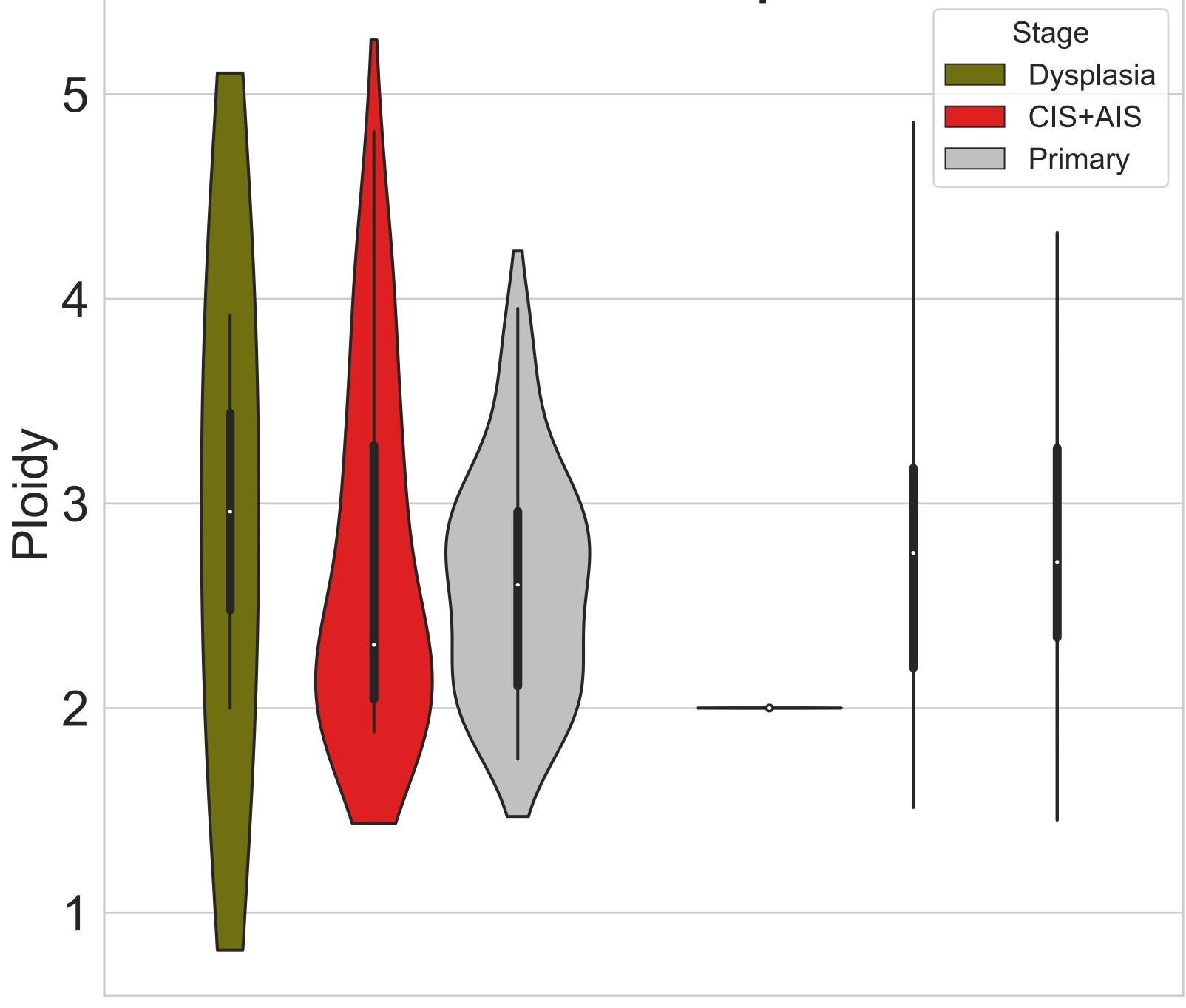
Kruskal-Wallis p=0.512



Lower Higher Mutation Shared Count (SYN) per TMB