RAP1GDS1: Kruskal-Wallis p=0.025 Stage Normal 17.5 CIS+AIS **Primary** 15.0 M.W.W. p ≤ 0.01 12.5 S1 expression M.W.W. p ≤ 0.05 10.0 RAP1GD 5.0 2.5 0.0 Higher Lower

Mutation Shared Count (SYN) per TMB