

Kruskal-Wallis p=0.000

1e9

Size of somatic CNV region (bp)

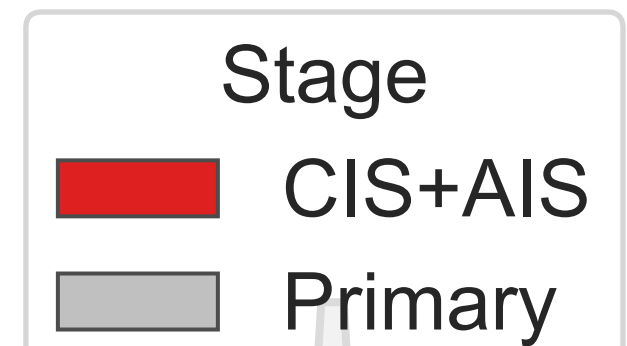
2.0

1.5

1.0

0.5

0.0



Lower Higher
Mutation Shared Count (SYN) per TMB

