SMAD2: Kruskal-Wallis p=0.019 Stage M.W.W. p = 0.46Normal CIS+AIS M.W.W. p  $\leq 0.05$ Primary 3000 M.W.W. p = 0.682500 M.W.W. p  $\leq 0.01$ M.W.W. p = 0.58M.W.W. p = 0.68M.W.W. p = 0.071000 500 Higher Lower

Mutation Shared Count per TMB