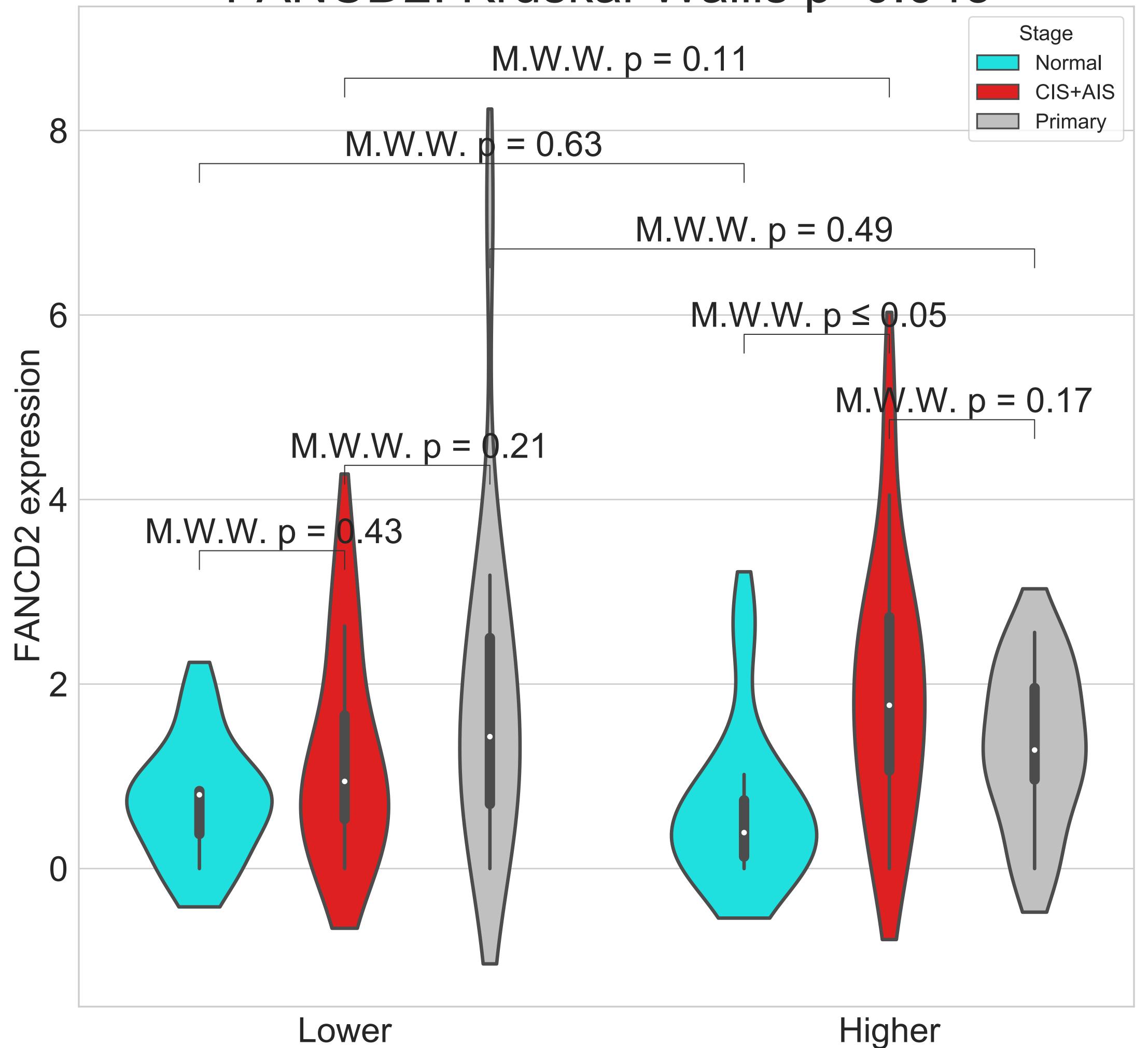
FANCD2: Kruskal-Wallis p=0.043



Mutation Shared Proportion