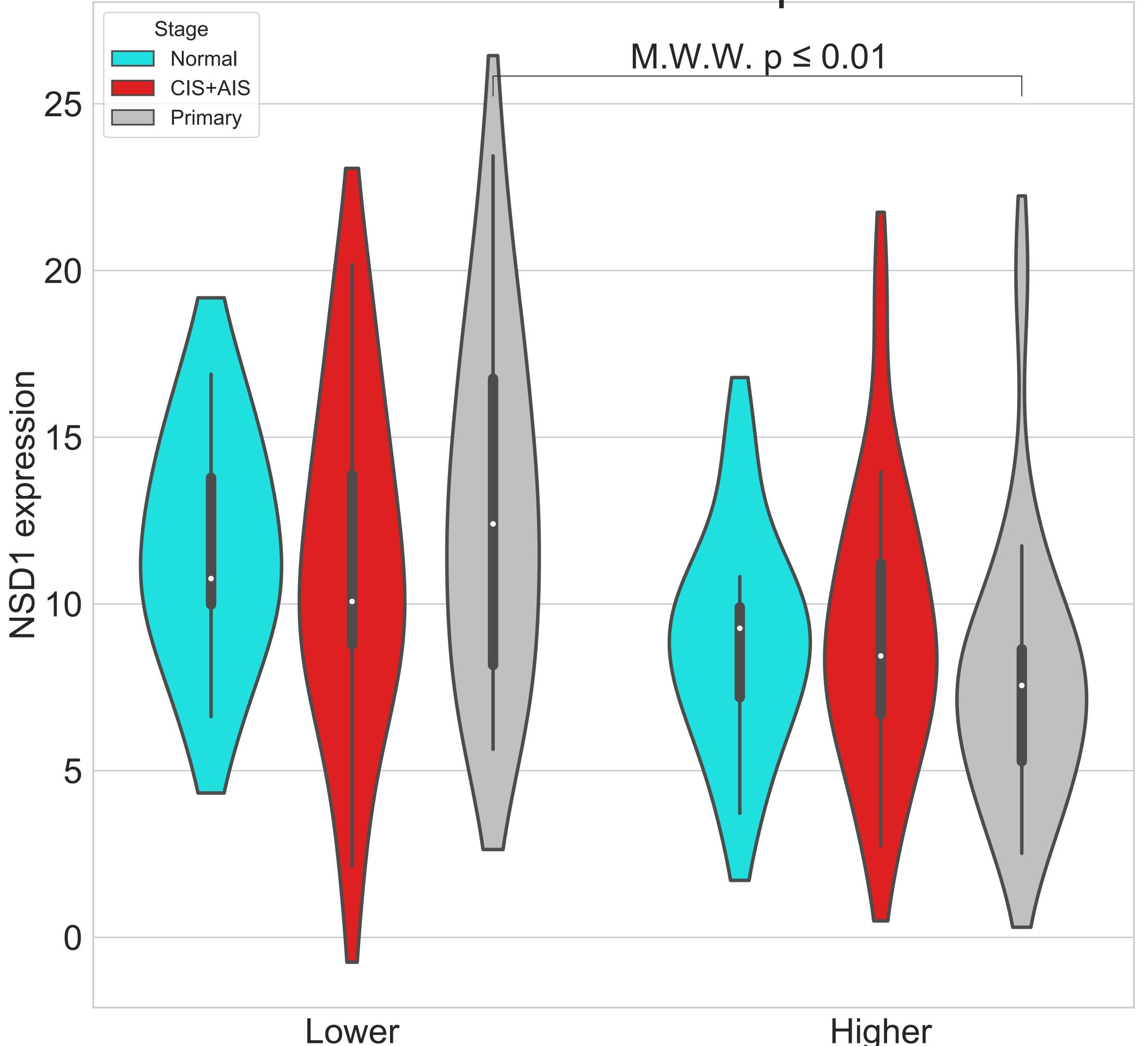
NSD1: Kruskal-Wallis p=0.013



wer
Mutation Shared Count per TMB