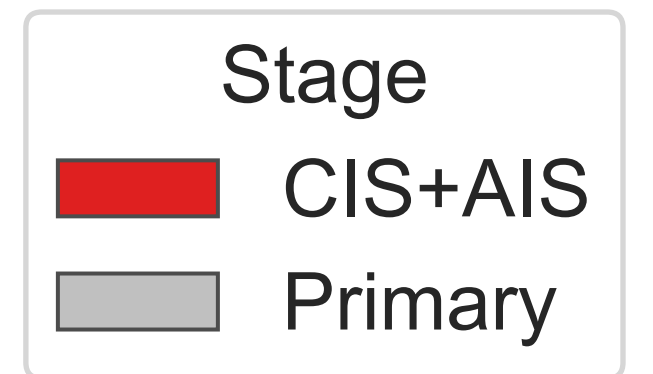


Kruskal-Wallis  $p=0.000$

Number of somatic CNV segment (count)



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

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

Lower Higher  
Mutation Shared Count (SYN)

