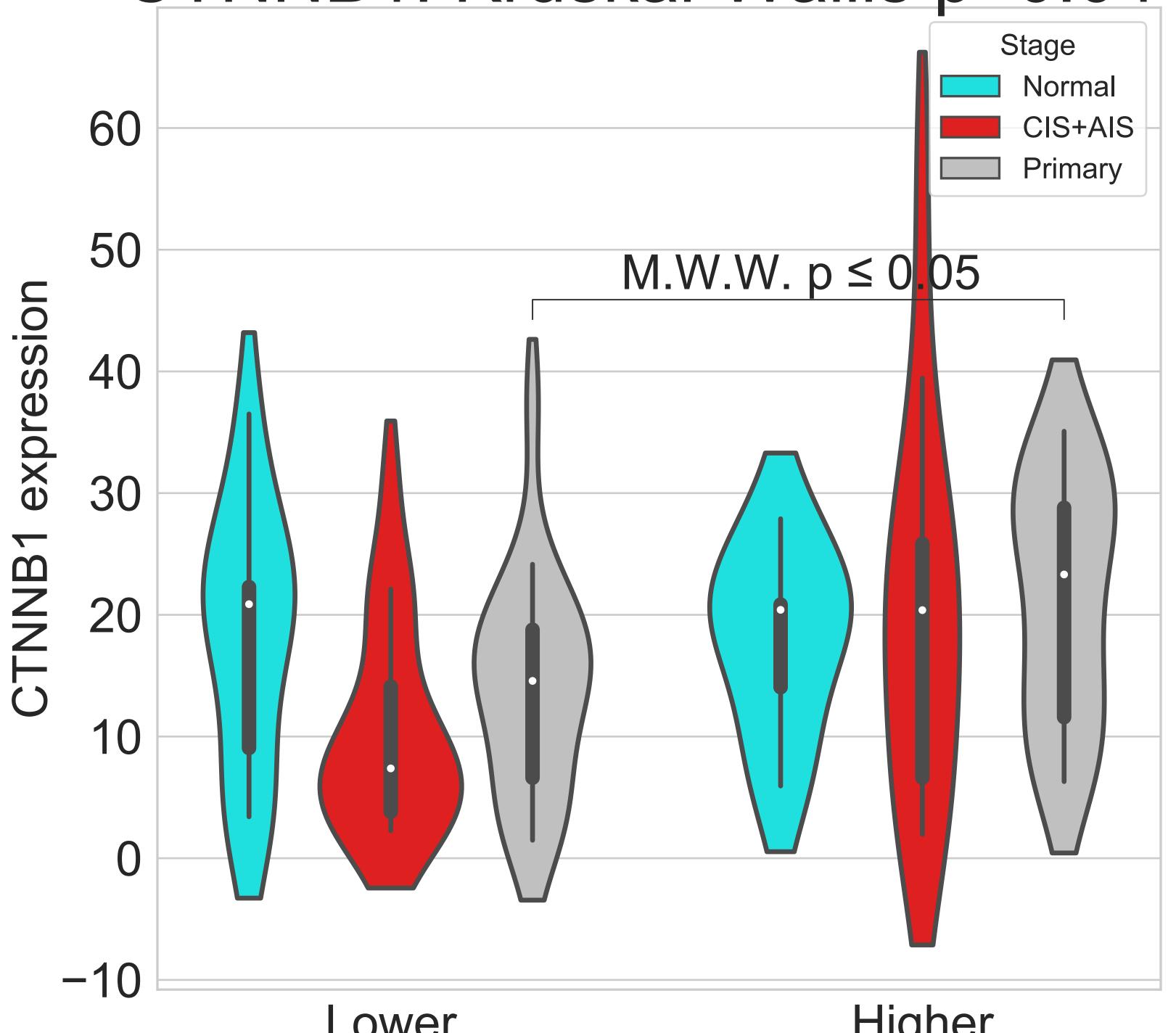
CTNNB1: Kruskal-Wallis p=0.044



Lower
Higher
Mutation Shared Proportion