GRIN2A: Kruskal-Wallis p=0.010 1.0 Stage M.W.W. p = 0.46Normal CIS+AIS Primary M.W.W. p = 0.378.0 M.W.W. p = 0.93M.W.W. $p \le 0.05$ GRIN2A expression M.W.W. p = 0.96M.W.W.p =M.W.W. $p \le 0.05$ 0.2 0.0 Higher Lower Mutation Shared Count per TMB