

Kruskal-Wallis  $p=0.000$

M.W.W.  $p \leq 1e-4$

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

Stage

CIS+AIS

Primary

Size of somatic CNV region (bp)

Lower Higher  
Mutation Shared Count (SYN)

1e9

3.0  
2.5  
2.0  
1.5  
1.0  
0.5  
0.0

