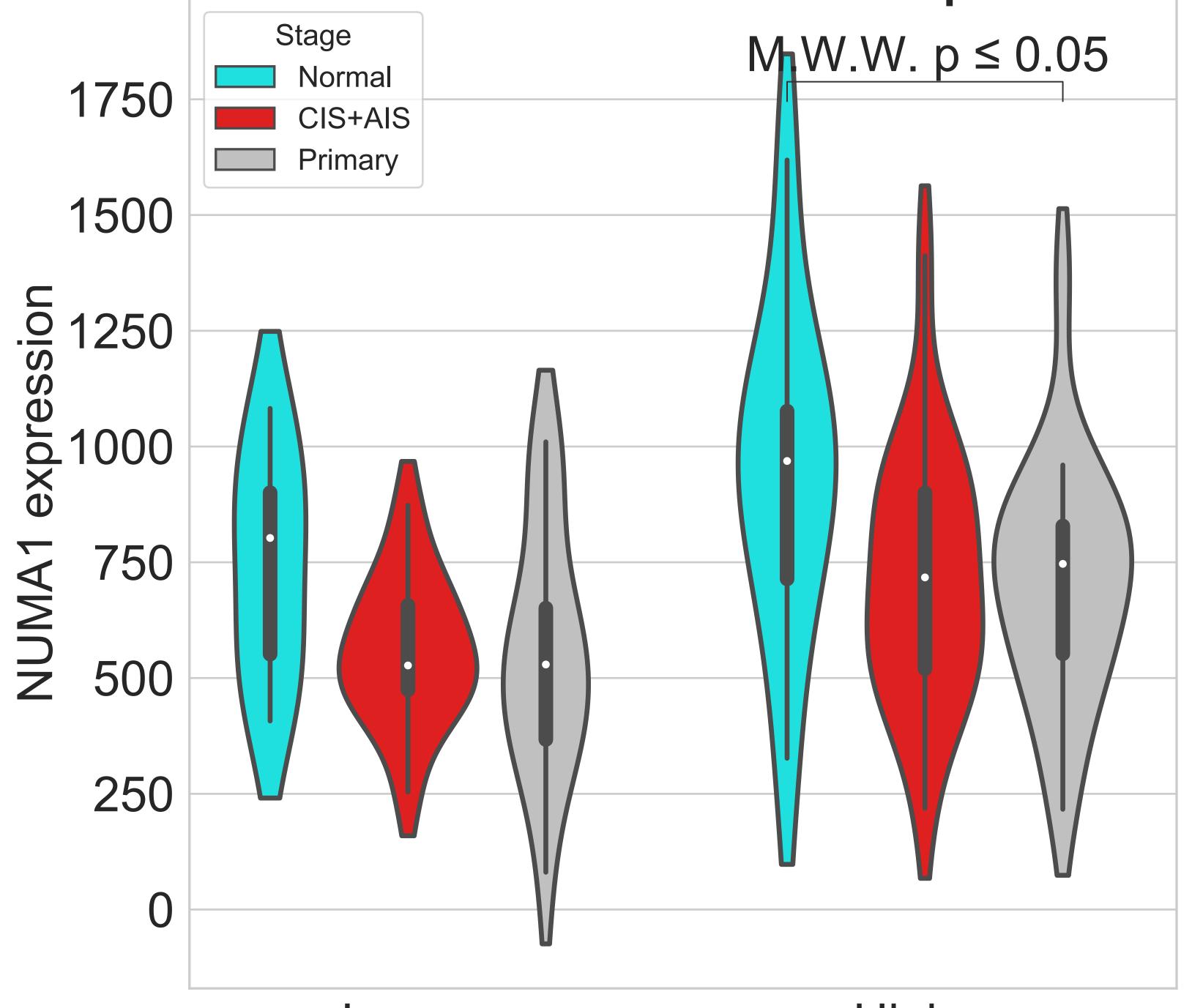
NUMA1: Kruskal-Wallis p=0.020



Lower Higher Mutation Shared Count (SYN) per TMB