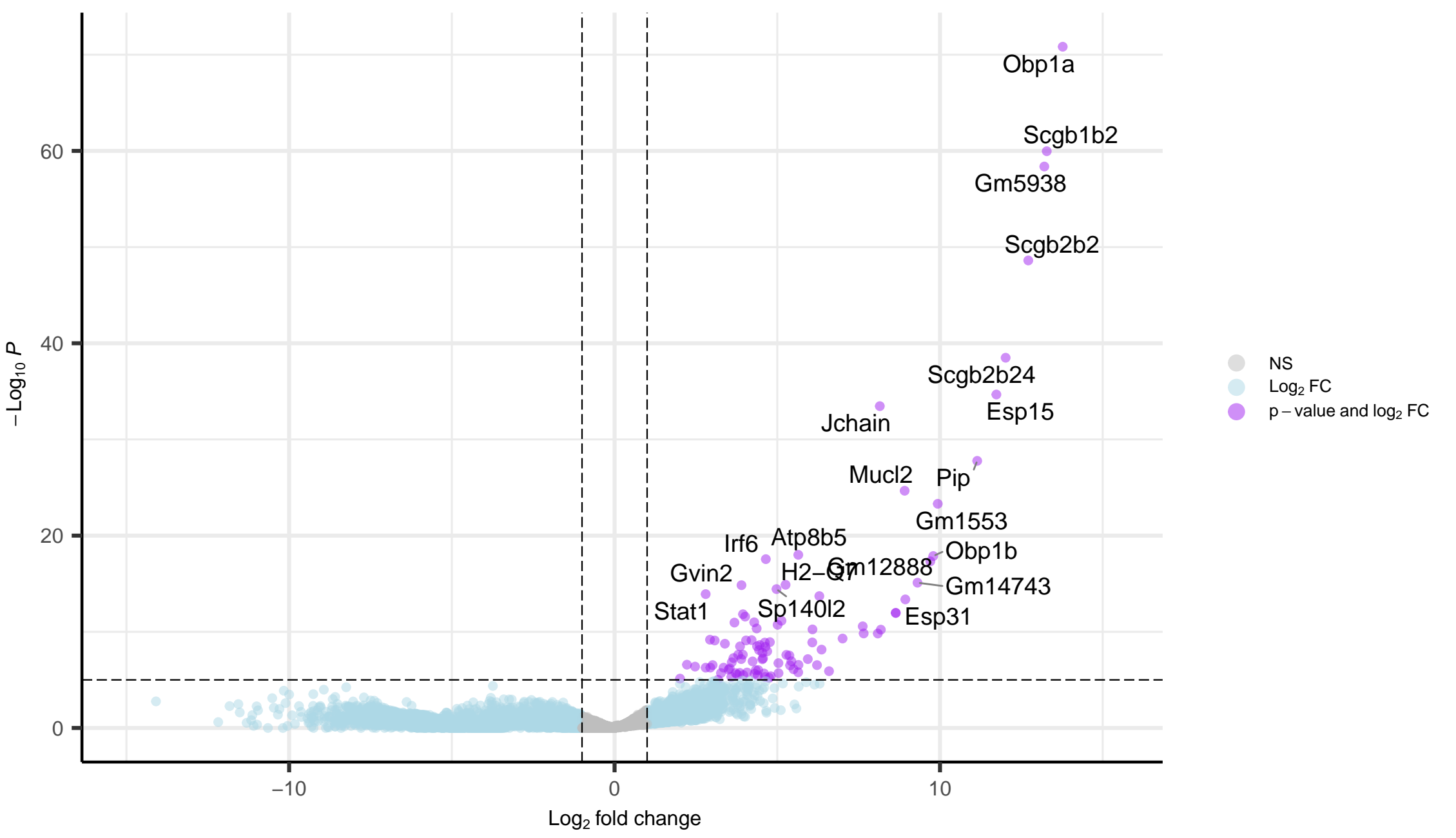


resultant_genotype: nf1 KO; pten KO; ink KO; atrx wt vs. nf1 wt; pten KO; ink KO; atrx KO



total = 16875 variables