TCL1A: K.W. p=0.048 Stage M.W. p ≤ 0.05 Normal CIS+AIS 2.5 Primary 2.0 rcL1A expression M.W.W. p ≤ 0.05 1.5 M.W.W. p≤ -0.5Higher Lower

Lower
Mutation Shared Count (SYN) per TMB