

Kruskal-Wallis p=0.000

1e9

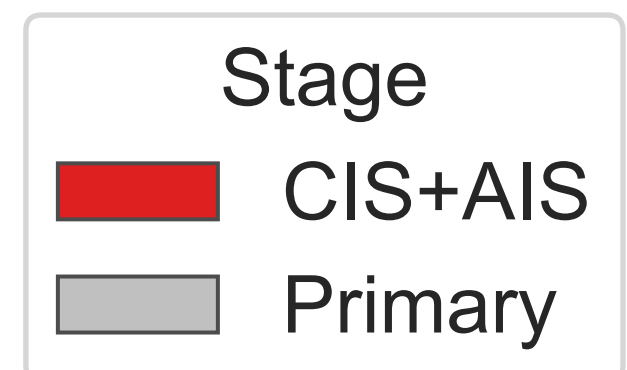
Size of somatic CNV region (bp)

2.9

2.8

2.7

2.6



Lower

Mutation Shared Count per TMB

Higher

