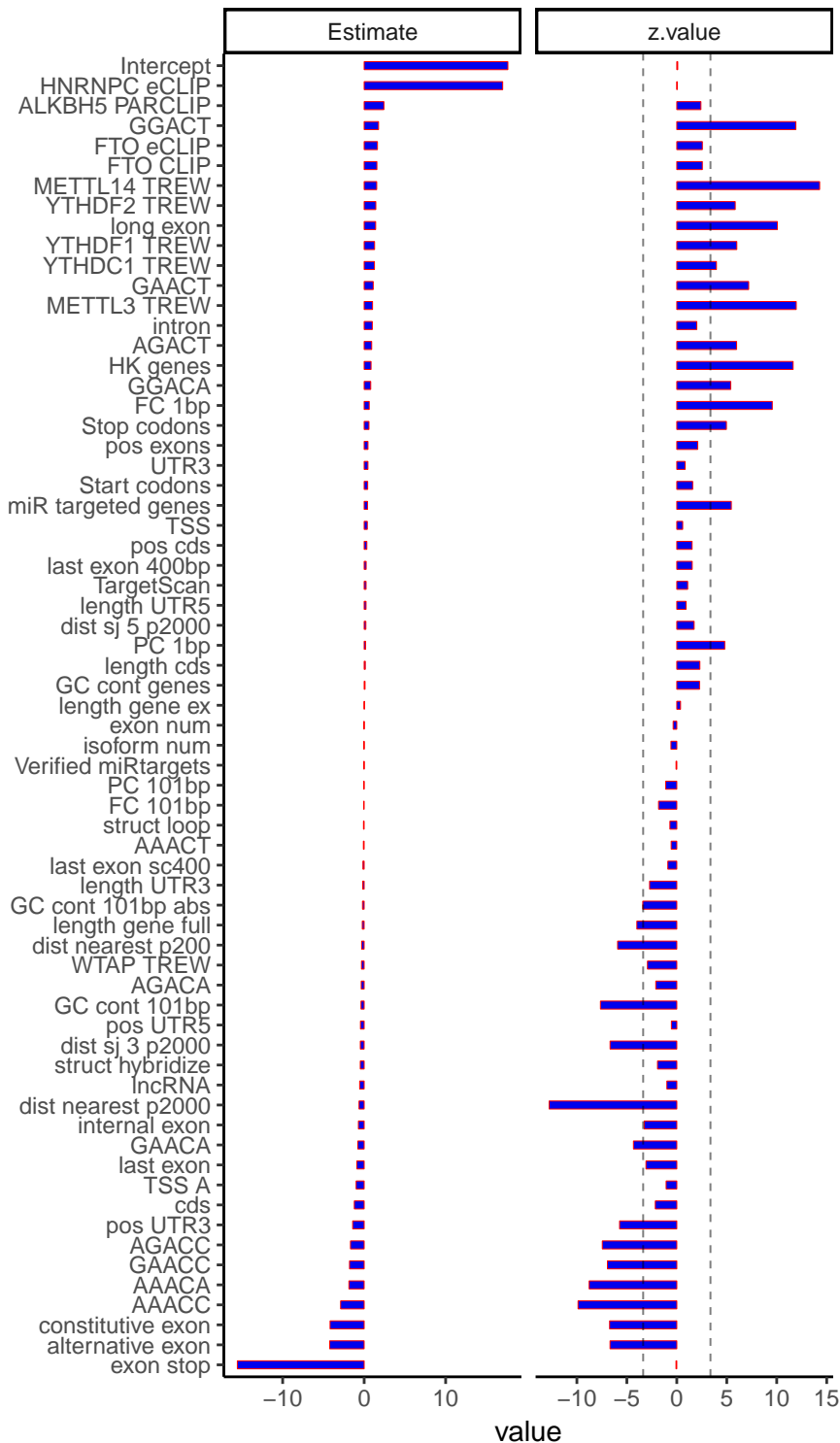binomial linear model on genomic features

glm