NUTM2D: Kruskal-Wallis p=0.003 Stage M.W.W. p = 0.07Normal CIS+AIS M.W. p = 0.68Primary M.W.W. p ≤ 0.05 M.W.W. p = 0.72M.W.W. p ≤ 0.05 M.W.W. p = 0.24

30

25

expression 5

NUTM2

Lower
Mutation Shared Count (SYN) per TMB