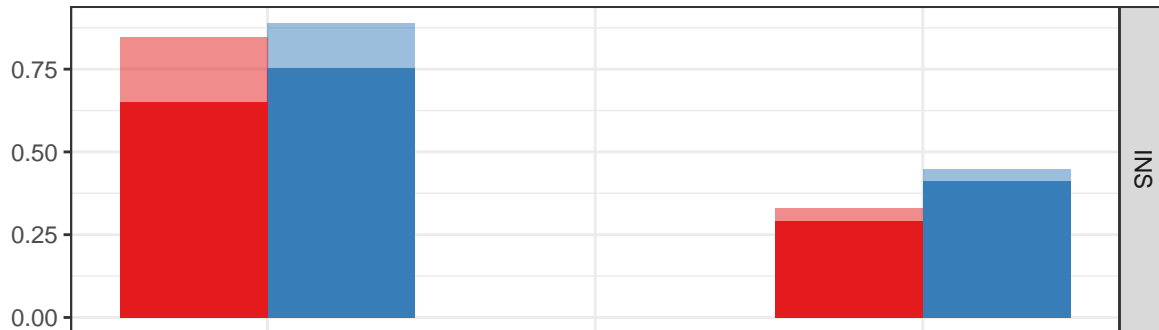
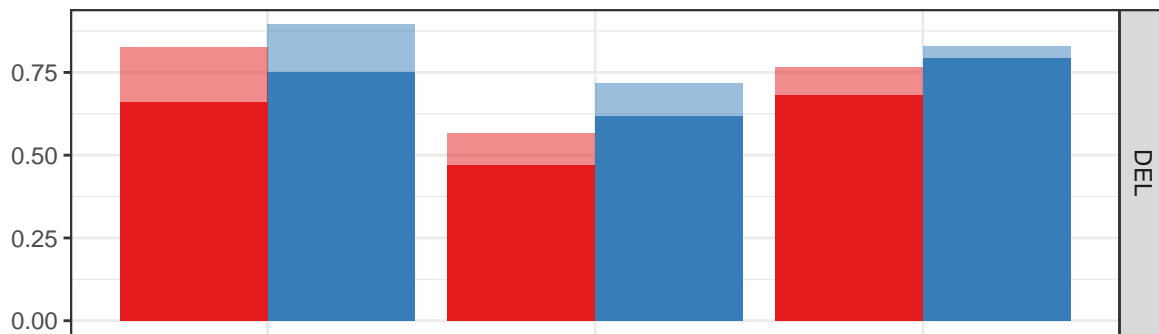
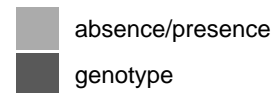


best F1



INS

SV evaluation



DEL

genomic regions

