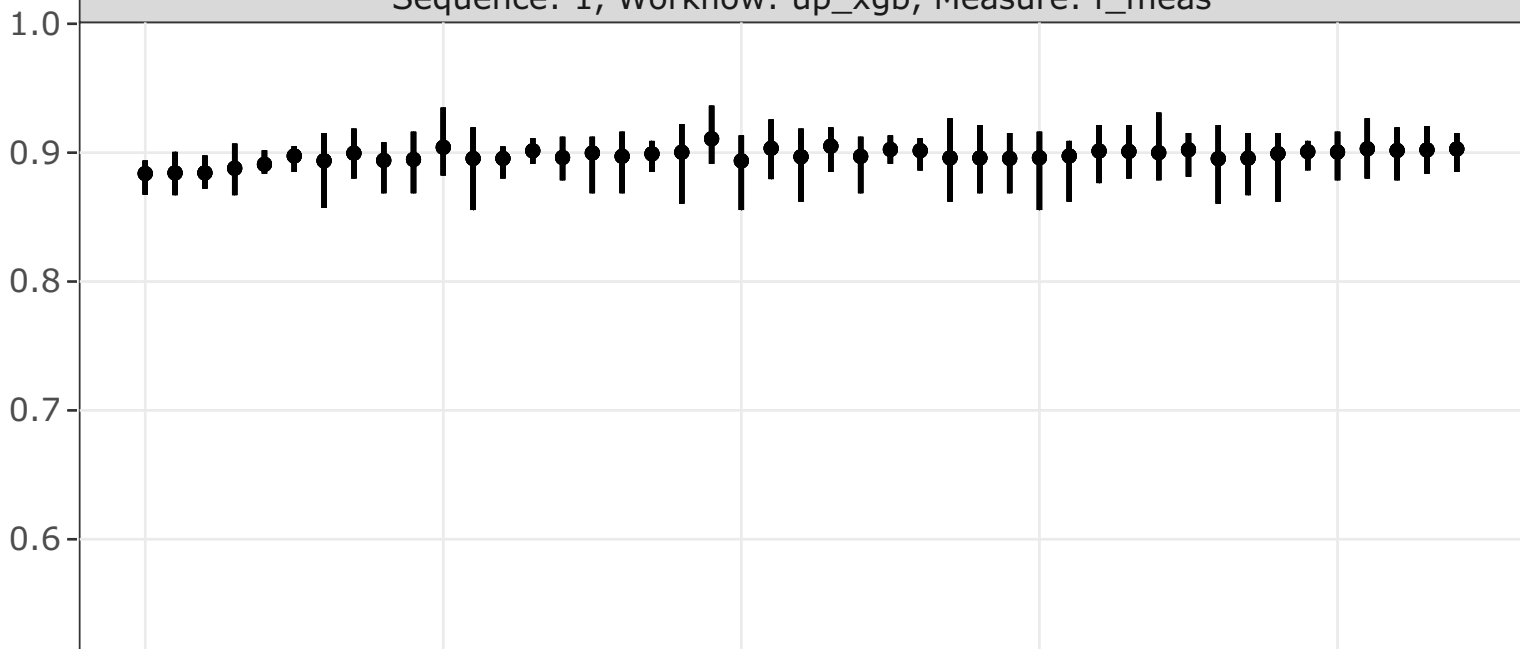
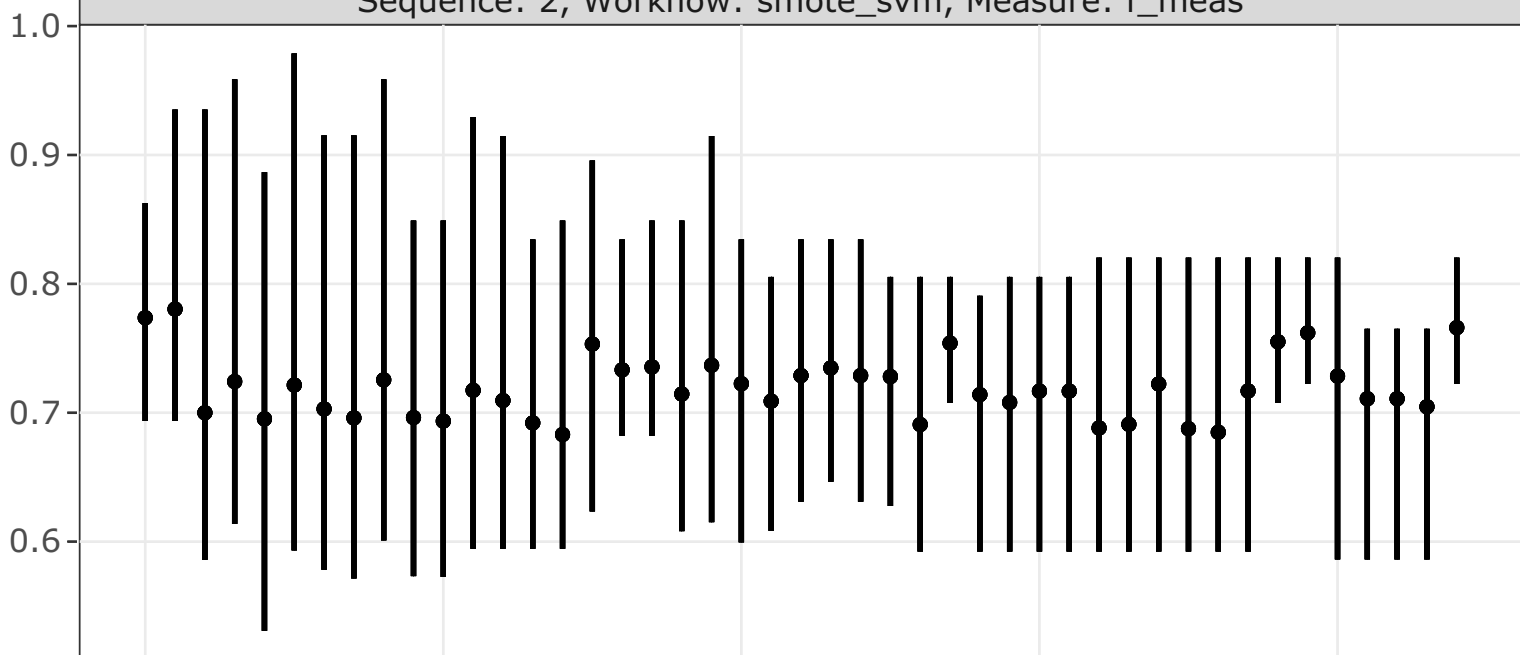


Gene Optimziation for 2-STEP Classifier

Sequence: 1, Workflow: up_xgb, Measure: f_meas



Sequence: 2, Workflow: smote_svm, Measure: f_meas



Genes Added