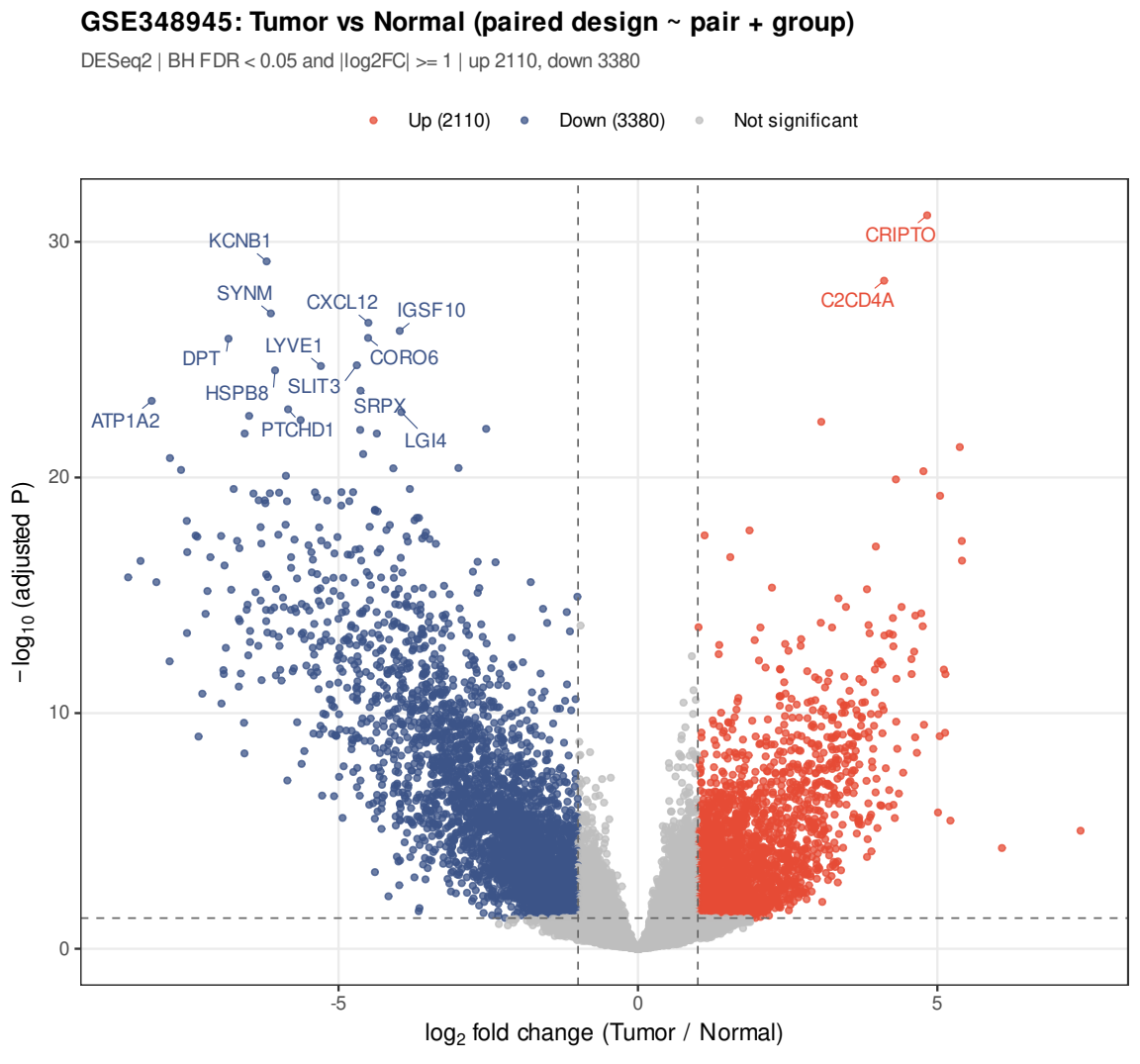
GSE348945: Tumor vs Normal (paired design ~ pair + group)
DESeq2 | BH FDR < 0.05 and |log2FC| >= 1 | up 2110, down 3380
Up (2110)
Down (3380)
Not significant
KCNB1
SYNM
CXCL12
IGSF10
DPT
LYVE1
CORO6
HSPB8
SLIT3
SRPX
ATP1A2
PTCHD1
LGI4
CRIPTO
C2CD4A
30
20
10
0
−log10 (adjusted P)
-5
0
5
log2 fold change (Tumor / Normal)