

Kruskal-Wallis p=0.000

1e9

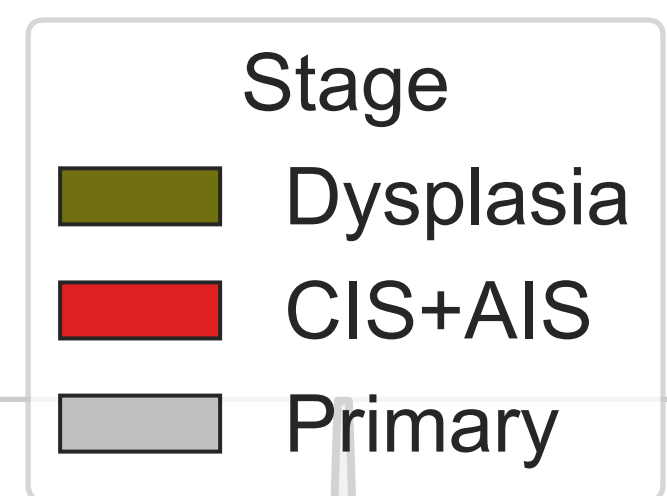
Size of somatic CNV region (bp)

2.9

2.8

2.7

2.6



**

Lower

Higher

Mutation Shared Count (SYN)

