Kruskal-Wallis p=0.000

1e9

Size of somatic CNV region (bp)

2.5
2.0
1.5
1.0
0.5
0.0

M.W.W. p ≤ 0.001

M.W.W. p ≤ 0.001

M.W.W. p ≤ 1e-5

Stage
CIS+AIS
Primary

Lower
Higher

Mutation Shared Count per TMB