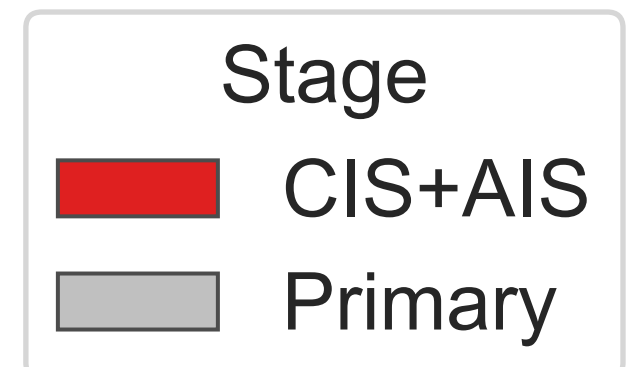


Kruskal-Wallis p=0.000

1e9

Size of somatic CNV region (bp)



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

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

Lower

Higher

Mutation Shared Proportion

