

# 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

