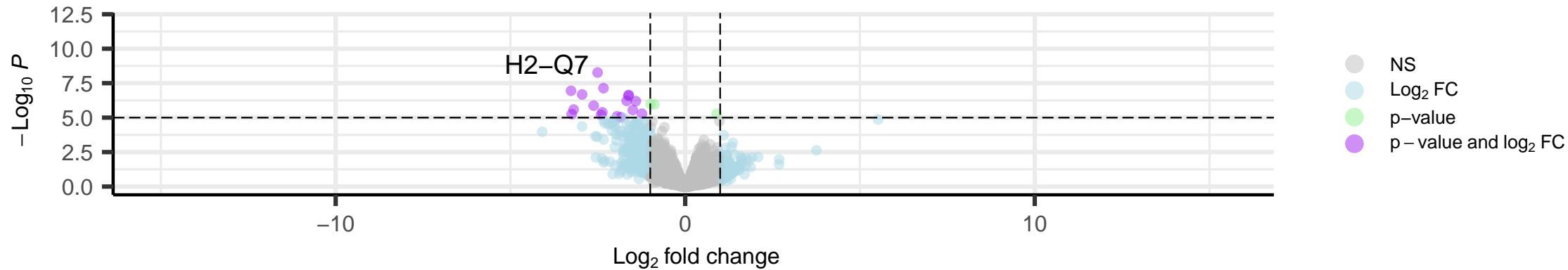


resultant_genotype: nf1 KO; pten KO; ink KO; atrx KO vs. nf1 KO; pten KO; ink KO; atrx wt



total = 14608 variables