PBRM1: Kruskal-Wallis p=0.000 Stage M.W.W. p = 0.25Normal CIS+AIS M.W.W. p = 0.07Primary 200 M.W.W. p = 0.62M.W.W. p ≤ 0.01 150 PBRM1 expression M.W.W. p = 0.32M.W.W.p = 0.93M.W.W. p ≤ 0.01 50 Higher Lower Mutation Shared Count (SYN) per TMB