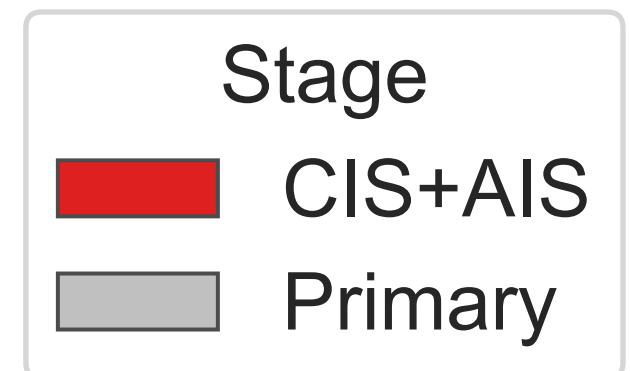


Kruskal-Wallis p=0.000

1e9

Size of somatic CNV region (bp)

2.9
2.8
2.7
2.6



Lower Higher
Mutation Shared Proportion (Union)