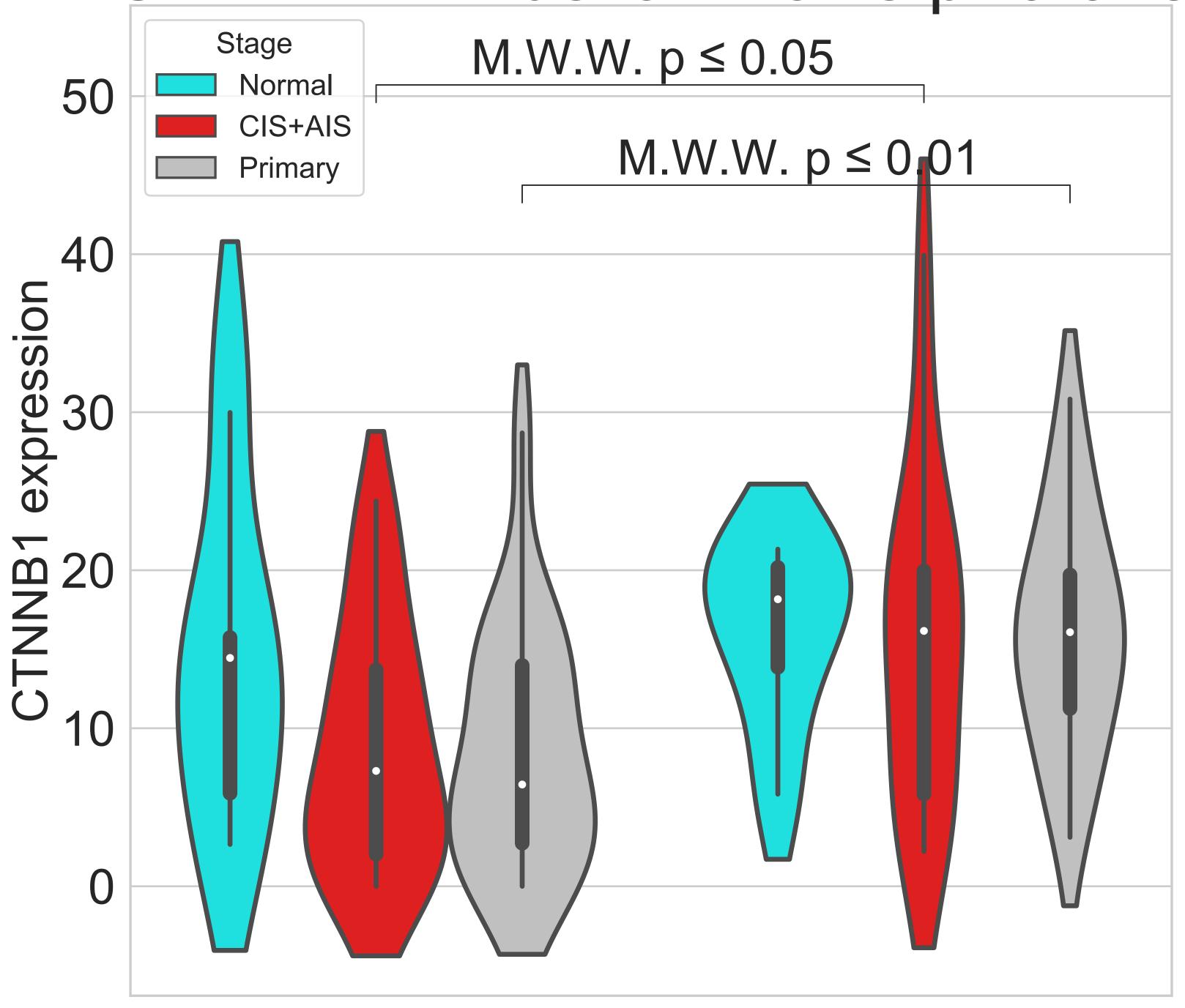
CTNNB1: Kruskal-Wallis p=0.015



Lower
Mutation Shared Count (SYN) per TMB