GRIN2A: Kruskal-Wallis p=0.007 1.0 Stage M.W.W. p = 0.84Normal CIS+AIS Primary M.W.W. p = 0.198.0 M.W.W. p = 0.76M.W.W.  $p \le 0.05$ GRIN2A expression M.W.W. p = 0.51M.W.W. p = 0.35M.W.W.  $p \le 0.05$ • 0.0 Higher Lower Mutation Shared Count (SYN) per TMB