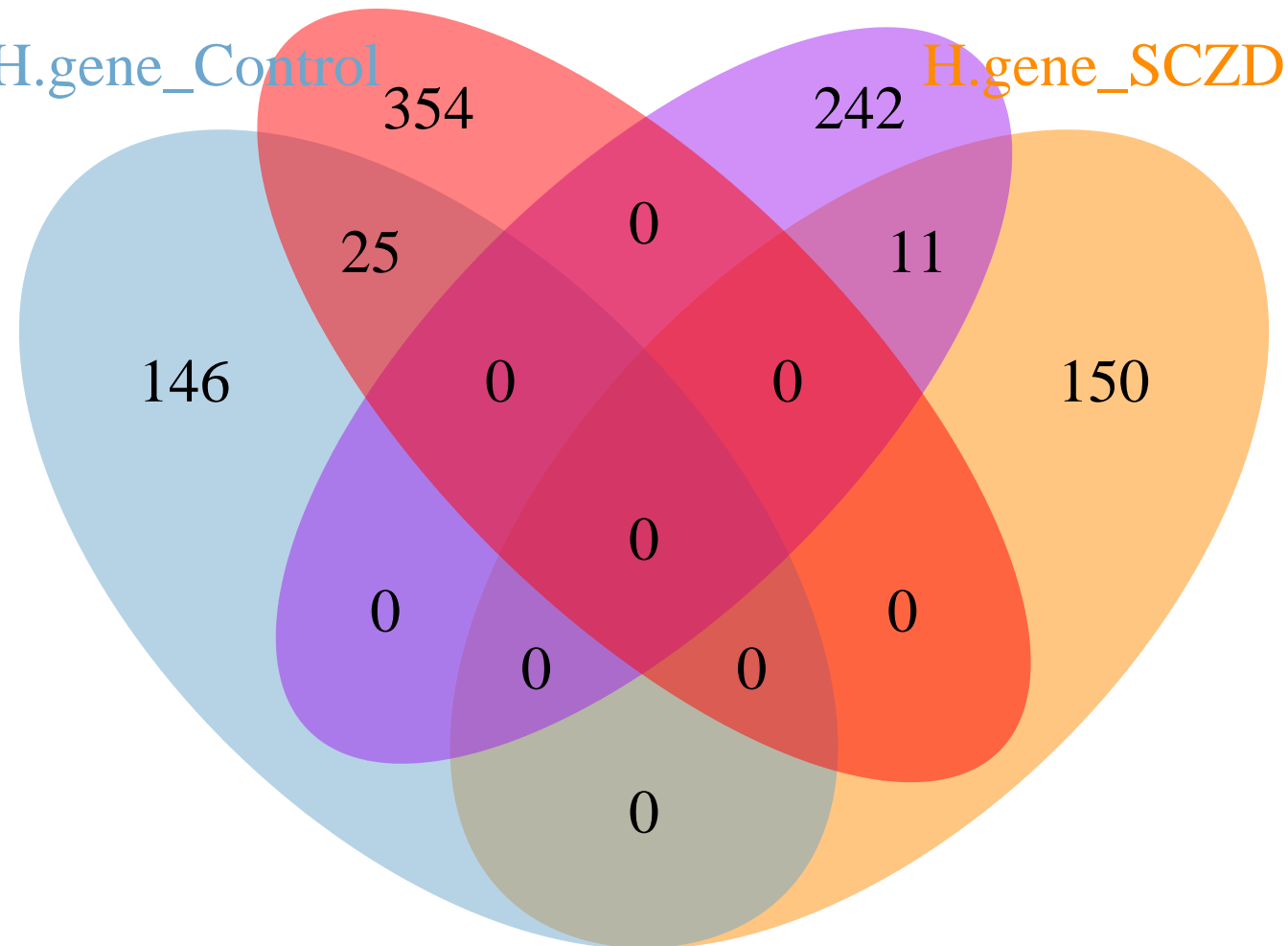


Gene DLPFC FDR10%, HIPPO FDR20%

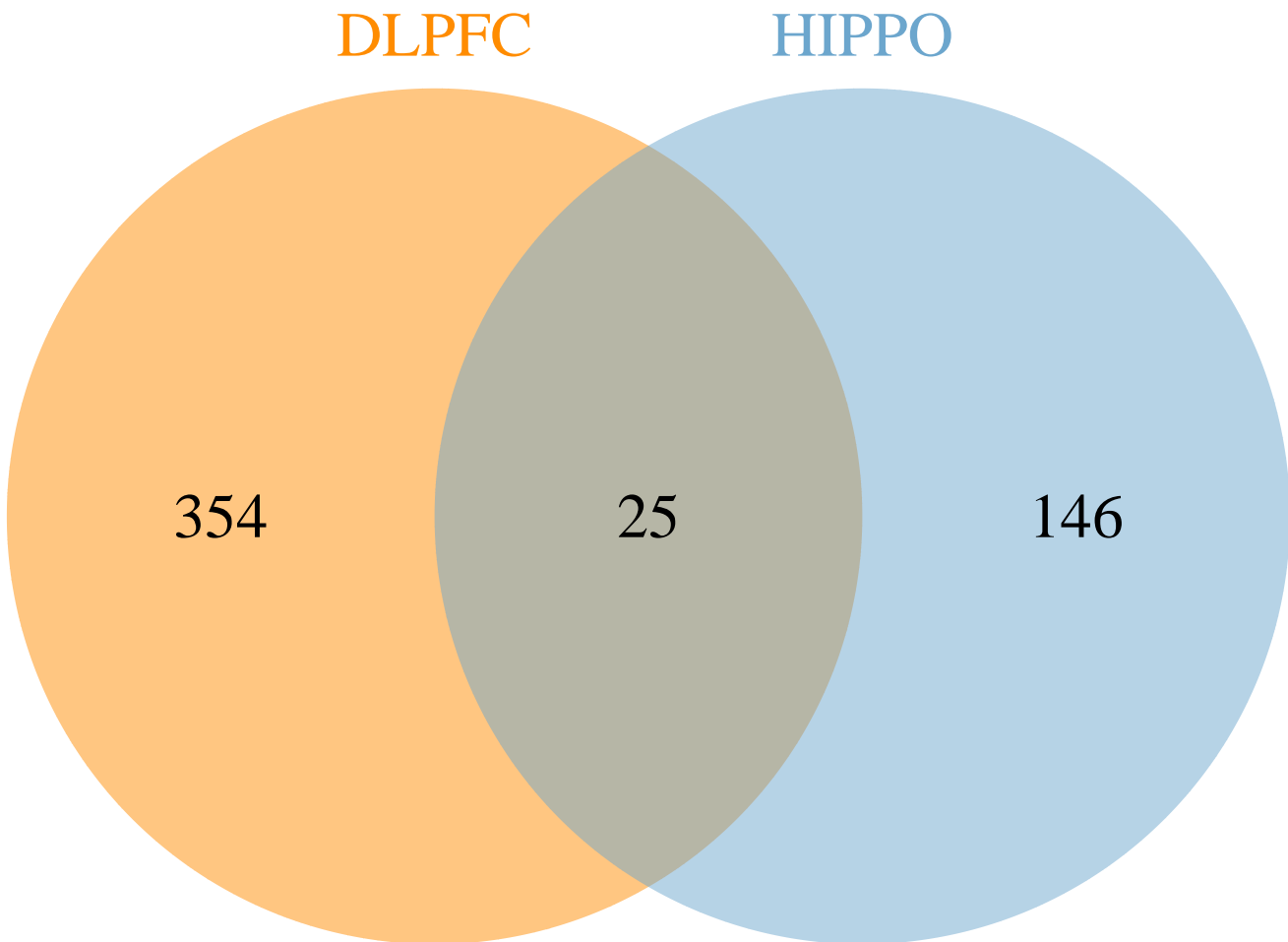
D.gene_Control D.gene_SCZD

H.gene_Control

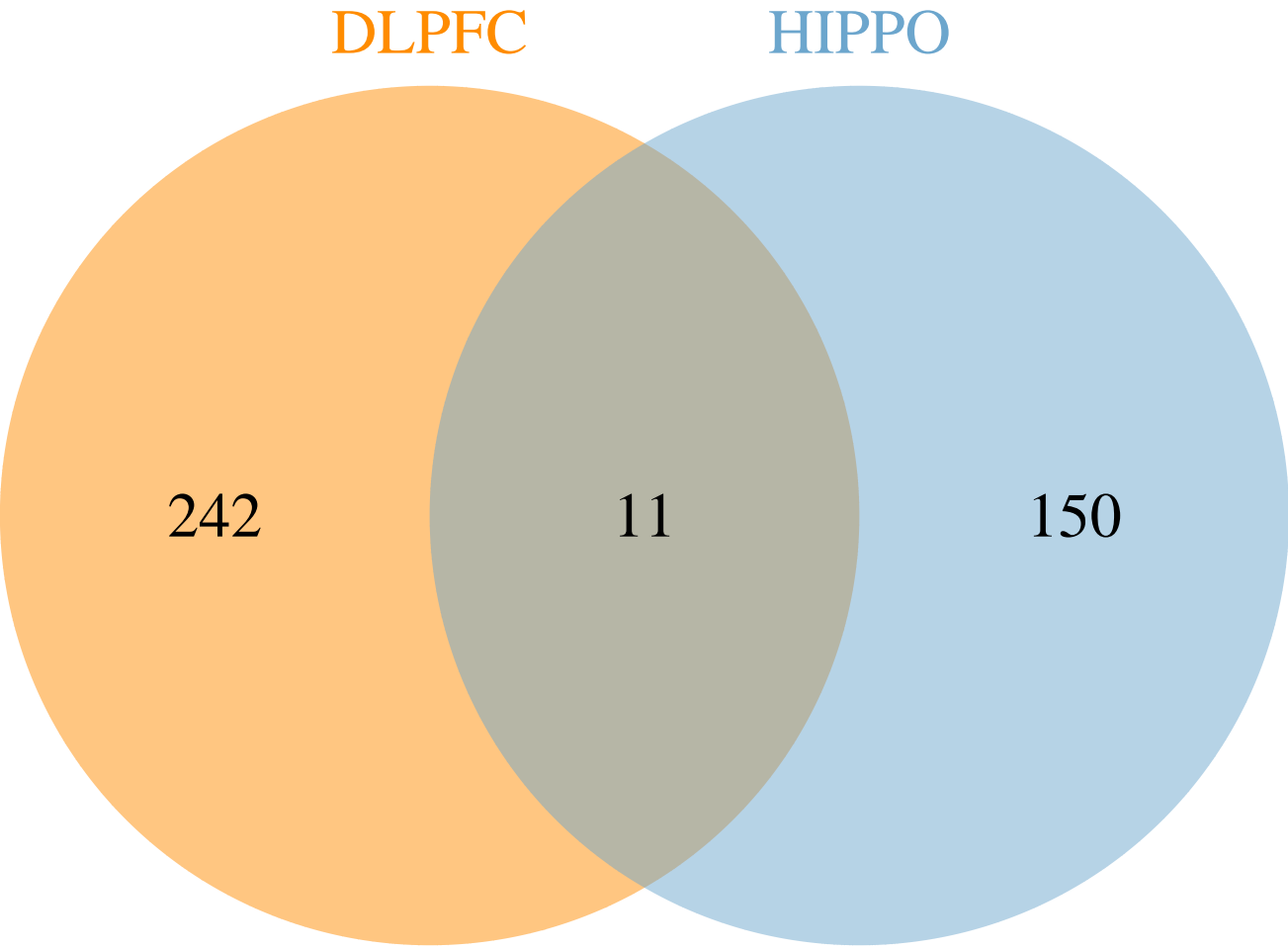
H.gene_SCZD



Gene Control (DLPFC FDR10%, HIPPO FDR20%)



Gene SCZD (DLPFC FDR10%, HIPPO FDR20%)

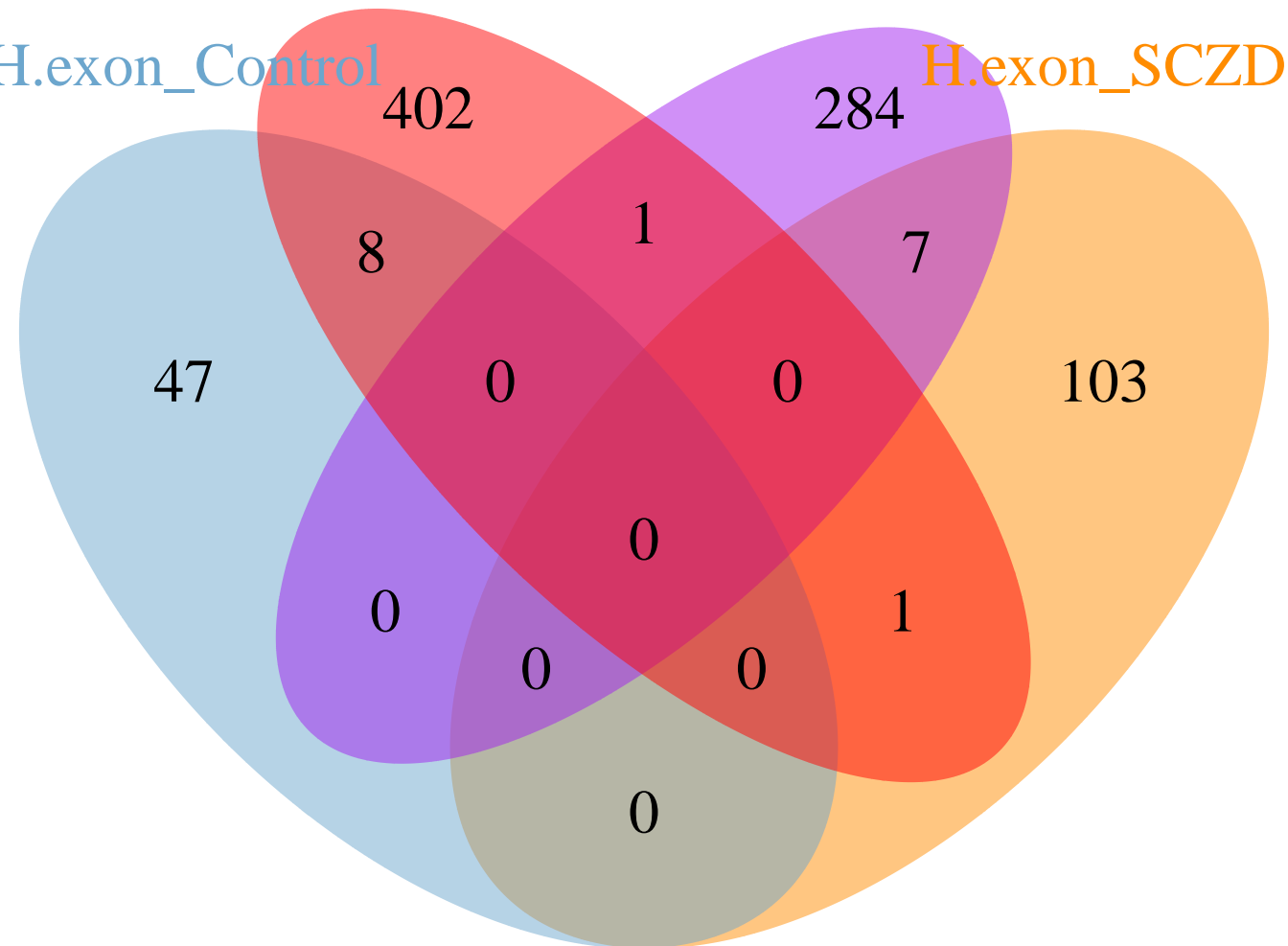


Exon DLPFC FDR10%, HIPPO FDR20%

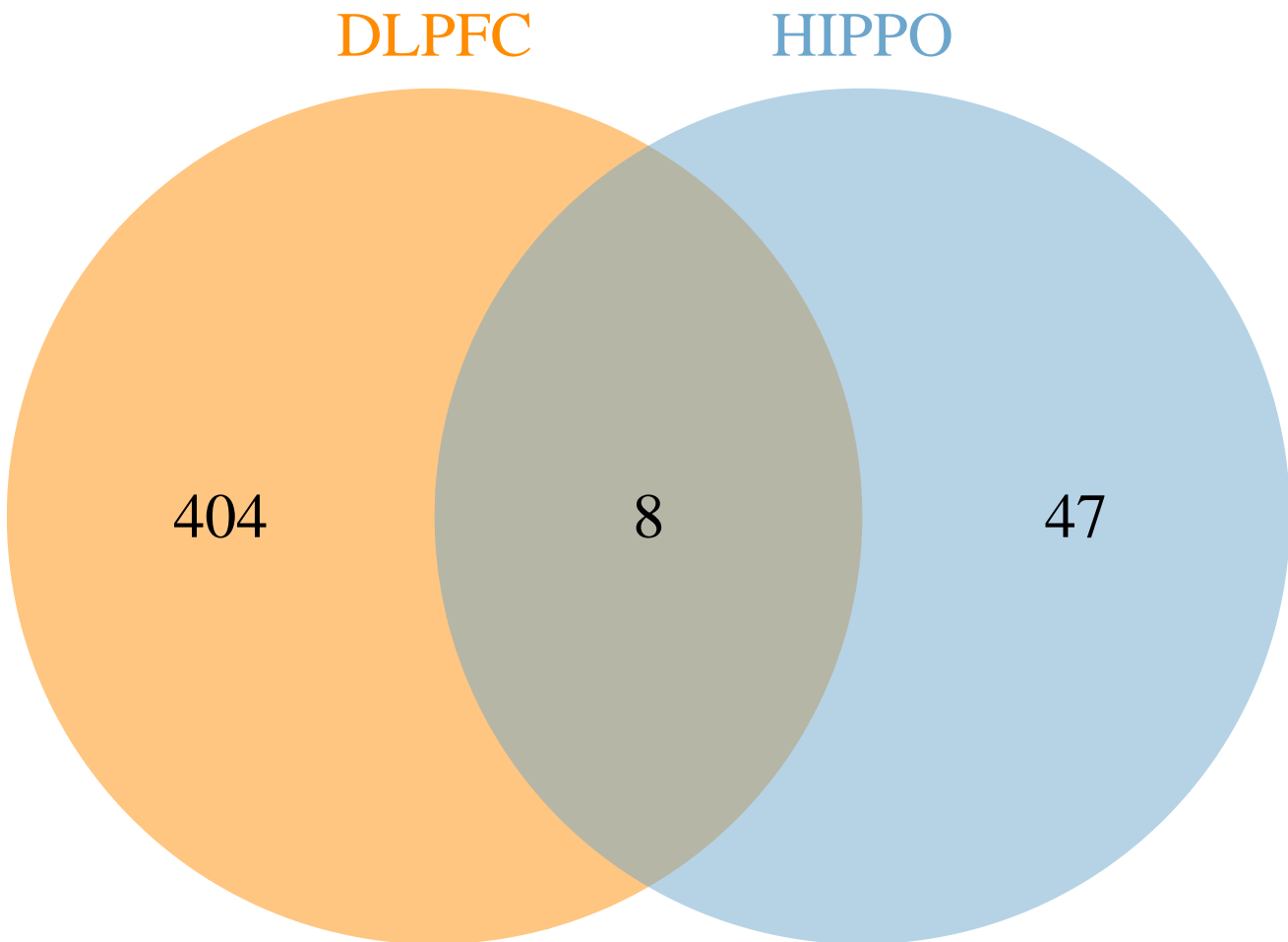
D.exon_Control D.exon_SCZD

H.exon_Control

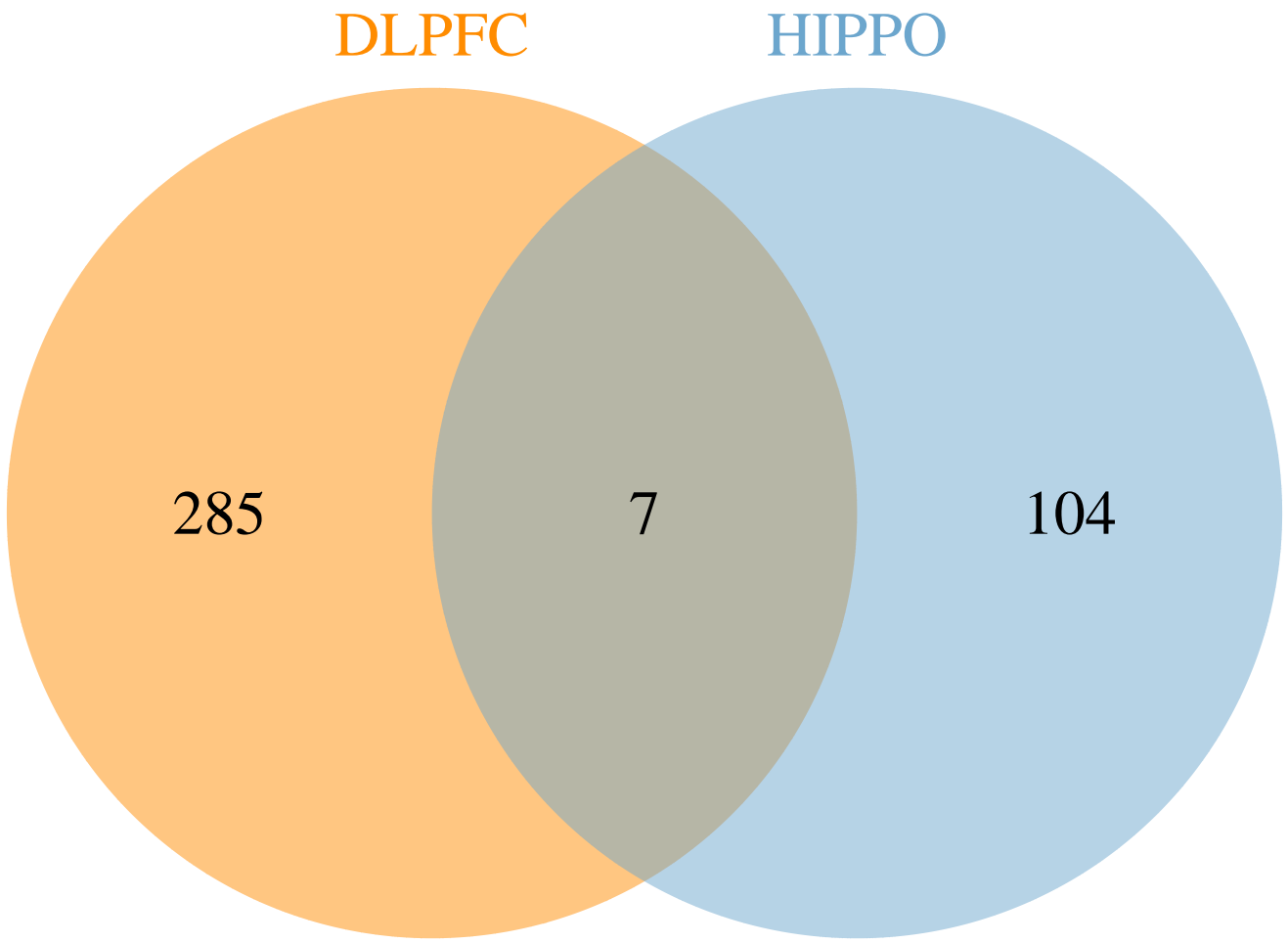
H.exon_SCZD



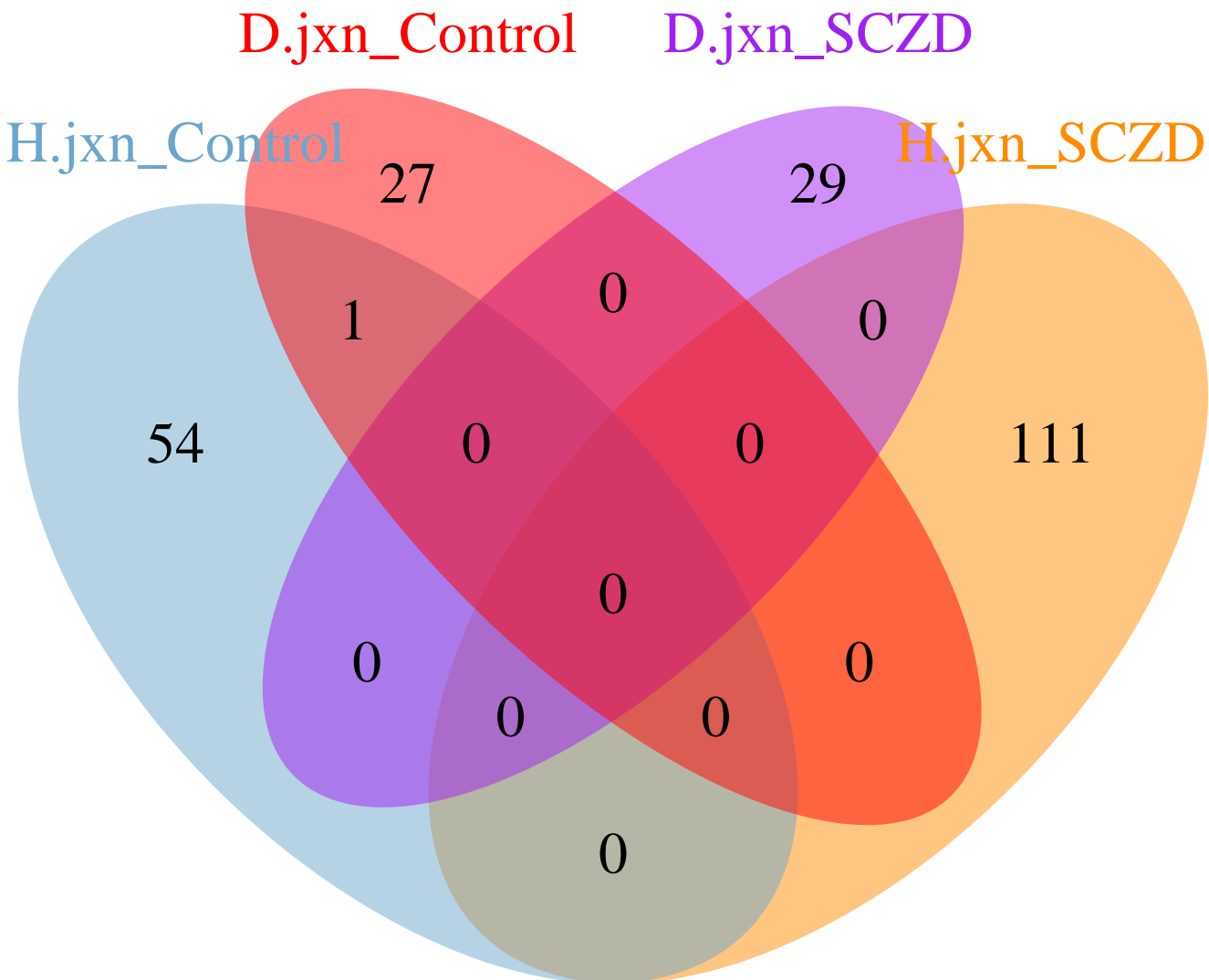
Exon Control (DLPFC FDR10%, HIPPO FDR20%)



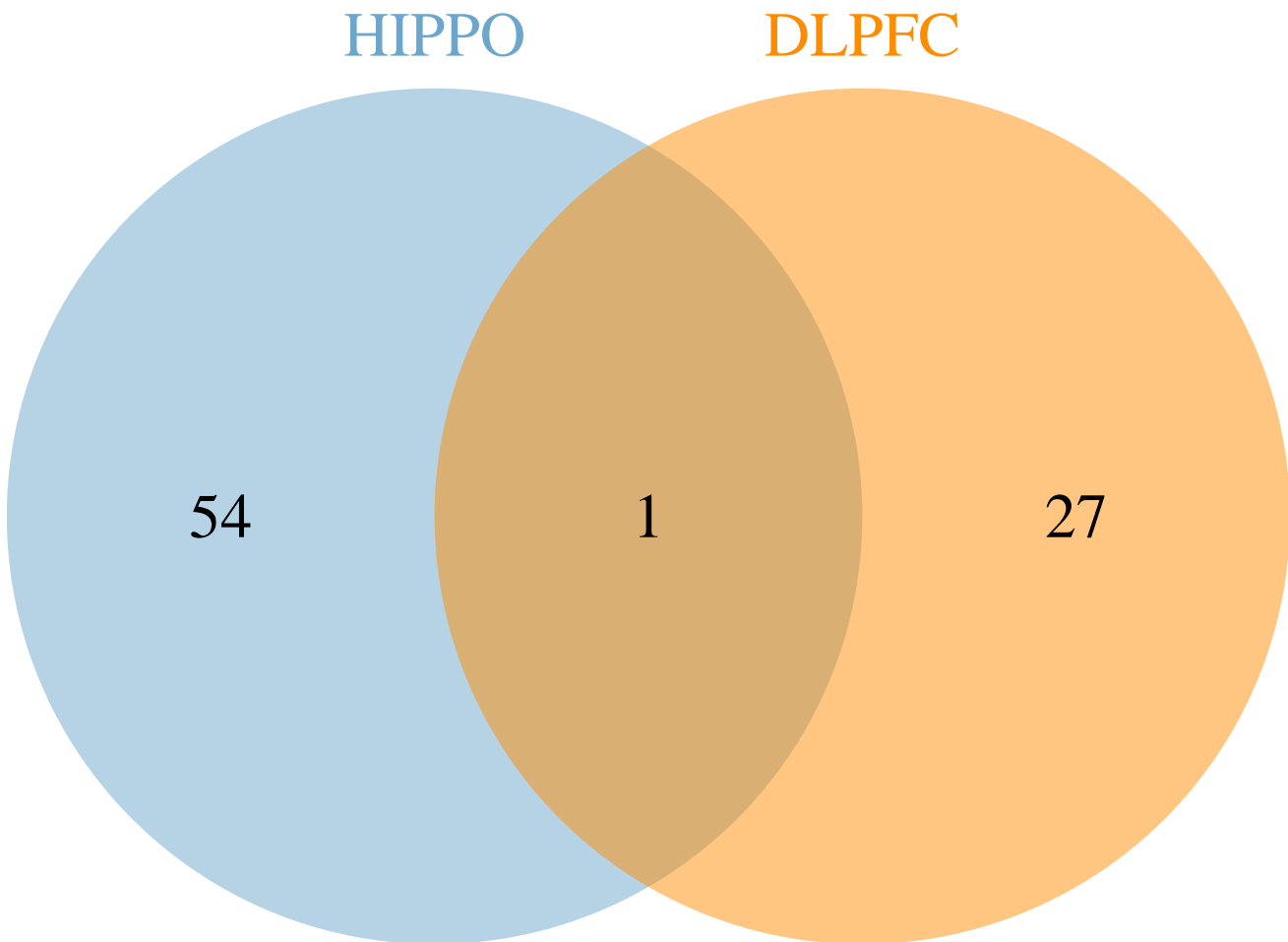
Exon SCZD (DLPFC FDR10%, HIPPO FDR20%)



JXN DLPFC FDR10%, HIPPO FDR20%



JXN Control (DLPFC FDR10%, HIPPO FDR20%)



JXN SCZD (DLPFC FDR10%, HIPPO FDR20%)

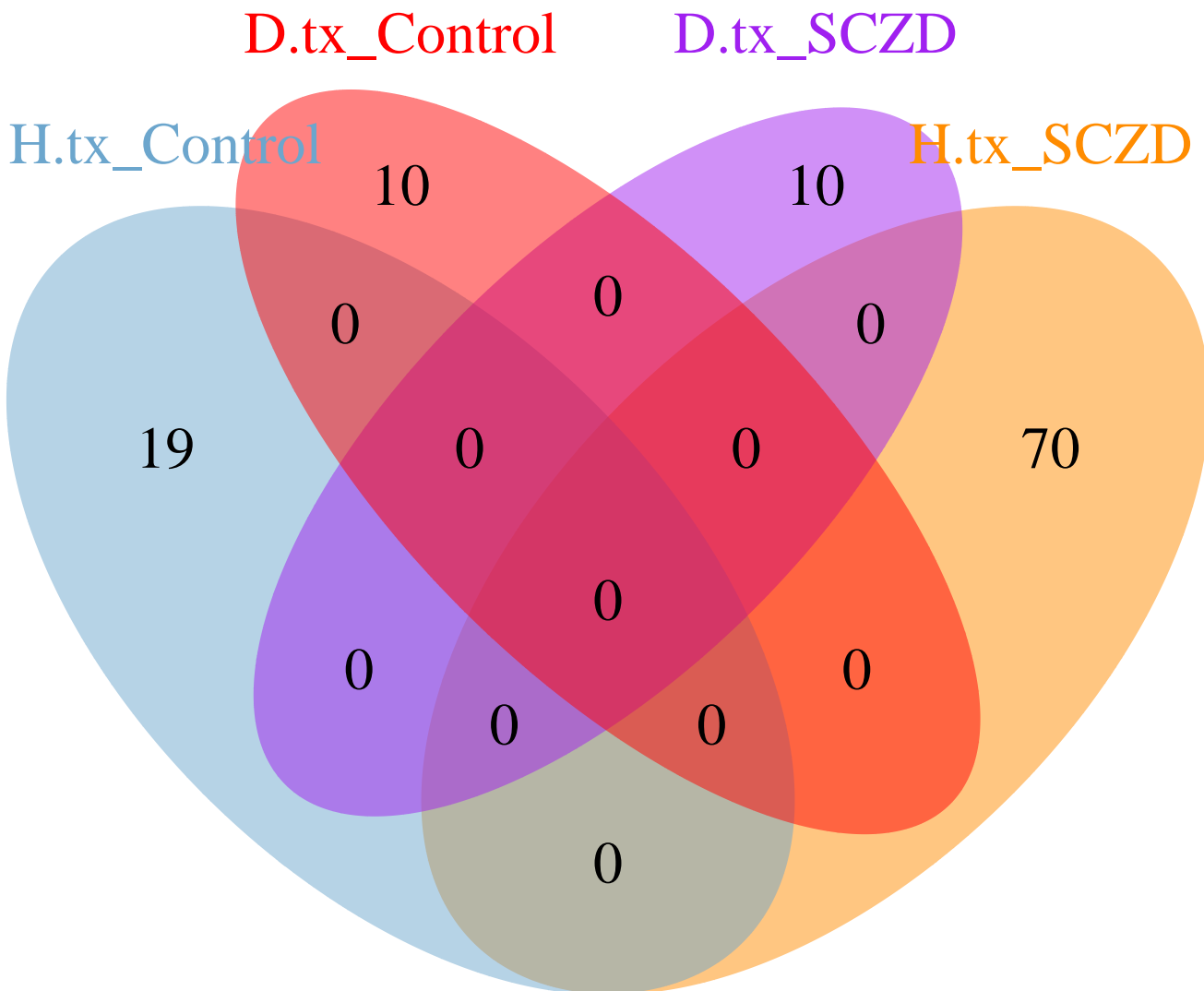
HIPPO

111

DLPFC

29

TX DLPFC FDR10%, HIPPO FDR20%



TX Control (DLPFC FDR10%, HIPPO FDR20%)

HIPPO

19

DLPFC

10

TX SCZD (DLPFC FDR10%, HIPPO FDR20%)

HIPPO

70

DLPFC

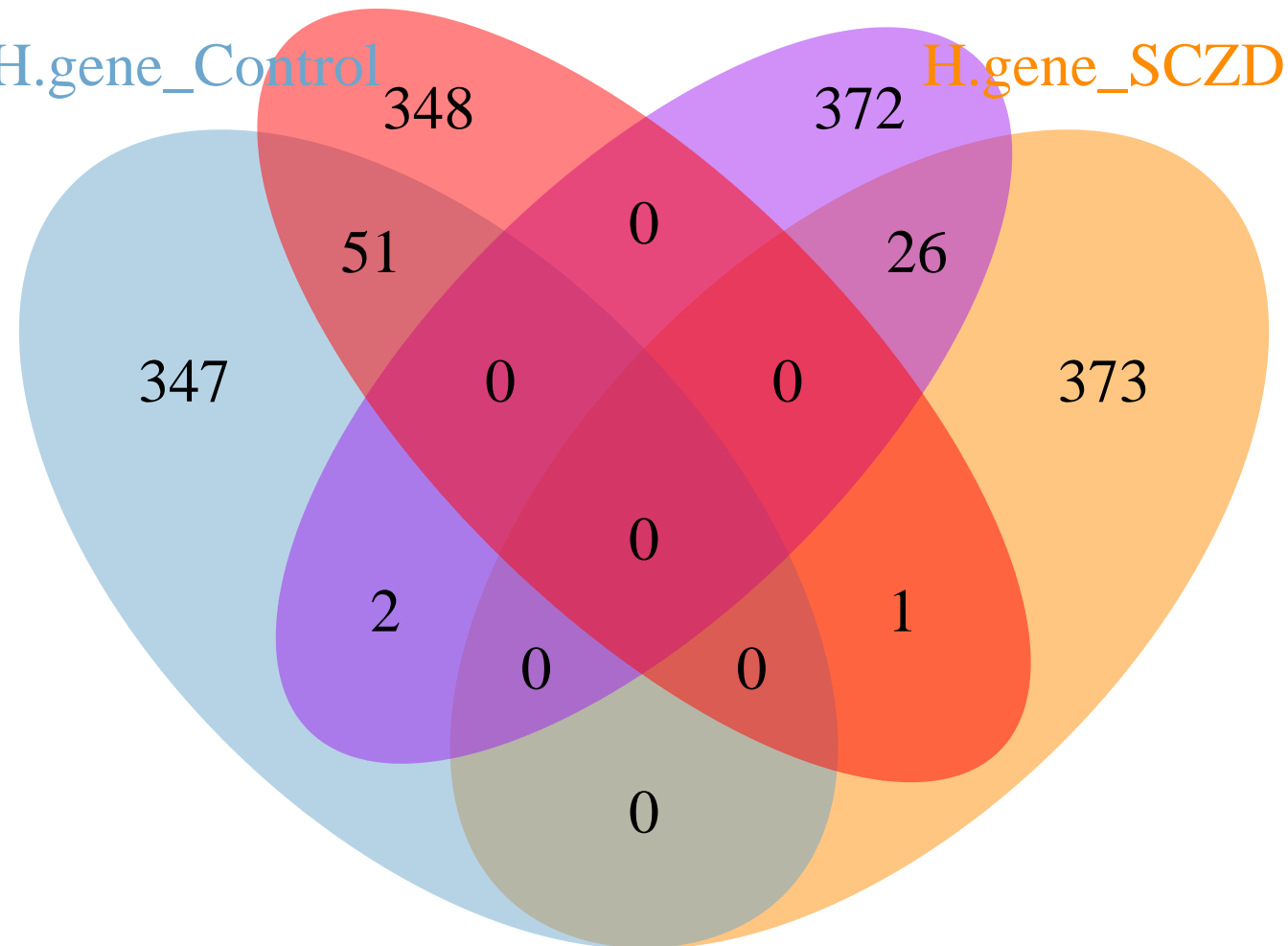
10

gene: top 400 in each group

D.gene_Control D.gene_SCZD

H.gene_Control

H.gene_SCZD



gene: Control (top 400)

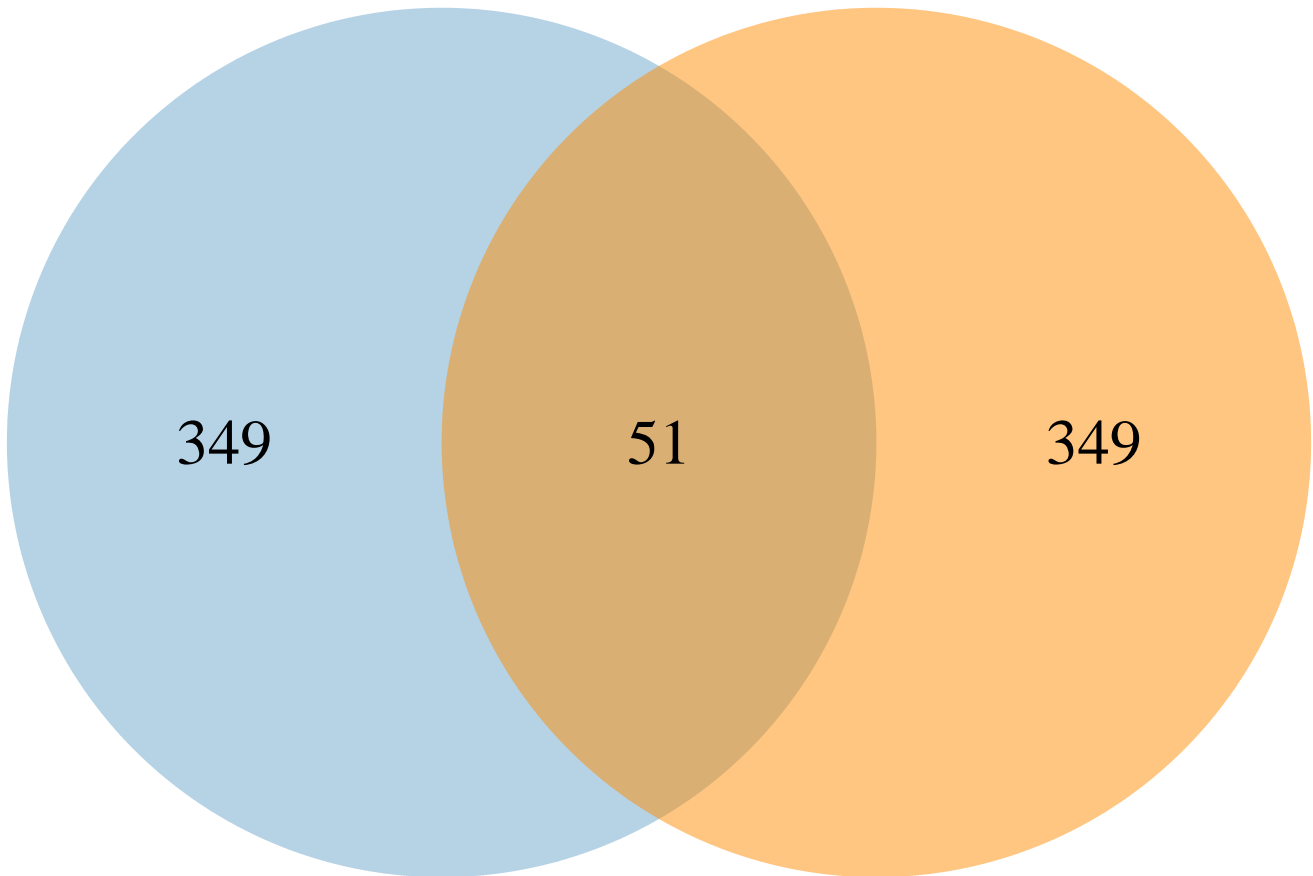
HIPPO

DLPFC

349

51

349



gene: SCZD (top 400)

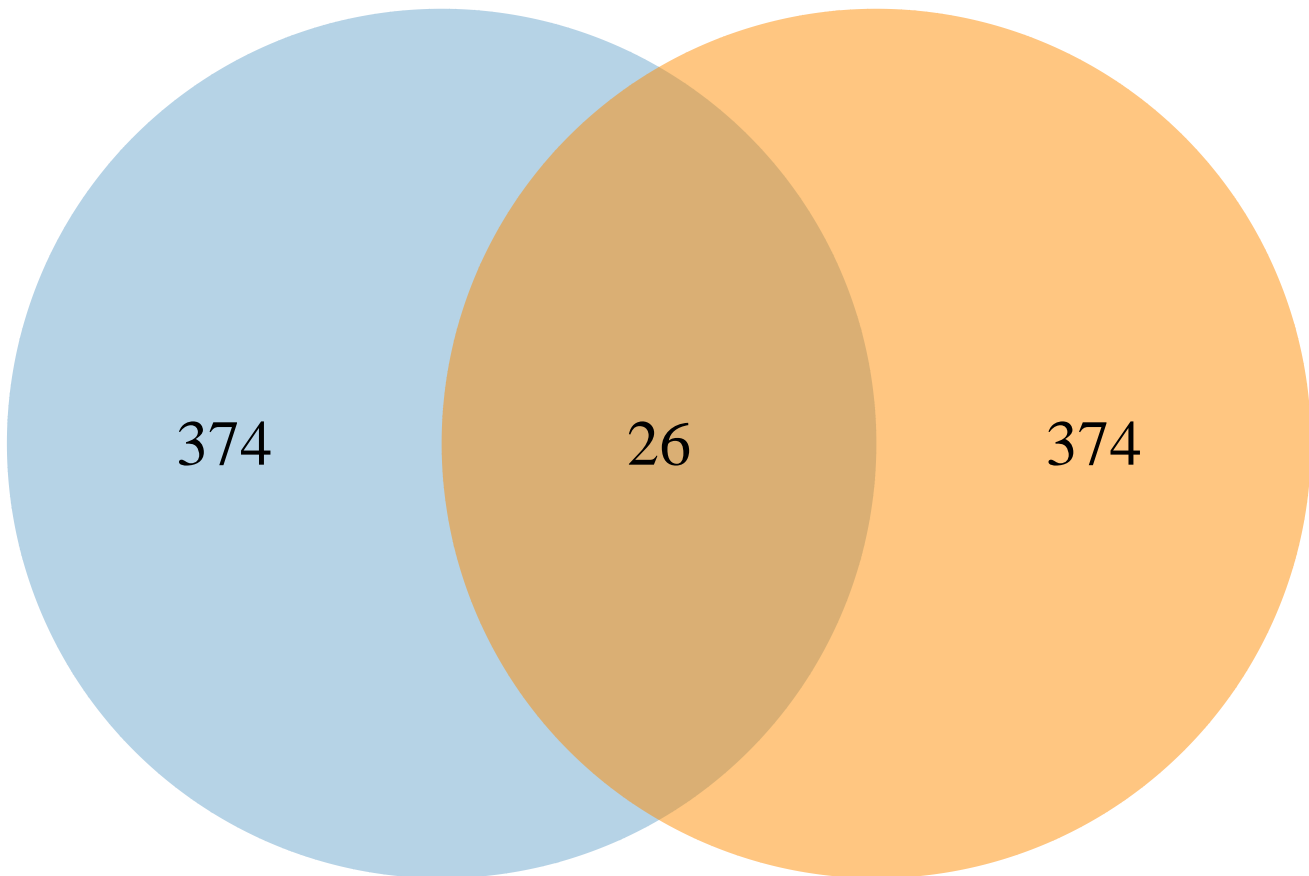
HIPPO

DLPFC

374

26

374

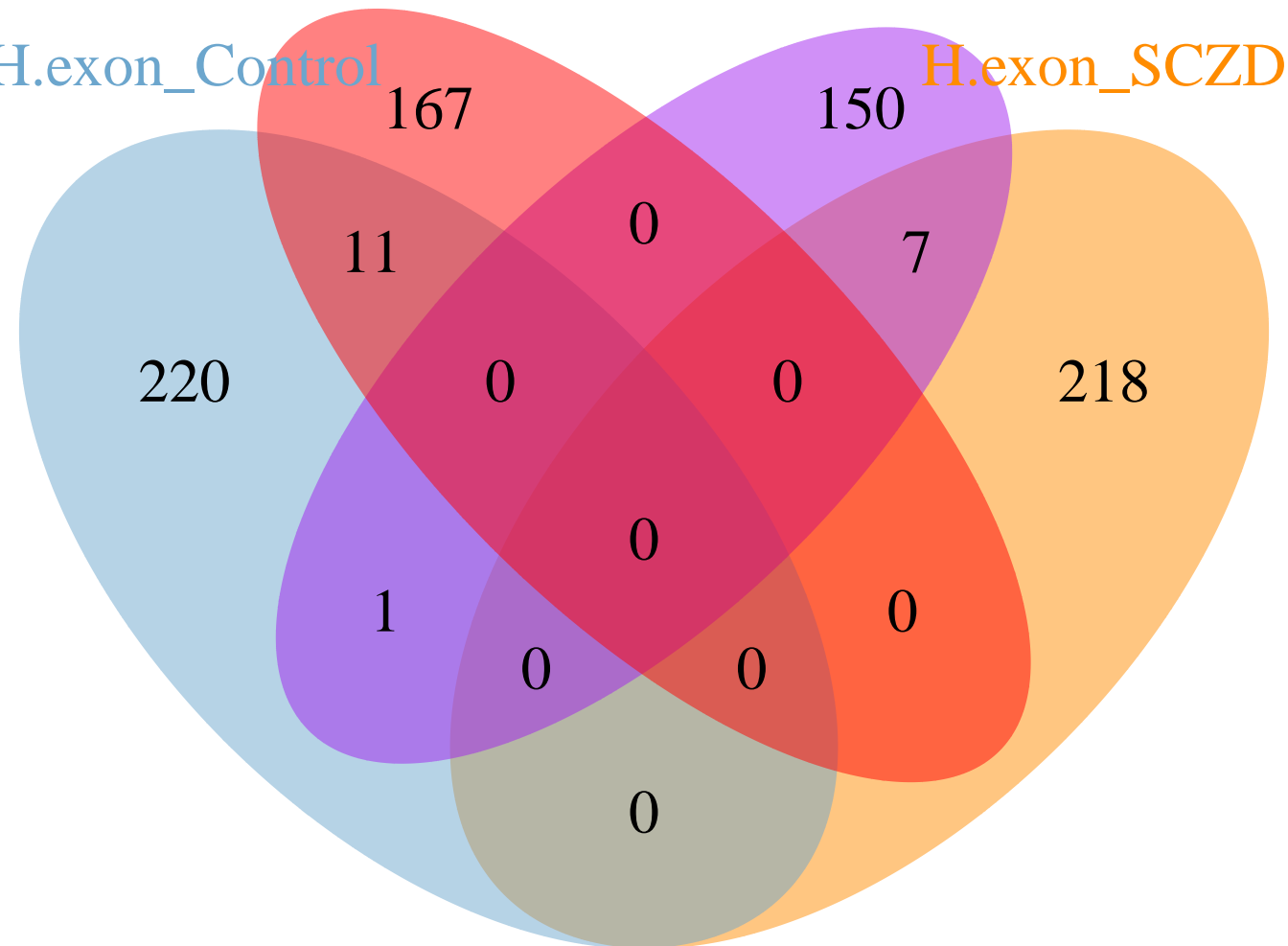


exon: top 400 in each group

D.exon_Control D.exon_SCZD

H.exon_Control

H.exon_SCZD



exon: Control (top 400)

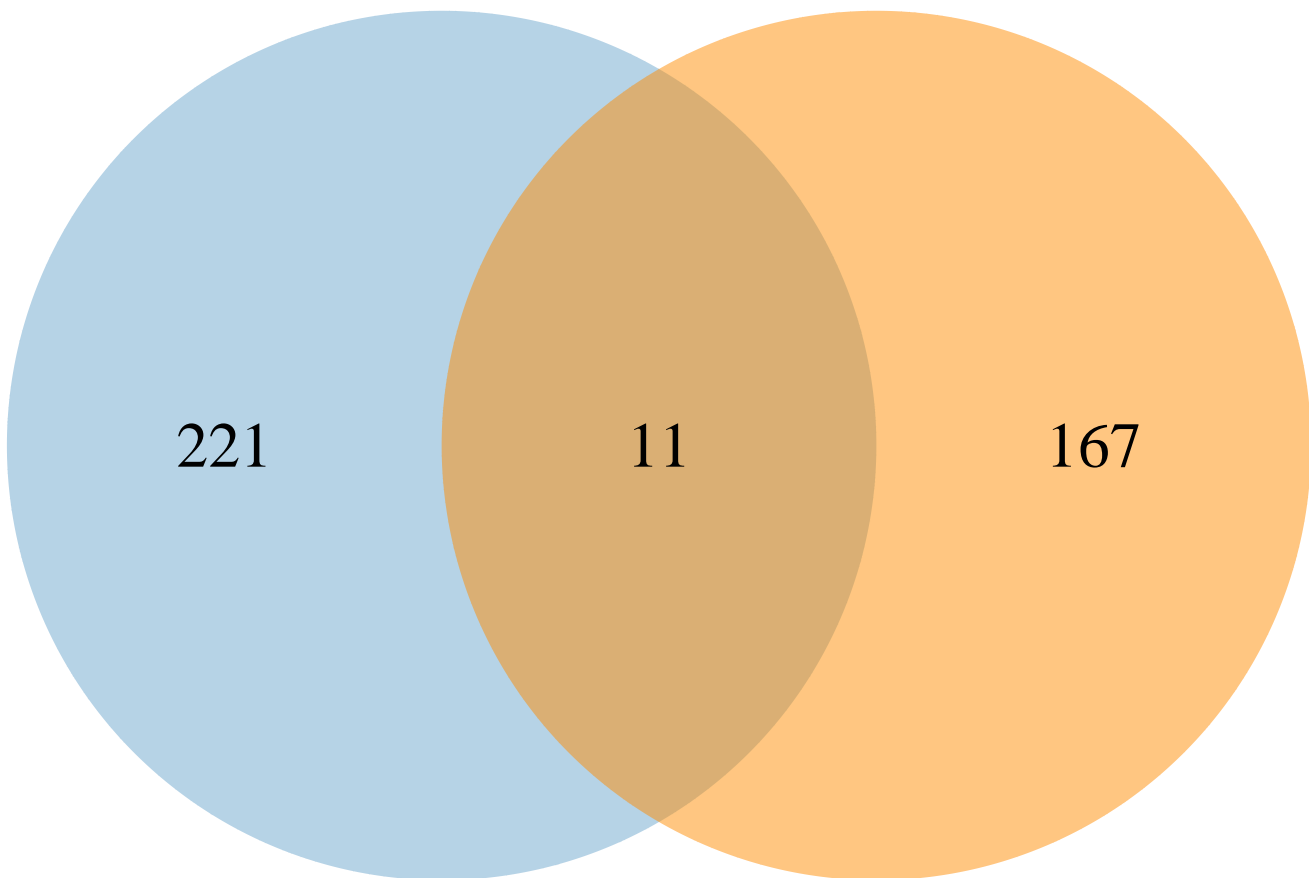
HIPPO

DLPFC

221

11

167



exon: SCZD (top 400)

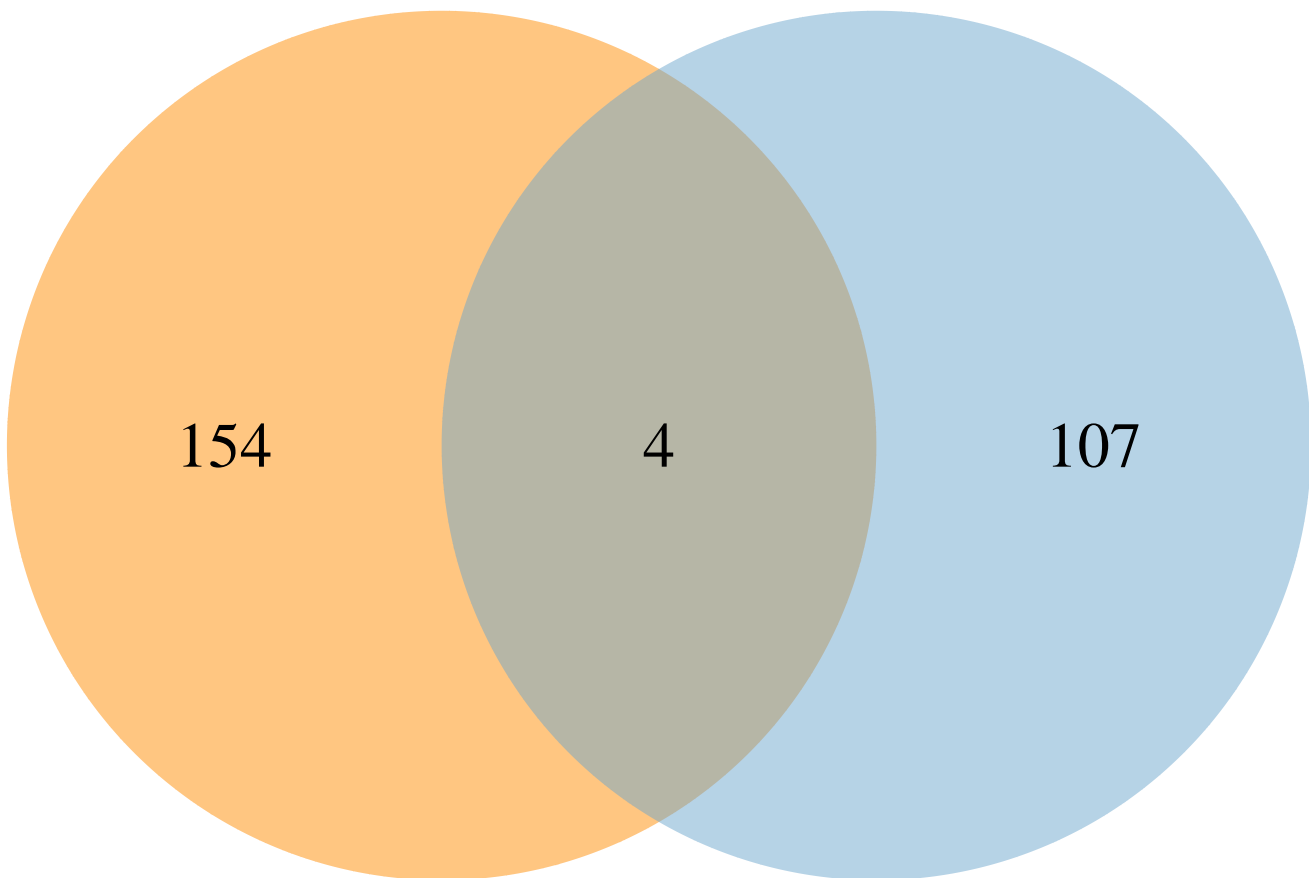
DLPFC

HIPPO

154

4

107



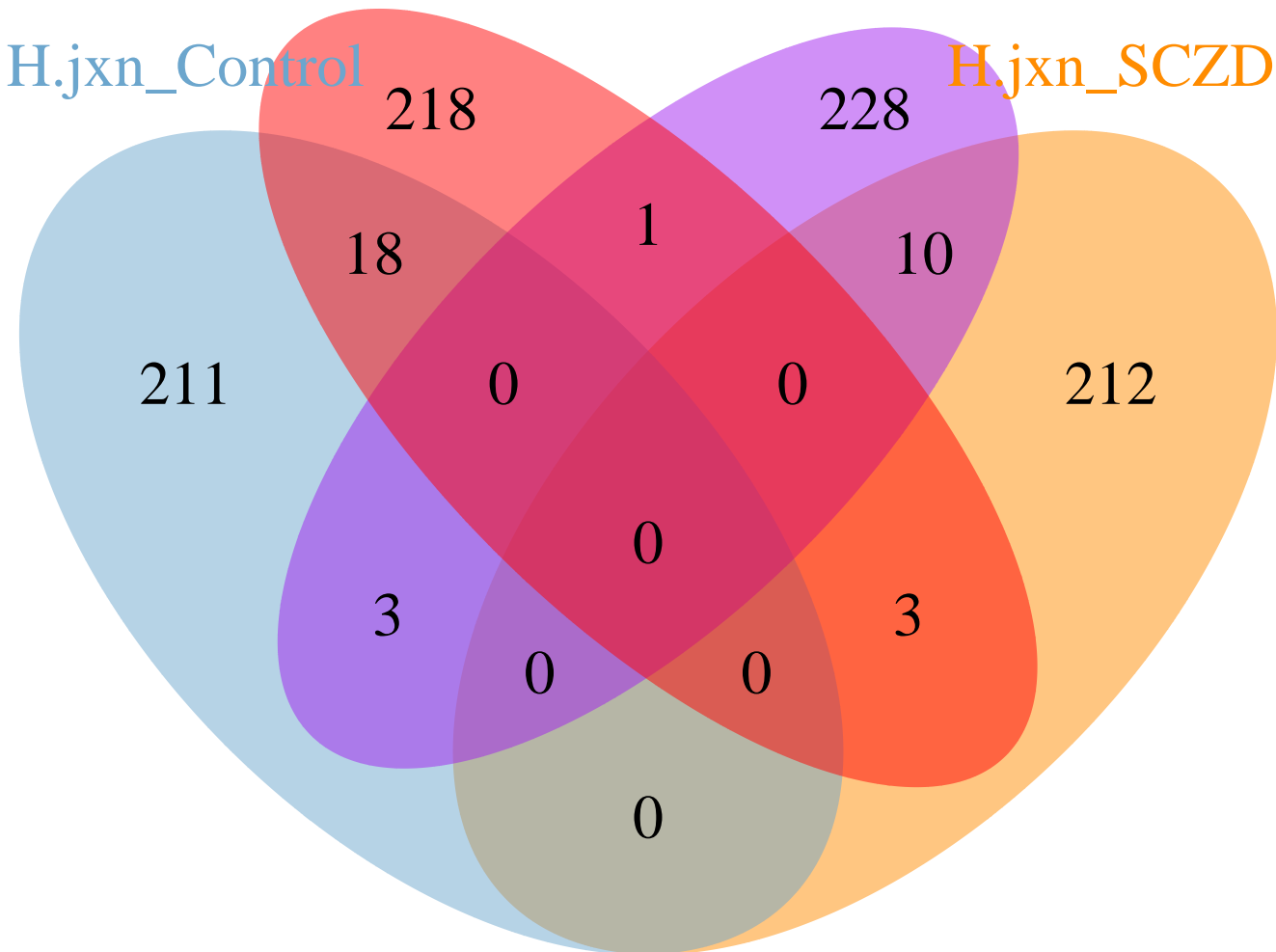
jxn: top 400 in each group

D.jxn_Control

D.jxn_SCZD

H.jxn_Control

H.jxn_SCZD



jxn: Control (top 400)

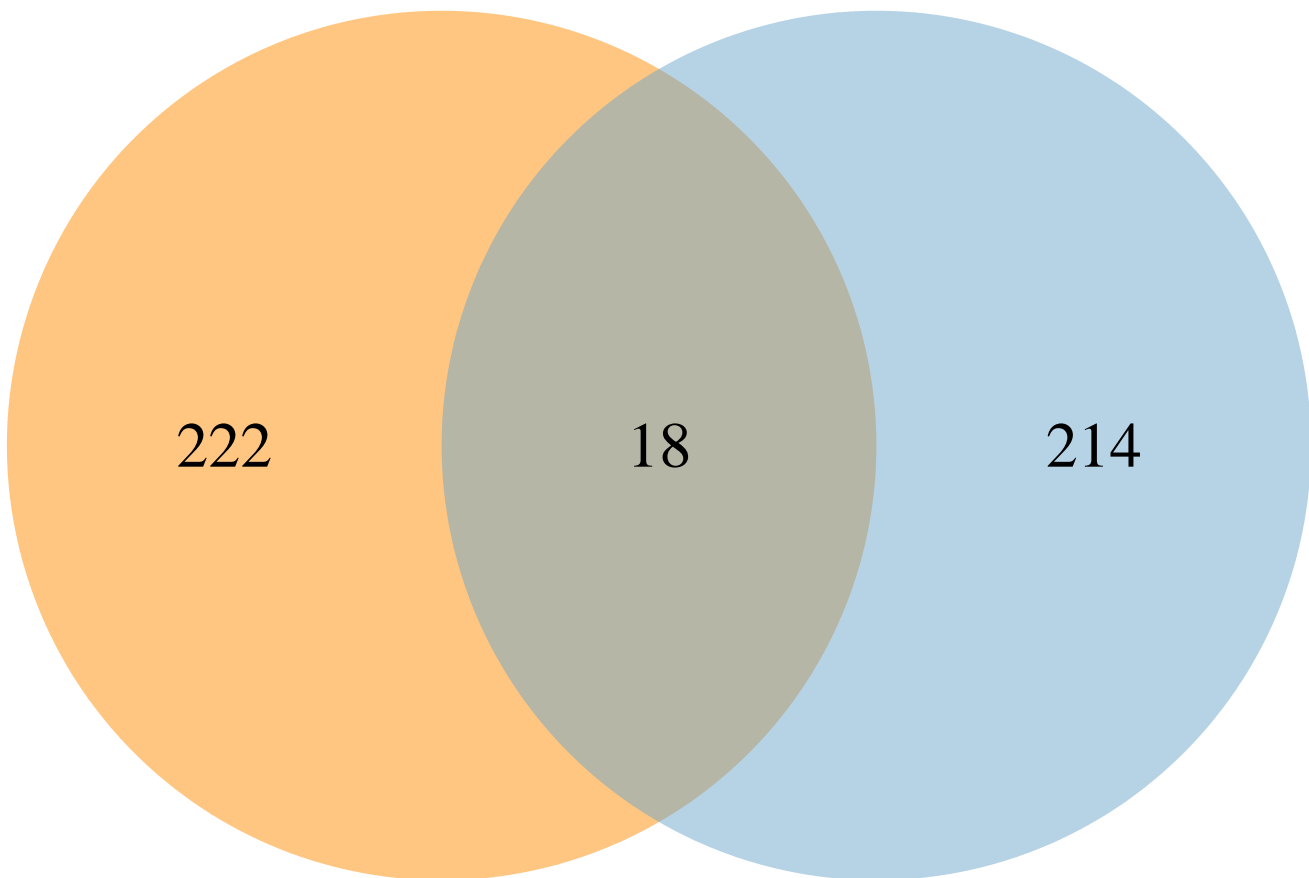
DLPFC

HIPPO

222

18

214



jxn: SCZD (top 400)

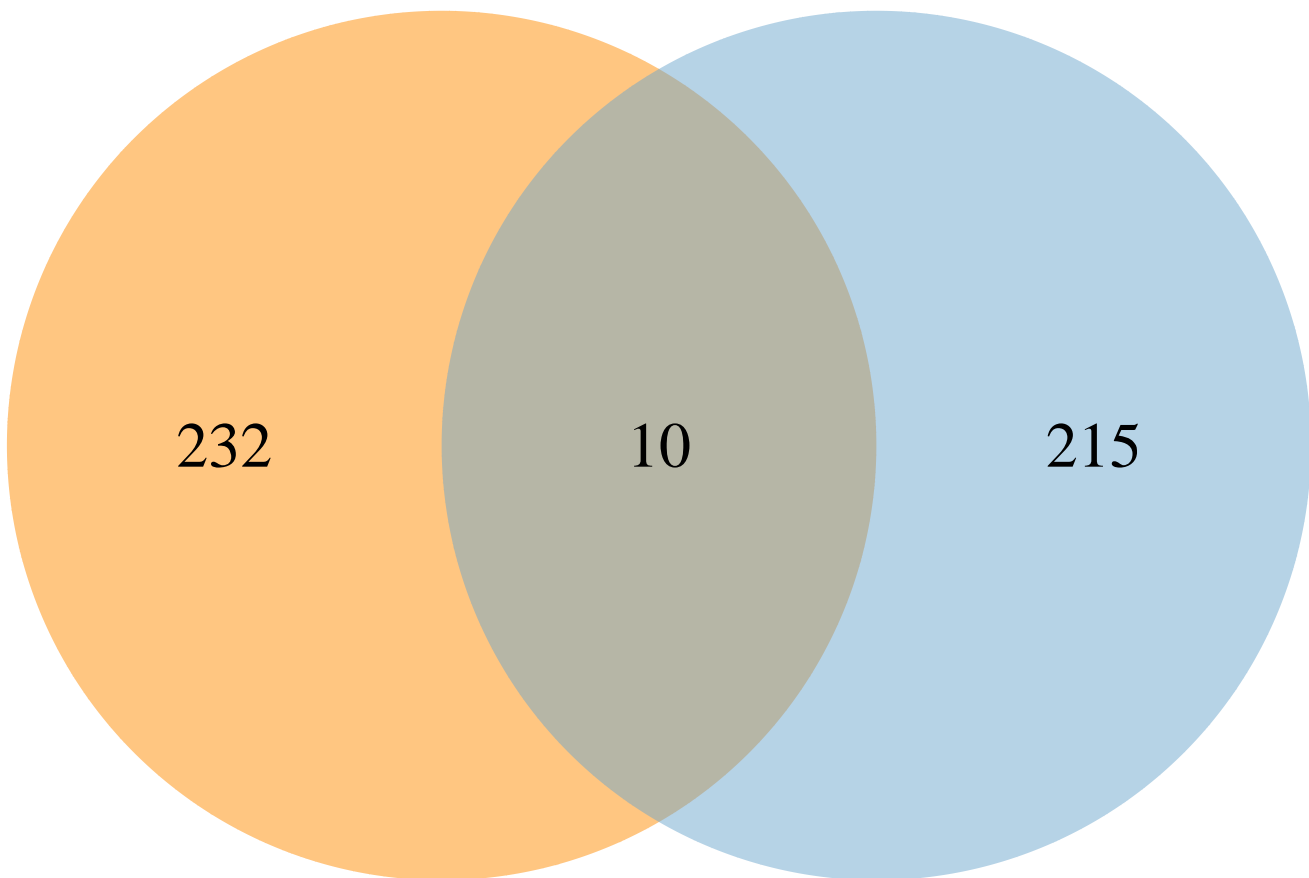
DLPFC

HIPPO

232

10

215



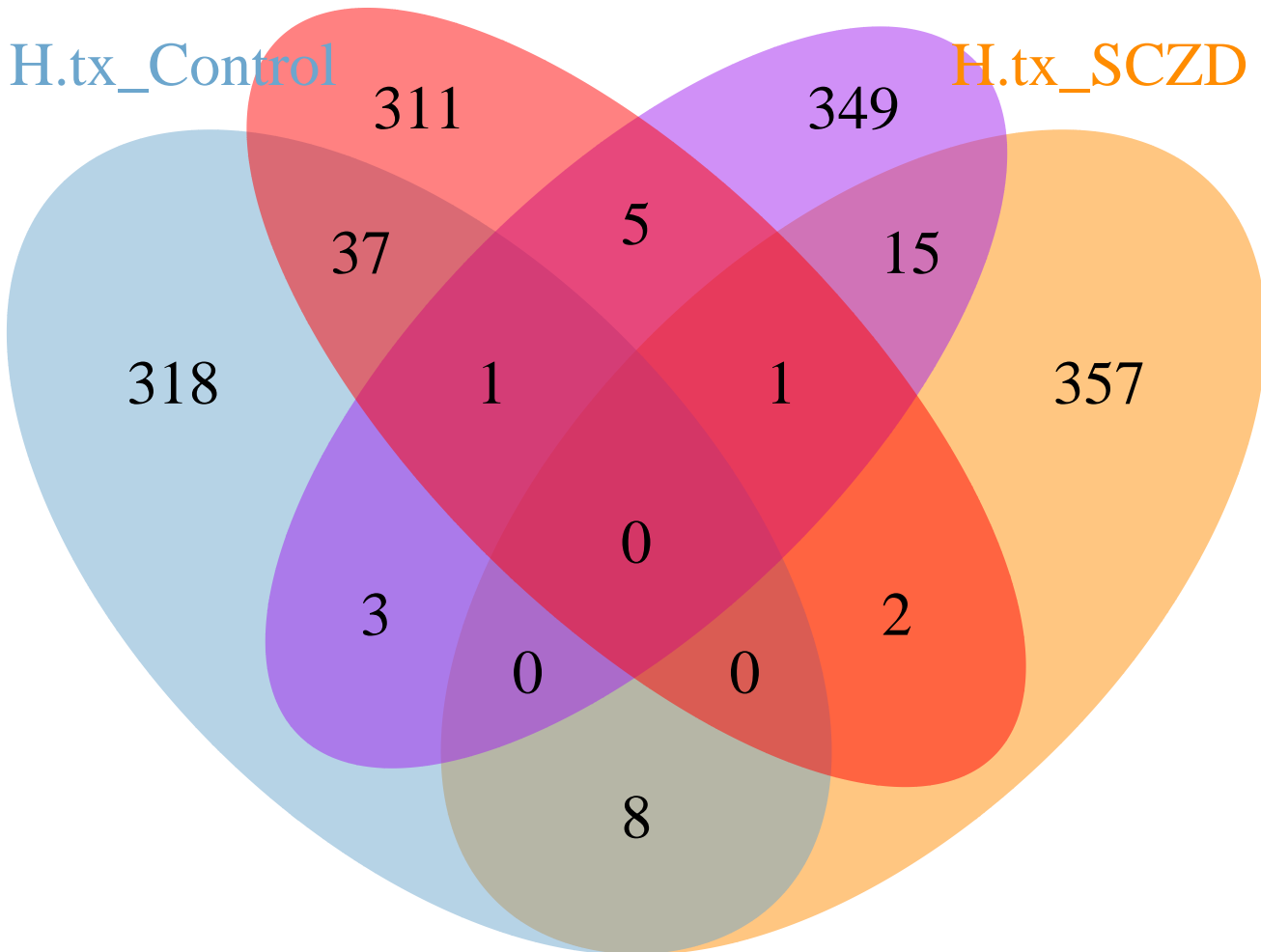
tx: top 400 in each group

D.tx_Control

D.tx_SCZD

H.tx_Control

H.tx_SCZD



tx: Control (top 400)

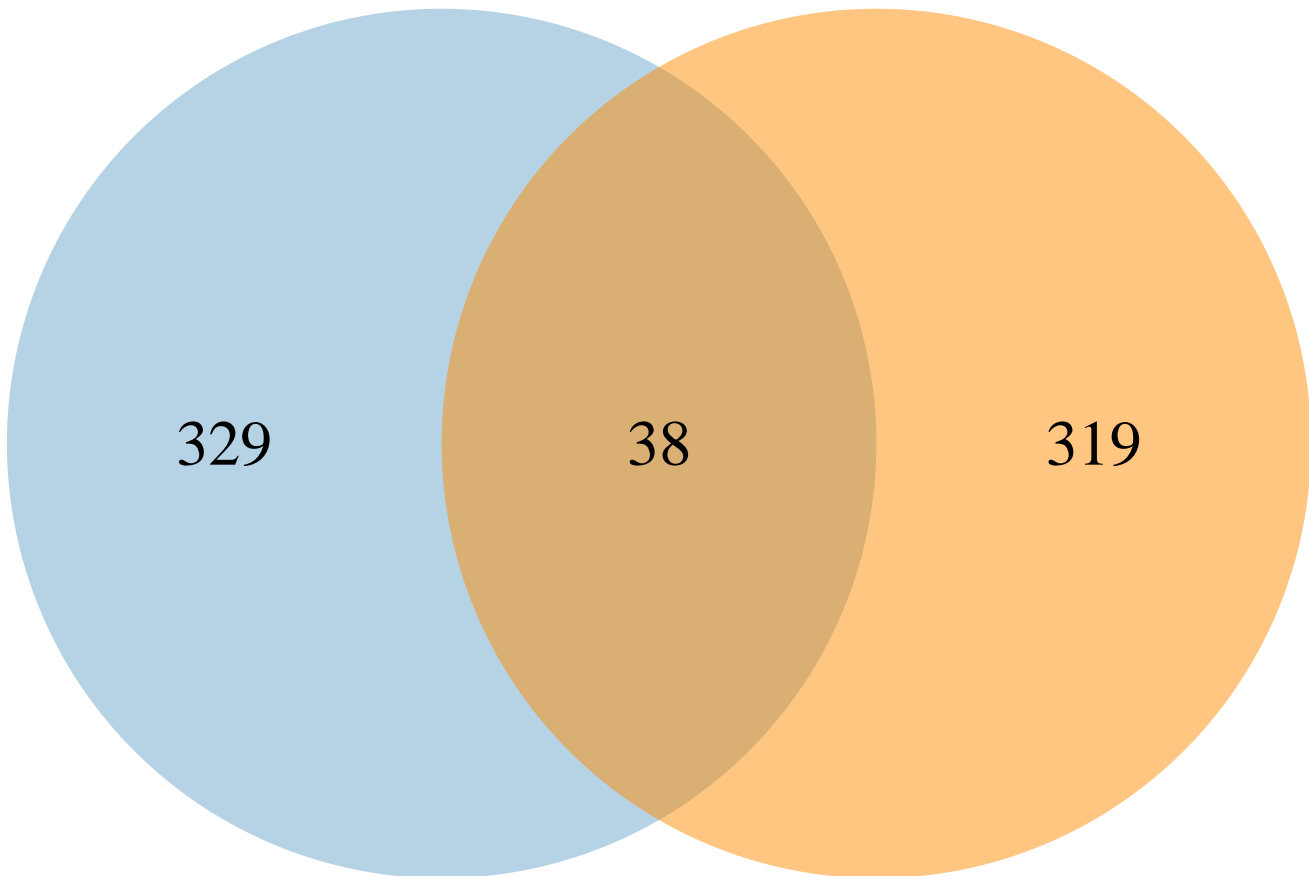
HIPPO

DLPFC

329

38

319



tx: SCZD (top 400)

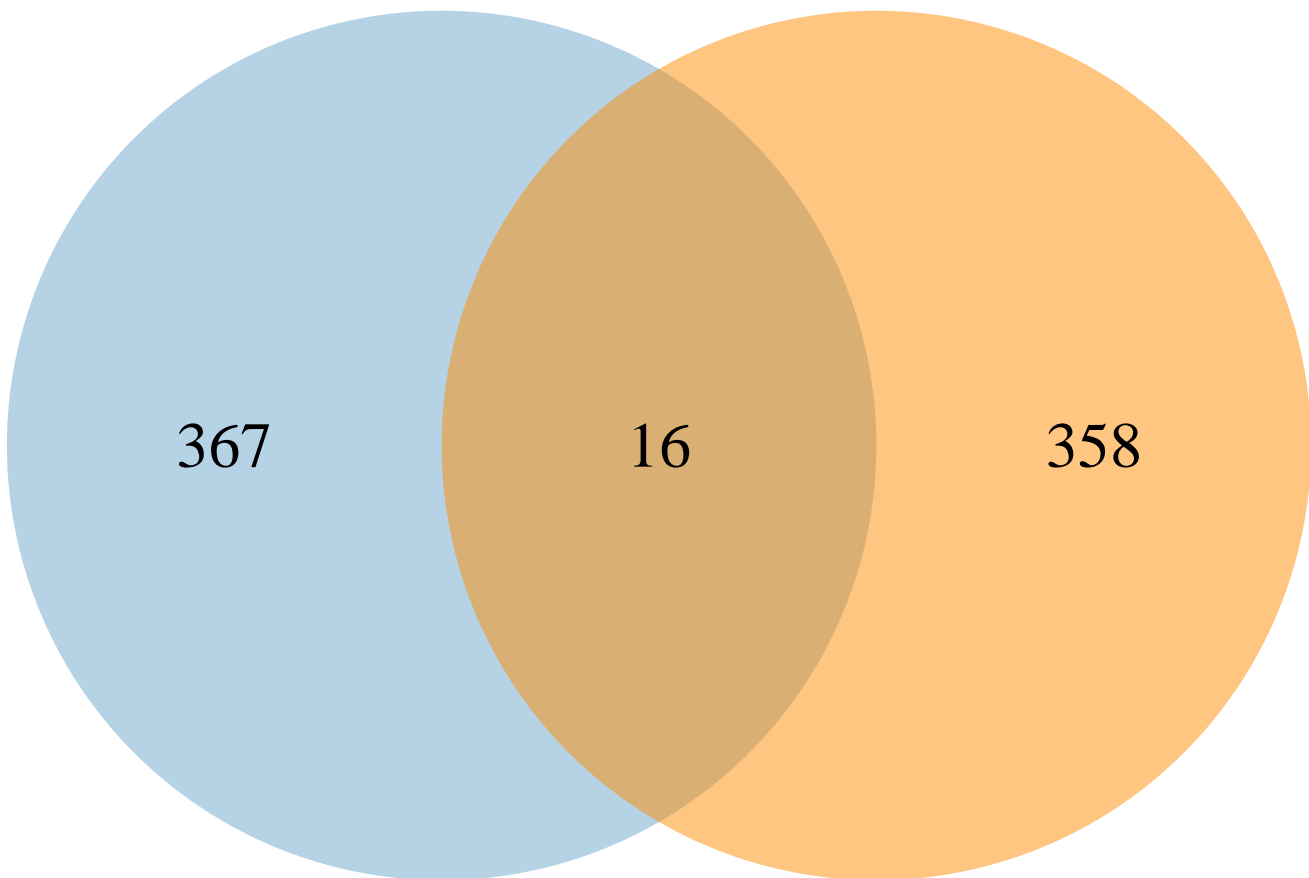
HIPPO

DLPFC

367

16

358

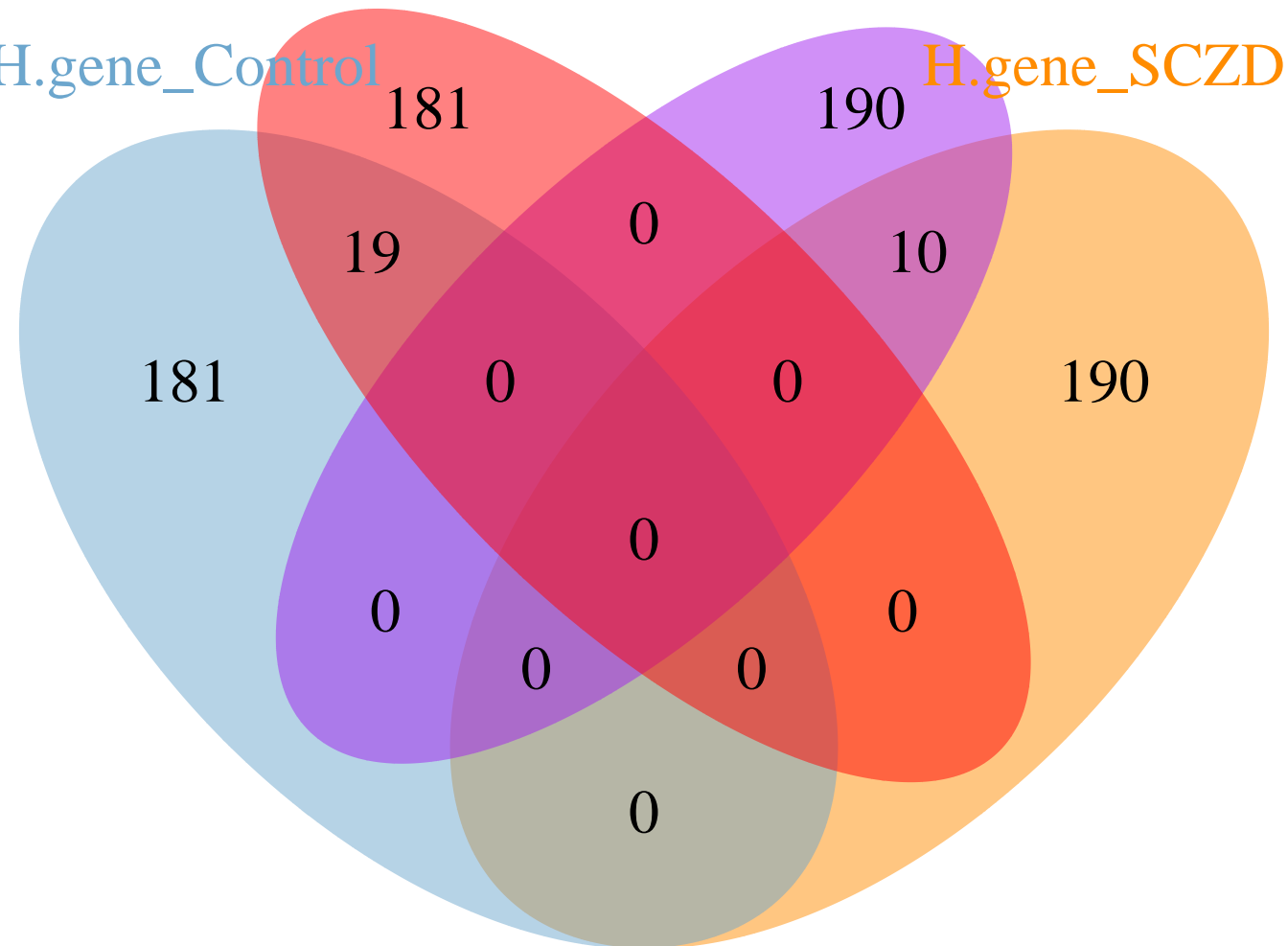


gene: top 200 in each group

D.gene_Control D.gene_SCZD

H.gene_Control

H.gene_SCZD



gene: Control (top 200)

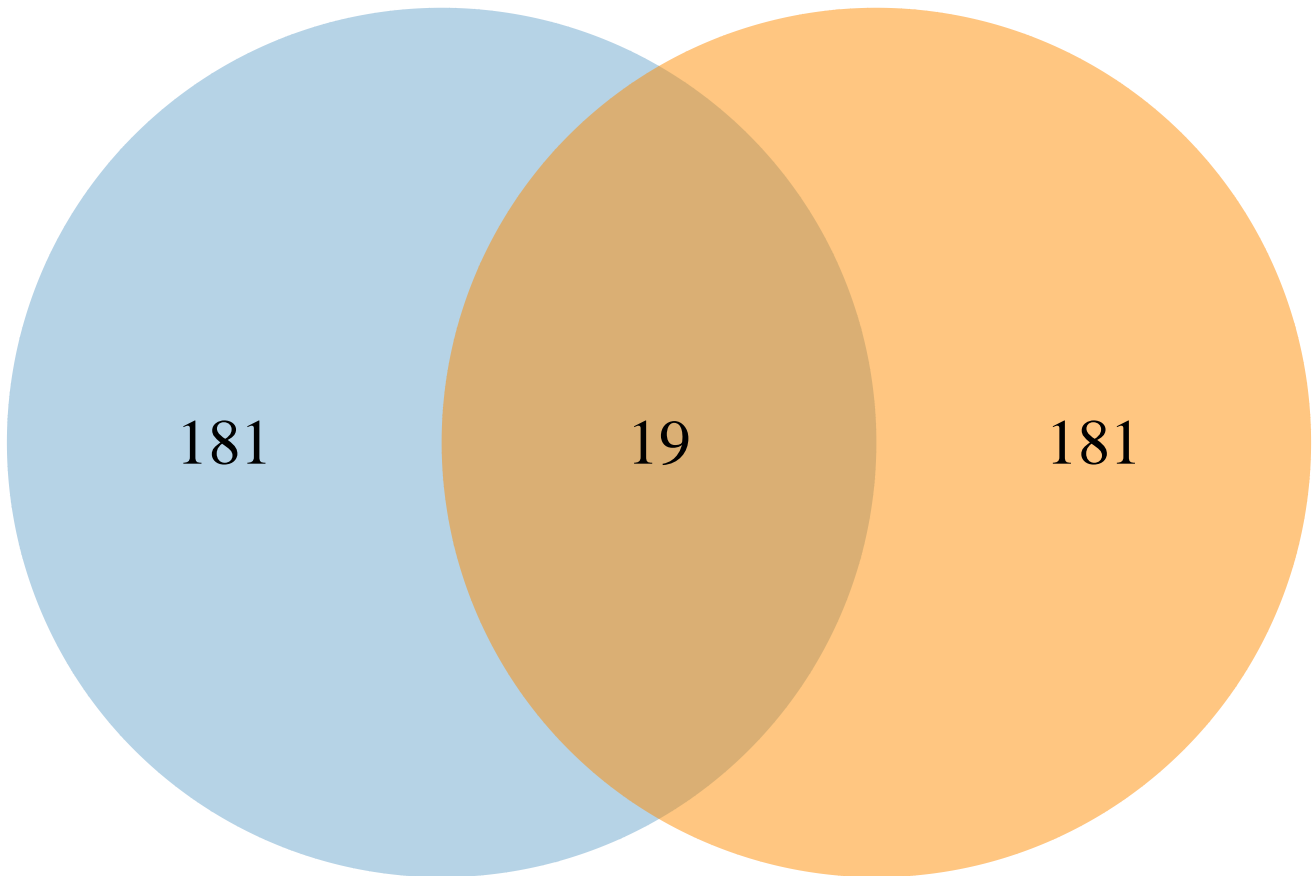
HIPPO

DLPFC

181

19

181



gene: SCZD (top 200)

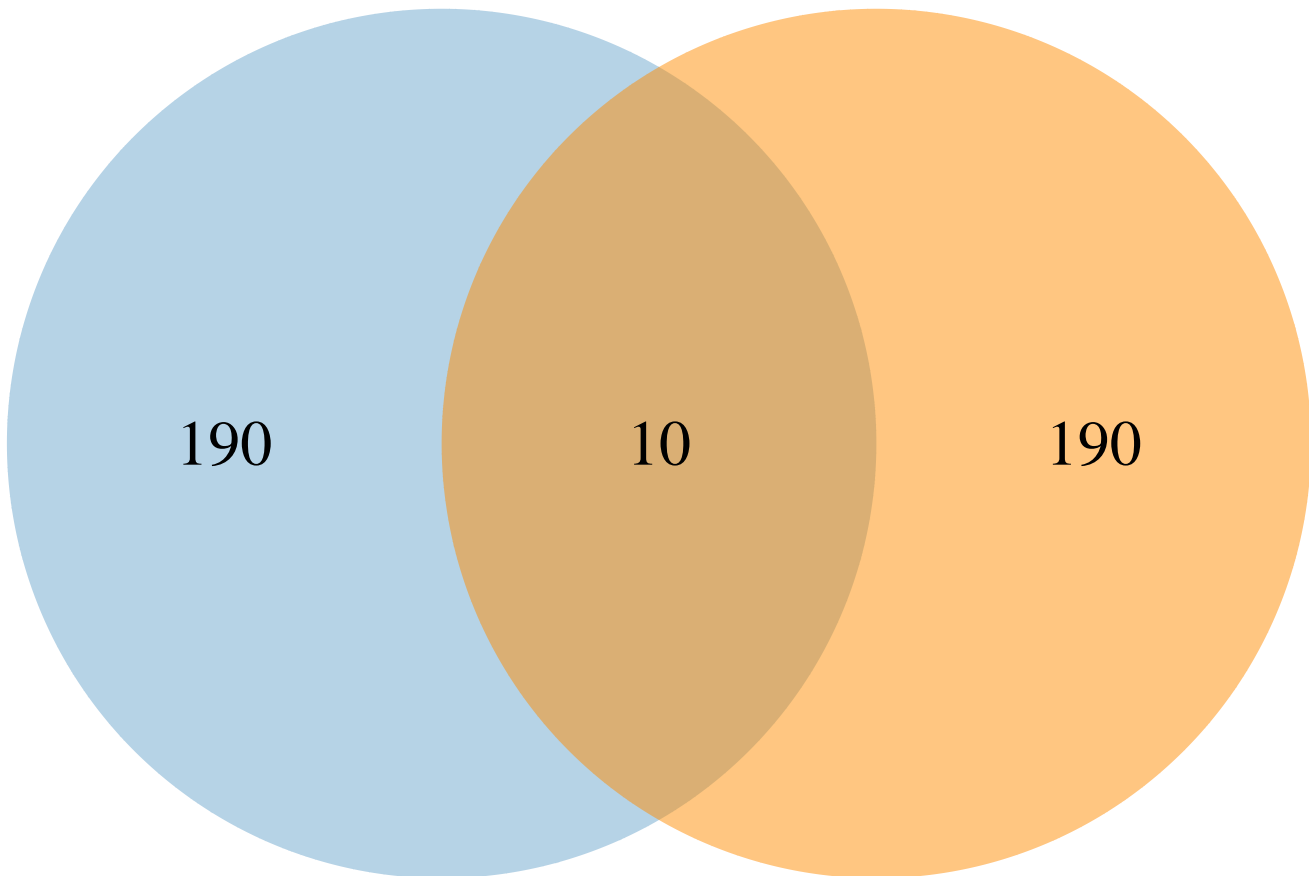
HIPPO

DLPFC

190

10

190

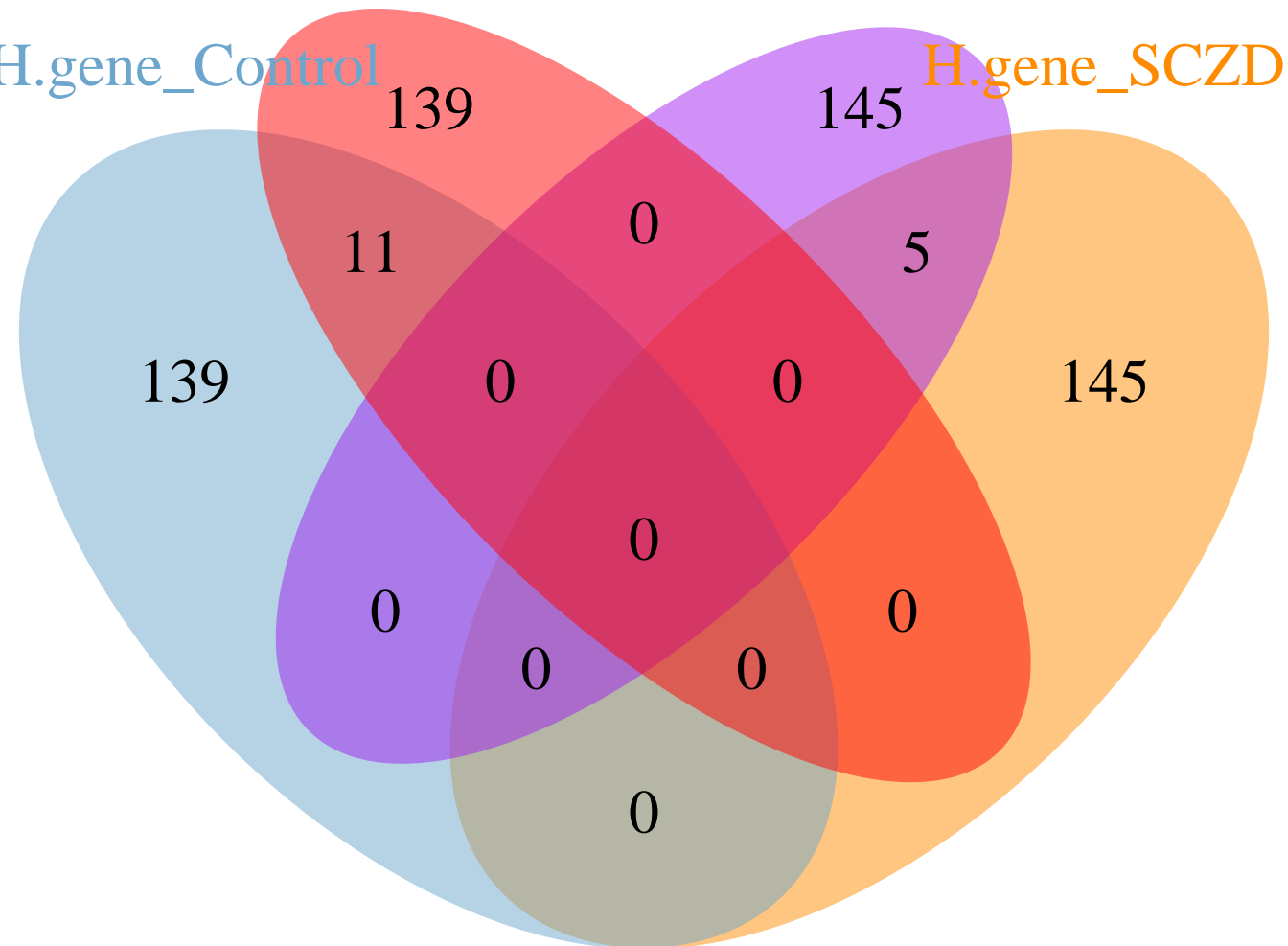


gene: top 150 in each group

D.gene_Control D.gene_SCZD

H.gene_Control

H.gene_SCZD



gene: Control (top 150)

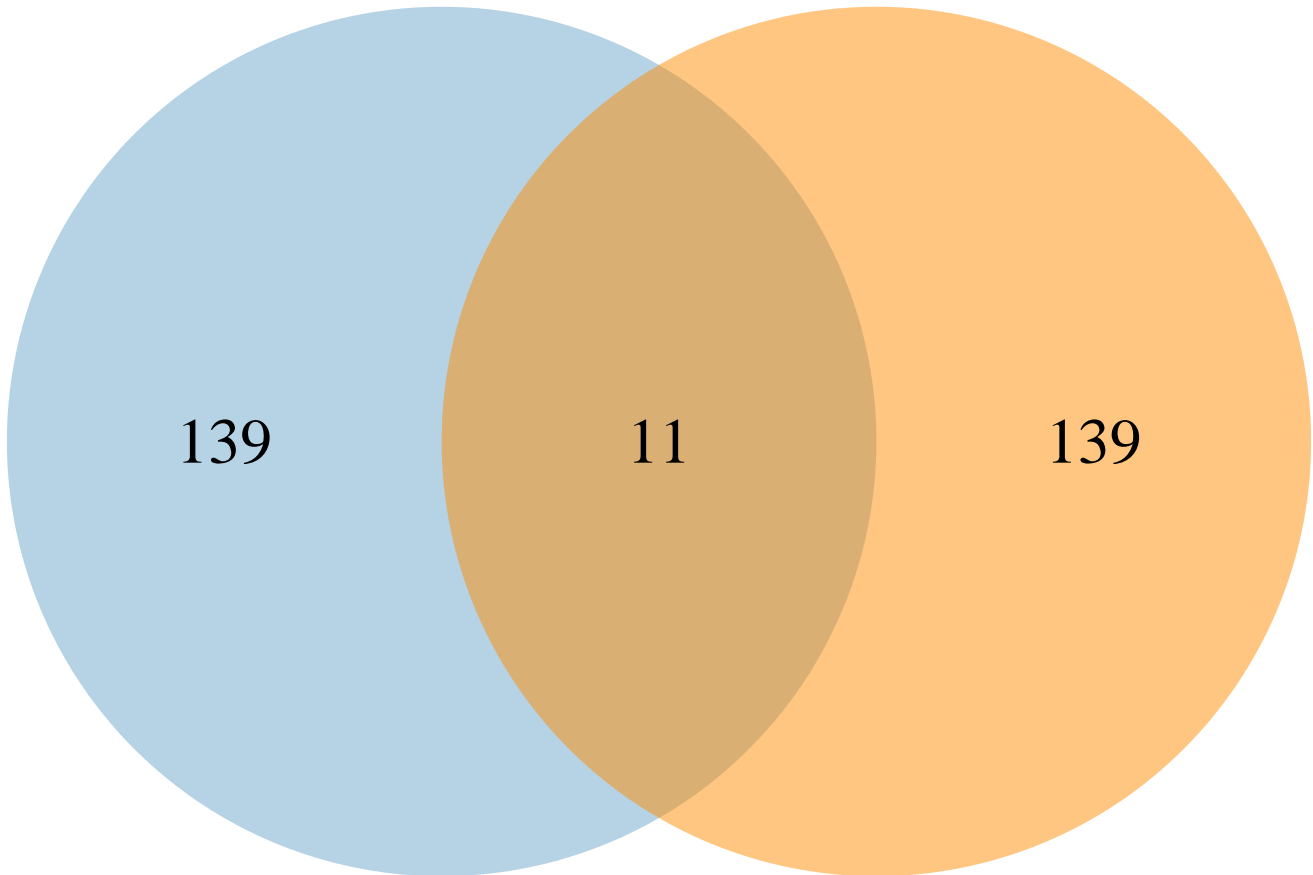
HIPPO

DLPFC

139

11

139



gene: SCZD (top 150)

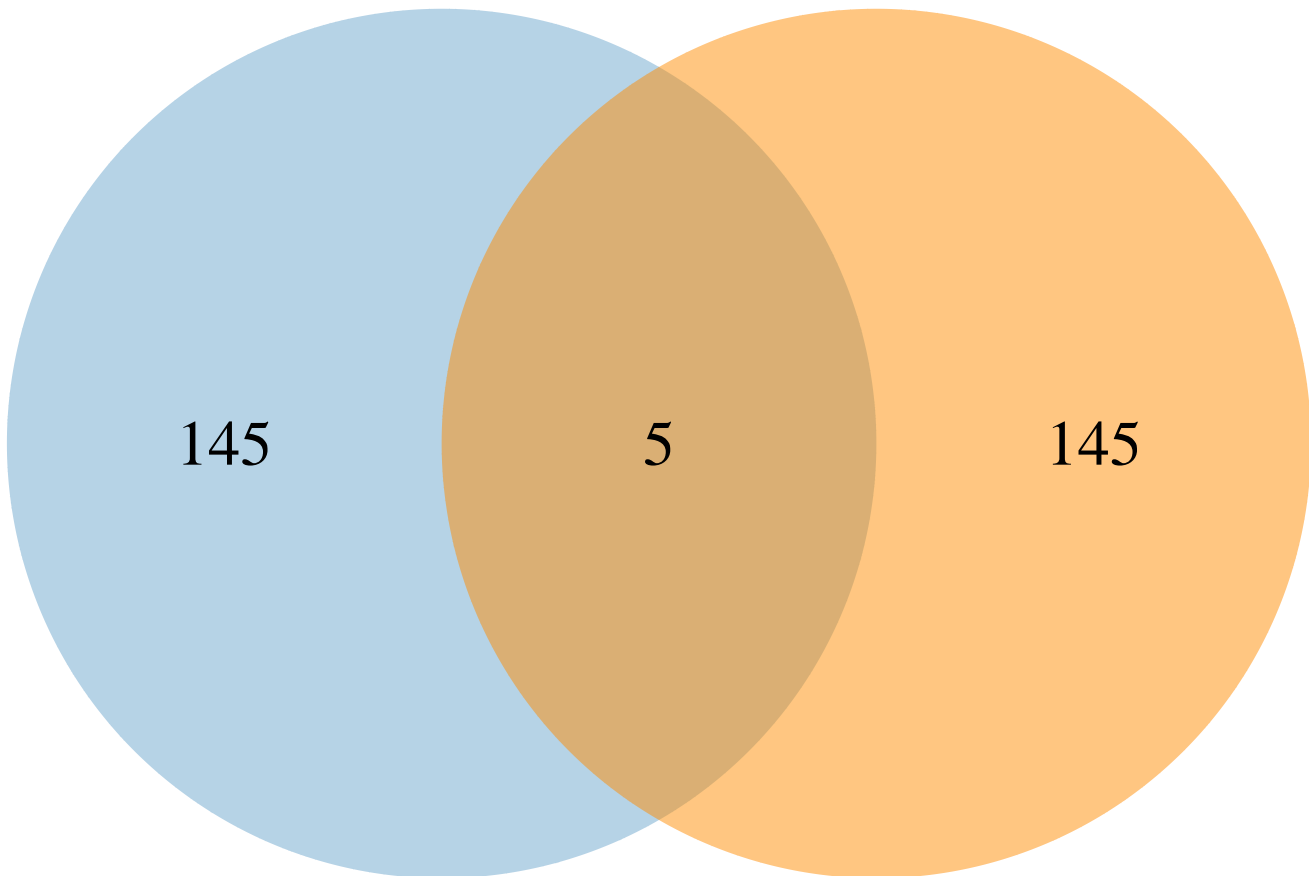
HIPPO

DLPFC

145

5

145

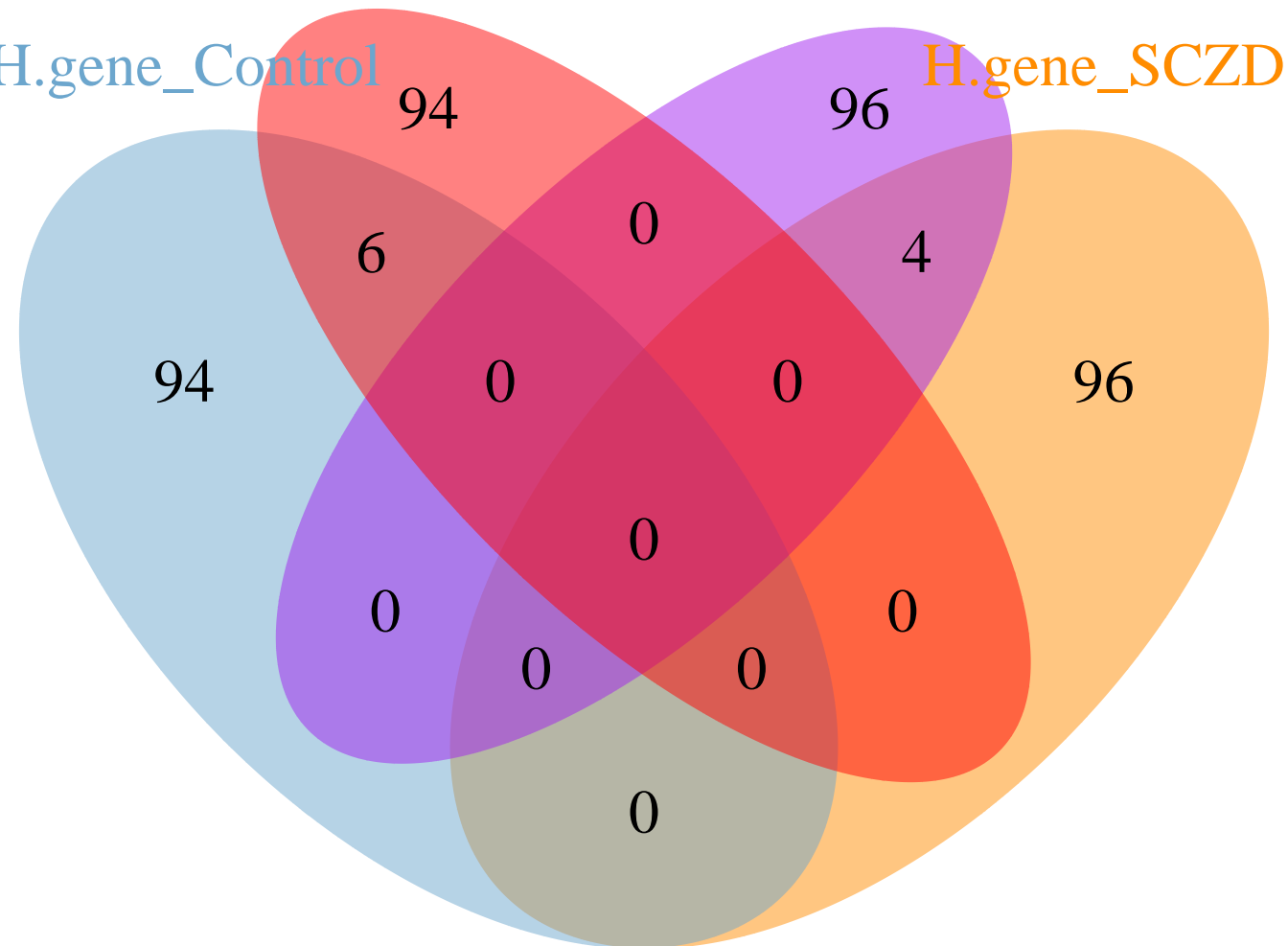


gene: top 100 in each group

D.gene_Control D.gene_SCZD

H.gene_Control

H.gene_SCZD



gene: Control (top 100)

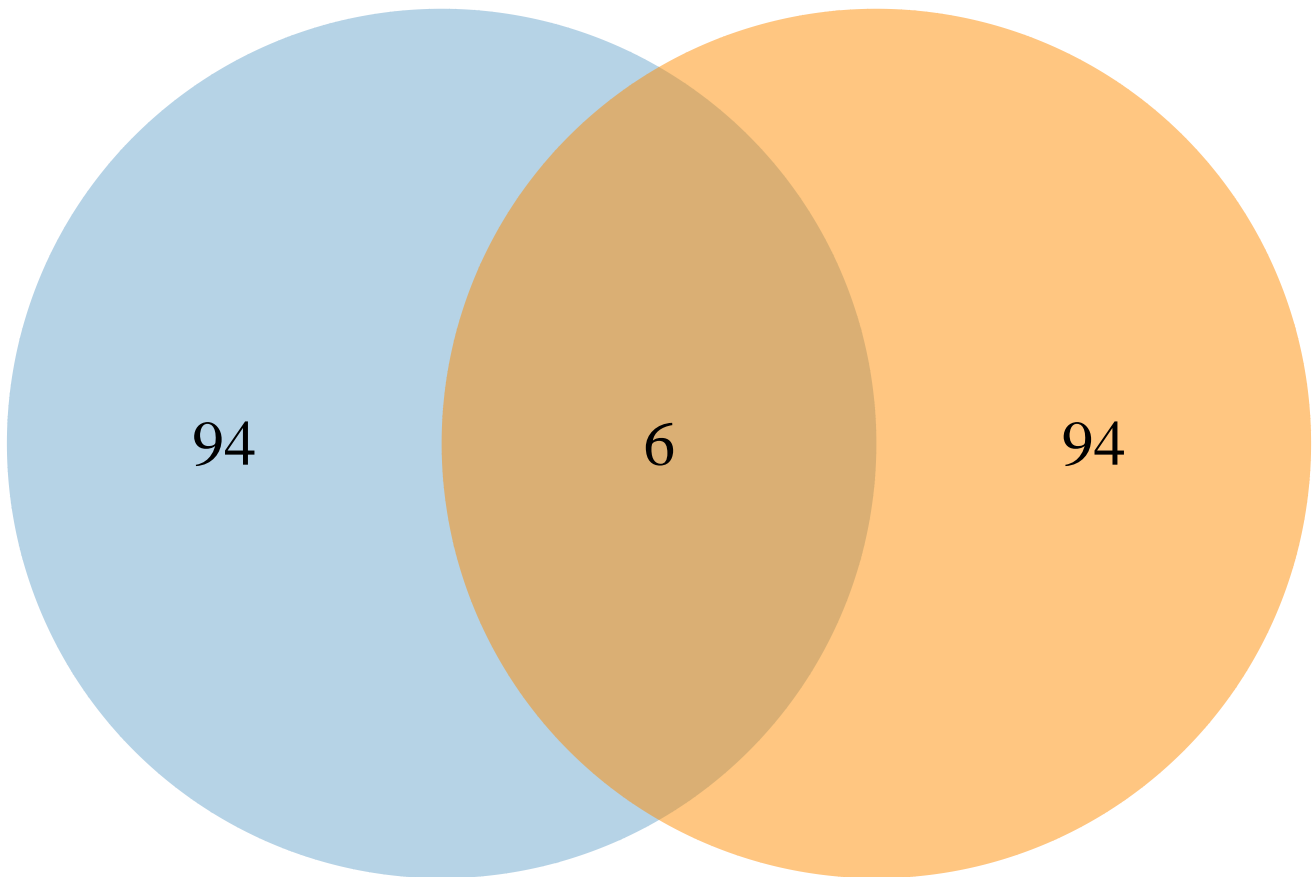
HIPPO

DLPFC

94

6

94



gene: SCZD (top 100)

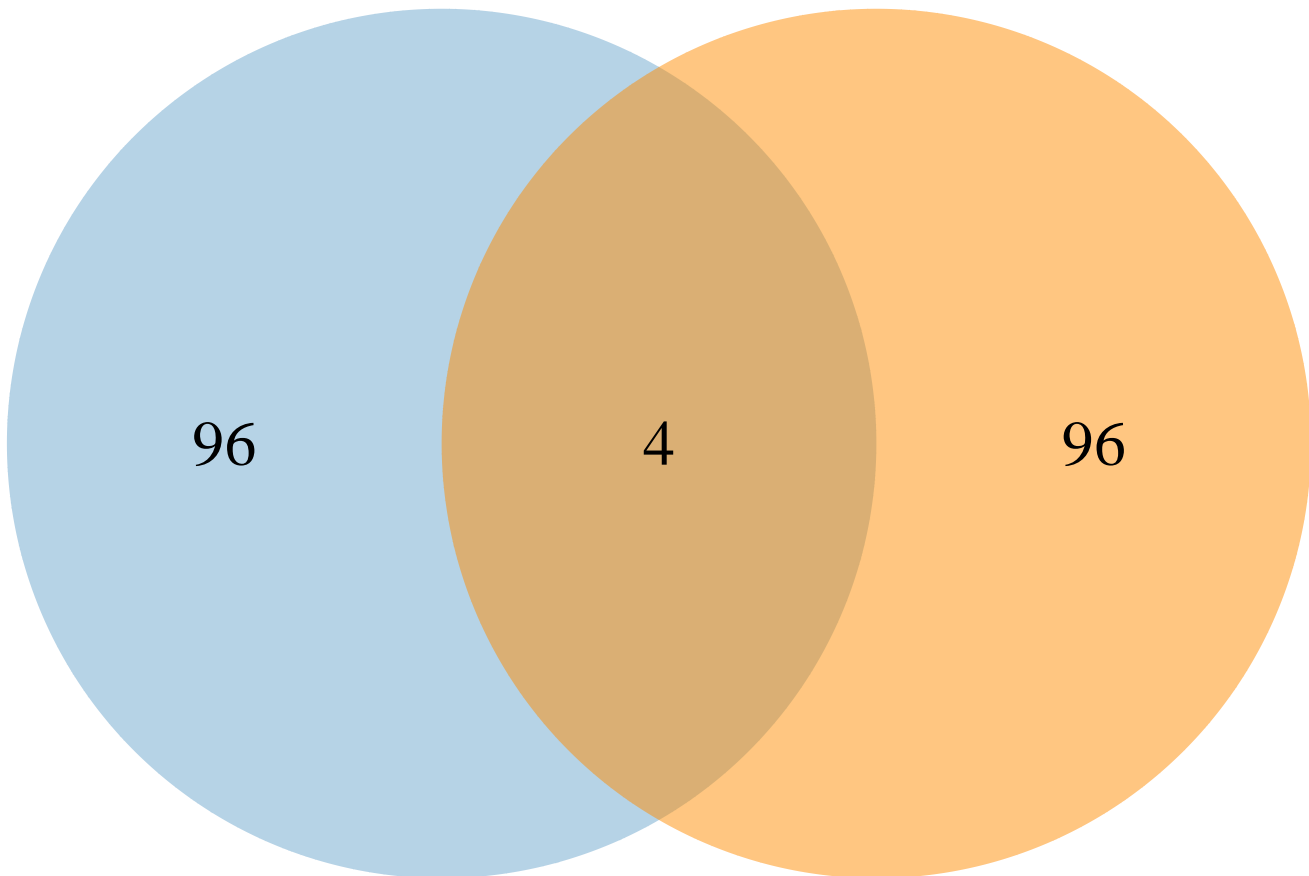
HIPPO

DLPFC

96

4

96

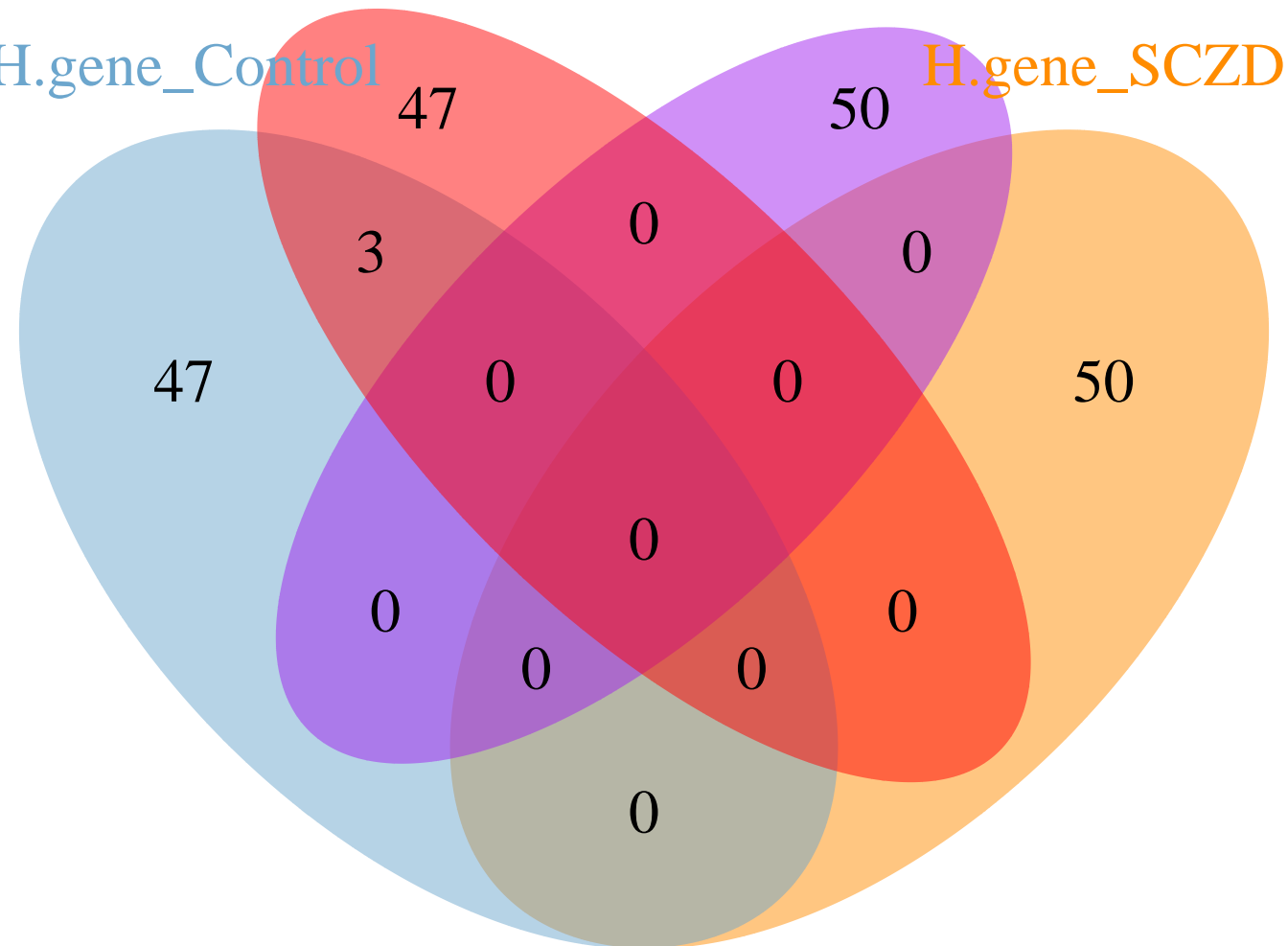


gene: top 50 in each group

D.gene_Control D.gene_SCZD

H.gene_Control

H.gene_SCZD



gene: Control (top 50)

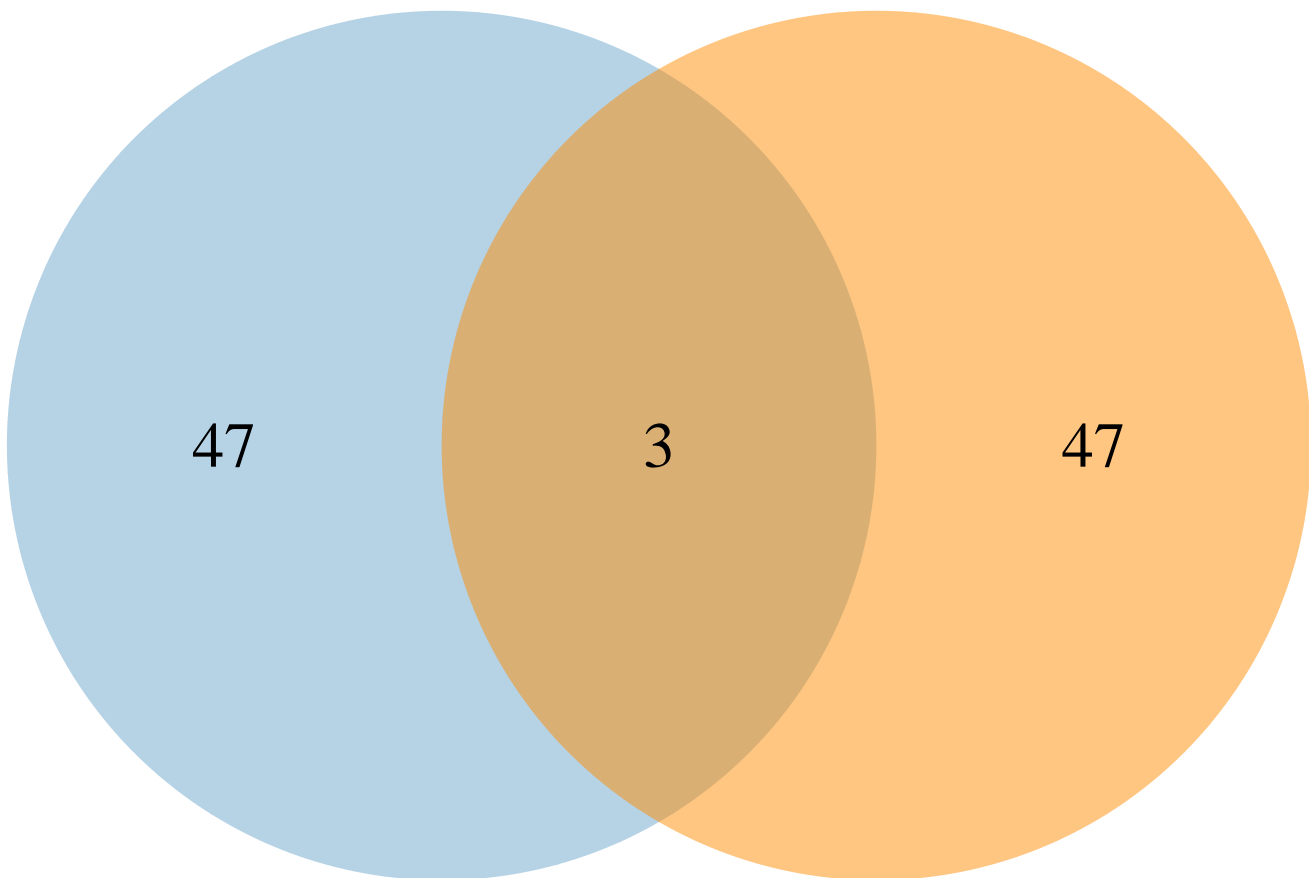
HIPPO

DLPFC

47

3

47



gene: SCZD (top 50)

HIPPO

50

DLPFC

50