Gene Optimziation for 2-STEP Classifier Sequence: 1, Workflow: up_xgb, Measure: f_meas 1.0 0.9 0.8 0.7 0.6 F1-Score Sequence: 2, Workflow: smote_svm, Measure: f_meas 1.0 0.9 0.8 0.7 0.6 20 0 10 30 40 Genes Added