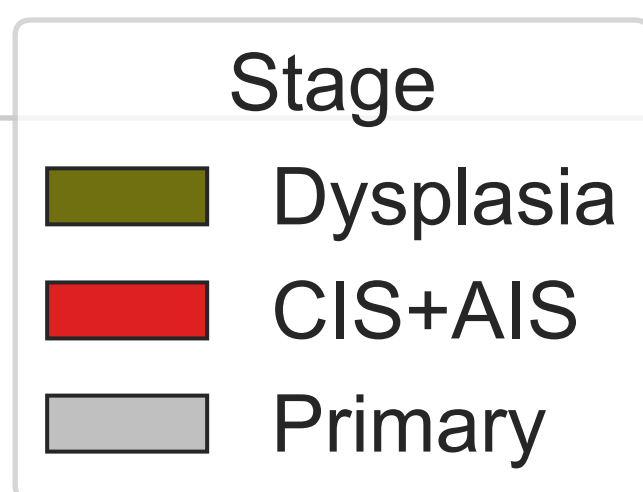


# Kruskal-Wallis p=0.000

1e9

Size of somatic CNV region (bp)



3.0  
2.5  
2.0  
1.5  
1.0  
0.5  
0.0

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Lower Higher  
Mutation Shared Proportion (SYN)

