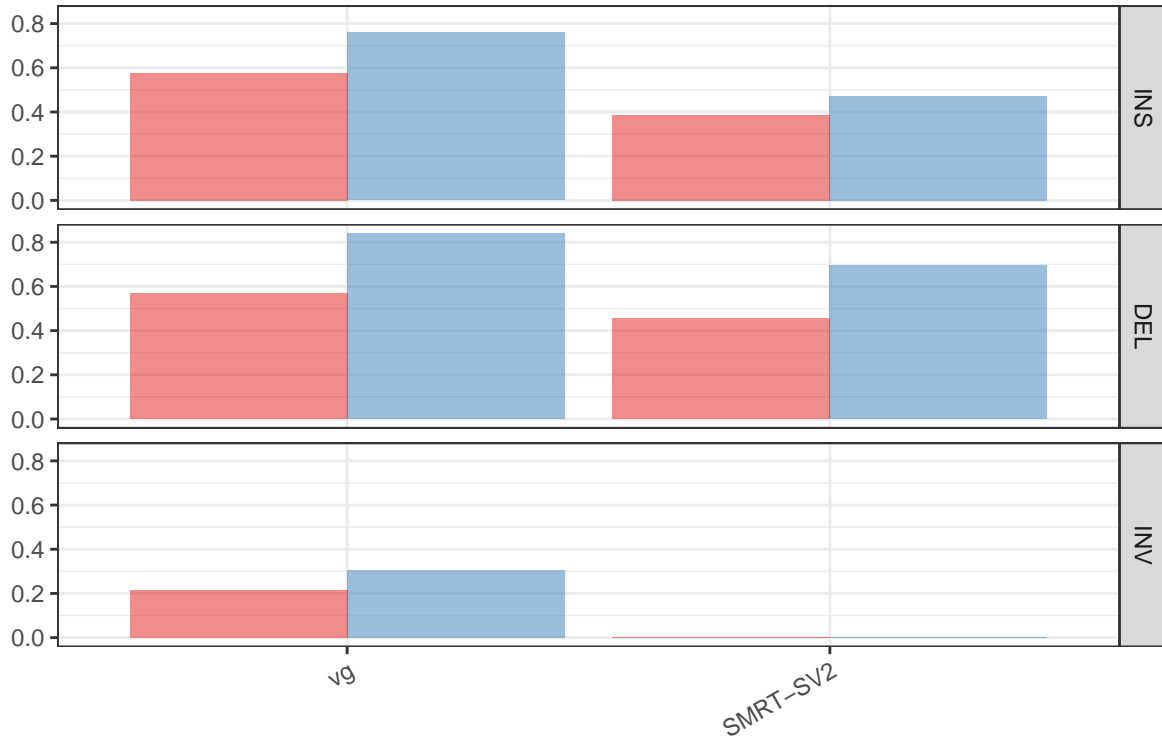


best F1



SV evaluation

absence/presence

genomic regions

all

non-repeat