RAP1GDS1: Kruskal-Wallis p=0.025 Stage M.W.W. p = 0.08Normal CIS+AIS 20 **Primary** M.W.W. p = 0.25M.W.W. p = 0.11M.W.W. p = 0.3415 S1 expression M.W.W. p = 0.76M.W.W. $p \le 0.05$ M.W.W. p = 0.34RAP1GD Higher Lower

Mutation Shared Count (SYN) per TMB