

Kruskal-Wallis p=0.000

M.W.W. $p \leq 0.001$

M.W.W. $p \leq 0.05$

Stage

CIS+AIS

Primary

Size of somatic CNV region (bp)

1e9

3.0

2.5

2.0

1.5

1.0

0.5

0.0

Lower

Higher

Mutation Shared Count (SYN) per TMB

