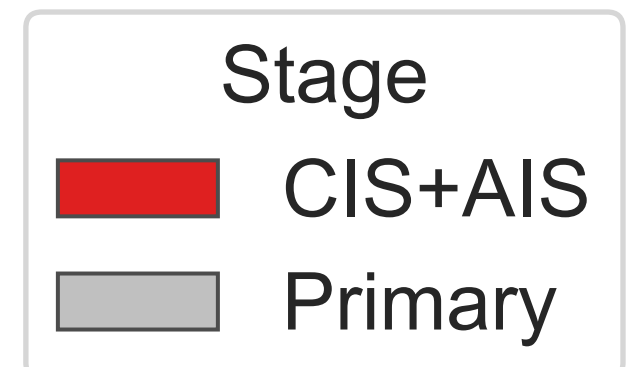


Kruskal-Wallis p=0.000

1e9

Size of somatic CNV region (bp)



M.W.W. $p \leq 1e-5$

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

Lower Higher
Mutation Shared Count (SYN) per TMB

