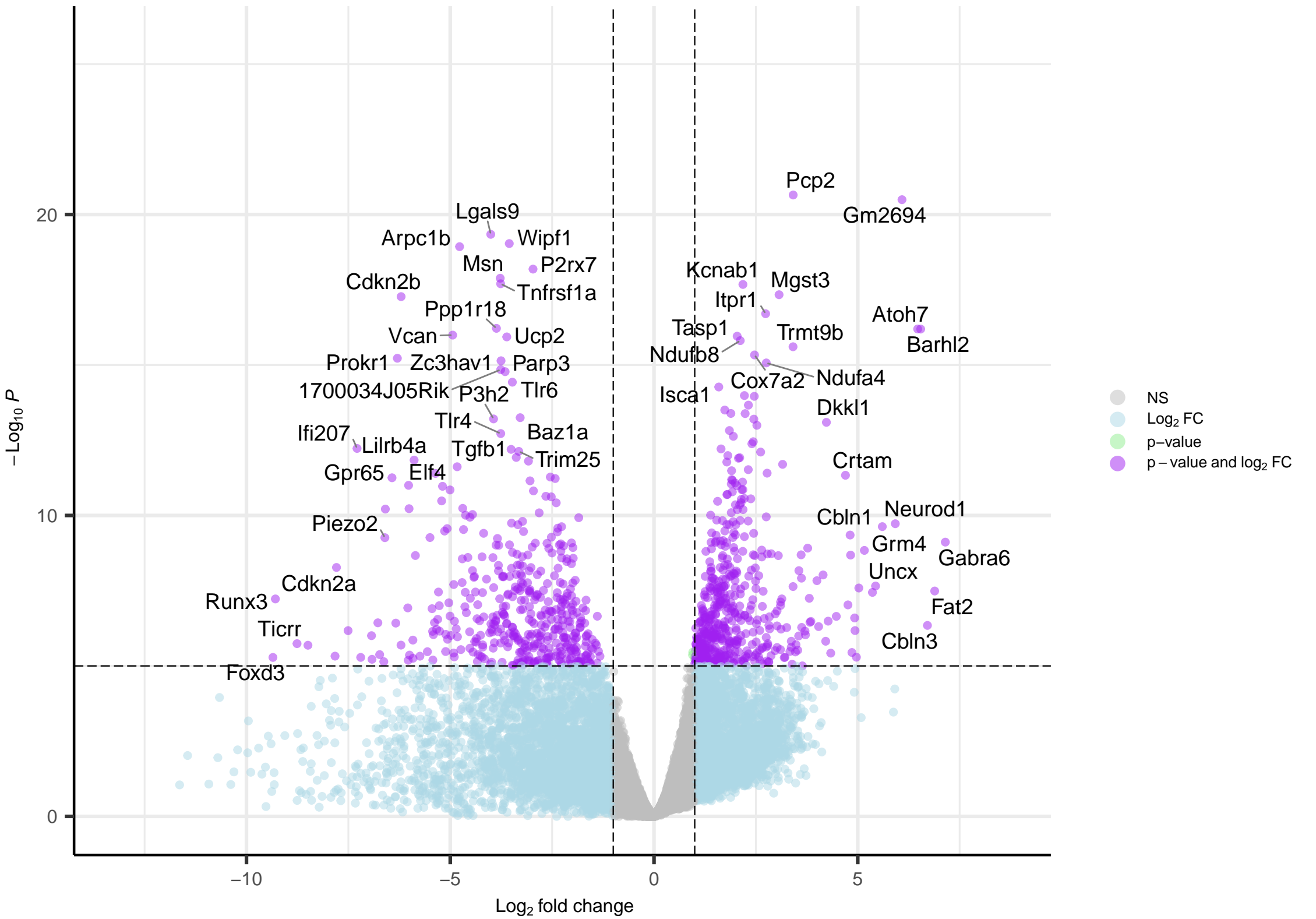


resultant_genotype: nf1 KO; pten KO; ink KO; atrx KO vs. nf1 wt; pten wt; ink wt; atrx wt



total = 16038 variables