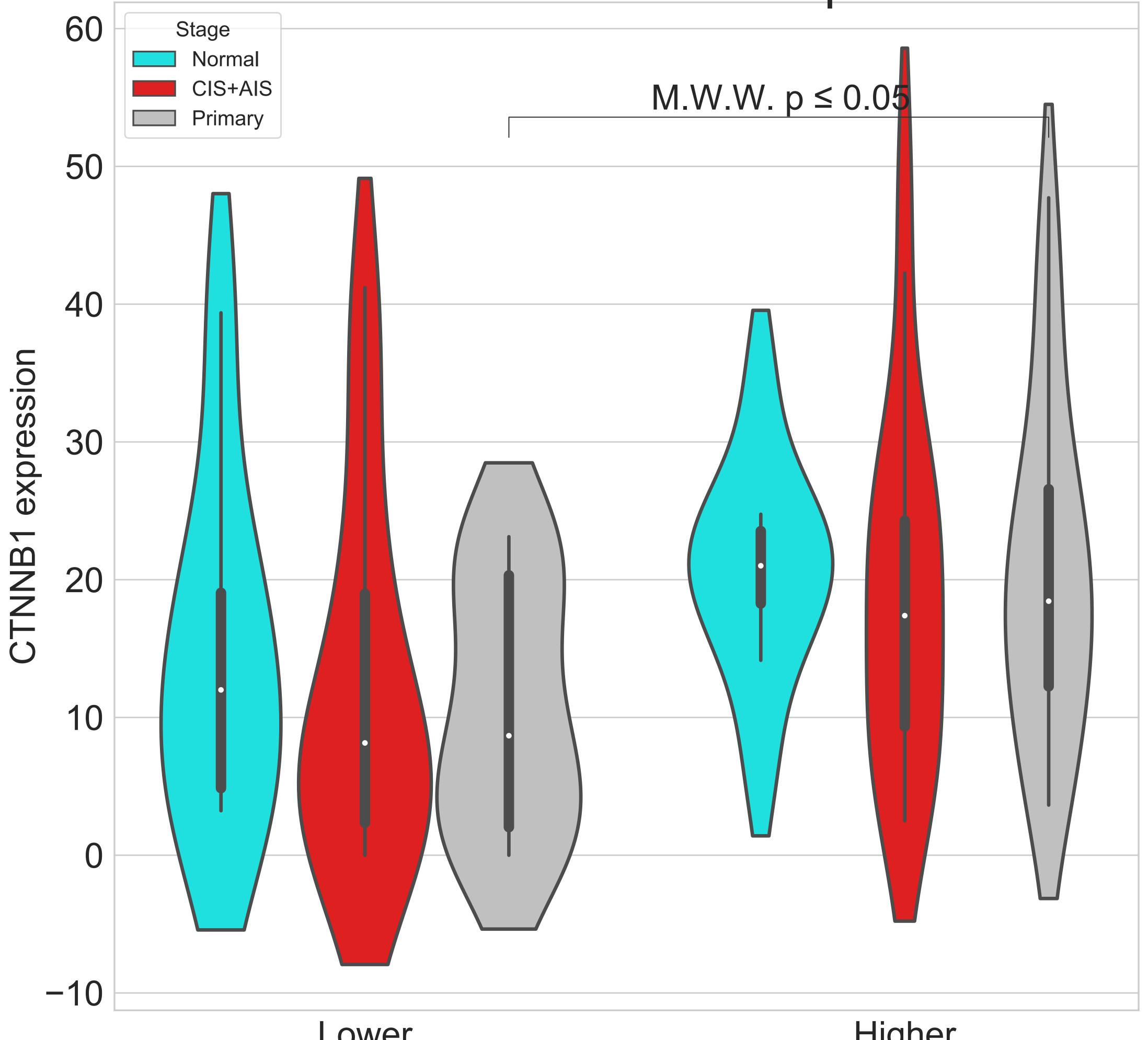
CTNNB1: Kruskal-Wallis p=0.037



Lower
Mutation Shared Proportion (SYN)