SMAD2: Kruskal-Wallis p=0.003 6000 Stage M.W.W. p = 1.00Normal CIS+AIS M.W.W. $p \le 0.05$ Primary M.W.W. p = 0.525000 M.W.W. $p \le 0.001$ M.W. p = 0.68M.W.W. p = 0.98M.W.W. p = 0.162000 1000 Higher Lower Mutation Shared Count (SYN) per TMB