

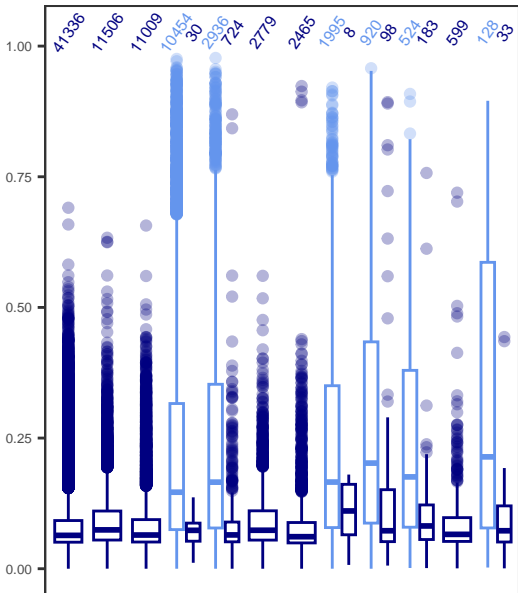
Effect Size (R-Squared)

type



cis

trans



equal_number_paralogs

hotspot

paralog_at_qtl

paralog_number_variation

single_copy_gene

Classification according to Gene and eQTL Characteristics