Gene Optimziation for Sequential Classifier Sequence: 1, Workflow: up_rf, Measure: f_meas 1.0-0.9 0.8 0.7 Sequence: 2, Workflow: smote_xgb, Measure: f_meas 1.0-0.9 0.8 F1-Score Sequence: 3, Workflow: hybrid_svm, Measure: f_meas 0.9 0.8 0.7 Sequence: 4, Workflow: hybrid_rf, Measure: f_meas 1.0 0.9 0.8 0.7 10 20 30 40 Genes Added