GRIN2A: Kruskal-Wallis p=0.008 1.0 Stage Normal CIS+AIS **Primary** 8.0 GRIN2A expression M.W.W.  $p \le 0.01$ M.W.W.  $p \le 0.01$ 0.2 0.0 Higher Lower Mutation Shared Count (SYN) per TMB