GRIN2A: Kruskal-Wallis p=0.015 Stage Normal 1.2 CIS+AIS **Primary** 1.0 8.0 GRIN2A expression 0.6 M.W.W. p ≤ 0.05 M.W. $p \le 0.01$ 0.4 0.2 0.0 -0.2Higher Lower Mutation Shared Count (SYN) per TMB