

Kruskal-Wallis p=0.000

1e9

Size of somatic CNV region (bp)

2.0  
1.5  
1.0  
0.5  
0.0

Stage  
CIS+AIS  
Primary

\*\*\*

\*\*\*\*

\*\*

Lower Higher  
Mutation Shared Count per TMB

