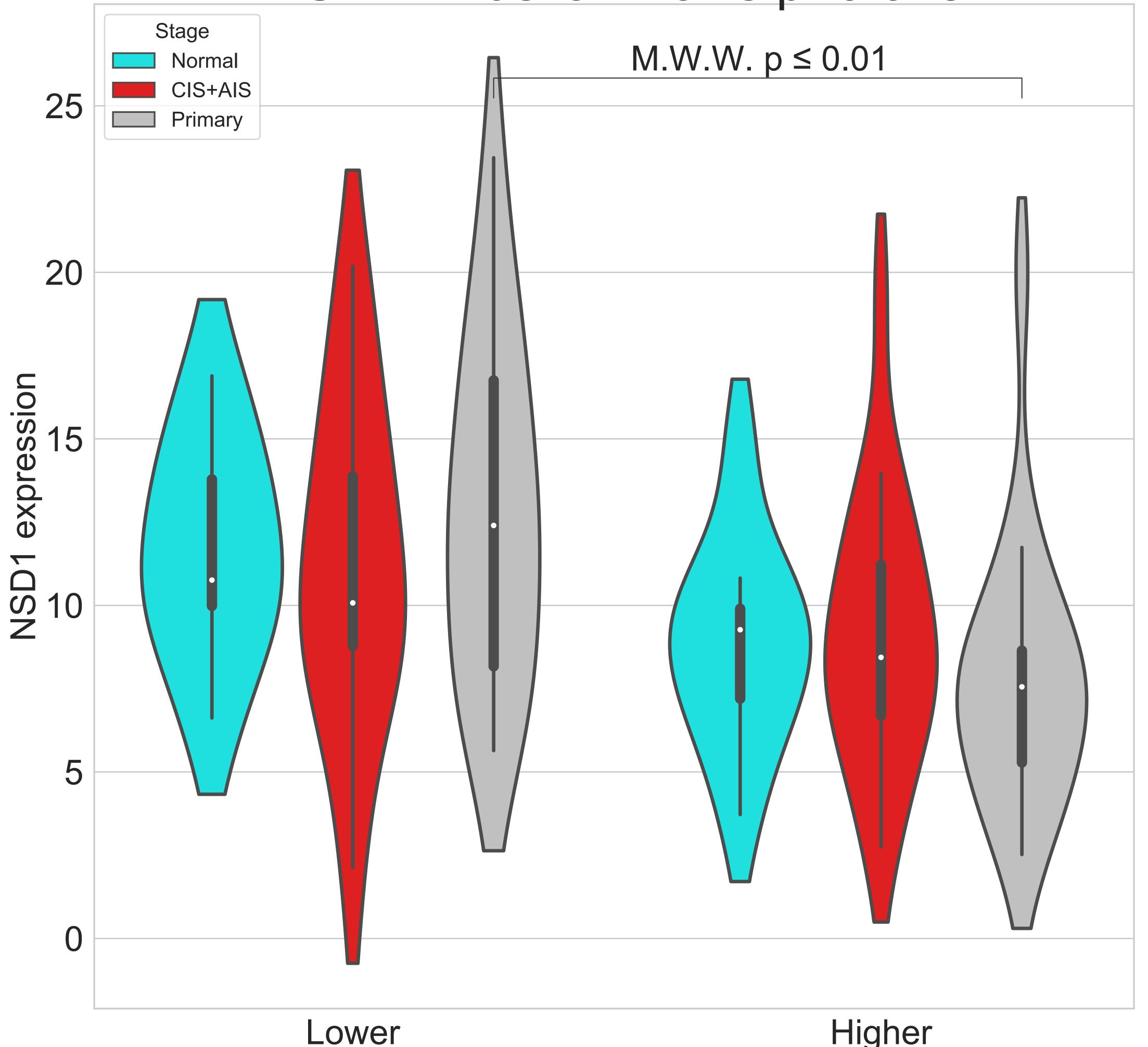
NSD1: Kruskal-Wallis p=0.013



er
Mutation Shared Proportion