NUTM2D: Kruskal-Wallis p=0.008 Stage 17.5 Normal CIS+AIS **Primary** M.W.W. $p \le 0.05$ 15.0 M.W.W. p ≤ 0.0512.5 expression 0 NUTM2 7.5 5.0 2.5 0.0 Higher Lower Mutation Shared Count per TMB