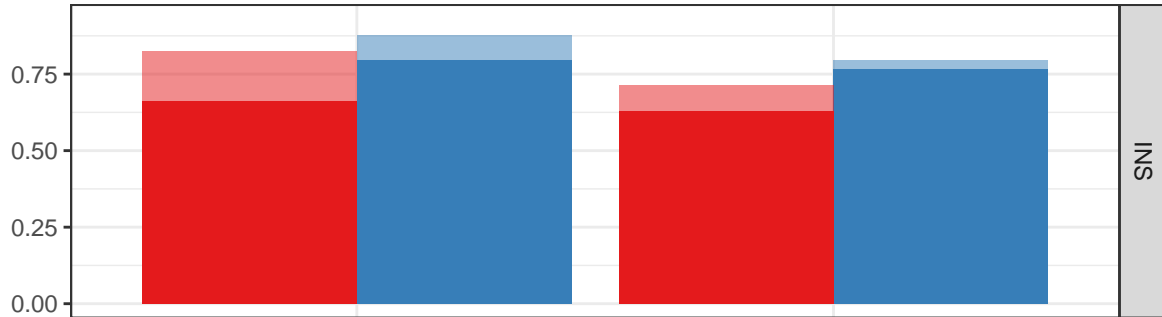


best F1



SV evaluation

absence/presence  
genotype

genomic regions

all  
non-repeat

vg

SMRT-SV2

INS

DEL