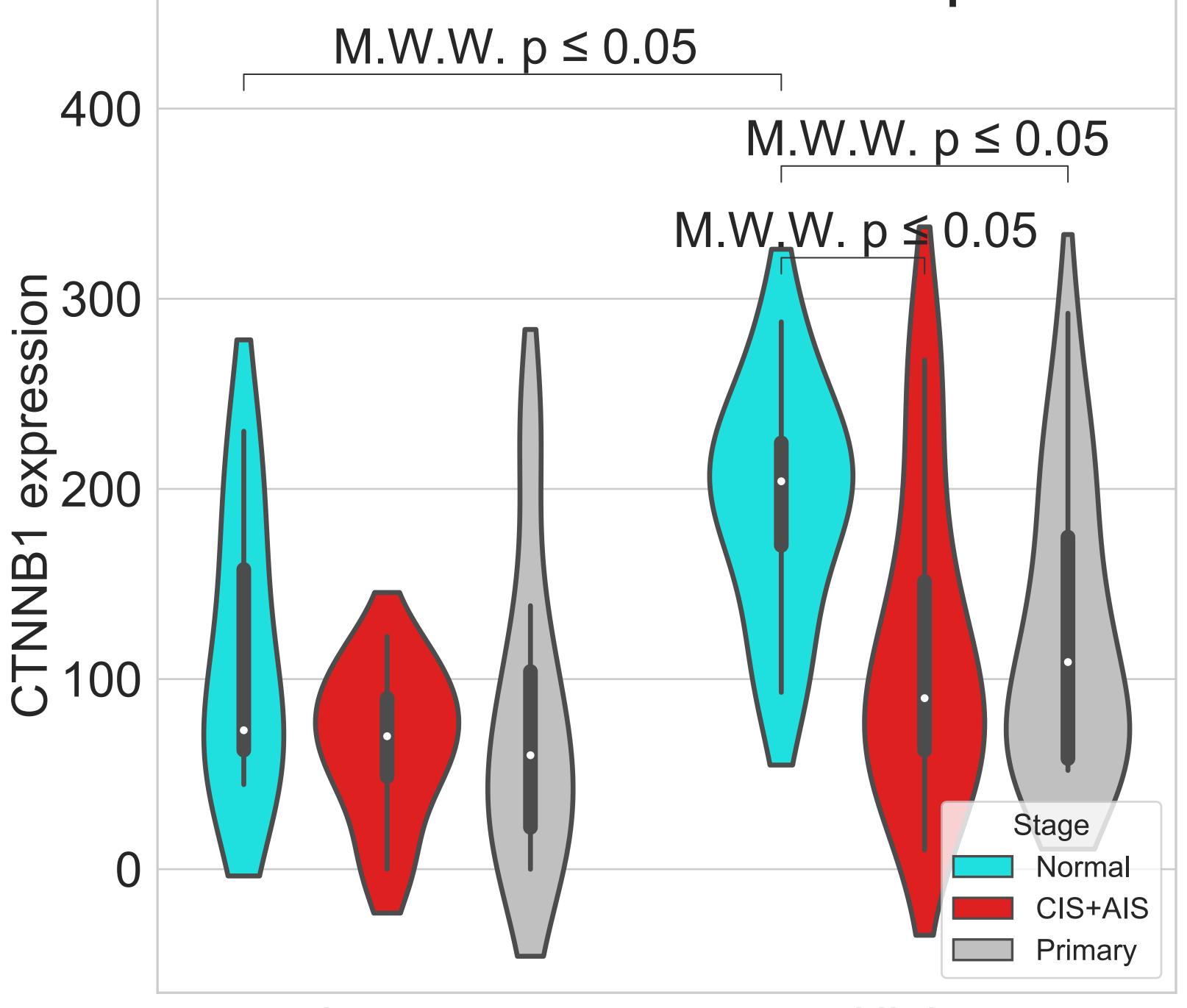
CTNNB1: Kruskal-Wallis p=0.005



Lower Higher Mutation Shared Proportion (SYN)