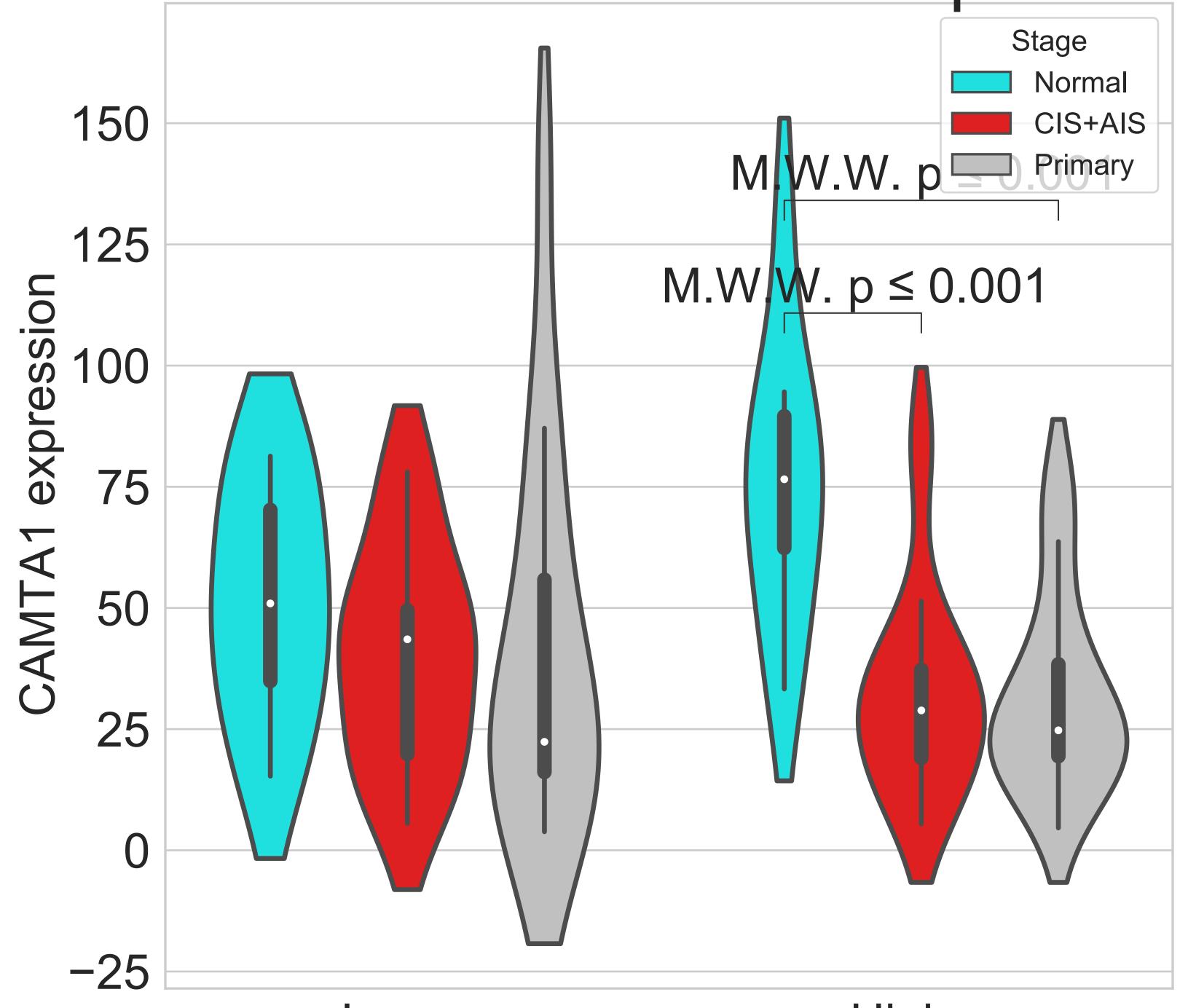
CAMTA1: Kruskal-Wallis p=0.009



Lower Higher Mutation Shared Count (SYN) per TMB