

# main

August 1, 2021

## 1 Check sex chromosomes for caudate nucleus

```
[1]: suppressMessages({library(dplyr)
                        library(ggpubr)})
```

### 1.1 Get and annotated sex from genotypes

```
[2]: sexcheck_file = paste0('/ceph/projects/brainseq/genotype/download/topmed/',
                             'convert2plink/filter_maf_01/check_sex/_m/
                             ↳LIBD_Brain_TopMed.sexcheck')
pheno_file = '/ceph/users/jbenja13/projects/sex_sz_ria/input/phenotypes/_m/
↳phenotypes.csv'

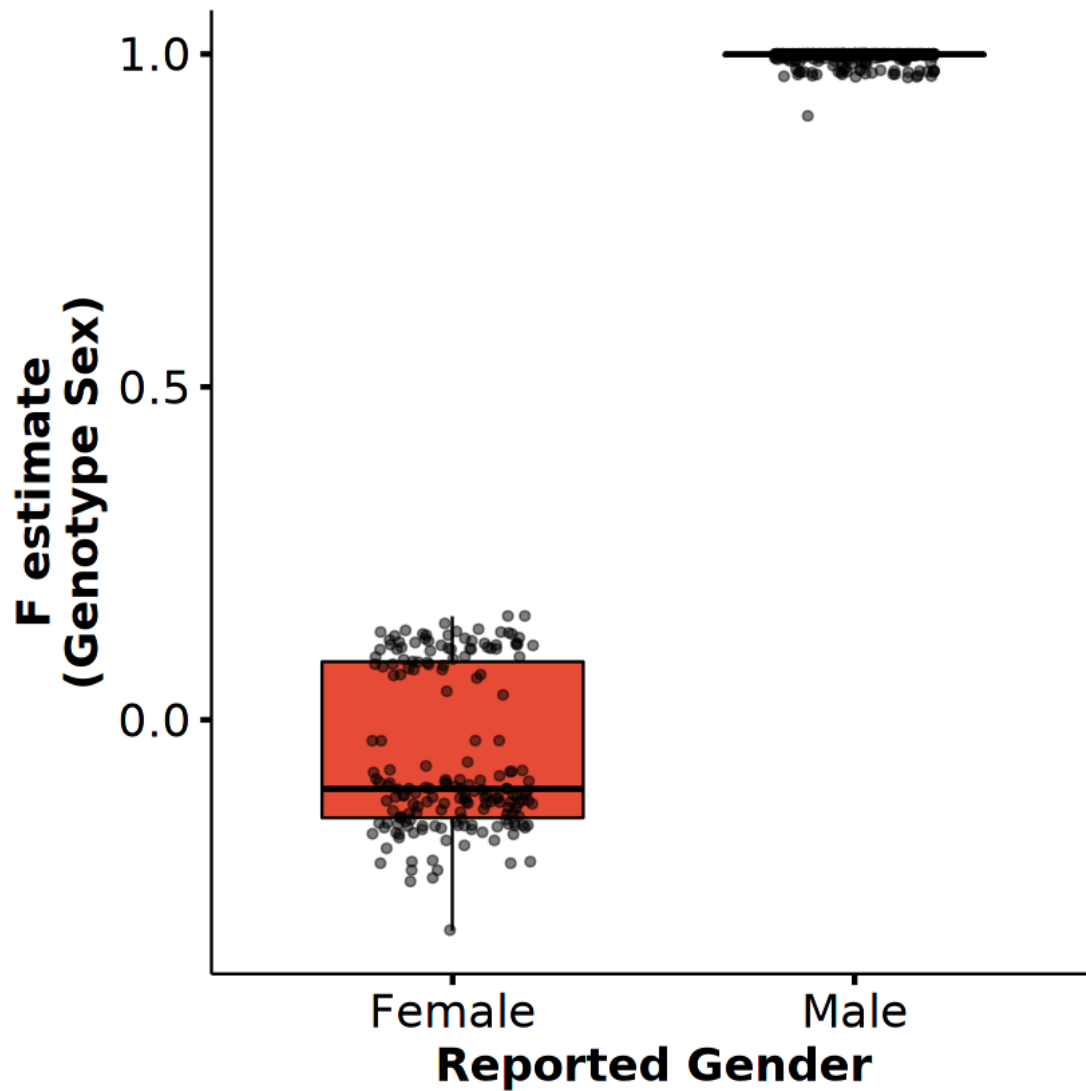
dt = data.table::fread(sexcheck_file) %>%
  inner_join(data.table::fread(pheno_file), by=c("FID"="BrNum")) %>%
  select(c("FID", "Dx", "Sex", "Age", "F", "SNPSEX")) %>%
  distinct() %>% mutate_if(is.character, as.factor)
levels(dt$Sex) <- c('Female', 'Male')
dt %>% head(2)
```

A data.table: 2 × 6

	FID <fct>	Dx <fct>	Sex <fct>	Age <dbl>	F <dbl>	SNPSEX <int>
	Br763	Schizo	Male	75.22	1	1
	Br822	Control	Male	26.73	1	1

### 1.2 Plot F estimate

```
[3]: fig <- ggboxplot(dt, x = "Sex", y = "F", fill='Sex', width=0.65, palette='npg',
                      xlab='Reported Gender', ylab='F estimate\n(Genotype Sex)',
                      ggtheme=theme_pubr(base_size=20), add="jitter",
                      add.params=list(alpha=0.5), legend="None") +
  font("xy.title", face='bold')
fig
```



```
[4]: ggsave("sex_check.png")  
ggsave("sex_check.pdf")  
ggsave("sex_check.svg")
```

Saving 6.67 x 6.67 in image

Saving 6.67 x 6.67 in image

Saving 6.67 x 6.67 in image

```
[5]: dt %>%  
group_by(Sex) %>%
```

```
summarize(mean_size = mean(F, na.rm = TRUE), max = max(F, na.rm=TRUE))
```

```

      Sex      mean_size      max
  <fct>    <dbl>         <dbl>
1 Female -0.05355674    0.156
2 Male   0.99623137    1.000

```

A tibble: 2 × 3

### 1.3 Check that reported gender matches genotype sex

```
[6]: sum(dt$Sex == "Male")
      sum(dt$SNPSEX == 1 & dt$Sex == "Male")
```

408

408

```
[7]: sum(dt$Sex == "Female")
      sum(dt$SNPSEX == 2 & dt$Sex == "Female")
```

187

187

#### 1.3.1 Examine number of X chromosomes

```
[8]: table(dt$SNPSEX)
```

```

  1  2
408 187

```

```
[9]: table(dt$Sex)
```

```

Female  Male
  187    408

```

Summary: All samples have reported gender matching genotype sex

### 1.4 Reproducibility Information

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

```
[1] "2021-08-01 15:56:07 EDT"
```

```

      user system elapsed
3.079   0.137   3.472

```

```

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     2021-08-01

```

```

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)
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)
fans          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)
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)
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)
magrittr	2.0.1	2020-11-17	[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)
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)
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)
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)
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)
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)
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