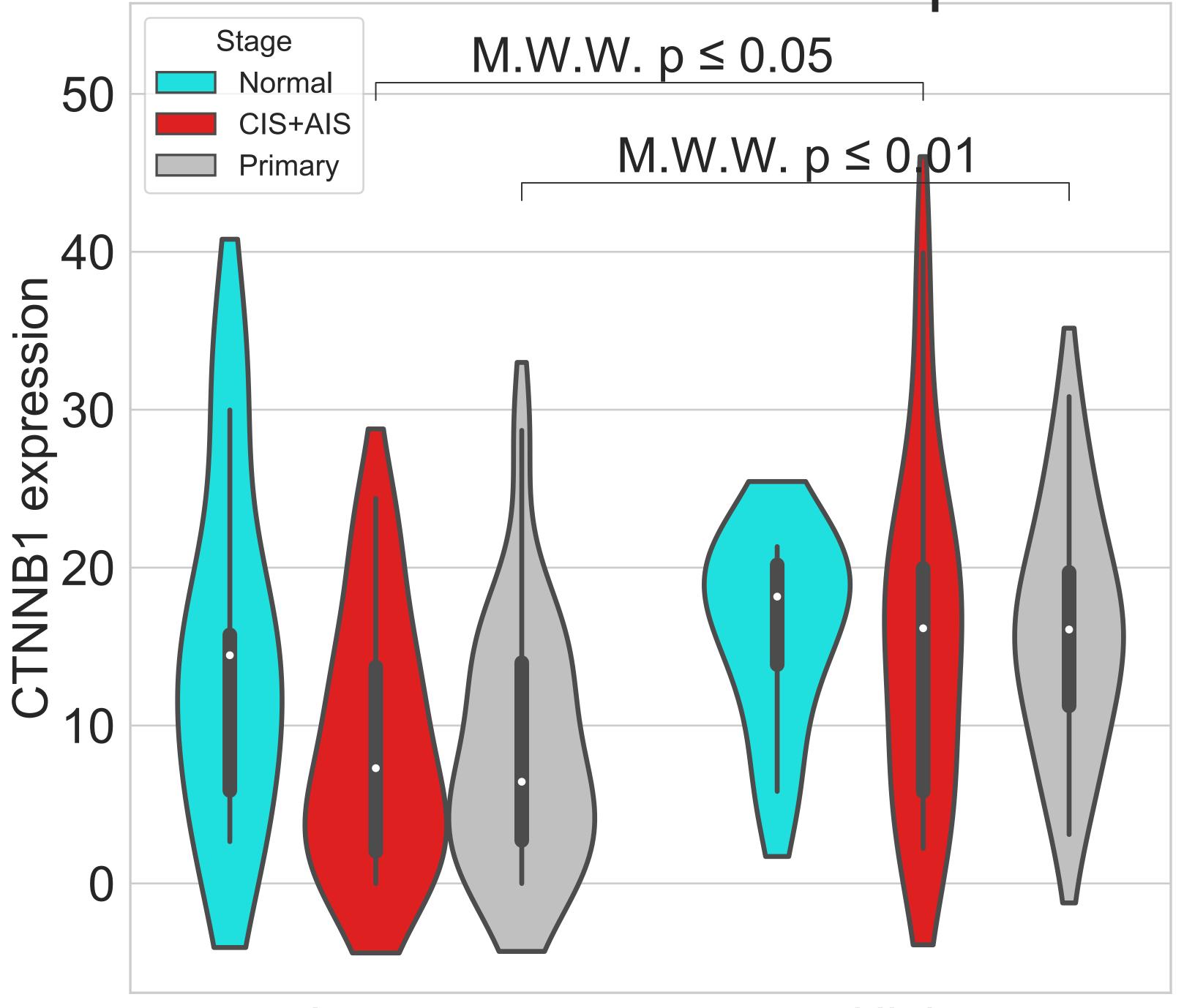
CTNNB1: Kruskal-Wallis p=0.015



Lower

Mutation Shared Proportion (Union & SYN)