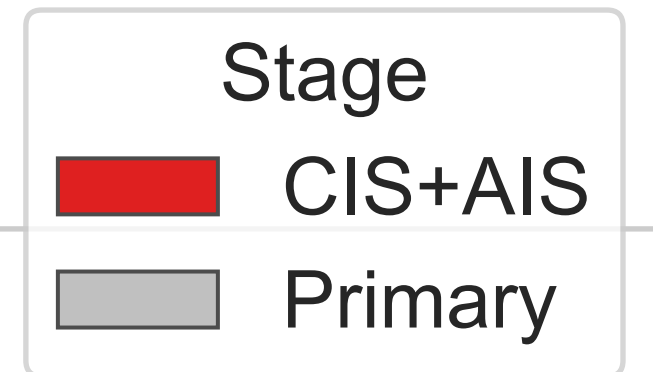


Kruskal-Wallis p=0.002

1e9

Size of somatic CNV region (bp)



**

**

Lower Higher
Mutation Shared Proportion (Union)

