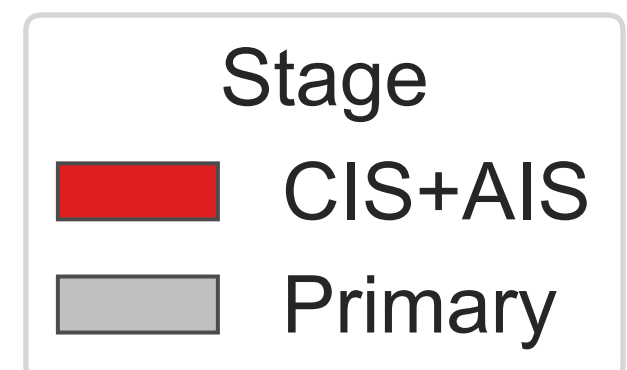


Kruskal-Wallis p=0.000

1e9

Size of somatic CNV region (bp)

2.9  
2.8  
2.7  
2.6



\*\*\*\*

\*\*\*\*

Lower Higher  
Mutation Shared Proportion (Union)

