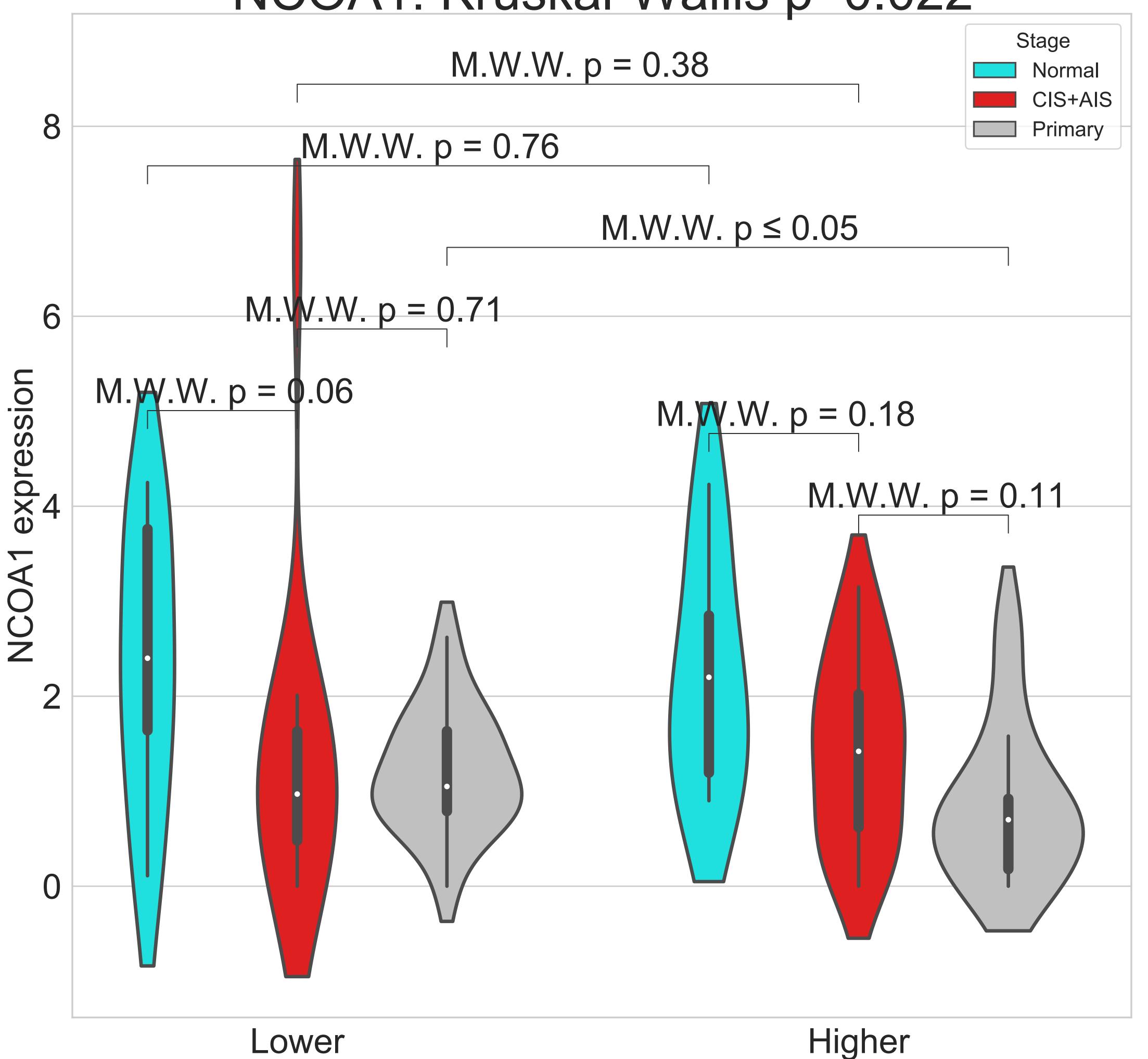
NCOA1: Kruskal-Wallis p=0.022



Higher Mutation Shared Count (SYN)