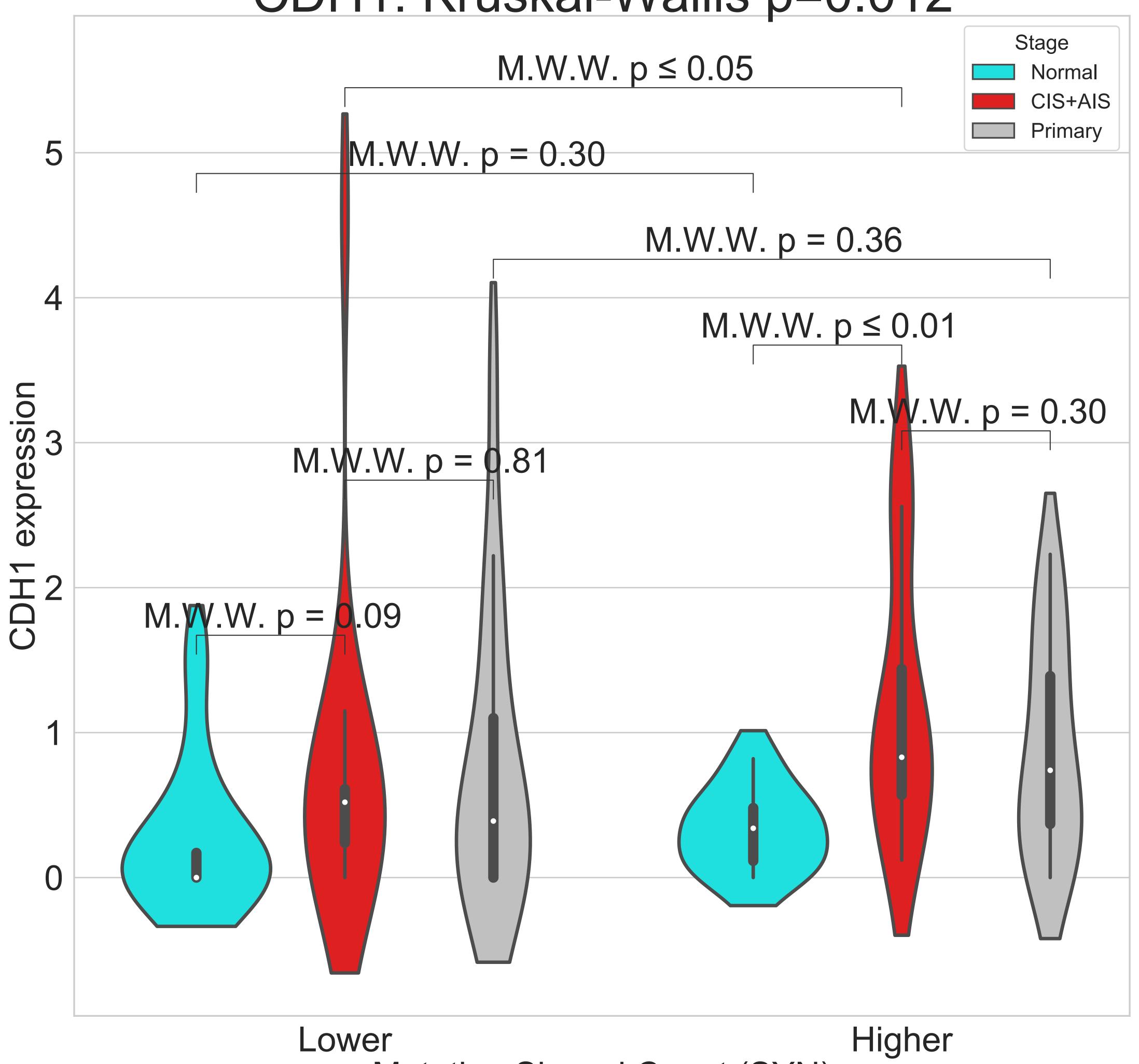
CDH1: Kruskal-Wallis p=0.012



Mutation Shared Count (SYN)