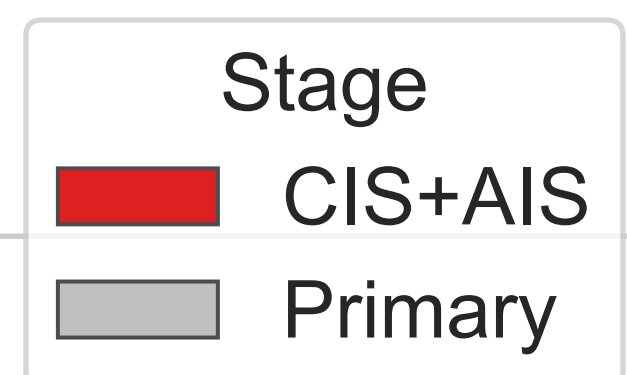


Kruskal-Wallis p=0.001

1e9

Size of somatic CNV region (bp)



**

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Lower Higher
Mutation Shared Proportion (Union & SYN)

