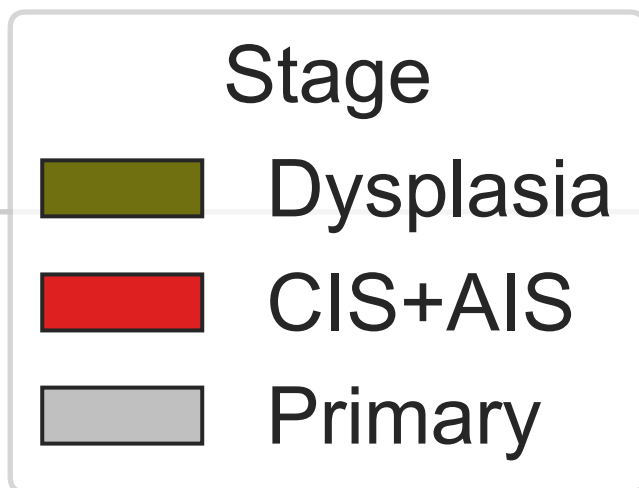


Kruskal-Wallis p=0.000

Number of somatic CNV segment (count)



M.W.W. $p \leq 0.001$

M.W.W. $p \leq 0.05$

M.W.W. $p \leq 0.05$

M.W.W. $p \leq 0.05$

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

M.W.W. $p \leq 0.05$

Lower Higher
Mutation Shared Count (SYN) per TMB

