ARHGAP26: Kruskal-Wallis p=0.005 Stage M.W. p = 0.30Normal CIS+AIS **Primary** M.W.W. p = 0.41M.W. p = 0.37M.W. p = 0.36M.W.W. p ≤ 0.01 M.W.N. p = 0.21M.W.W. p = 0.35

25

20

26 expression

Lower
Mutation Shared Count (SYN) per TMB