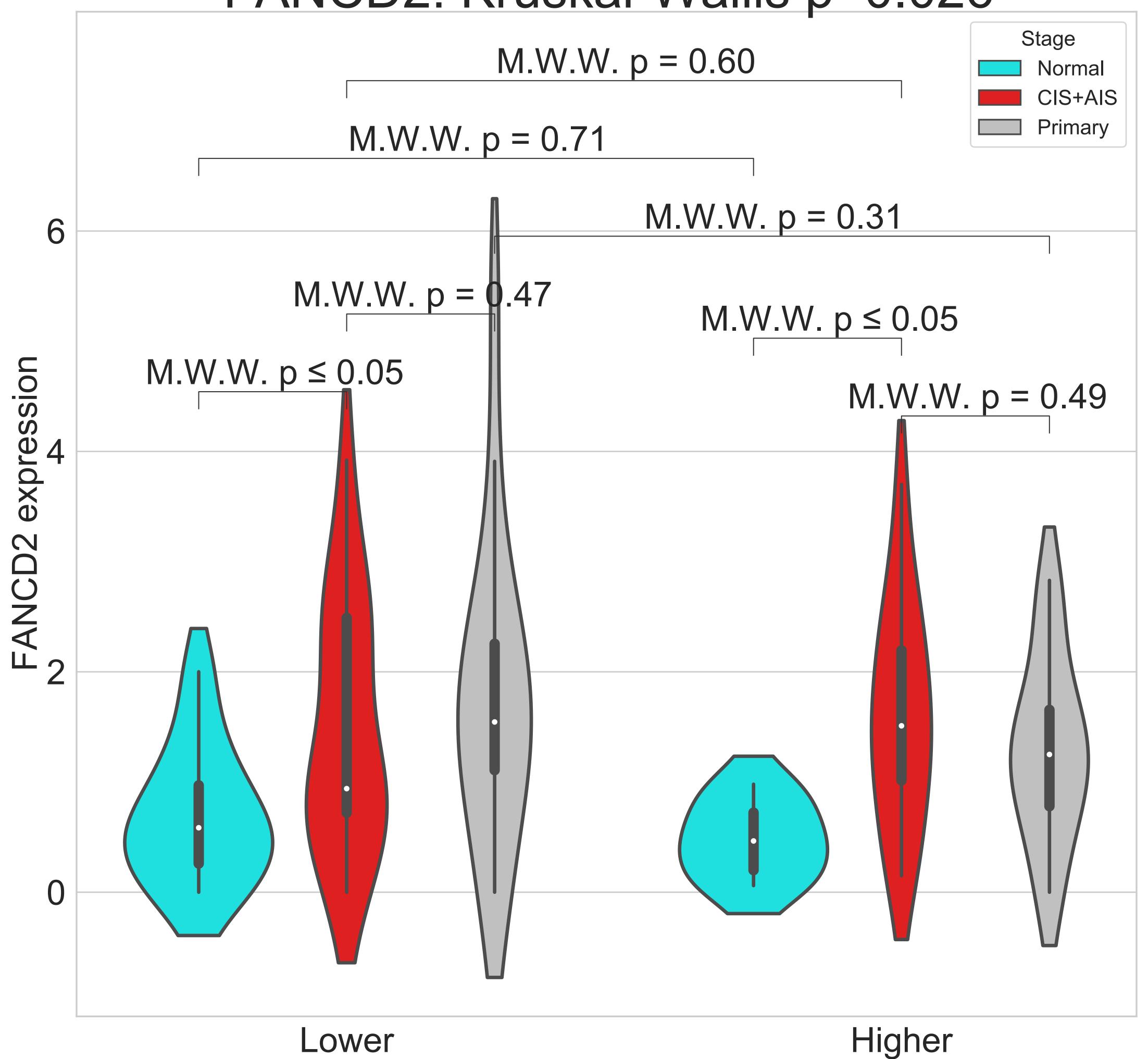
FANCD2: Kruskal-Wallis p=0.026



Mutation Shared Proportion (SYN)