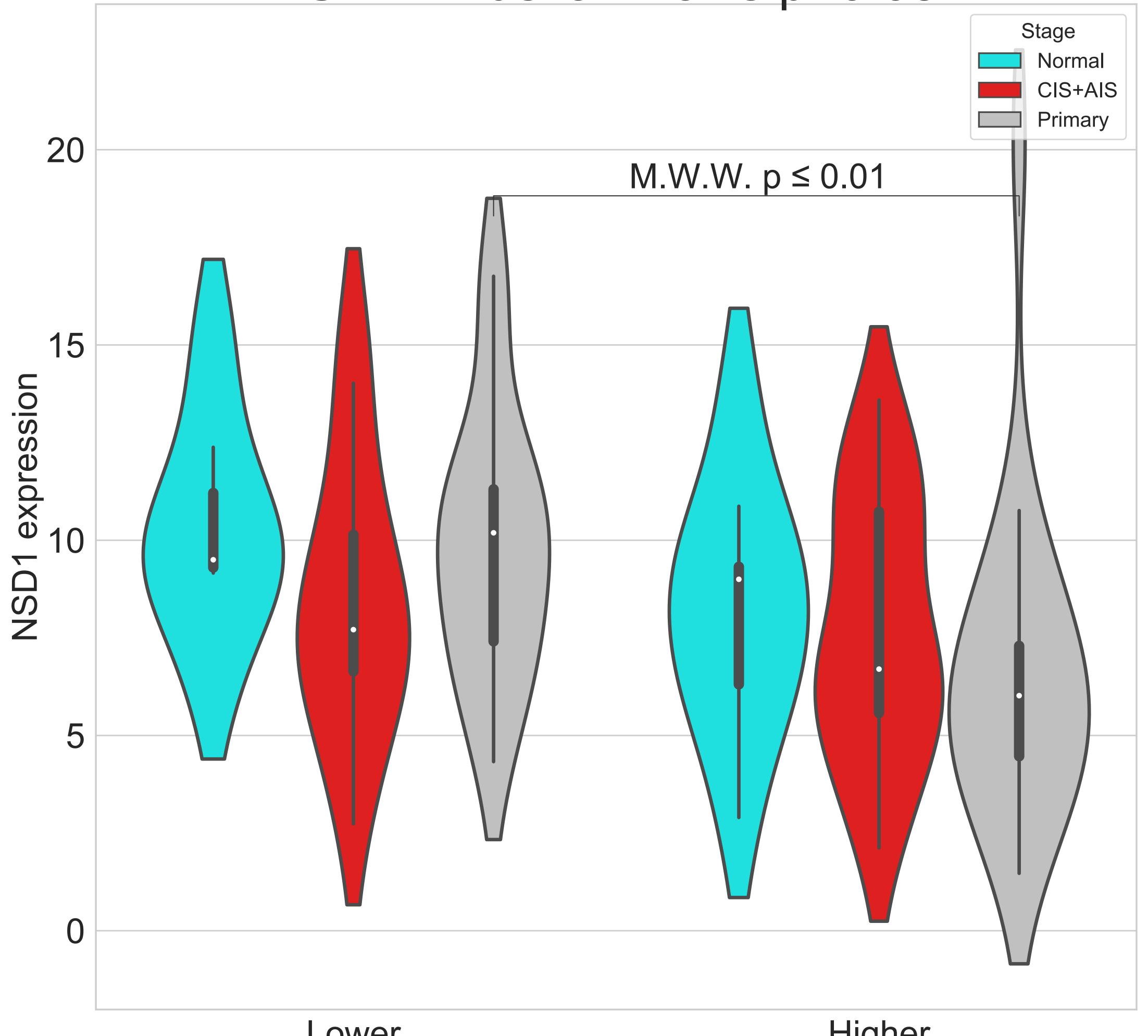
NSD1: Kruskal-Wallis p=0.034



Lower
Mutation Shared Count per TMB