(R 4.0.3)
rstudioapi	0.13	2020-11-12	[1]	CRAN	(R 4.0.2)
rvest	1.0.0	2021-03-09	[1]	CRAN	(R 4.0.3)
scales	1.1.1	2020-05-11	[1]	CRAN	(R 4.0.2)
sessioninfo	1.1.1	2018-11-05	[1]	CRAN	(R 4.0.2)
stringi	1.7.3	2021-07-16	[1]	CRAN	(R 4.0.3)
stringr	* 1.4.0	2019-02-10	[1]	CRAN	(R 4.0.2)
svglite	2.0.0	2021-02-20	[1]	CRAN	(R 4.0.3)
systemfonts	1.0.2	2021-05-11	[1]	CRAN	(R 4.0.3)
tibble	* 3.1.2	2021-05-16	[1]	CRAN	(R 4.0.3)
tidyr	* 1.1.3	2021-03-03	[1]	CRAN	(R 4.0.3)
tidyselect	1.1.1	2021-04-30	[1]	CRAN	(R 4.0.3)
tidyverse	* 1.3.1	2021-04-15	[1]	CRAN	(R 4.0.3)
utf8	1.2.1	2021-03-12	[1]	CRAN	(R 4.0.3)
uuid	0.1-4	2020-02-26	[1]	CRAN	(R 4.0.2)
vctrs	0.3.8	2021-04-29	[1]	CRAN	(R 4.0.3)
withr	2.4.2	2021-04-18	[1]	CRAN	(R 4.0.3)
xml2	1.3.2	2020-04-23	[1]	CRAN	(R 4.0.2)
zip	2.2.0	2021-05-31	[1]	CRAN	(R 4.0.3)

[1] /home/jbenja13/R/x86_64-pc-linux-gnu-library/4.0

[2] /usr/lib/R/library