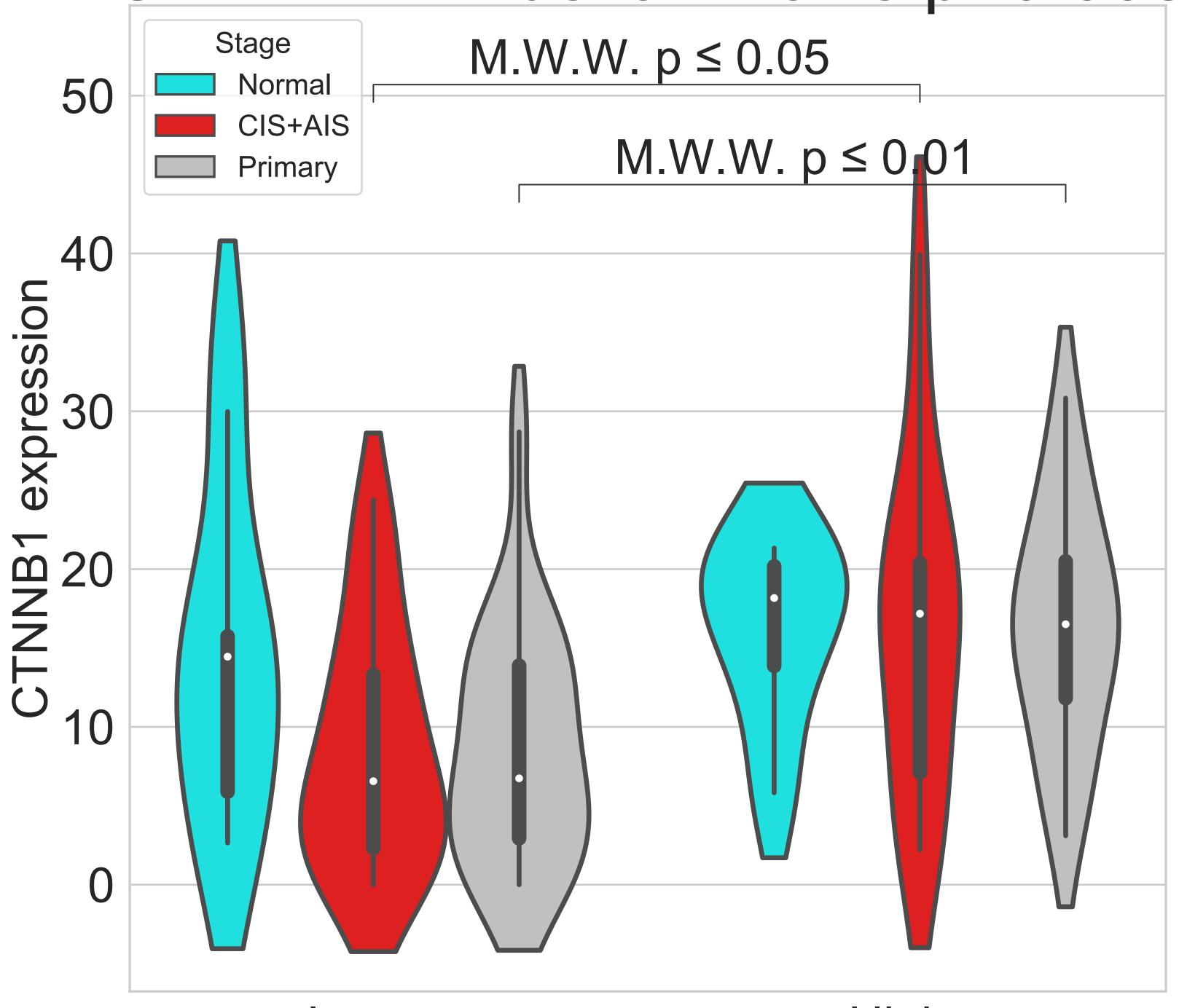
CTNNB1: Kruskal-Wallis p=0.008



Lower

Mutation Shared Proportion