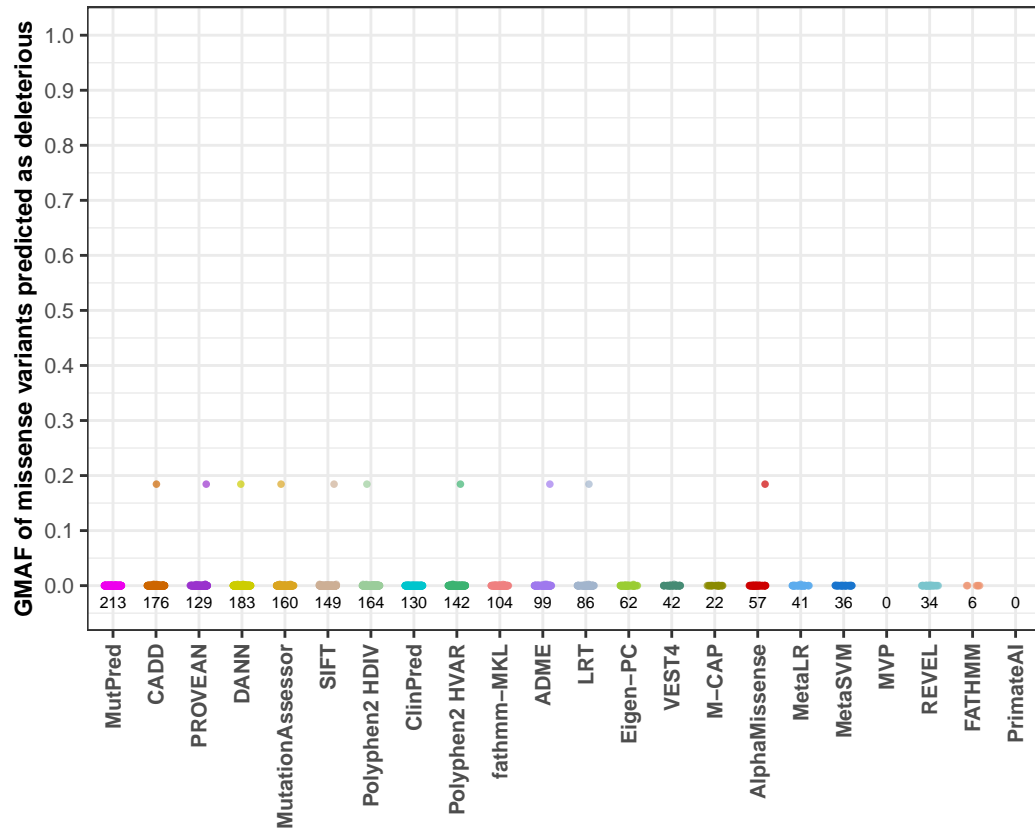


## UGT3A1

277 variants predicted as deleterious in total



## UGT3A2

252 variants predicted as deleterious in total

