NUTM2D: Kruskal-Wallis p=0.003 30 Stage M.W. p = 0.08Normal CIS+AIS M.W.W. p = 0.31Primary 25 M.W.W. p ≤ 0.05 M.W.W. p = 0.75M.W. p = 0.1120 expression M.W.W. p = 0.21NUTM2 M.W.W. p = 0.26Higher Lower Mutation Shared Count (SYN) per TMB