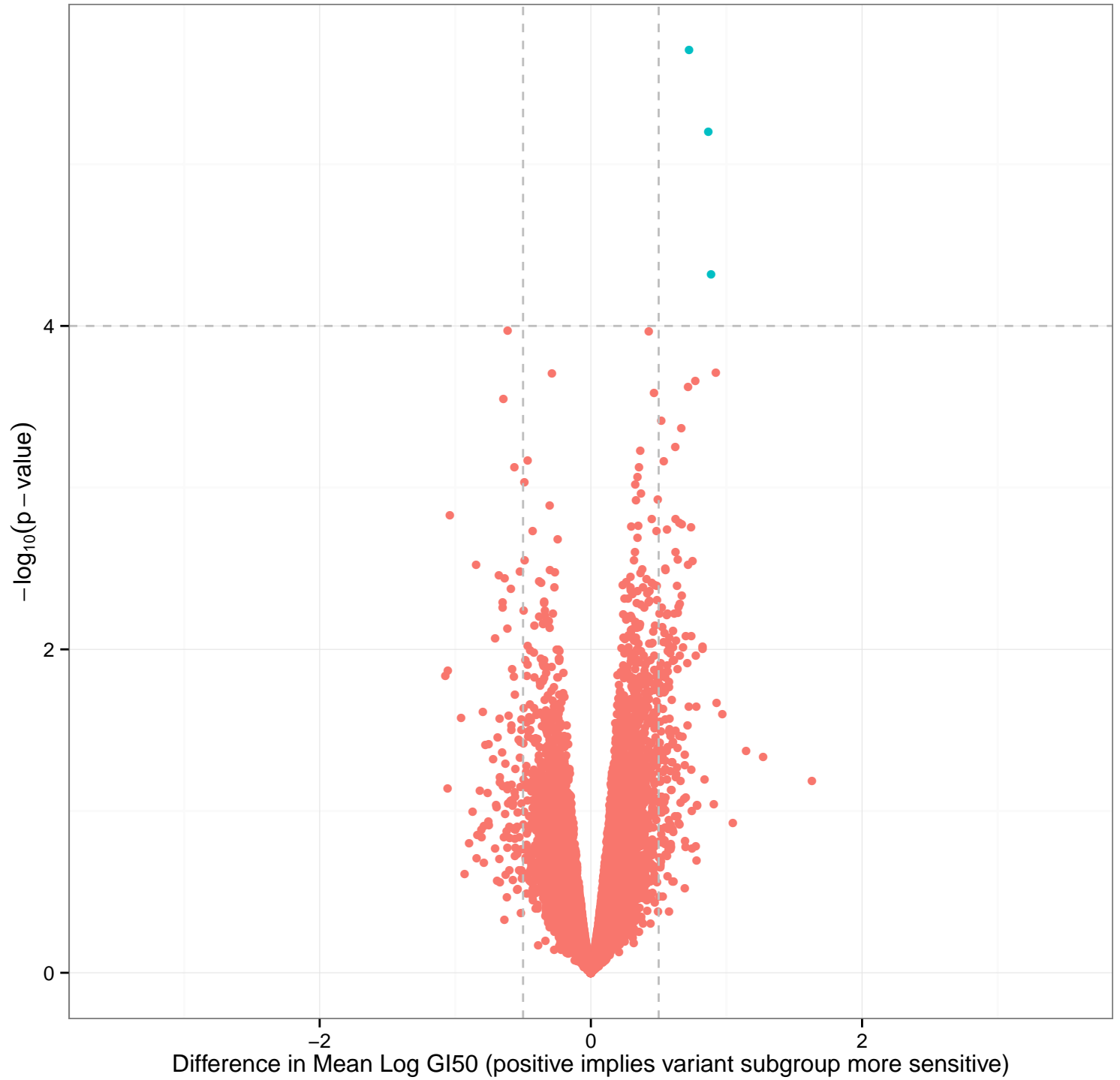
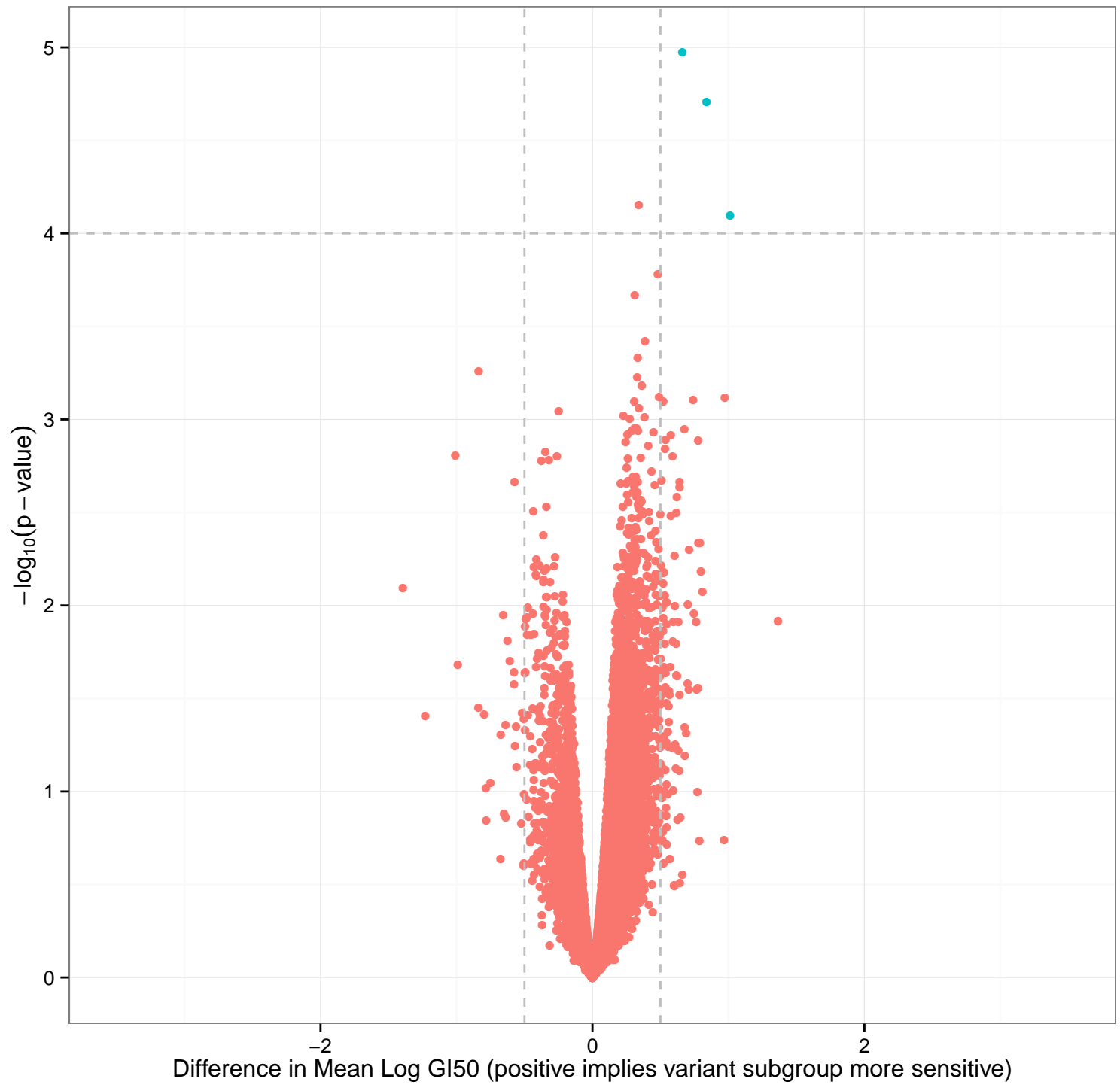


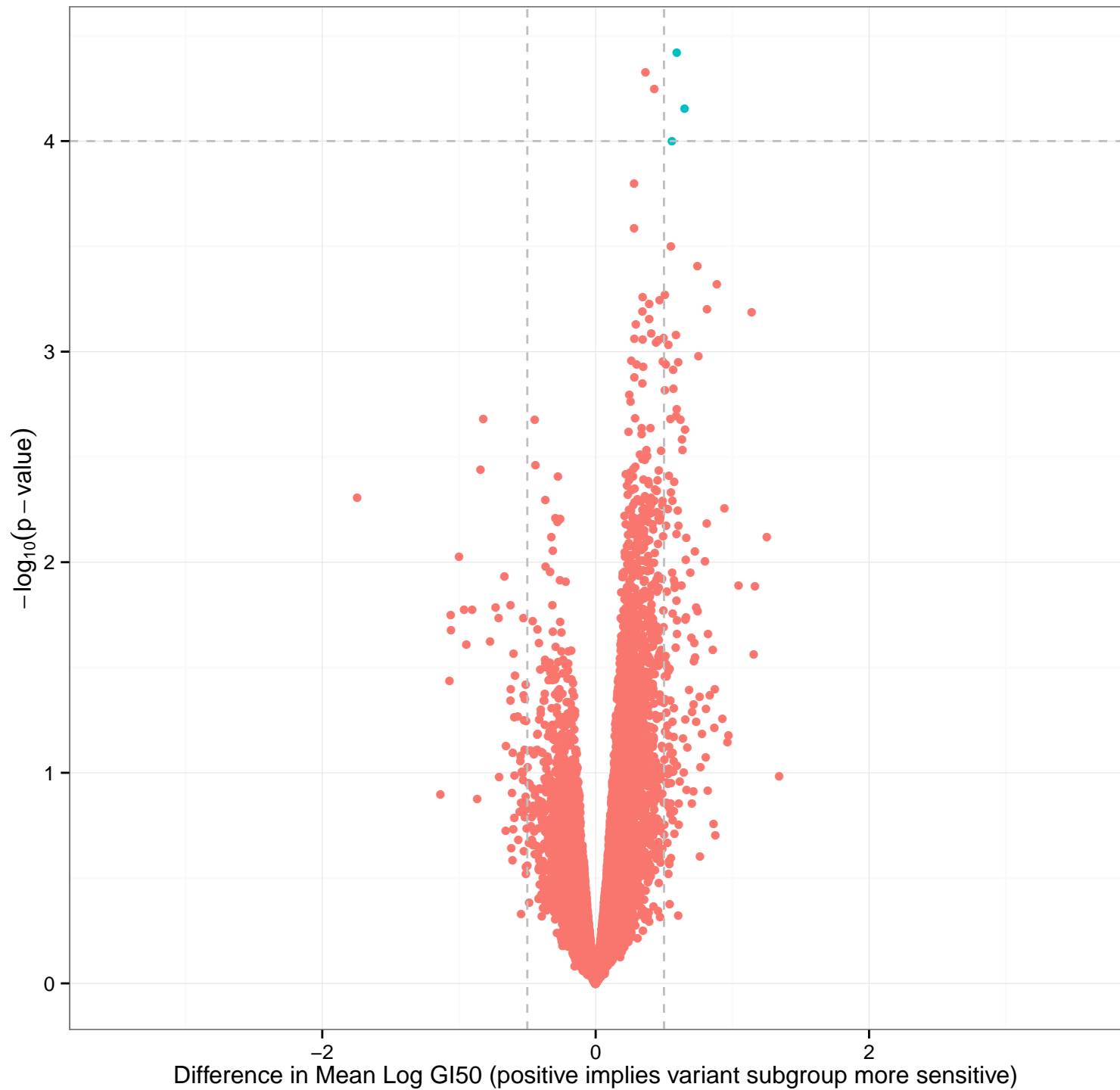
ALK



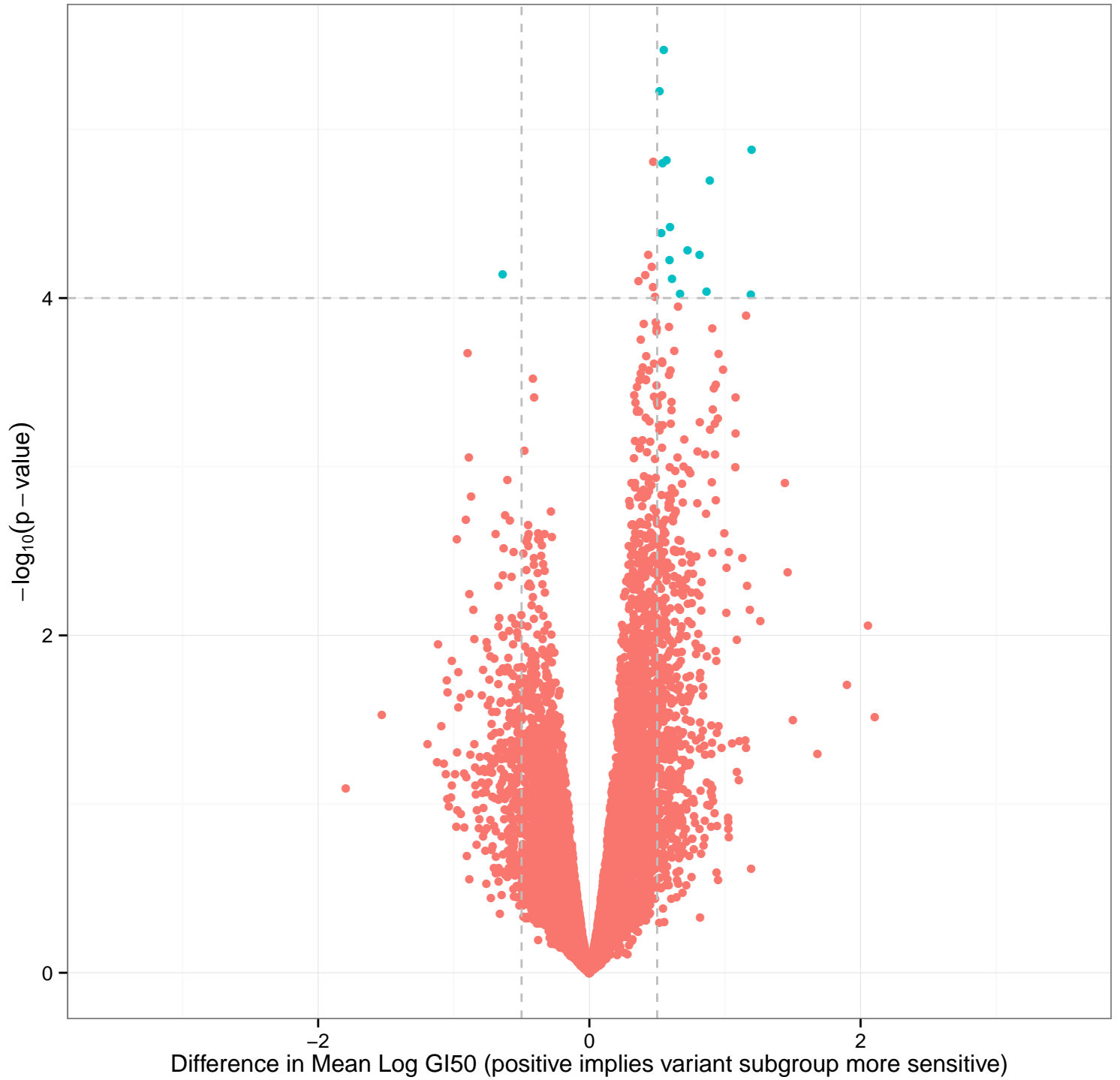
ARID1A



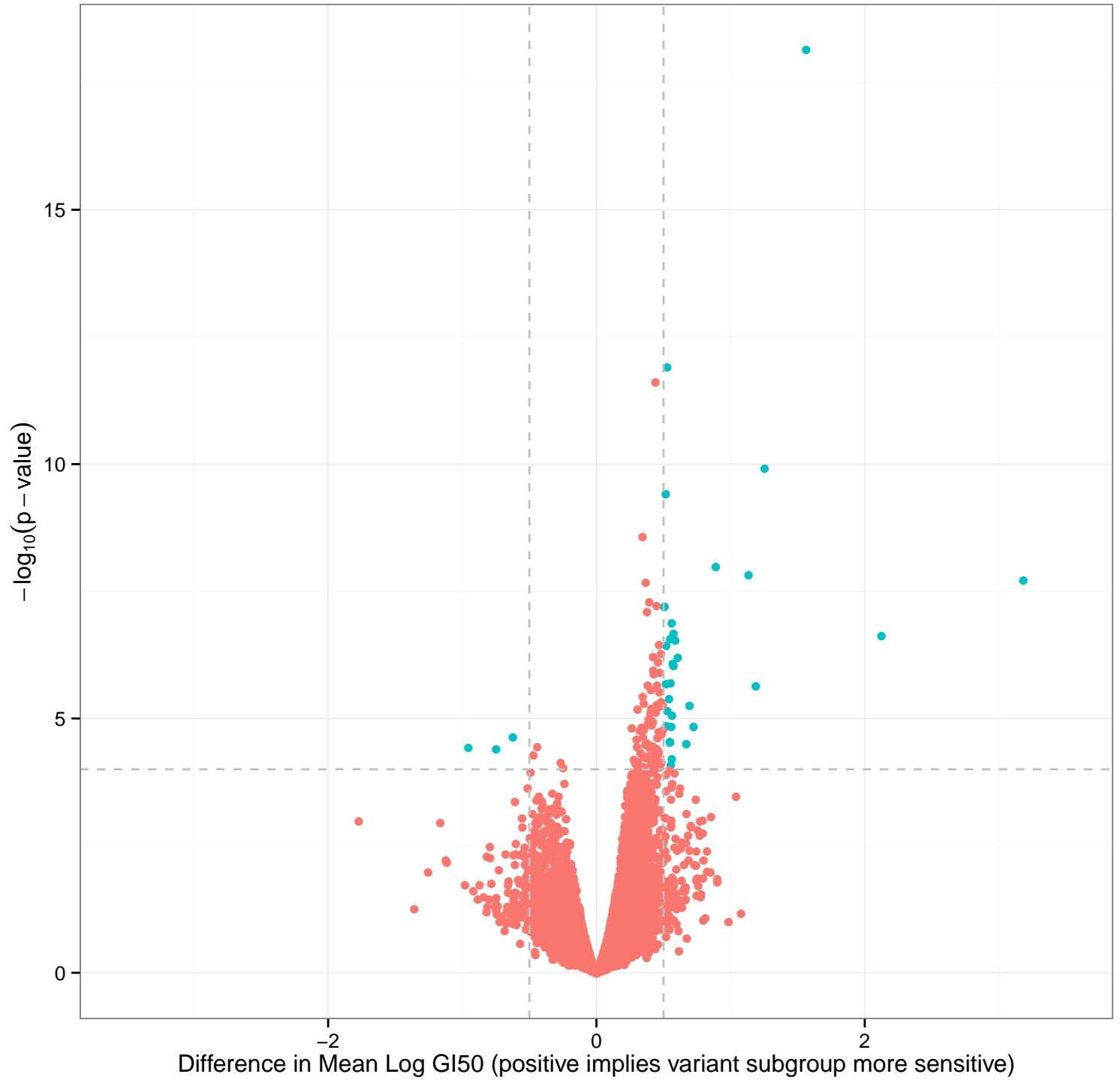
ATM



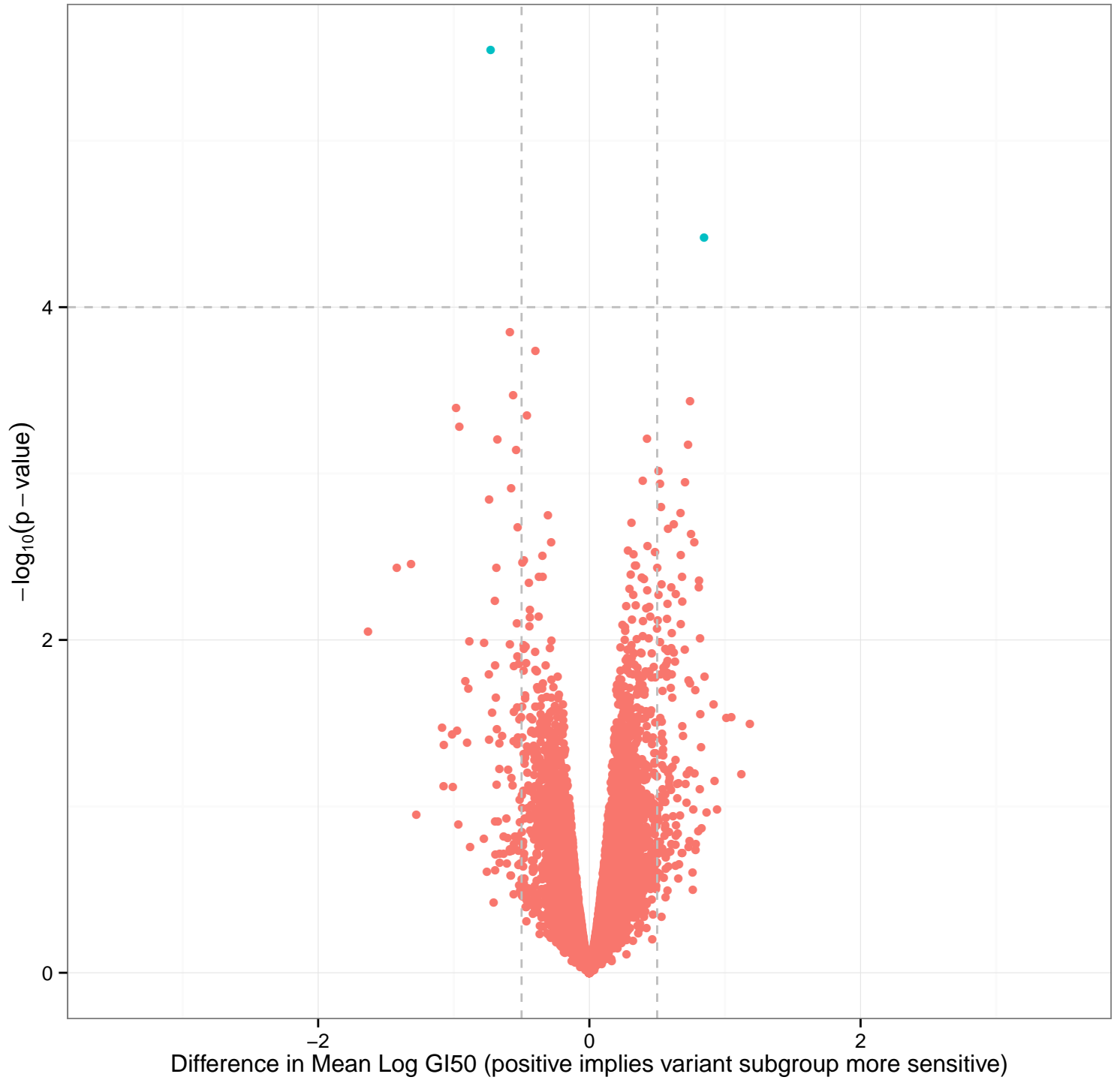
ATR



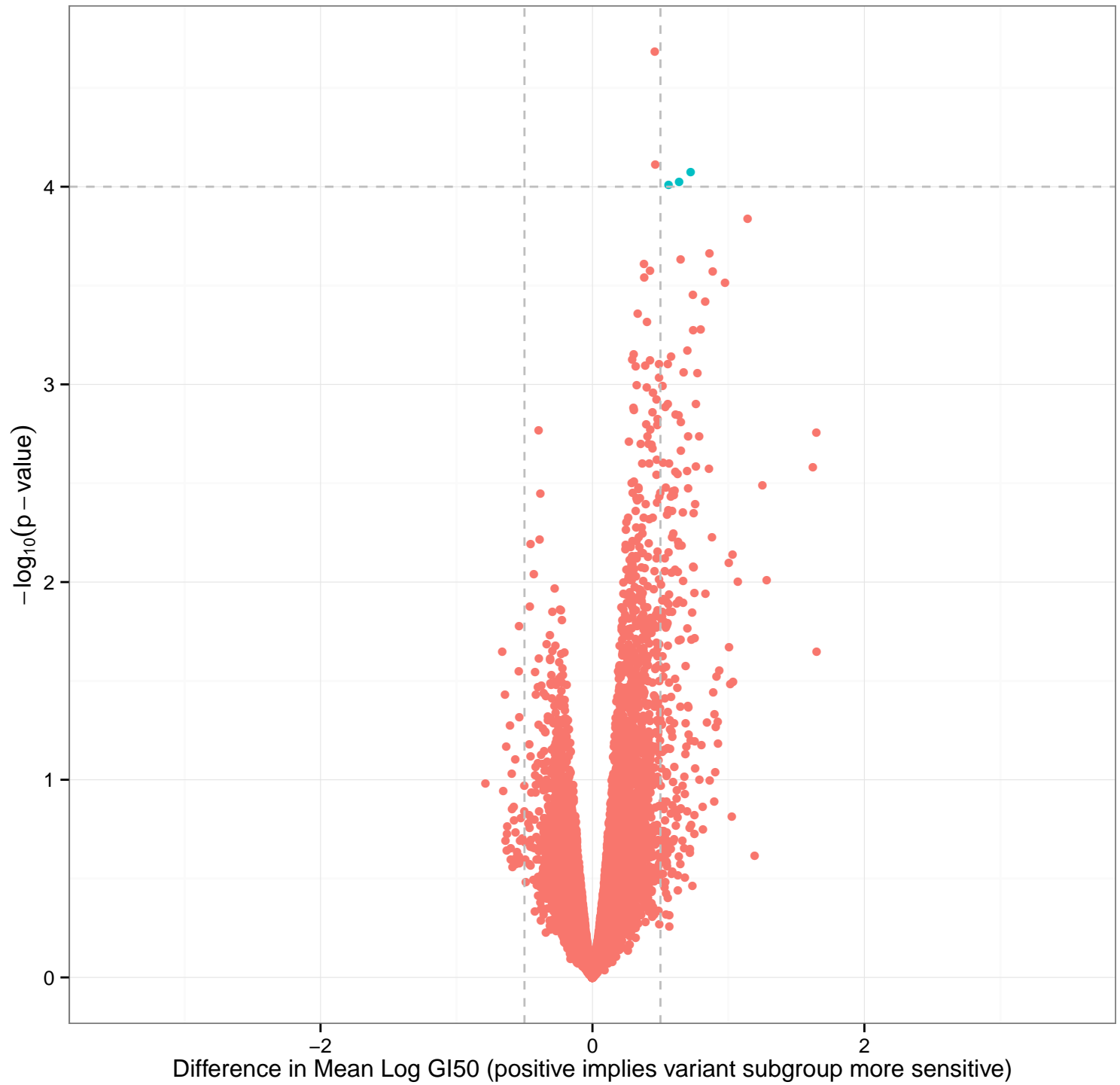
BRAF



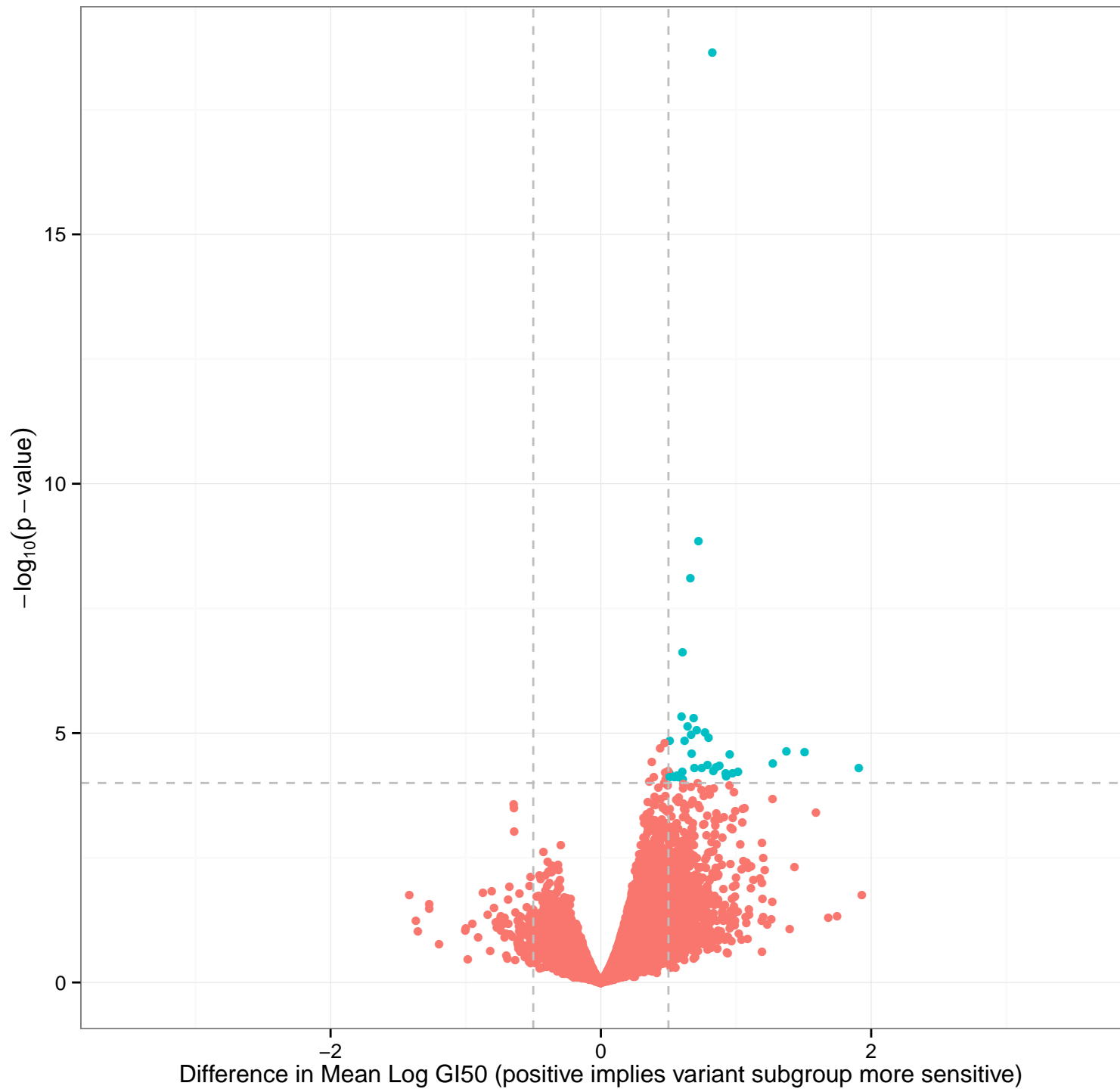
BRCA1



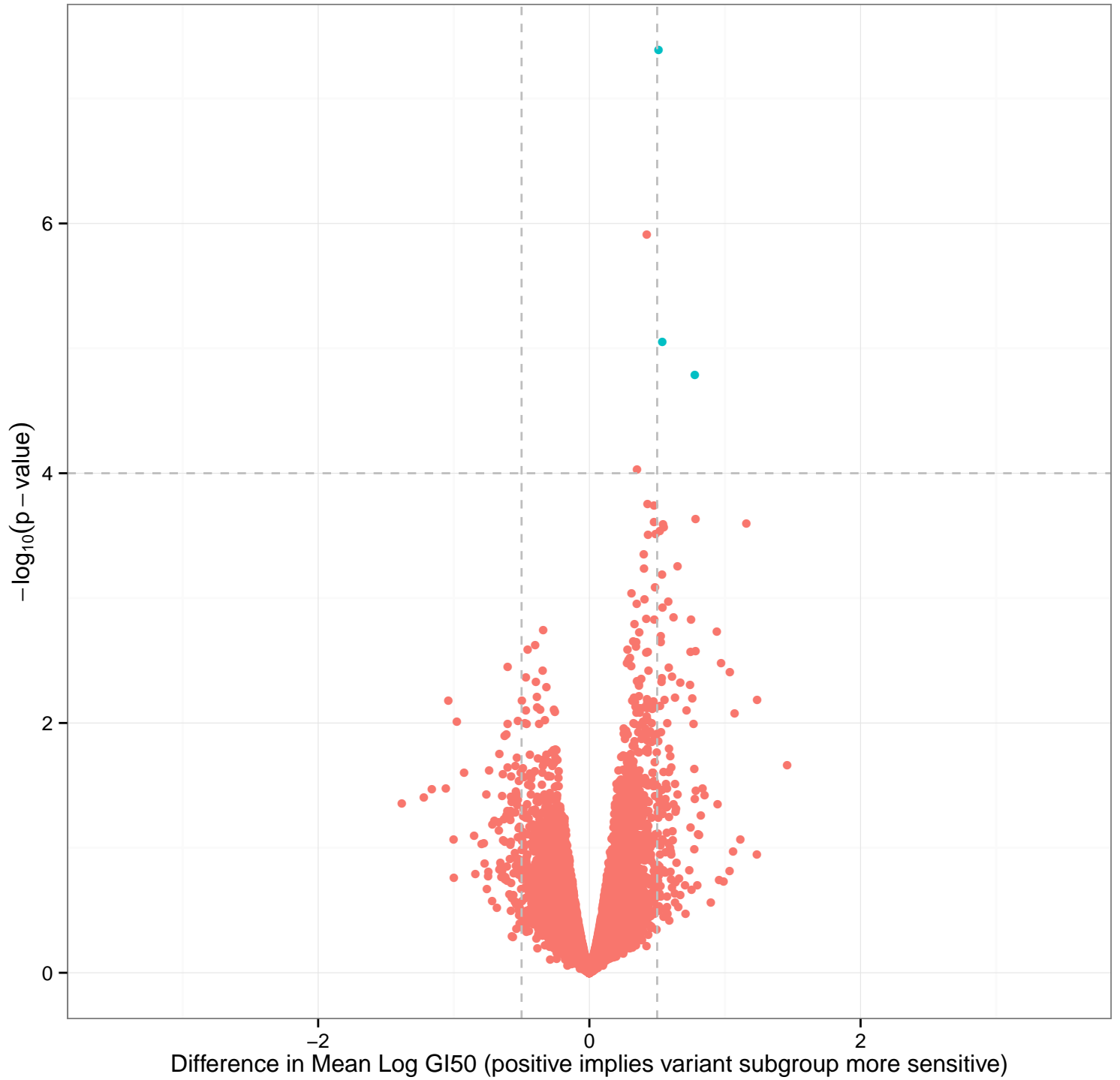
CARD11



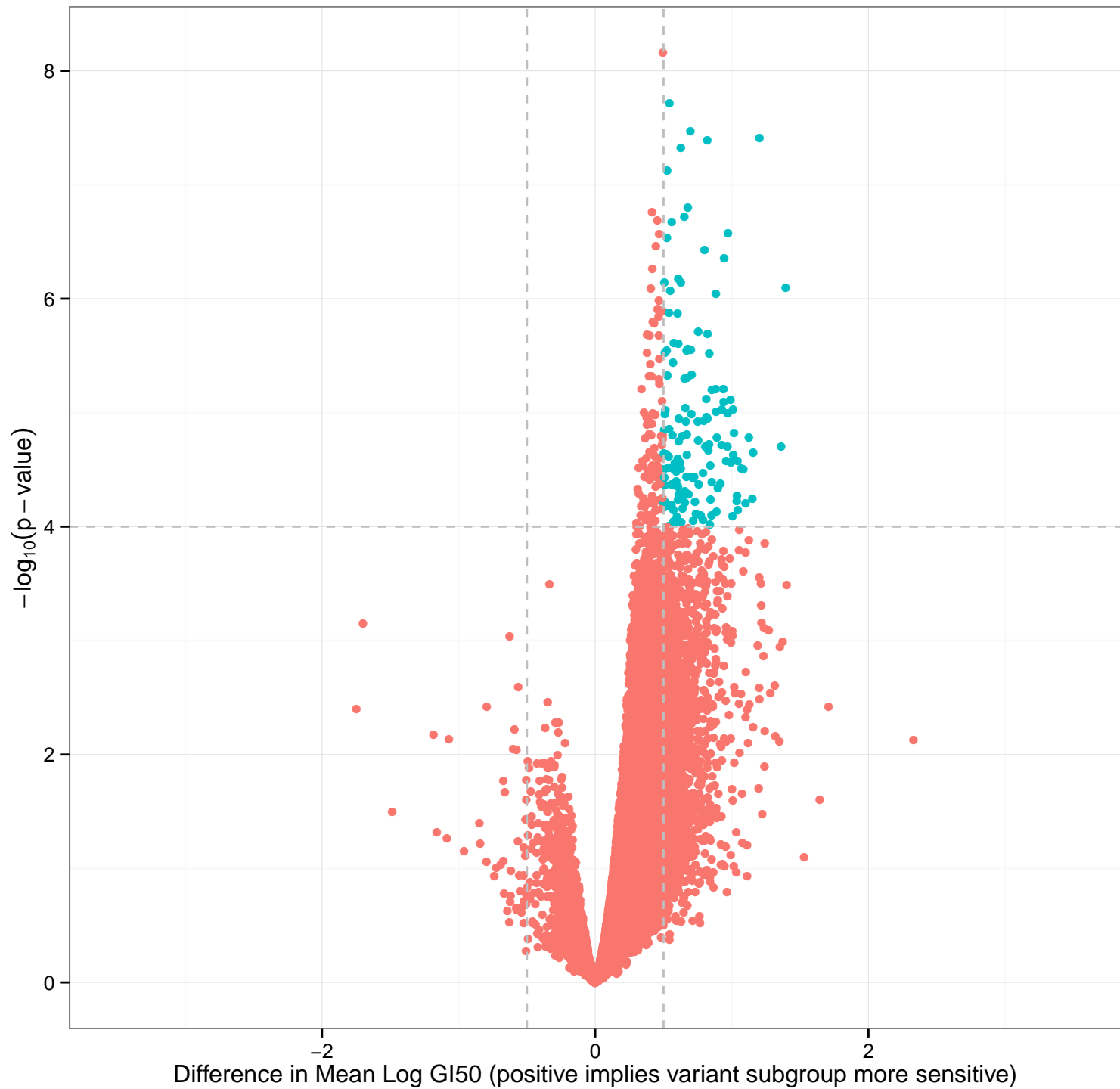
CDH1



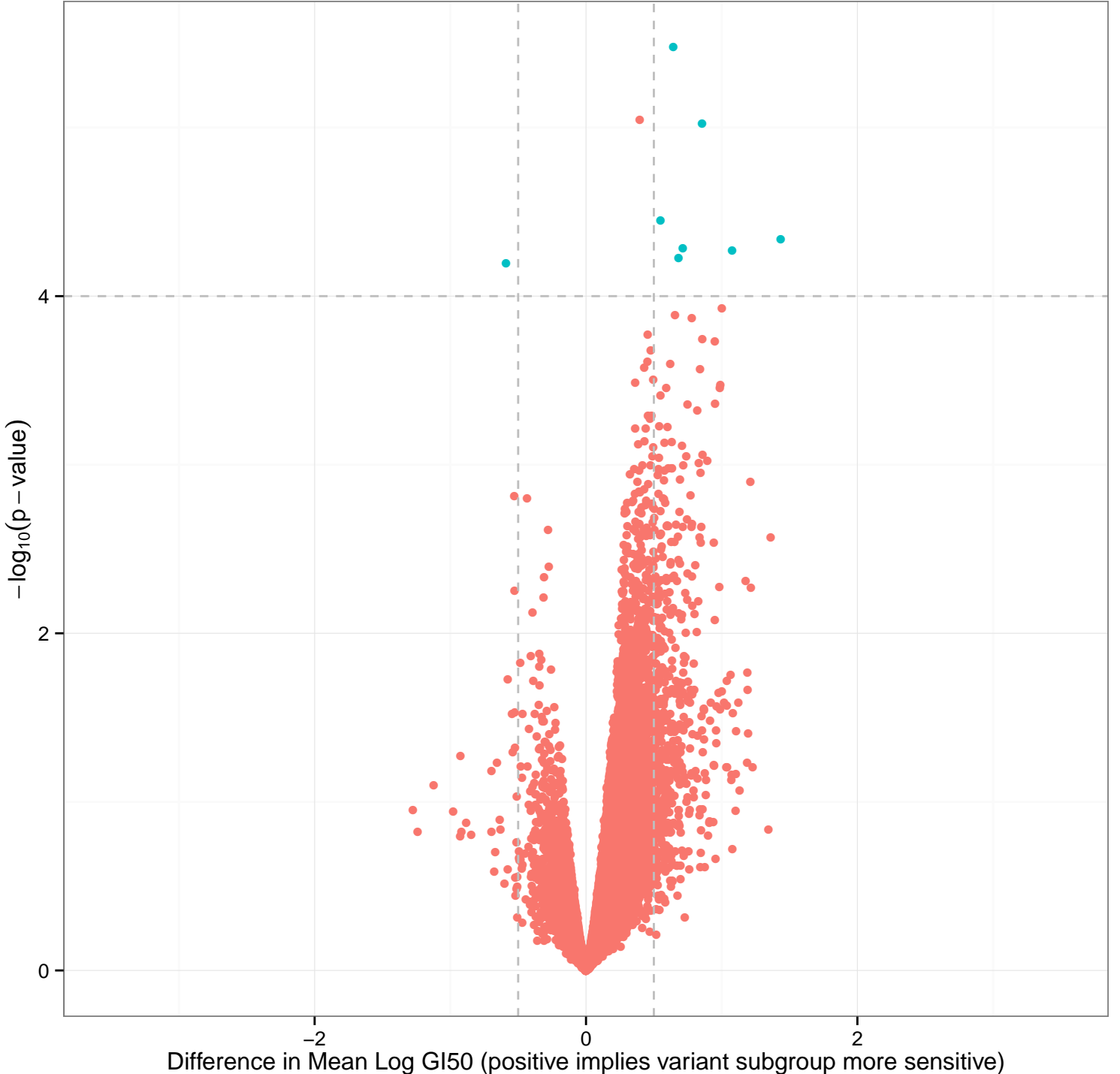
CDH5



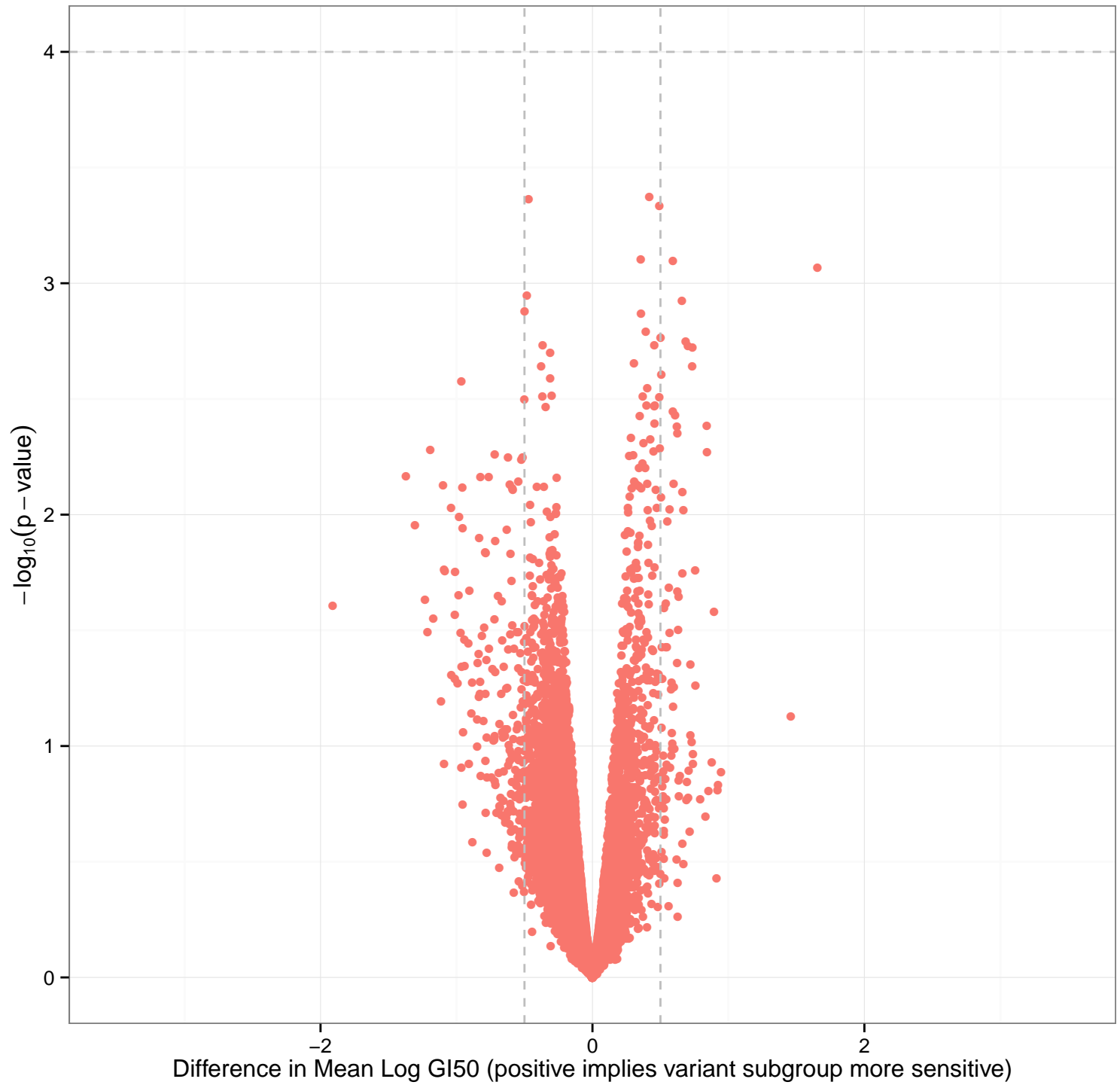
CHD4



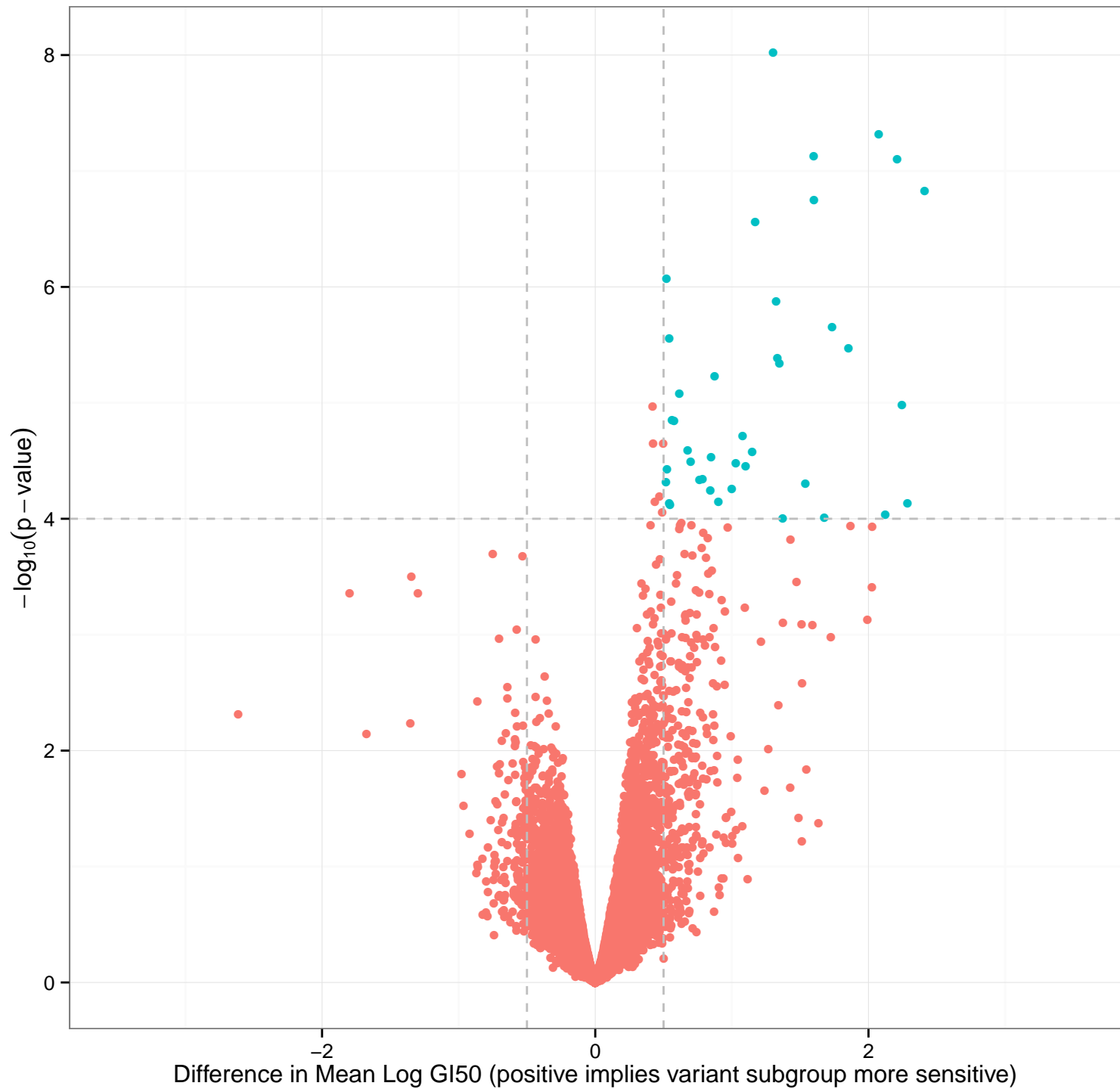
CNTRL



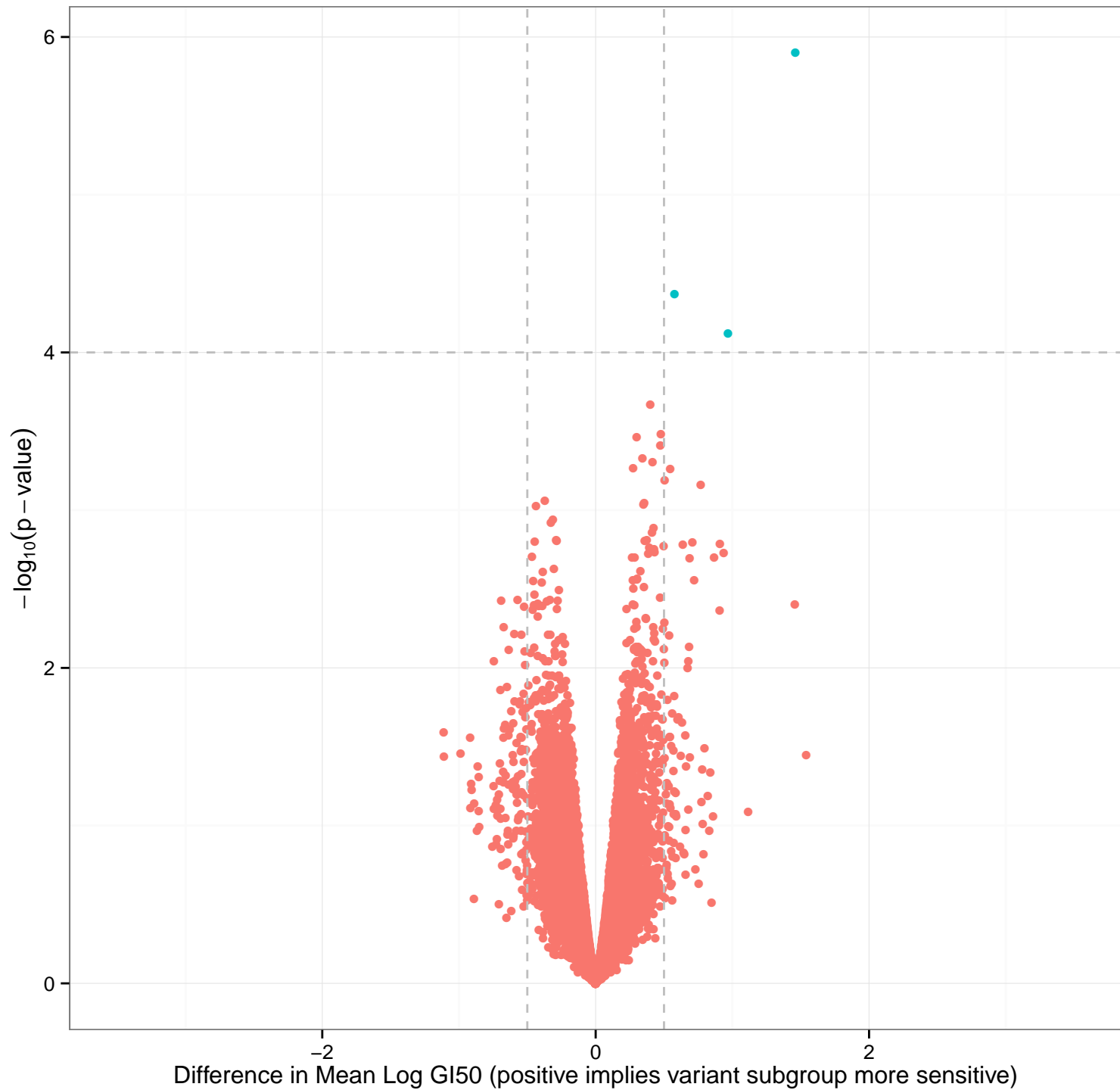
DNMT3A



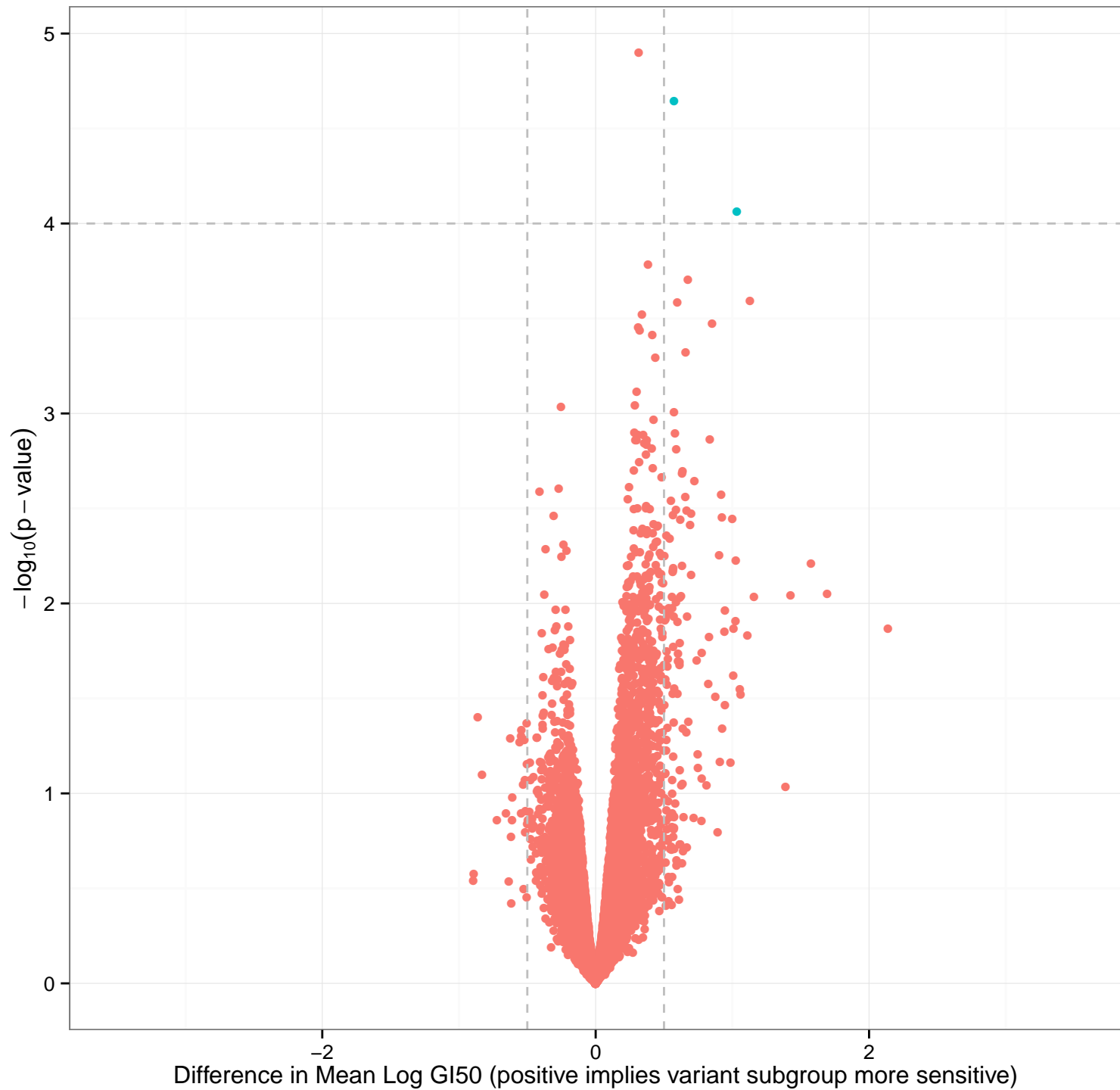
DST



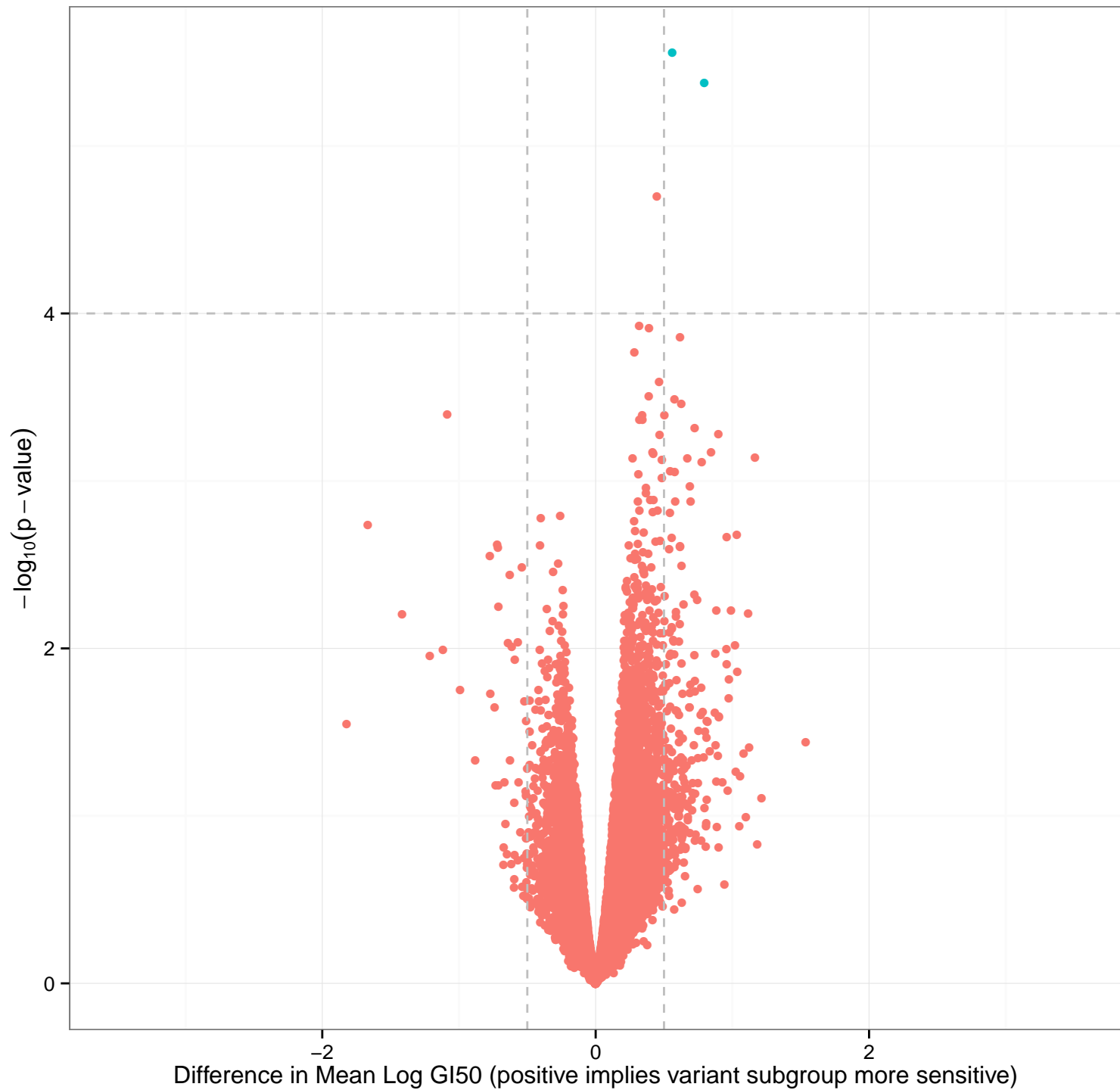
EPHA3



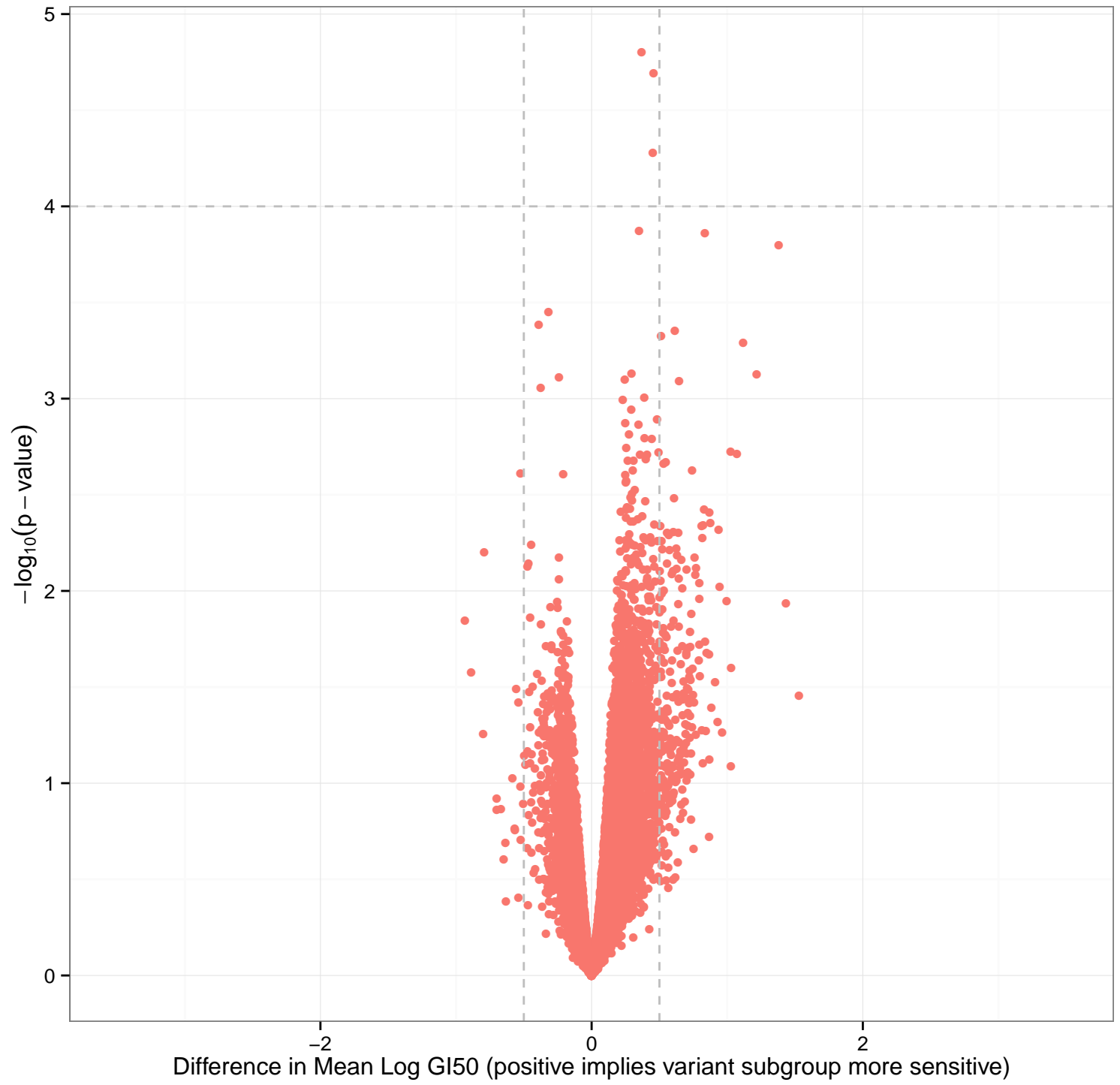
EPHA5



FLT1

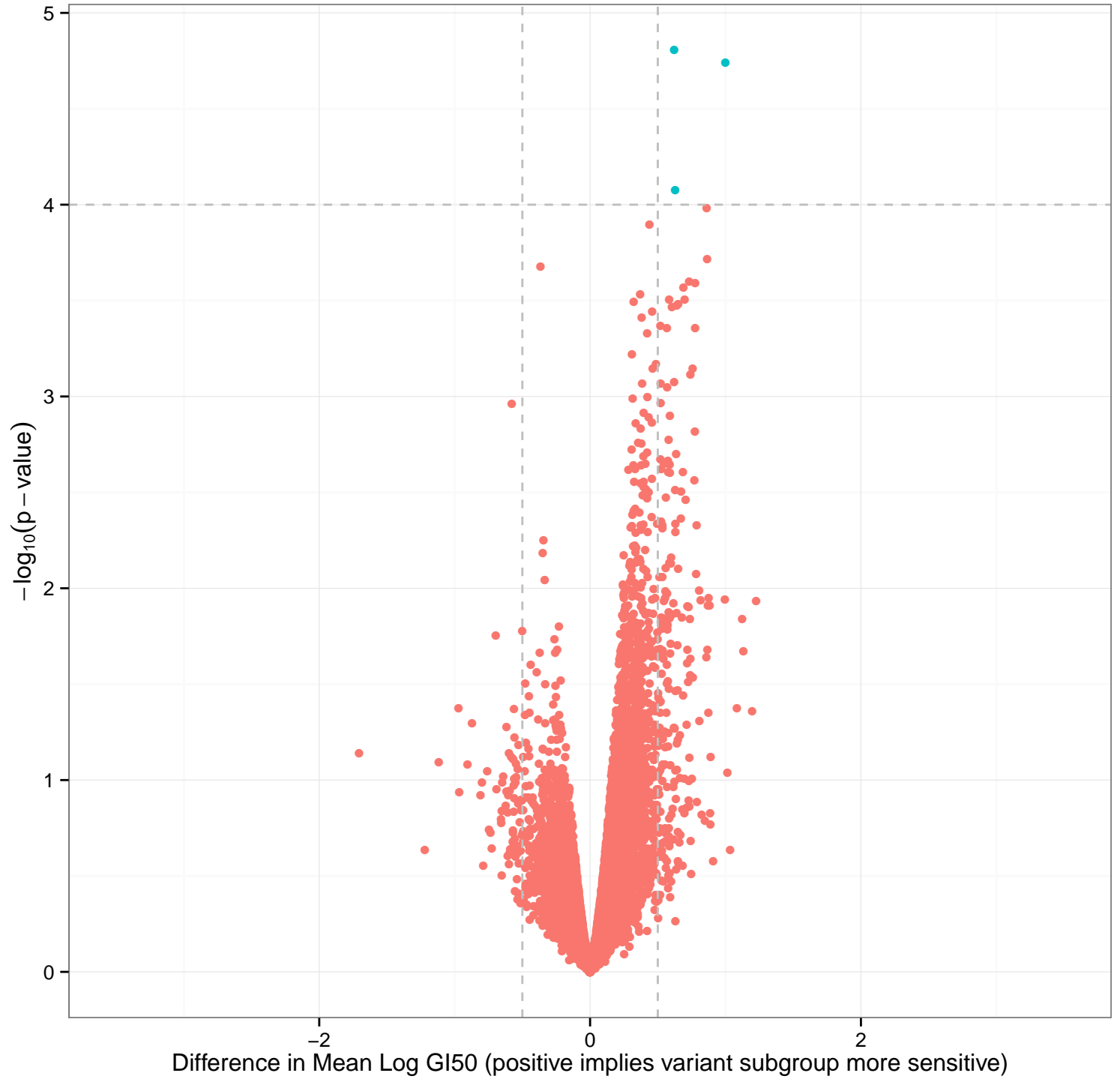


FLT3

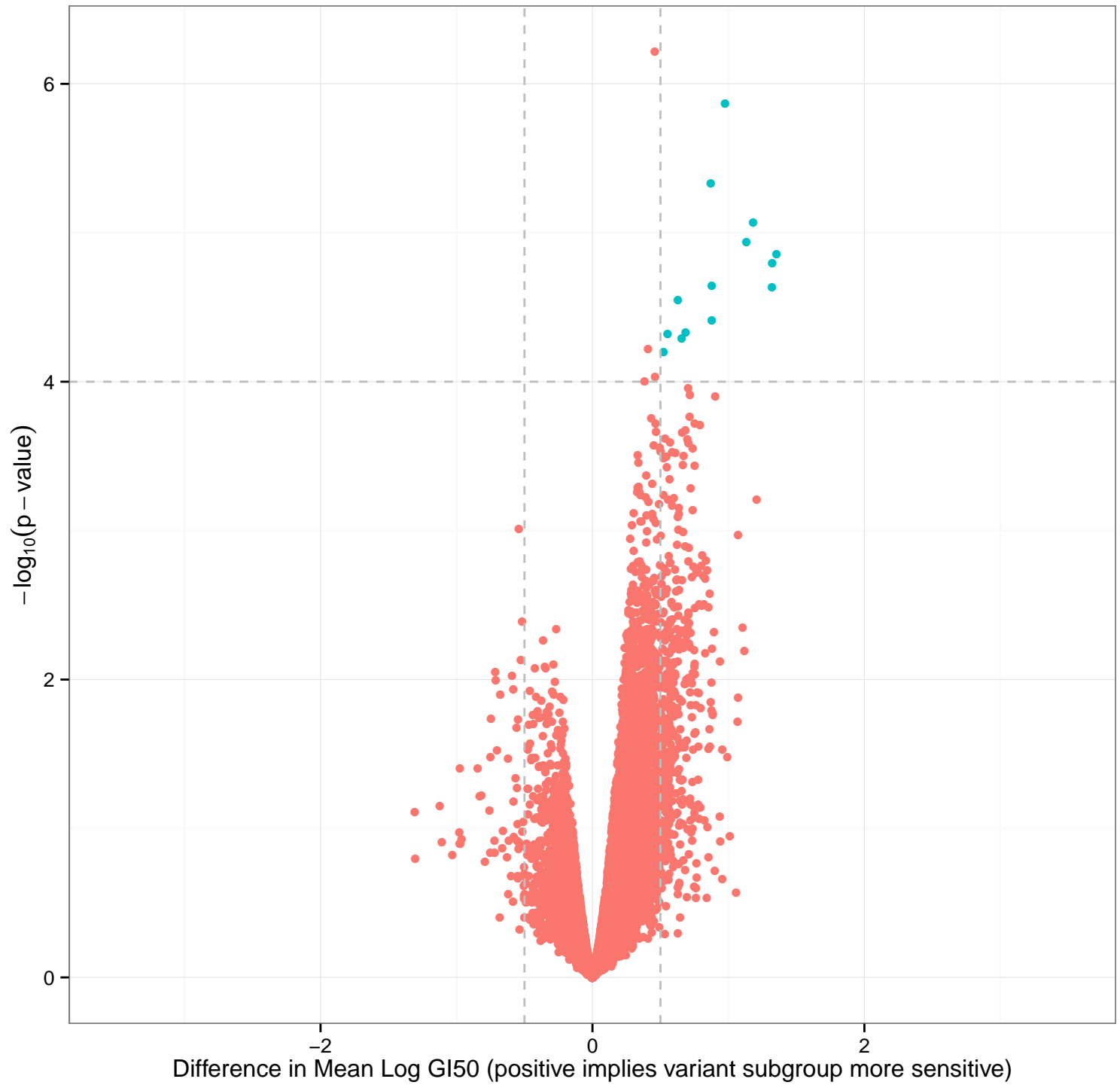


Difference in Mean Log GI50 (positive implies variant subgroup more sensitive)

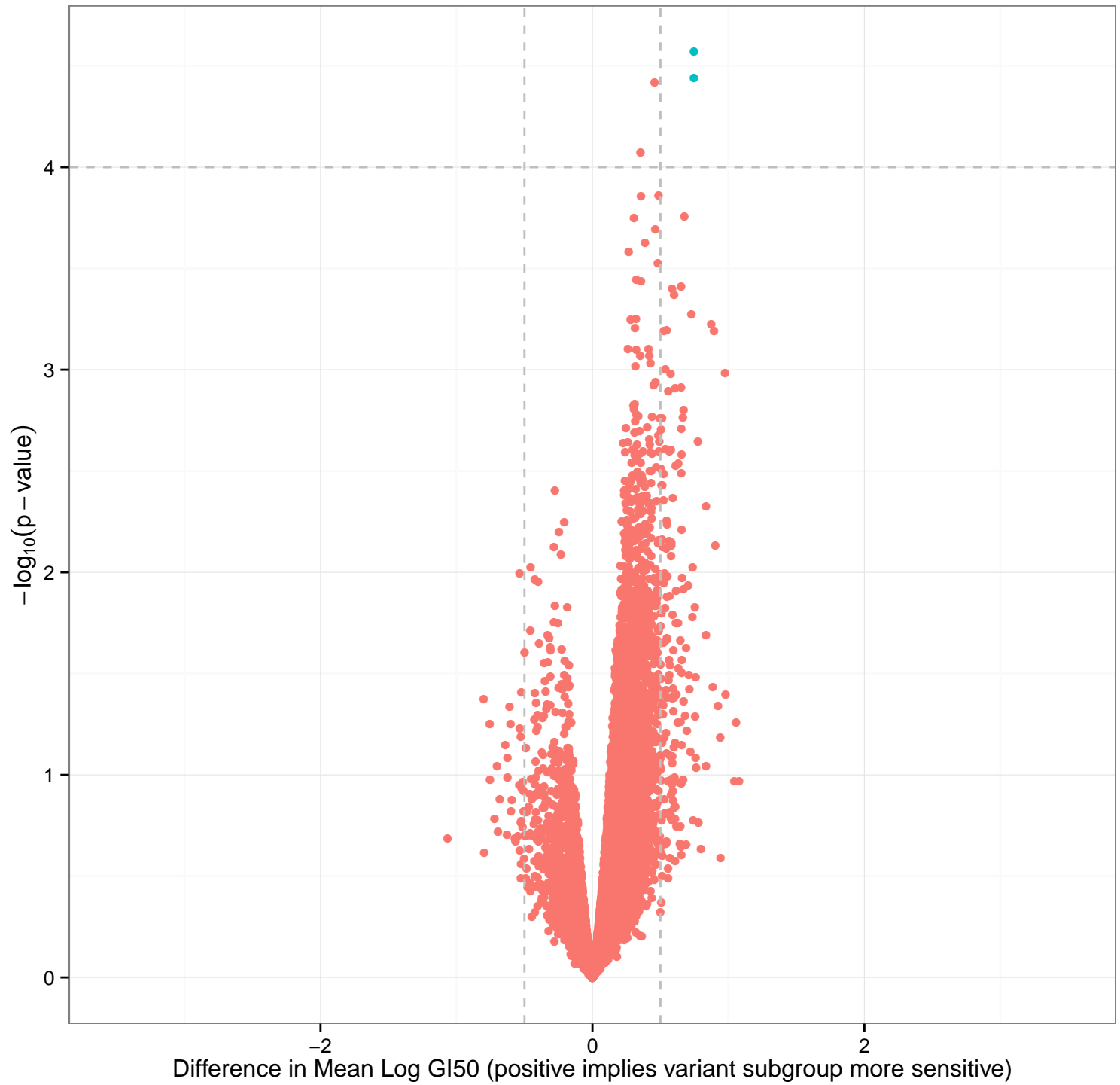
GNAS



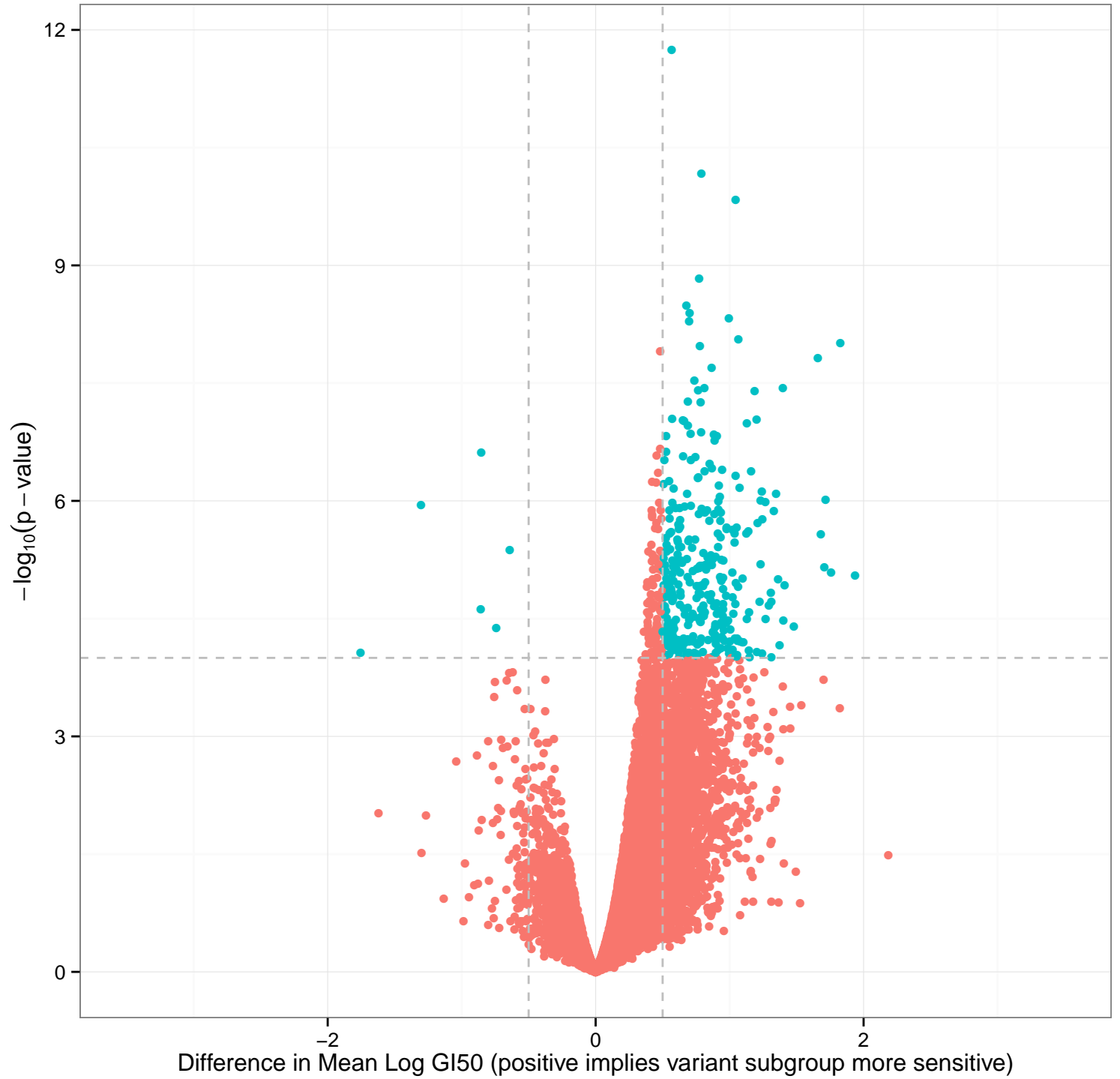
GRIN2A



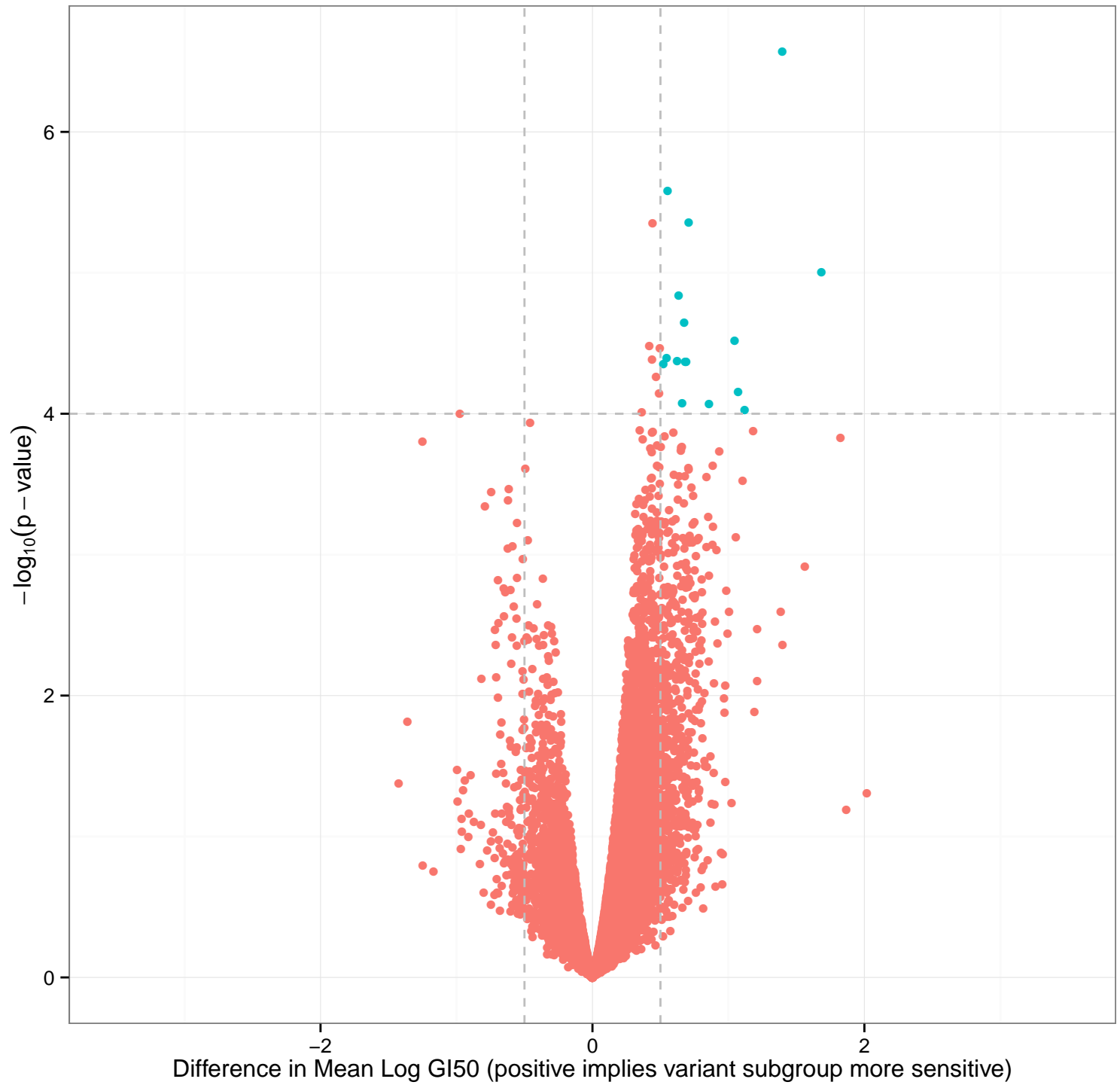
GUCY1A2



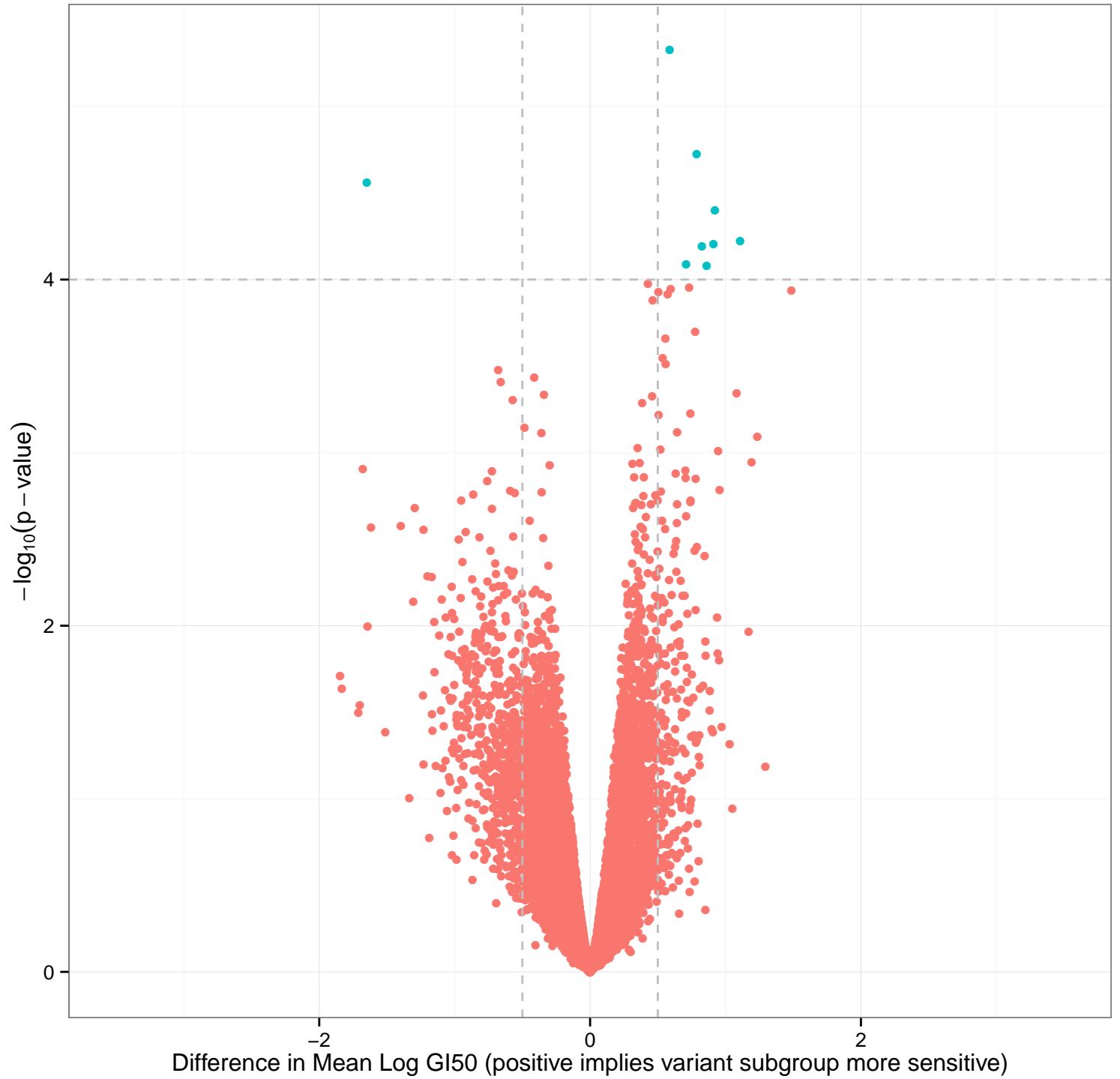
IGF1R



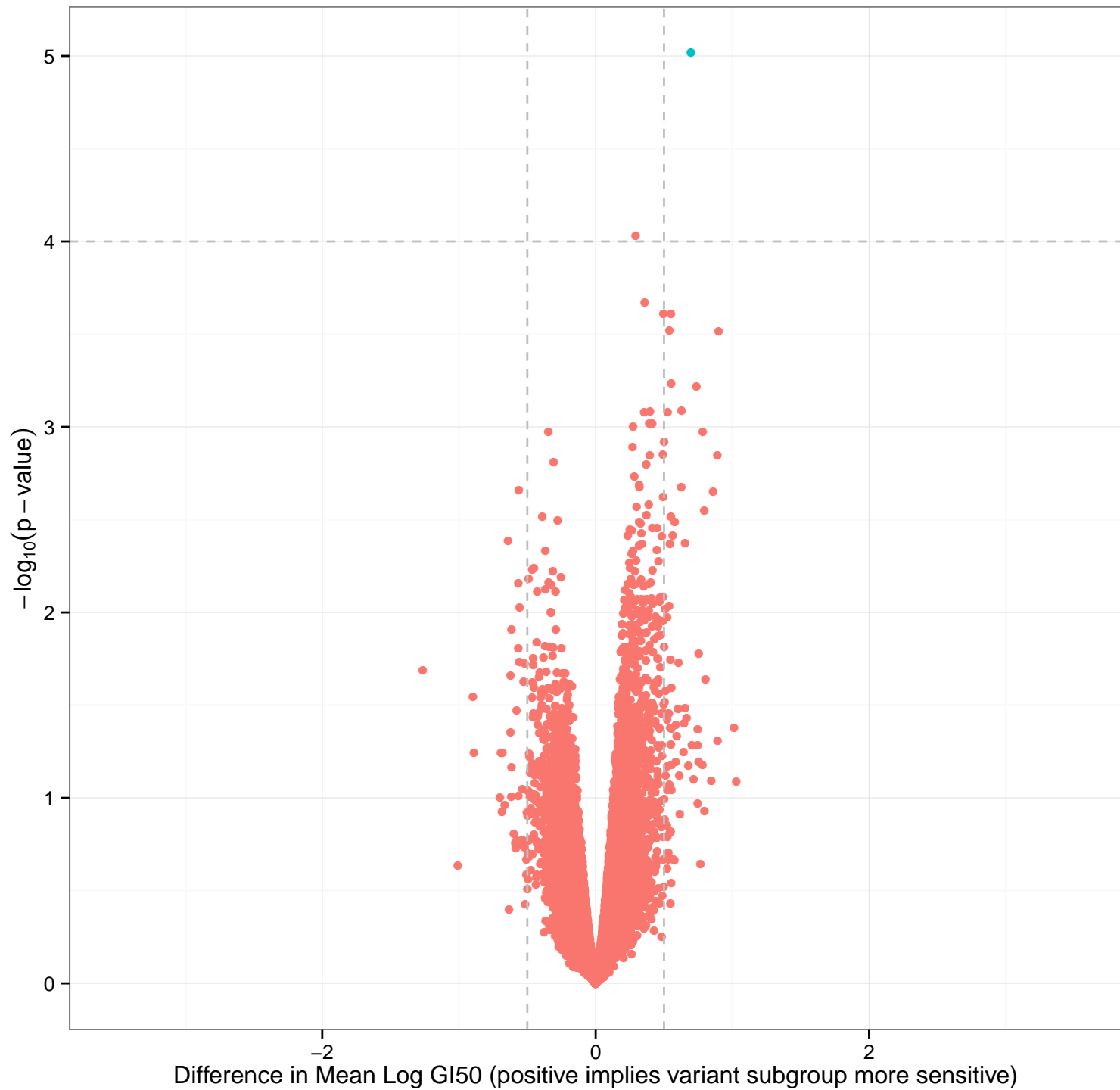
IGF2R



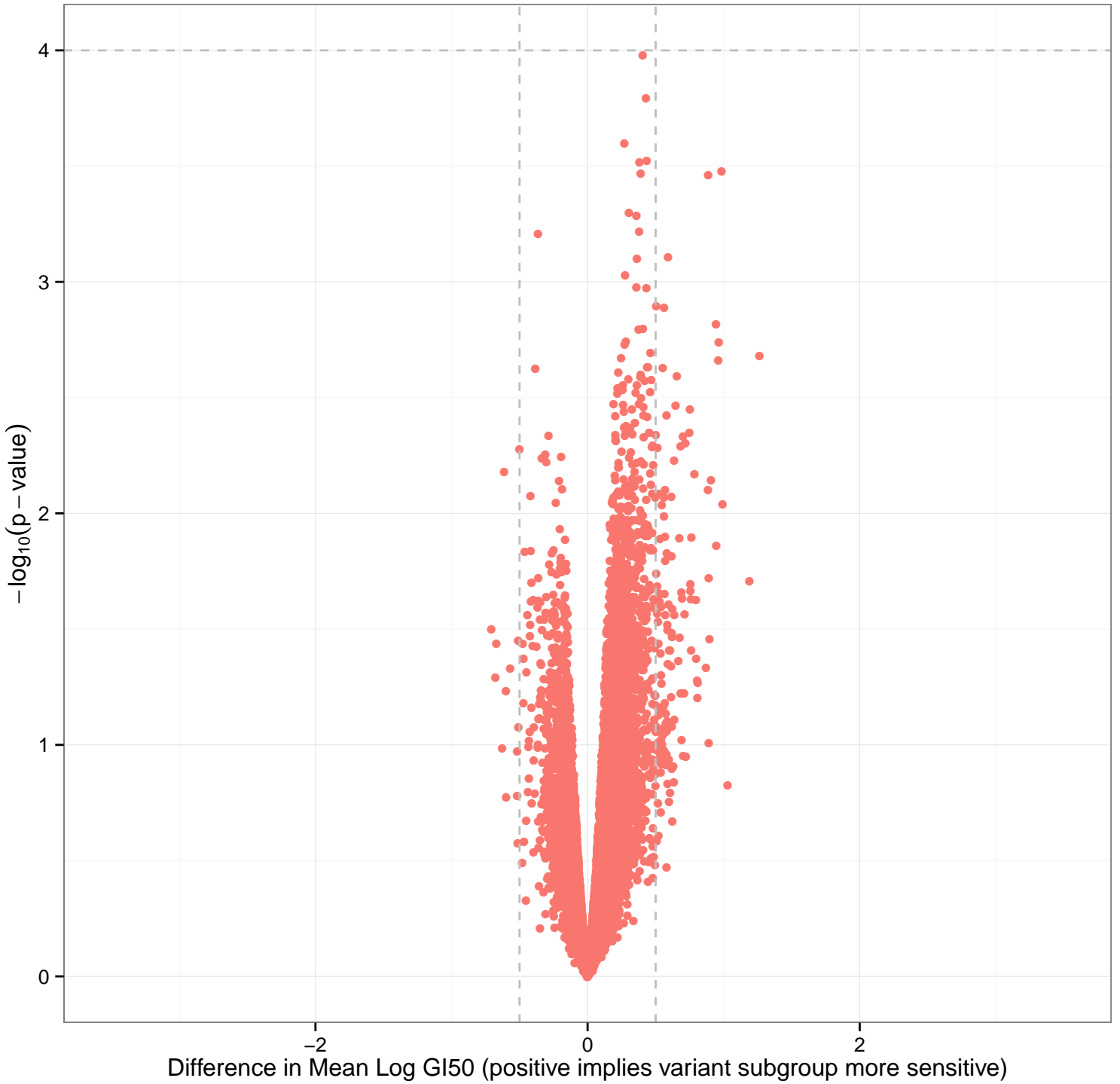
IL4R



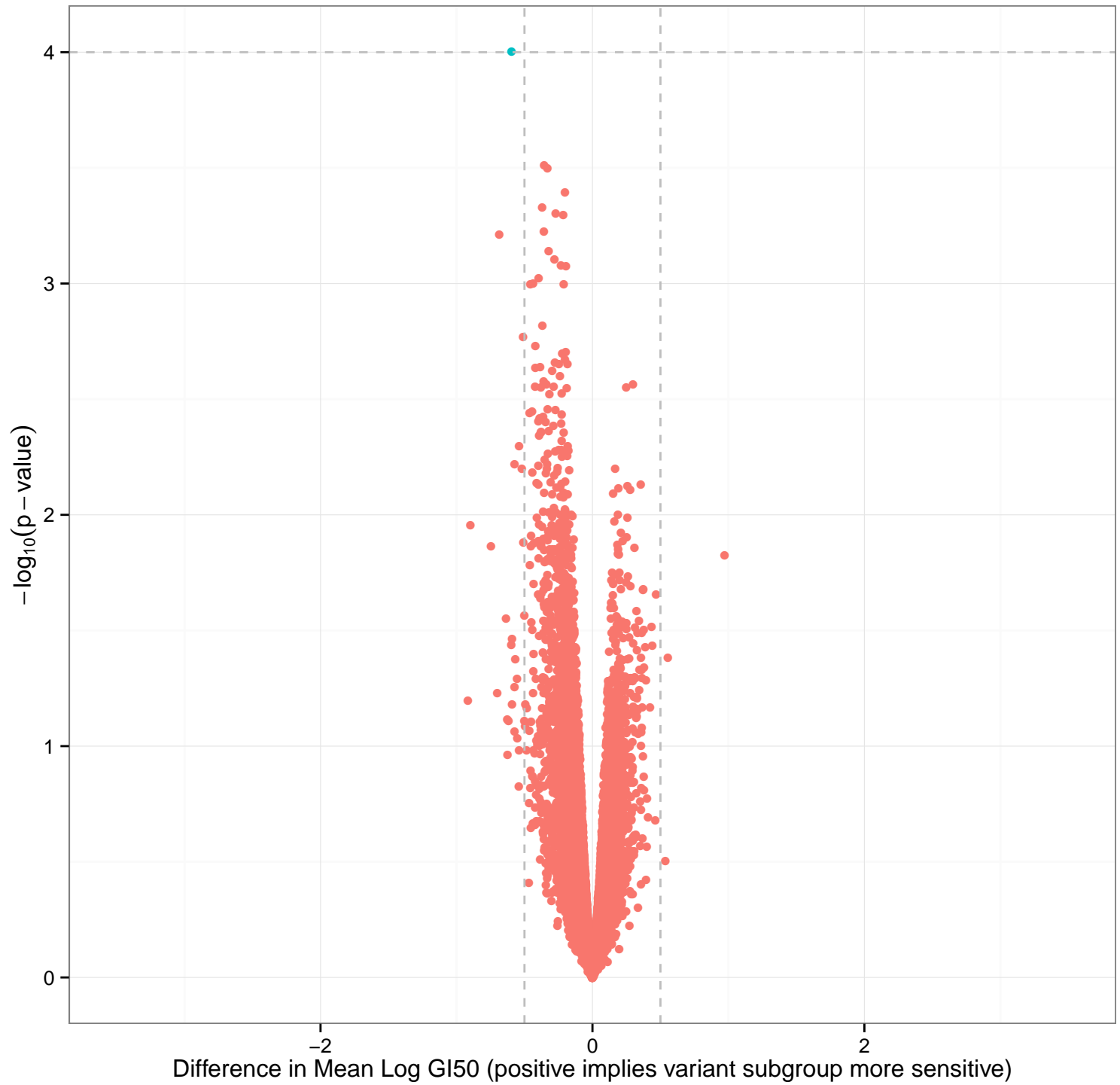
KDR



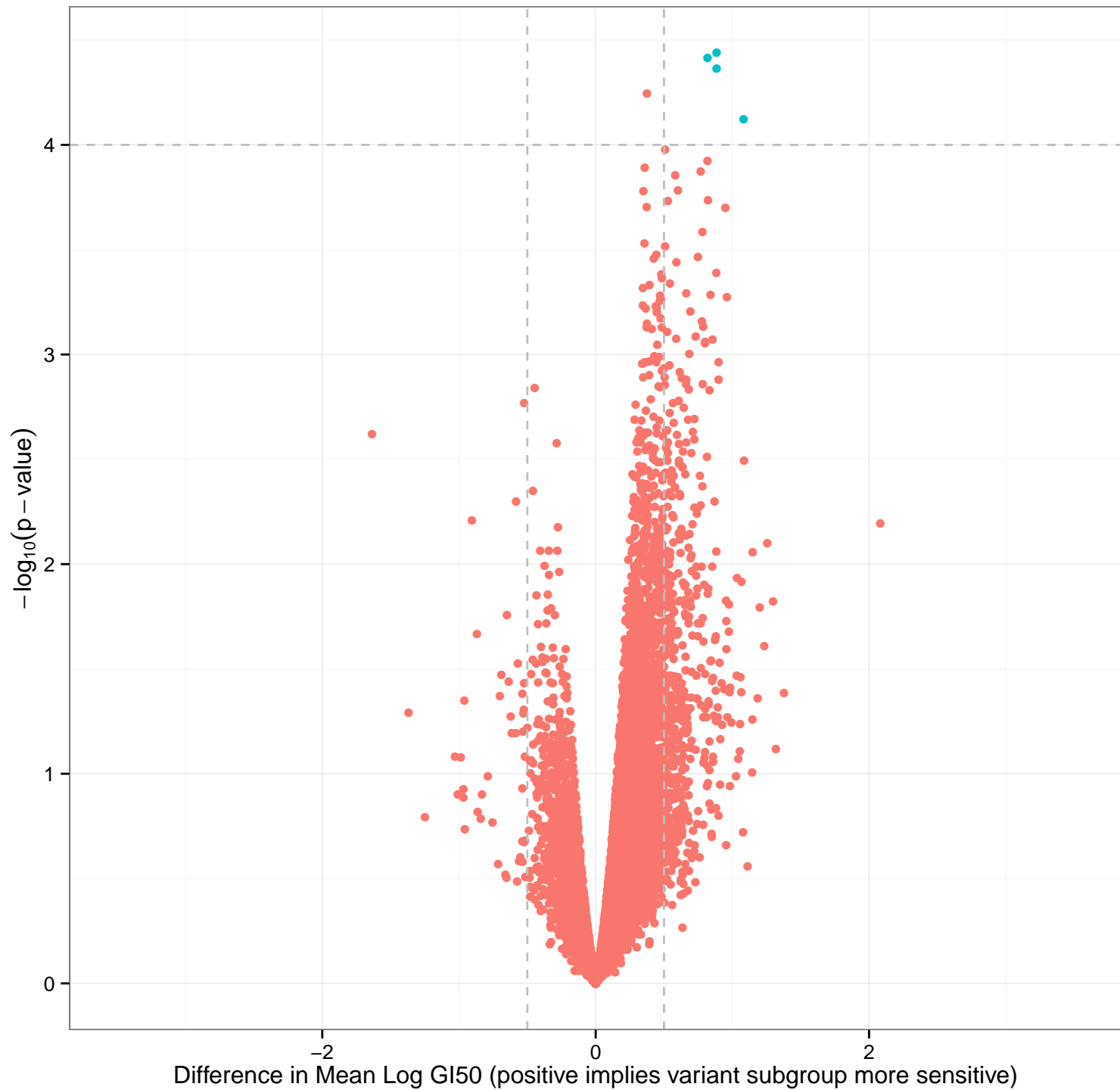
KRAS



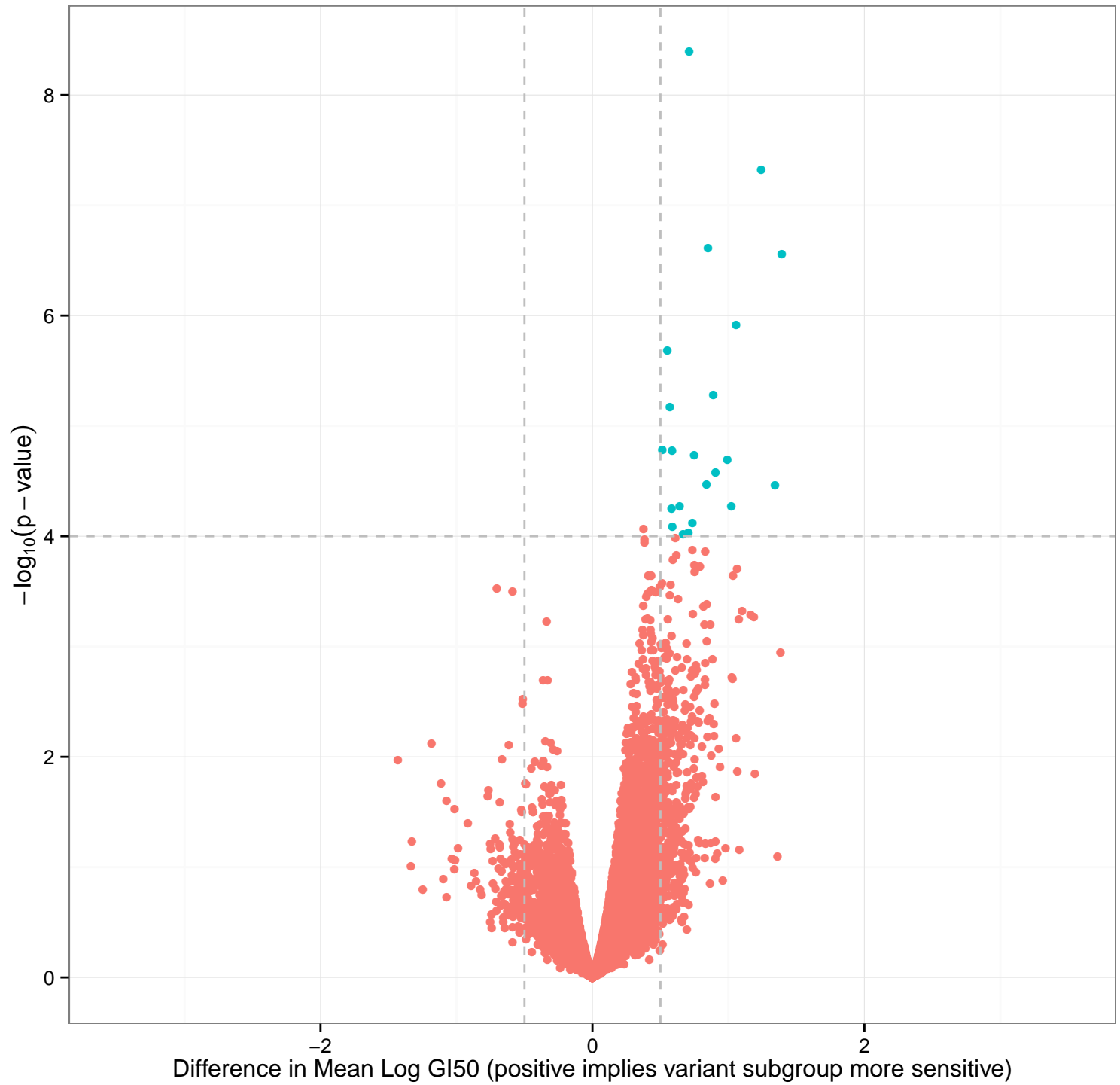
LRP1B



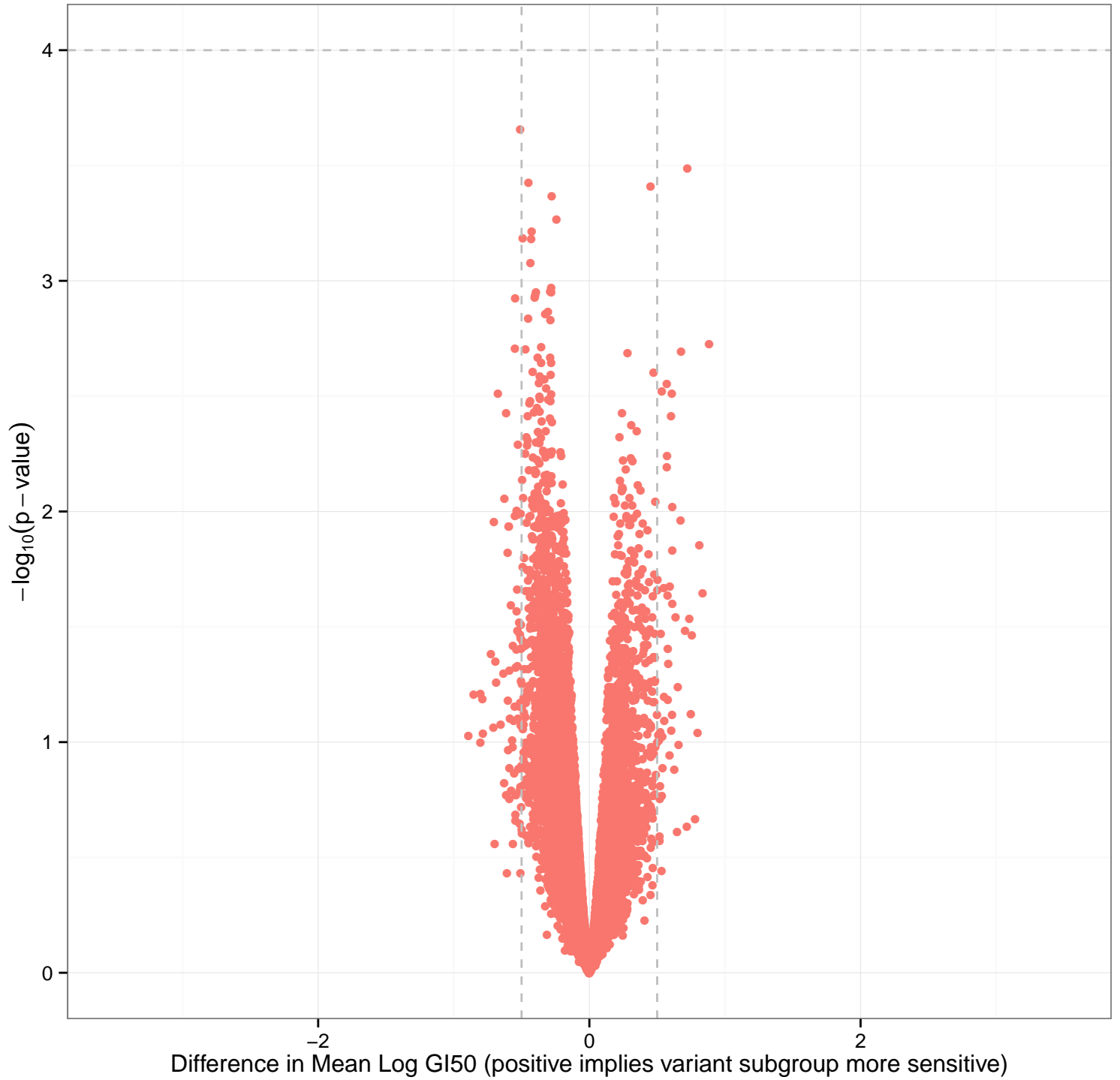
MLH1



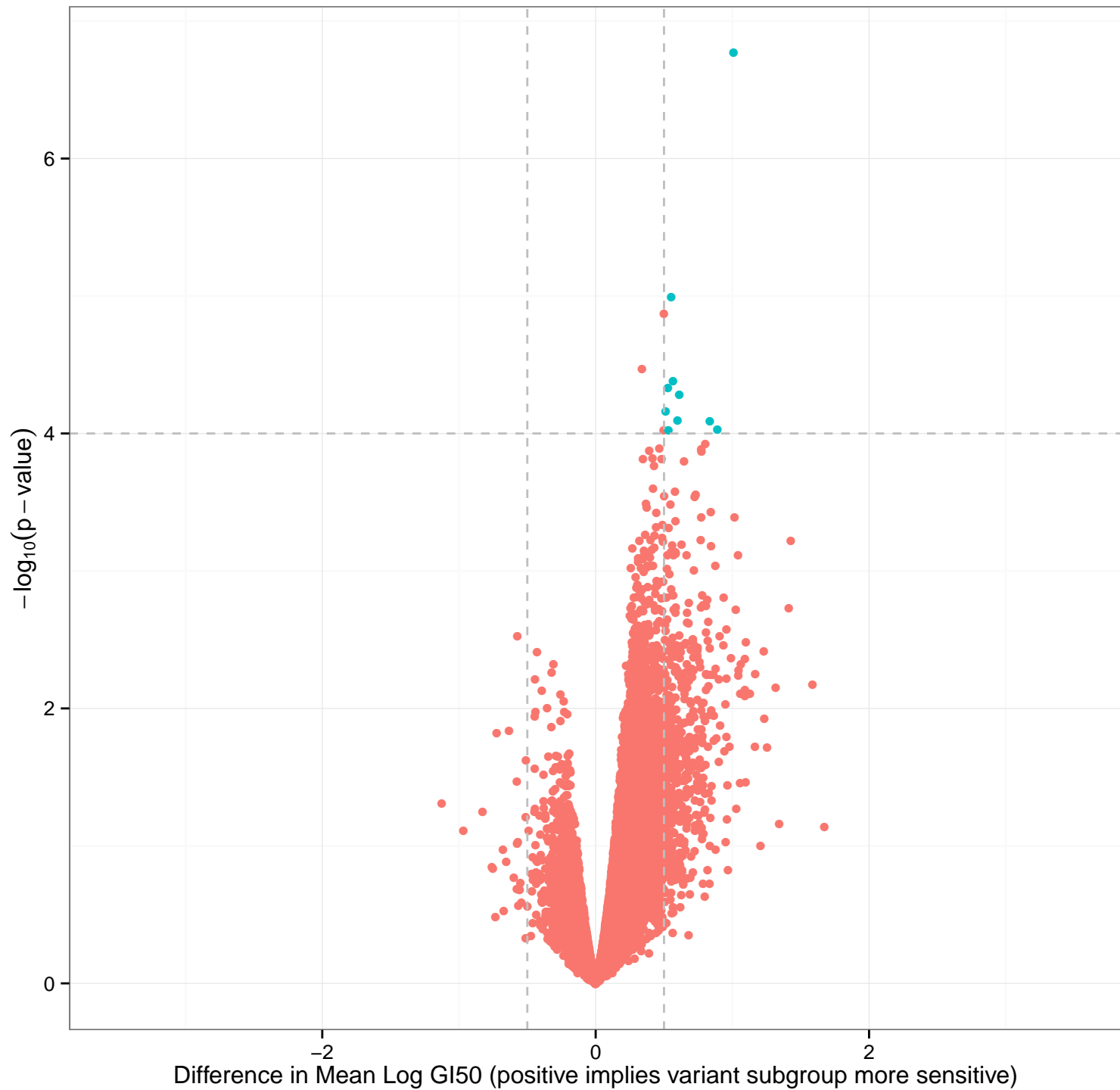
MSH2



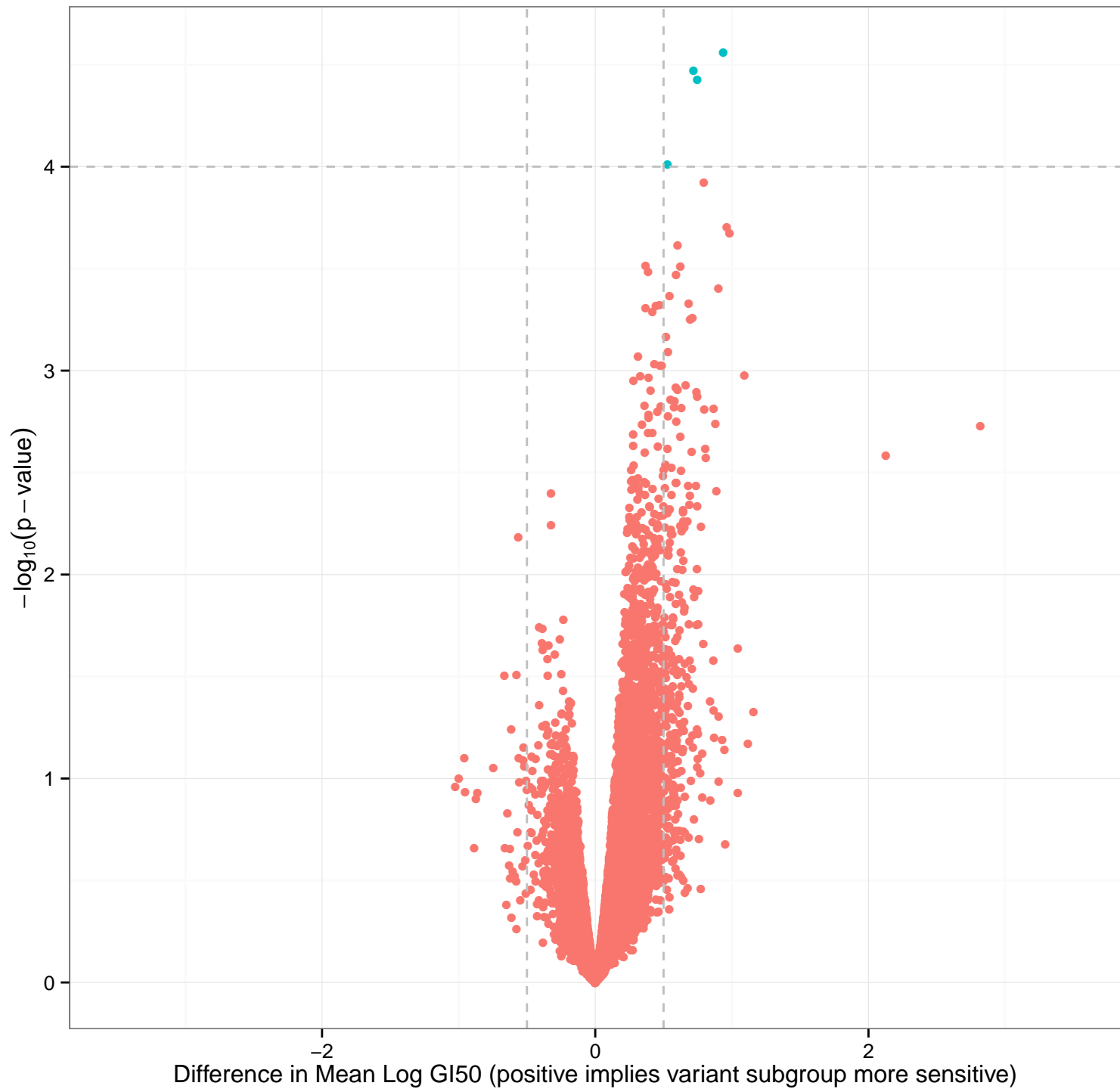
NF1



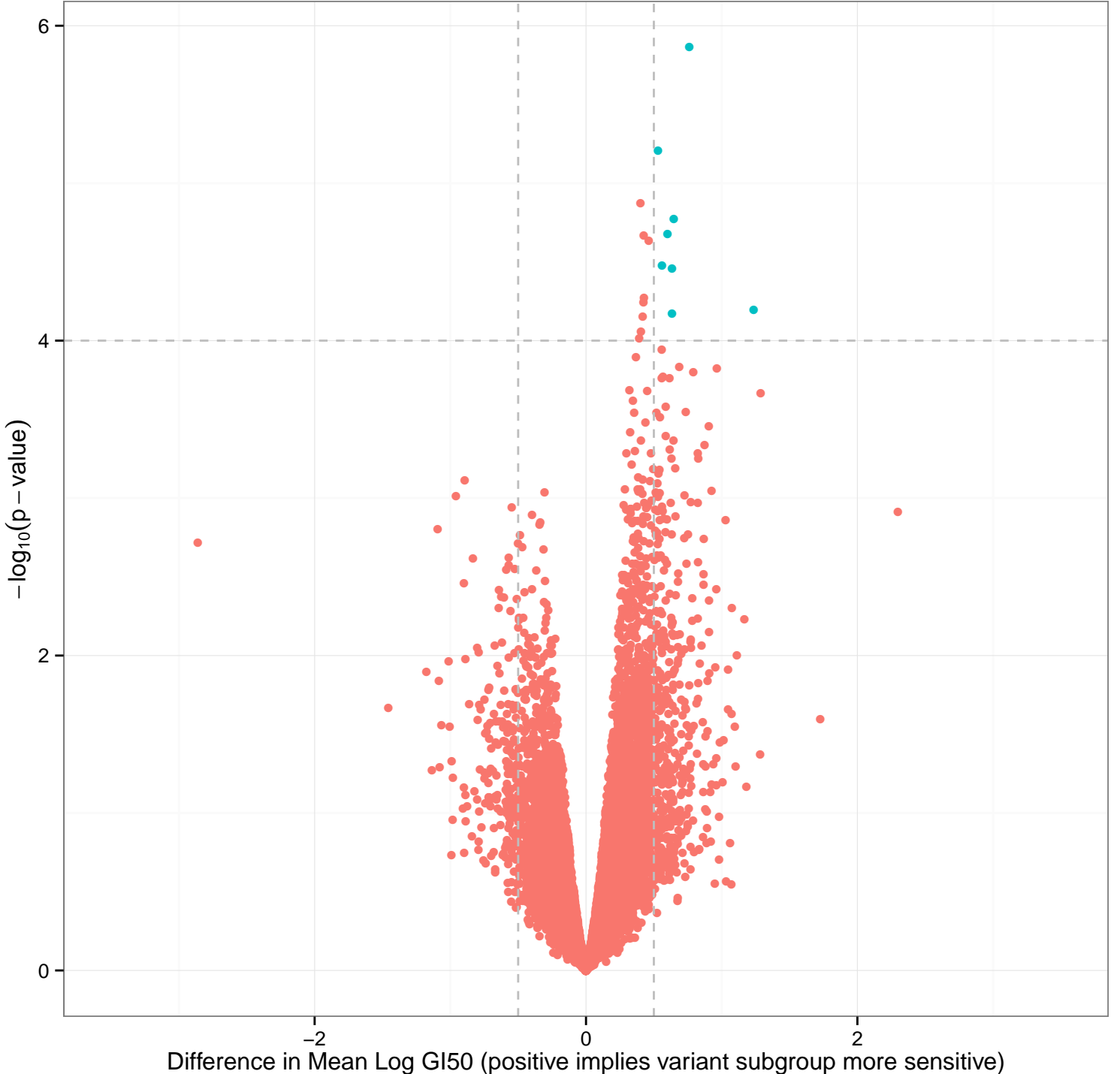
NOTCH1



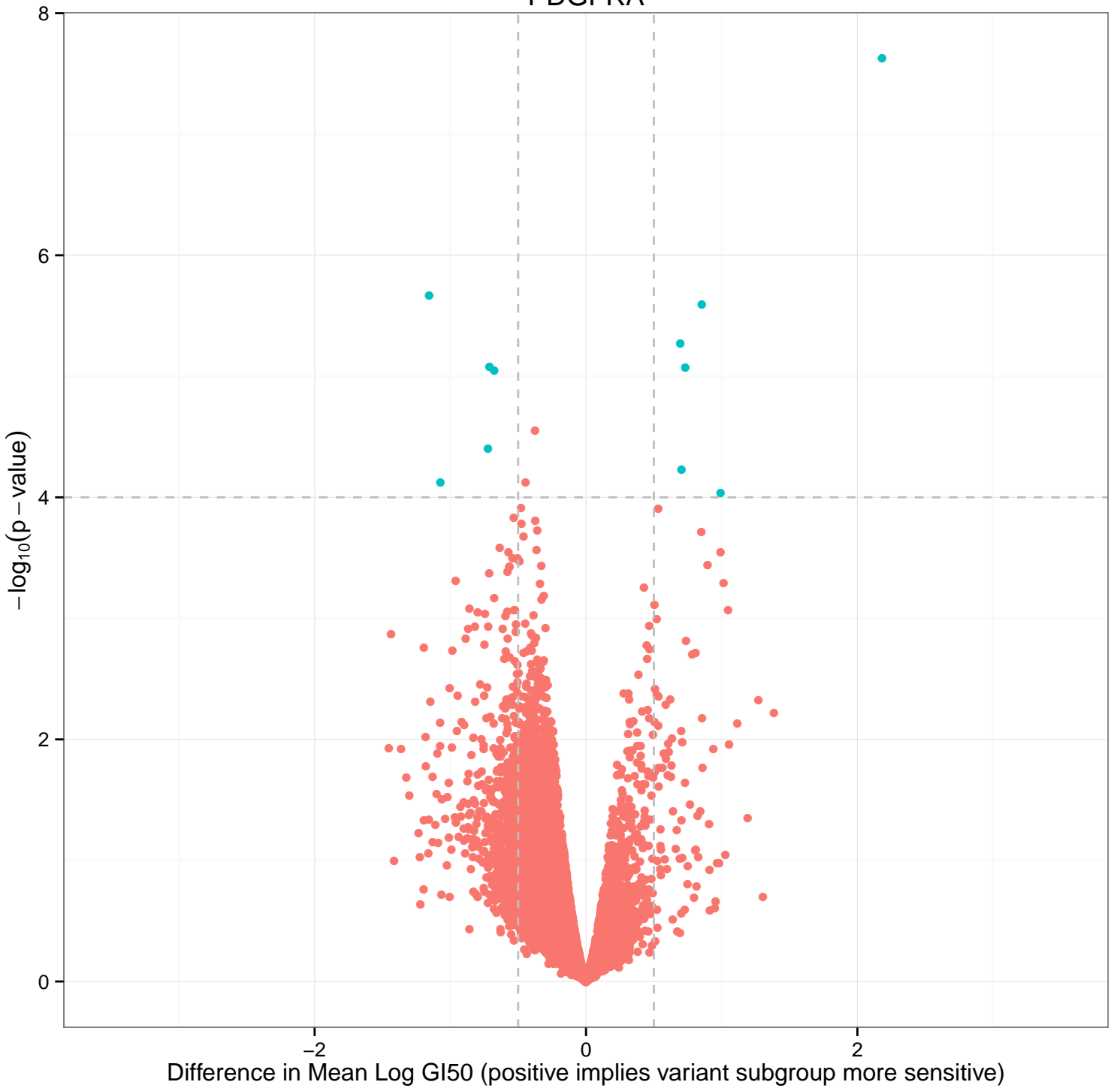
NOTCH3



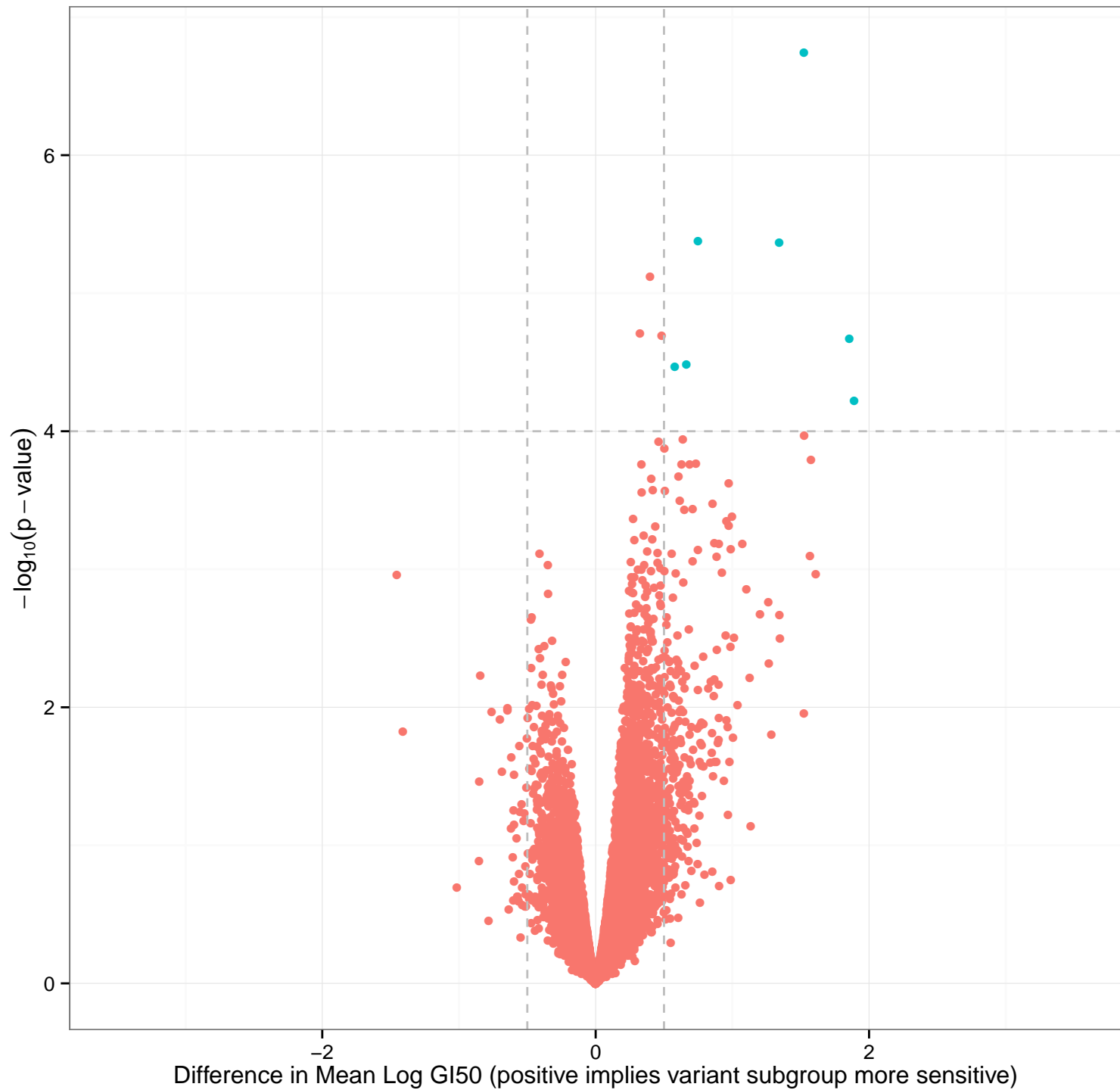
NTRK2



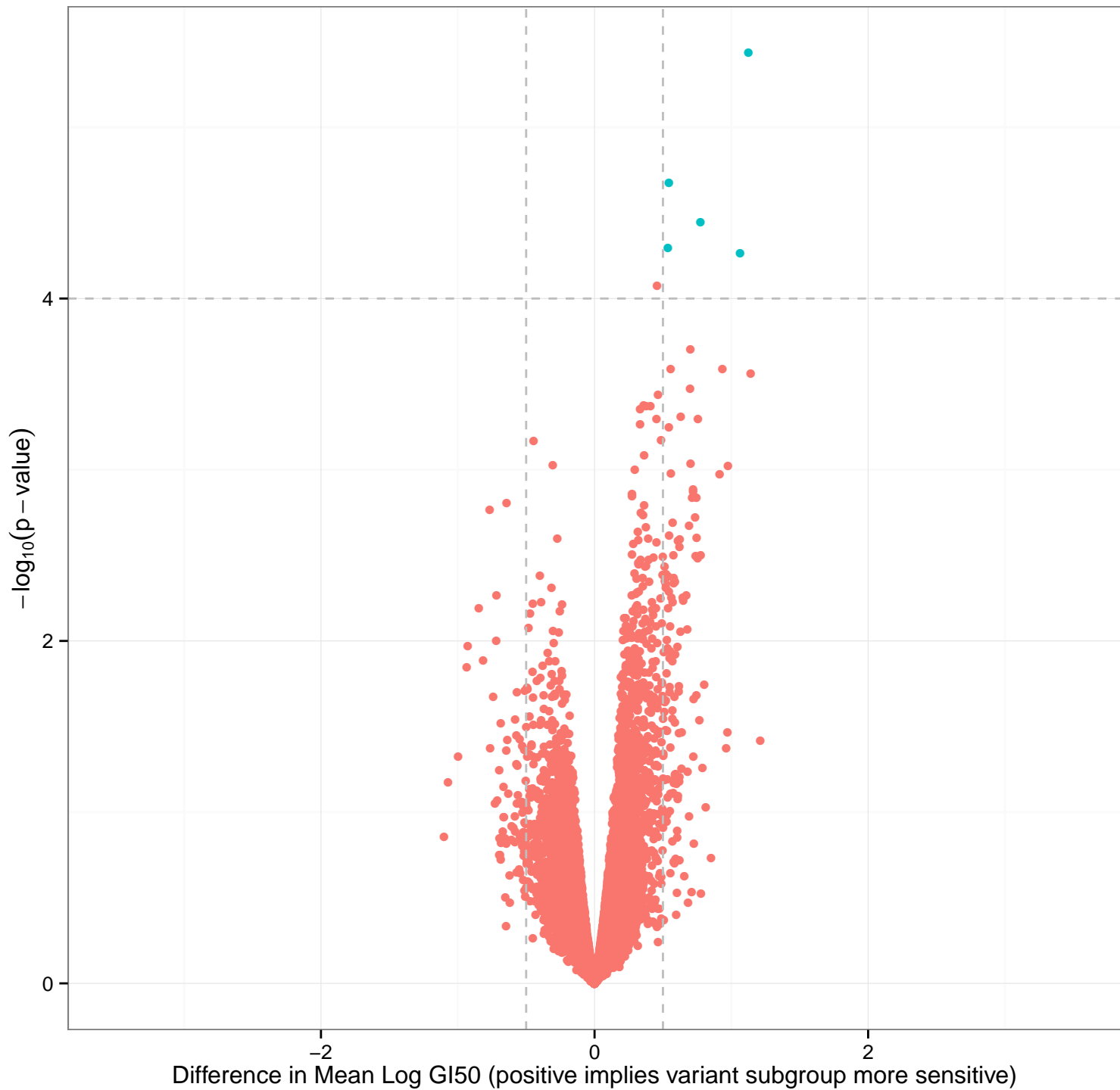
PDGFRA



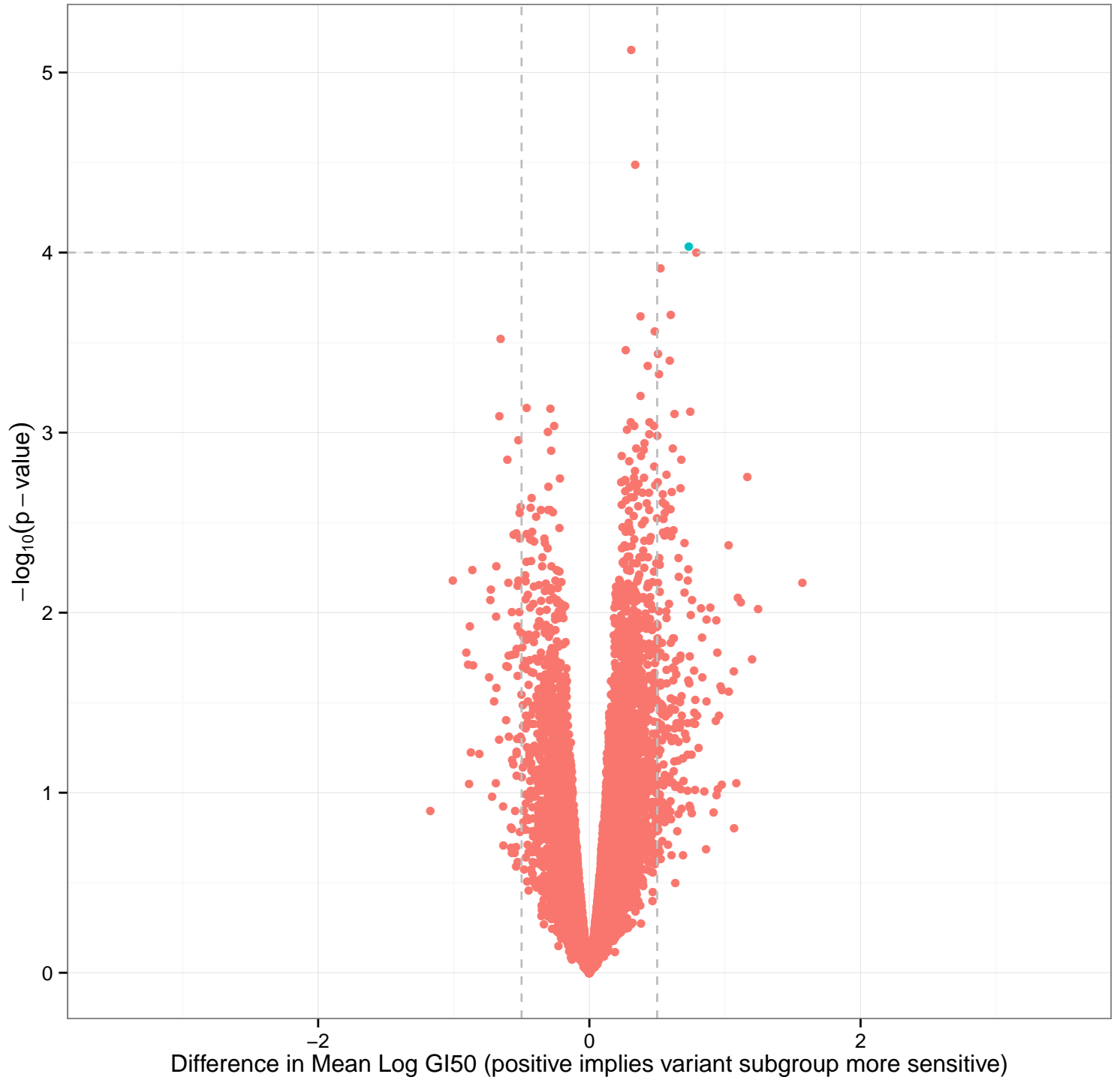
PIK3CA



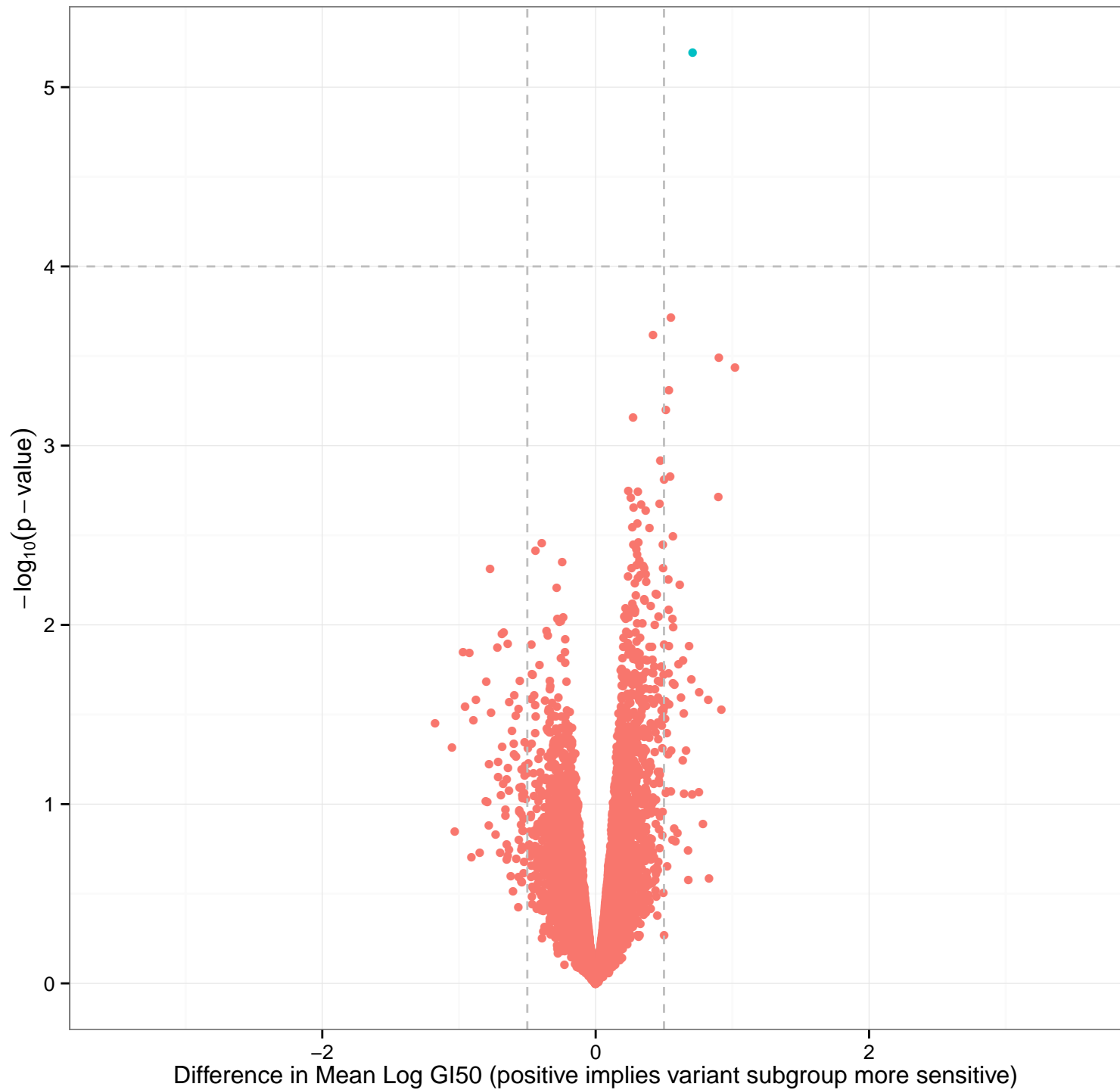
PIK3CB



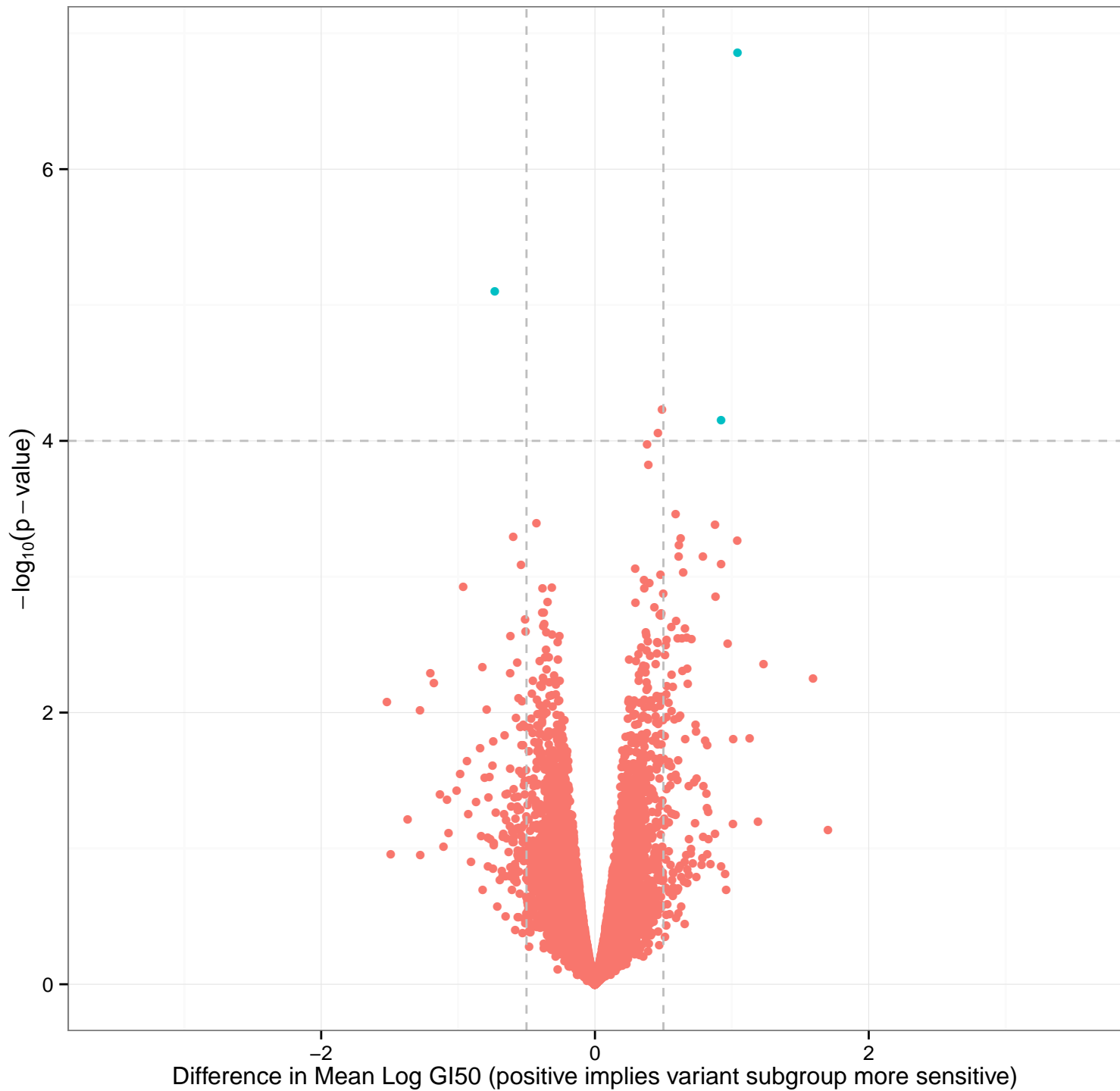
PKHD1



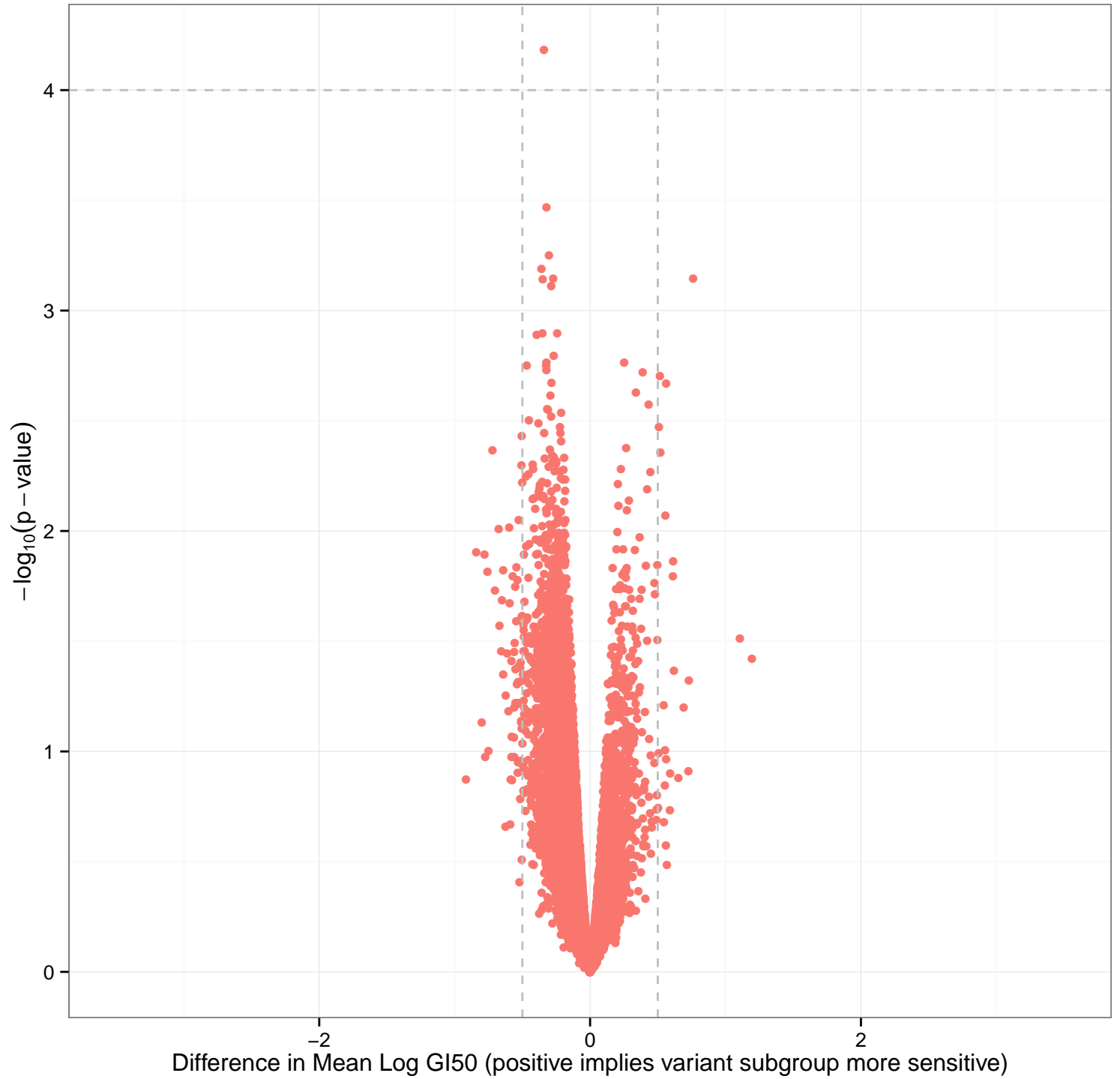
POLE



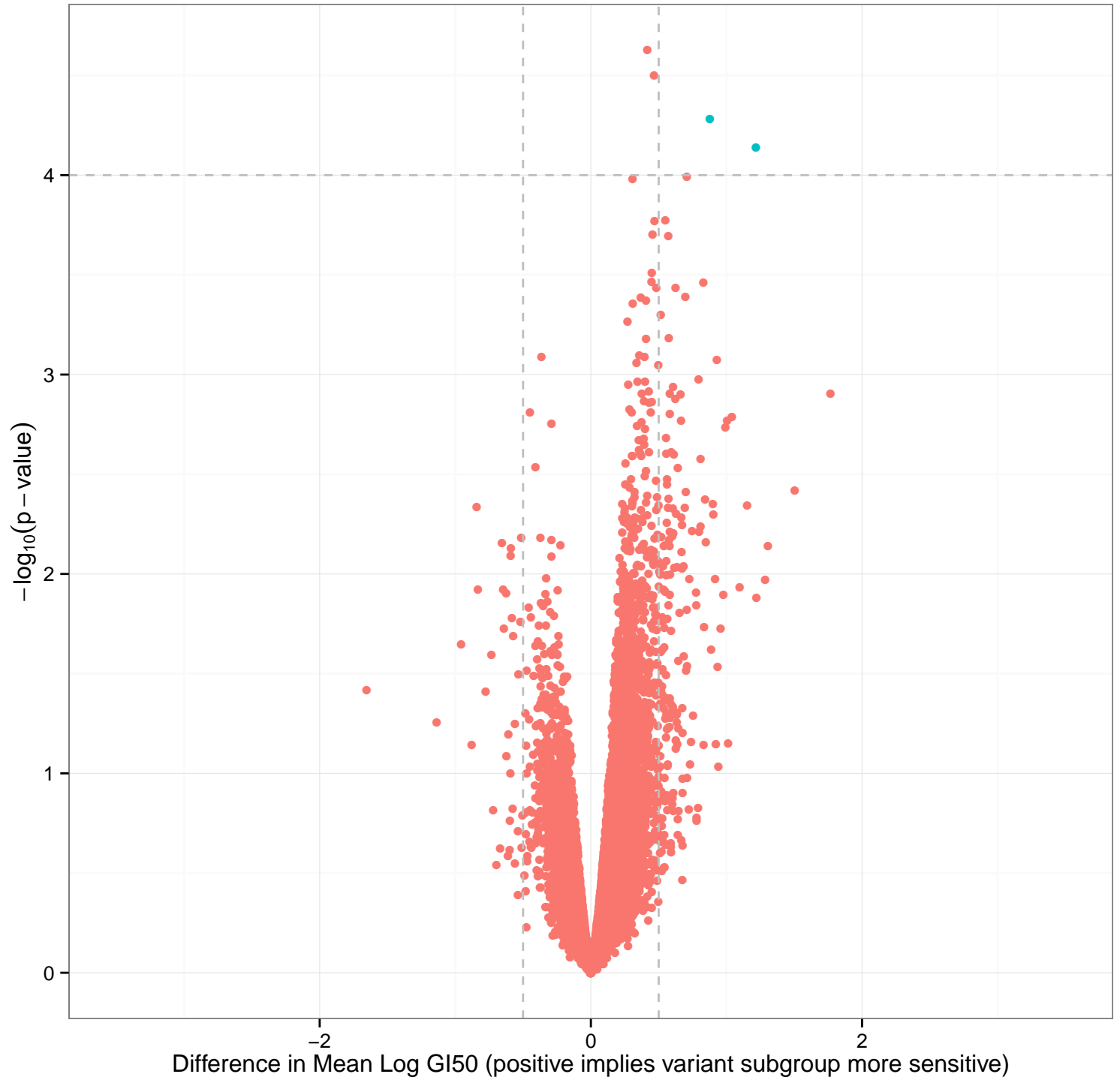
RET



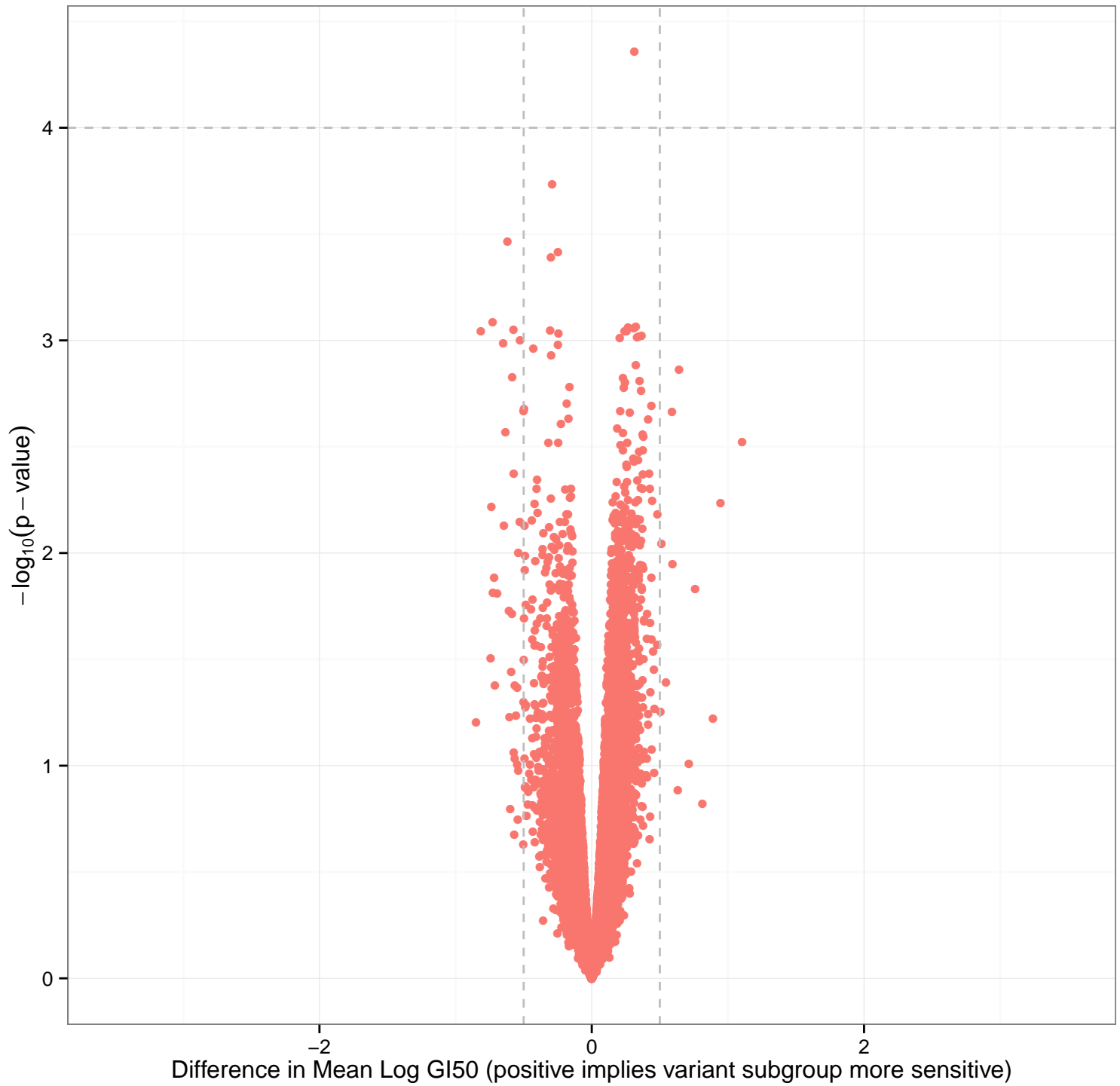
RETNLB



SMARCA4



TP53



TSC2

