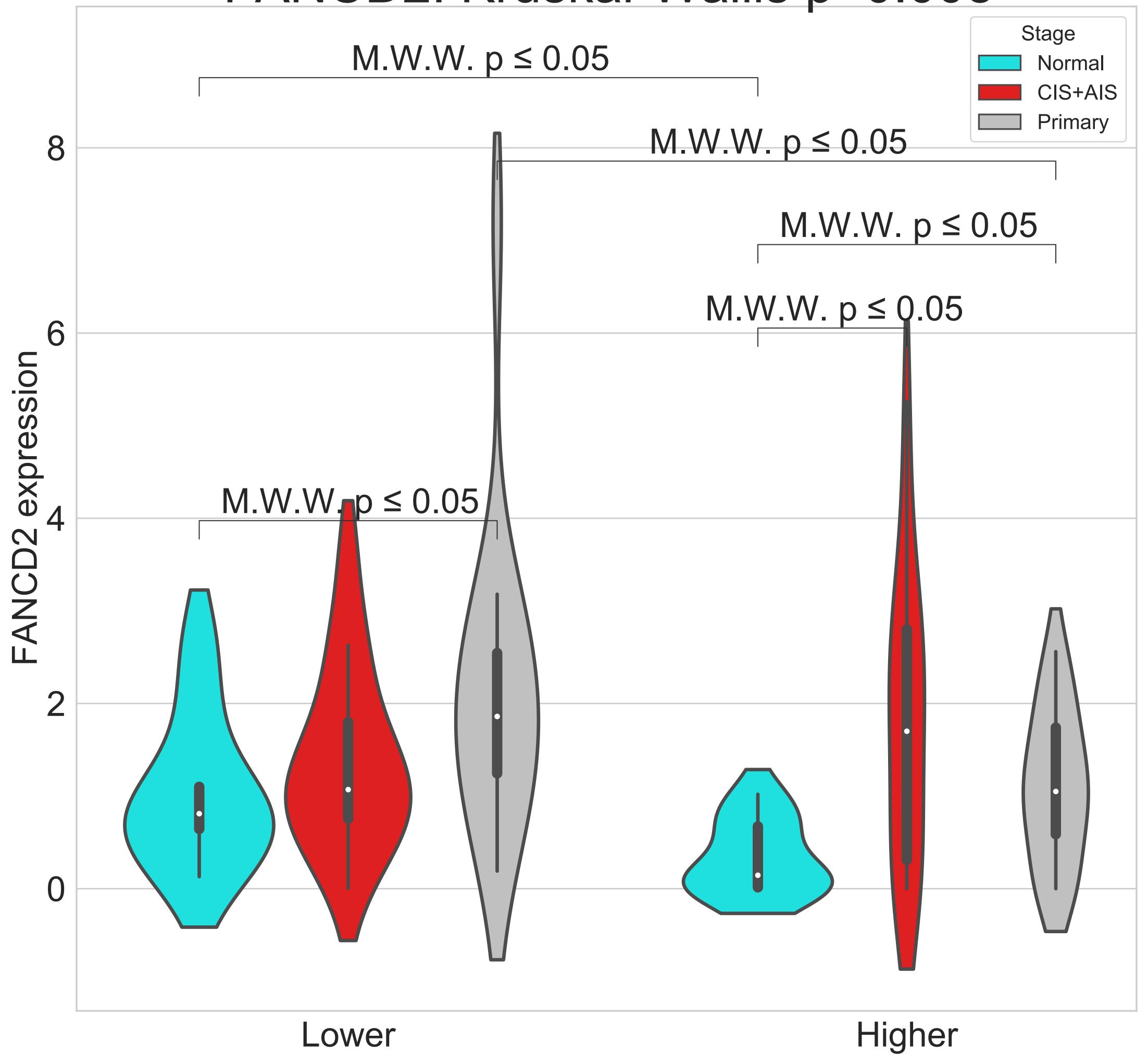
FANCD2: Kruskal-Wallis p=0.008



Mutation Shared Count (SYN)