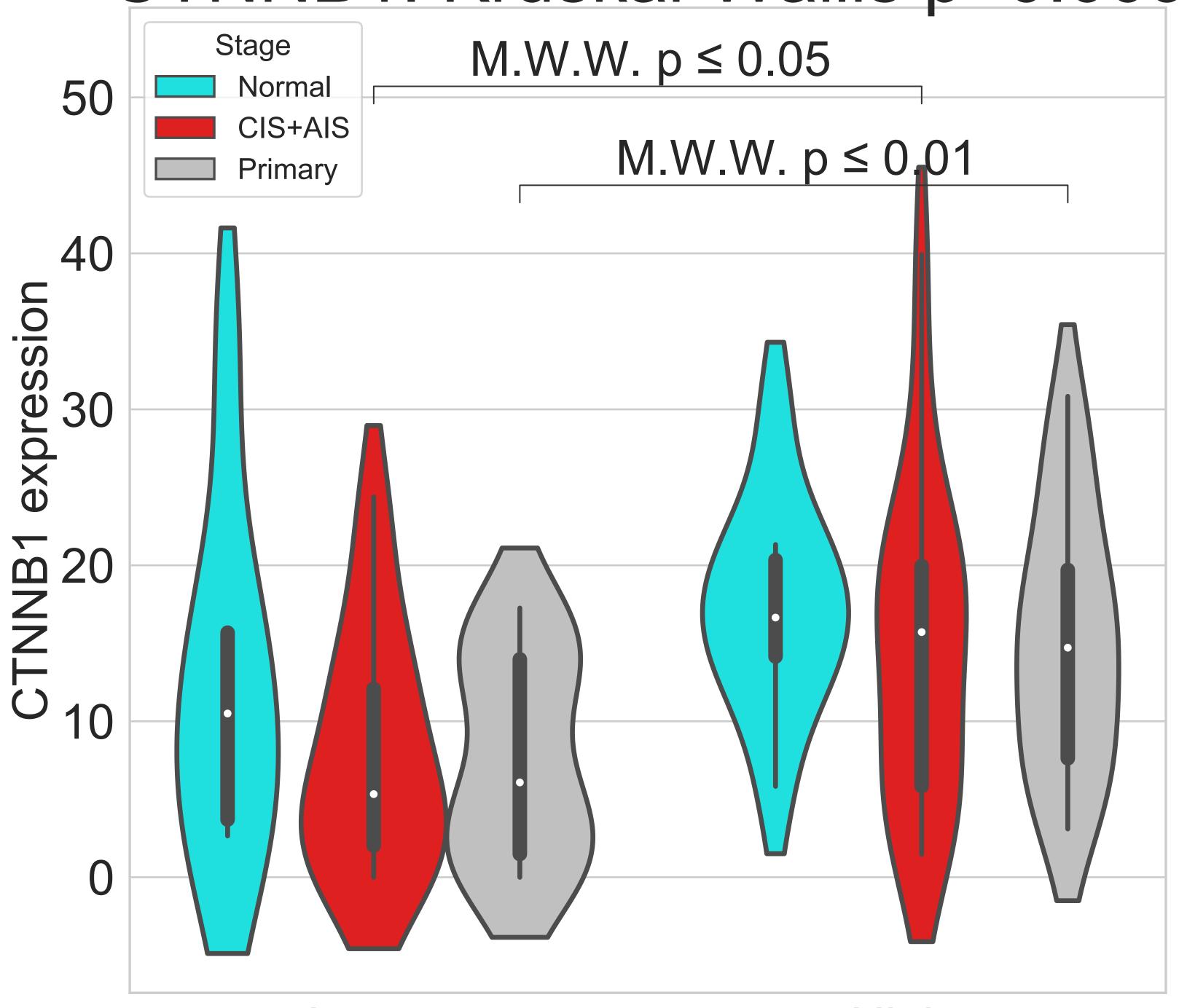
CTNNB1: Kruskal-Wallis p=0.009



Lower
Mutation Shared Proportion (SYN)