SMAD2: Kruskal-Wallis p=0.002 6000 Stage M.W.W. p = 0.66Normal CIS+AIS M.W. p ≤ 0.05 Primary M.W.W. p = 0.295000 M.W.W. p ≤ 0.01 M.W.W. p = 0.62M.W. p = 0.98M.W.W. p = 0.112000 1000 Higher Lower Mutation Shared Count per TMB