

Gene Optimziation for Sequential Class

F1-Score

Sequence: 1, Workflow: smote_rf, Measure: f_meas

Sequence: 2, Workflow: hybrid_xgb, Measure: f_meas

Sequence: 3, Workflow: up_svm, Measure: f_meas

Sequence: 4, Workflow: hybrid_mr, Measure: f_meas

0 10 20 30 40

Genes Added

