SMAD3: Kruskal-Wallis p=0.034 Stage M.W.W. p = 0.36Normal CIS+AIS **Primary** M.W.W. p ≤ 0.05 100 M.W.W. p = 0.96M.W.W. p = 0.4280 M.W.W. p = 0.23M.W.W. p = 0.46expression M.W.W. p = 0.1060 SMAD3 40 20 Higher Lower Mutation Shared Count (SYN) per TMB