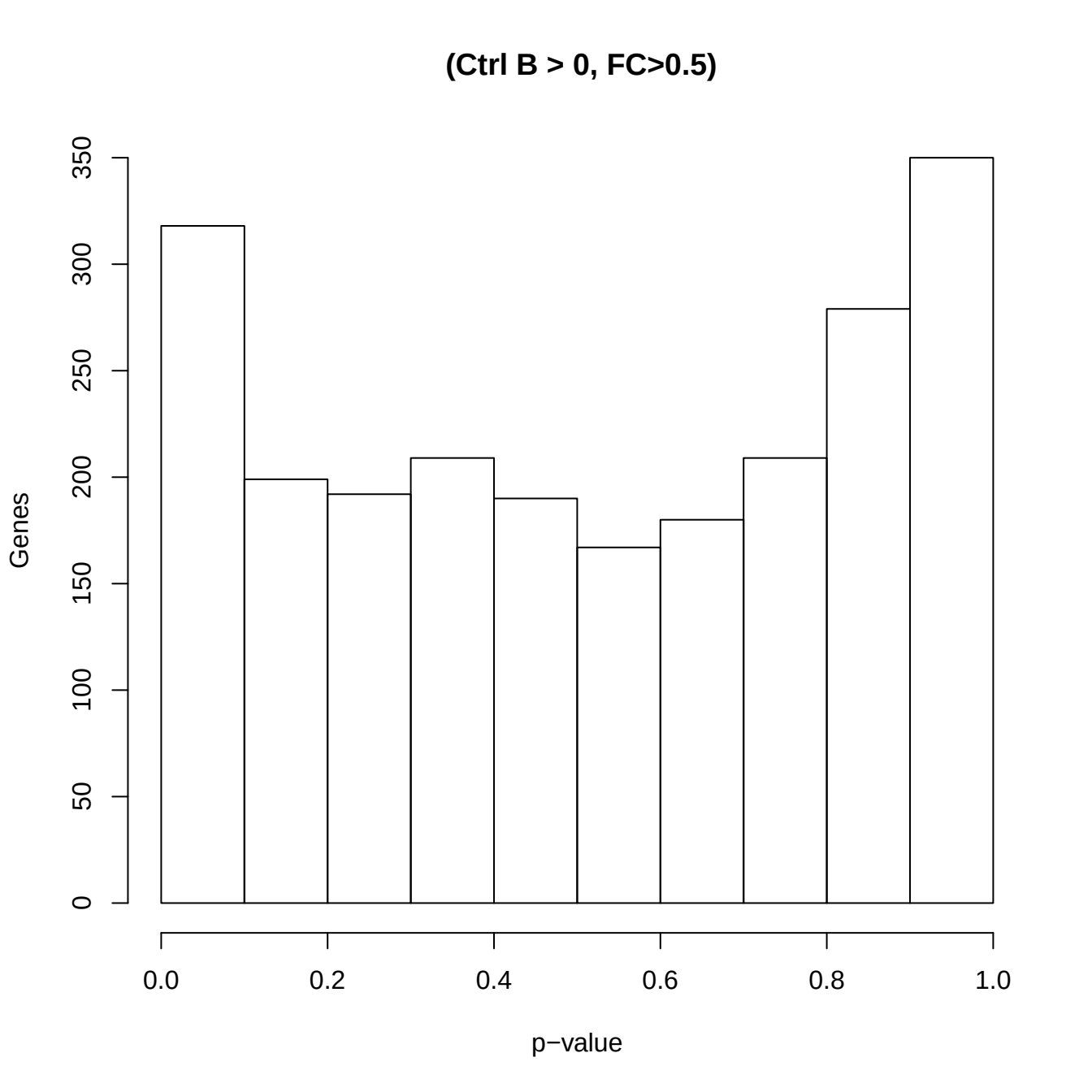
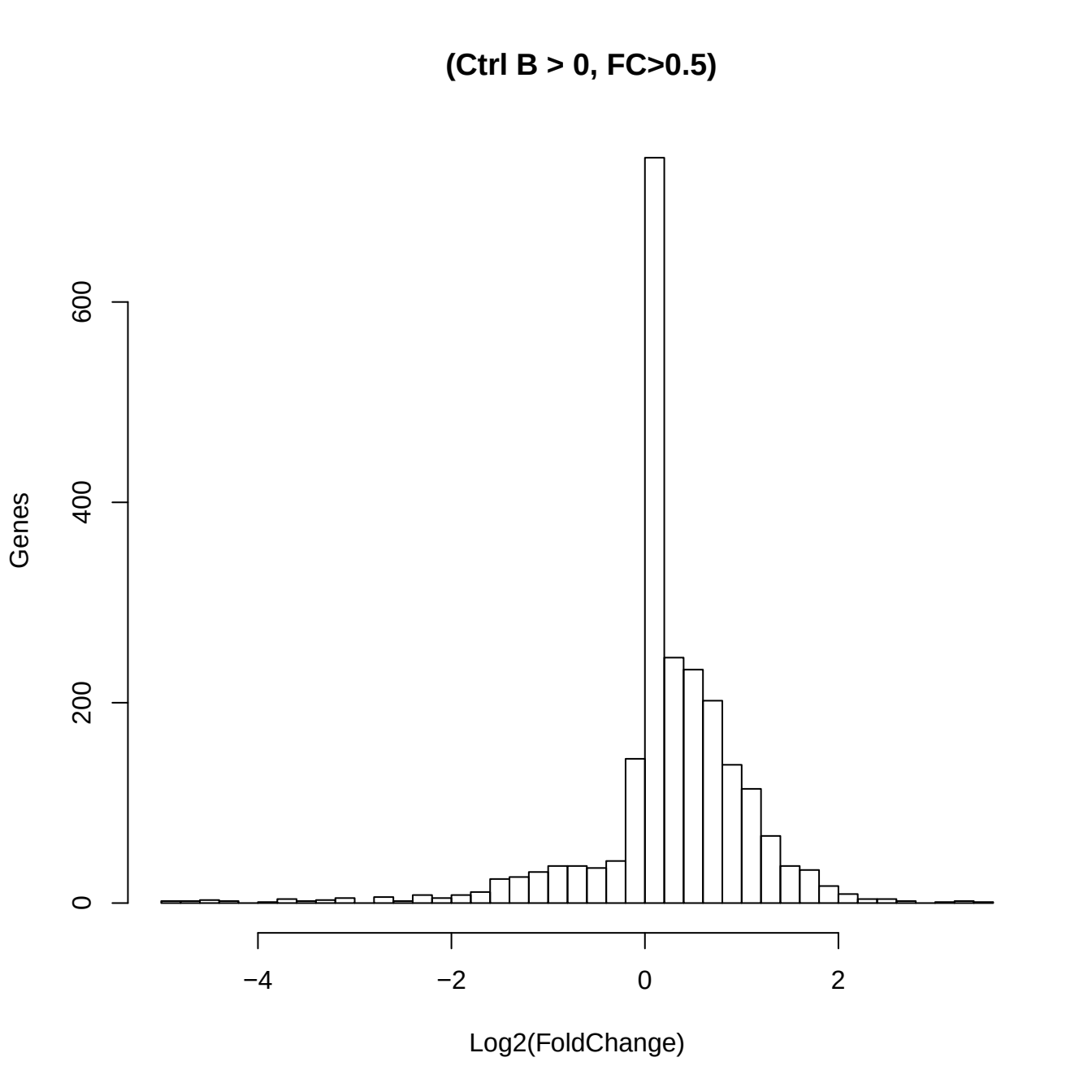
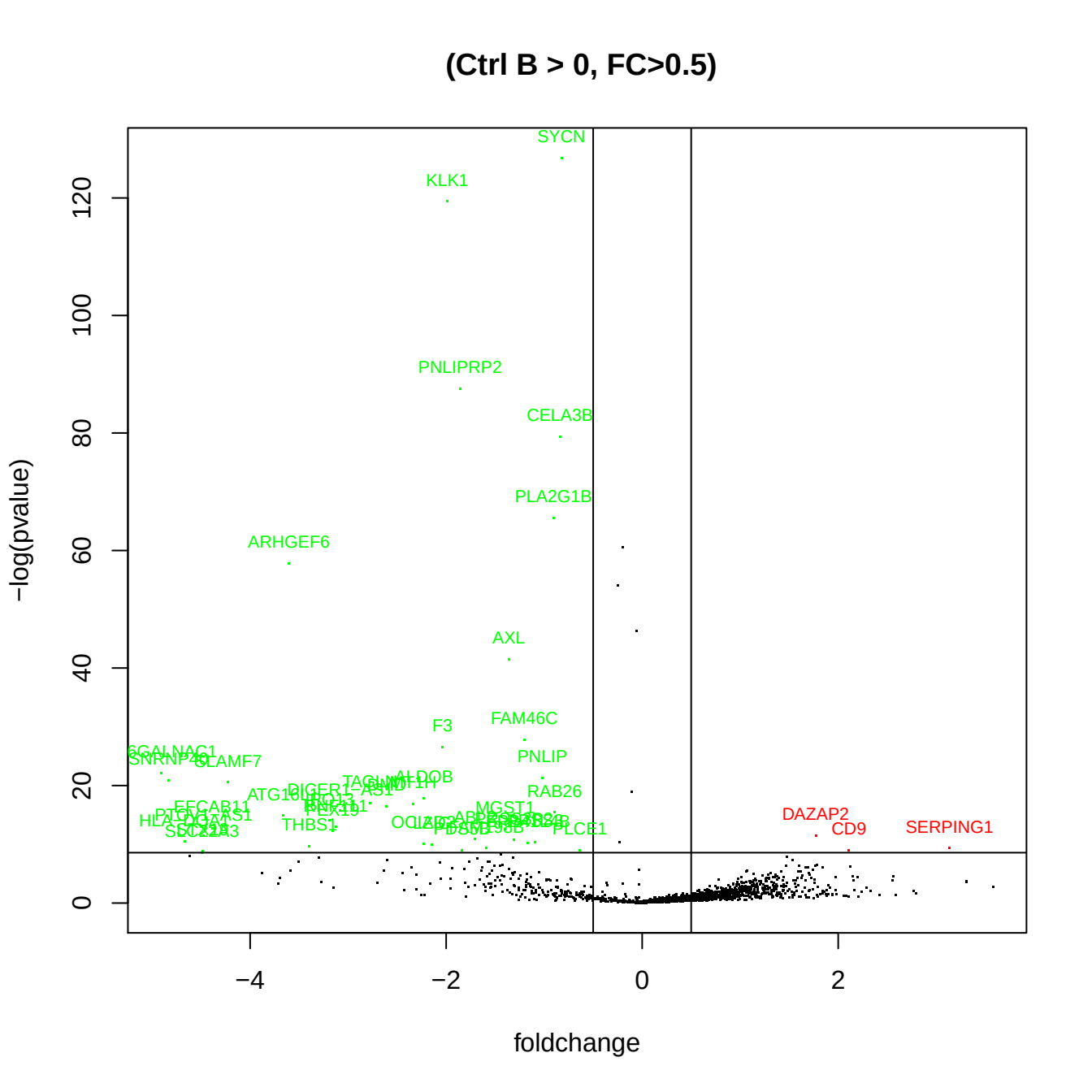

(Ctrl B > 0, FC>0.5)
Genes
0
50
100
150
200
250
300
350
0.0
0.2
0.4
0.6
0.8
1.0
p−value


(Ctrl B > 0, FC>0.5)
Genes
0
200
400
600
−4
−2
0
2
Log2(FoldChange)


(Ctrl B > 0, FC>0.5)
SYCN
KLK1
PNLIPRP2
CELA3B
PLA2G1B
ARHGEF6
AXL
F3
FAM46C
PNLIP
RAB26
SLAMF7
THBS1
PLCE1
DAZAP2
CD9
SERPING1
−log(pvalue)
foldchange
0
20
40
60
80
100
120
−4
−2
0
2