GRIN2A: Kruskal-Wallis p=0.007 1.0 Stage M.W.W. p = 0.82Normal CIS+AIS Primary M.W.W. p = 0.198.0 M.W.W. p = 0.60M.W.W. $p \le 0.05$ GRIN2A expression M.W.W. p = 0.54M.W.W. p = 0.34M.W.W. $p \le 0.05$ 0.0 Higher Lower Mutation Shared Count per TMB