

Kruskal-Wallis p=0.000

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

M.W.W.  $p \leq 1e-5$

M.W.W.  $p \leq 0.01$

Stage

CIS+AIS

Primary

Size of somatic CNV region (bp)

1e9

2.0

1.5

1.0

0.5

0.0

Lower

Mutation Shared Count per TMB

Higher

