SHAP Waterfall: Highest−Scoring gRNA

Feature contributions to prediction

f(x)=104

102 other features — +17.6

freq_TT = 0 — +7.01

freq_GA = 0.158 — +4.4

pos19_G = 1 — +3.56

pos15_A = 1 — +3.52

pos4_G = 1 — +3.35

pos16_A = 1 — +2.82

freq_GT = 0.158 — +2.77

freq_AG = 0.158 — +2.48

pos14_G = 0 — +2.24

exon_rank = 5 — +2.06

pos10_A = 1 — +2.02

E[f(x)]=50

Prediction