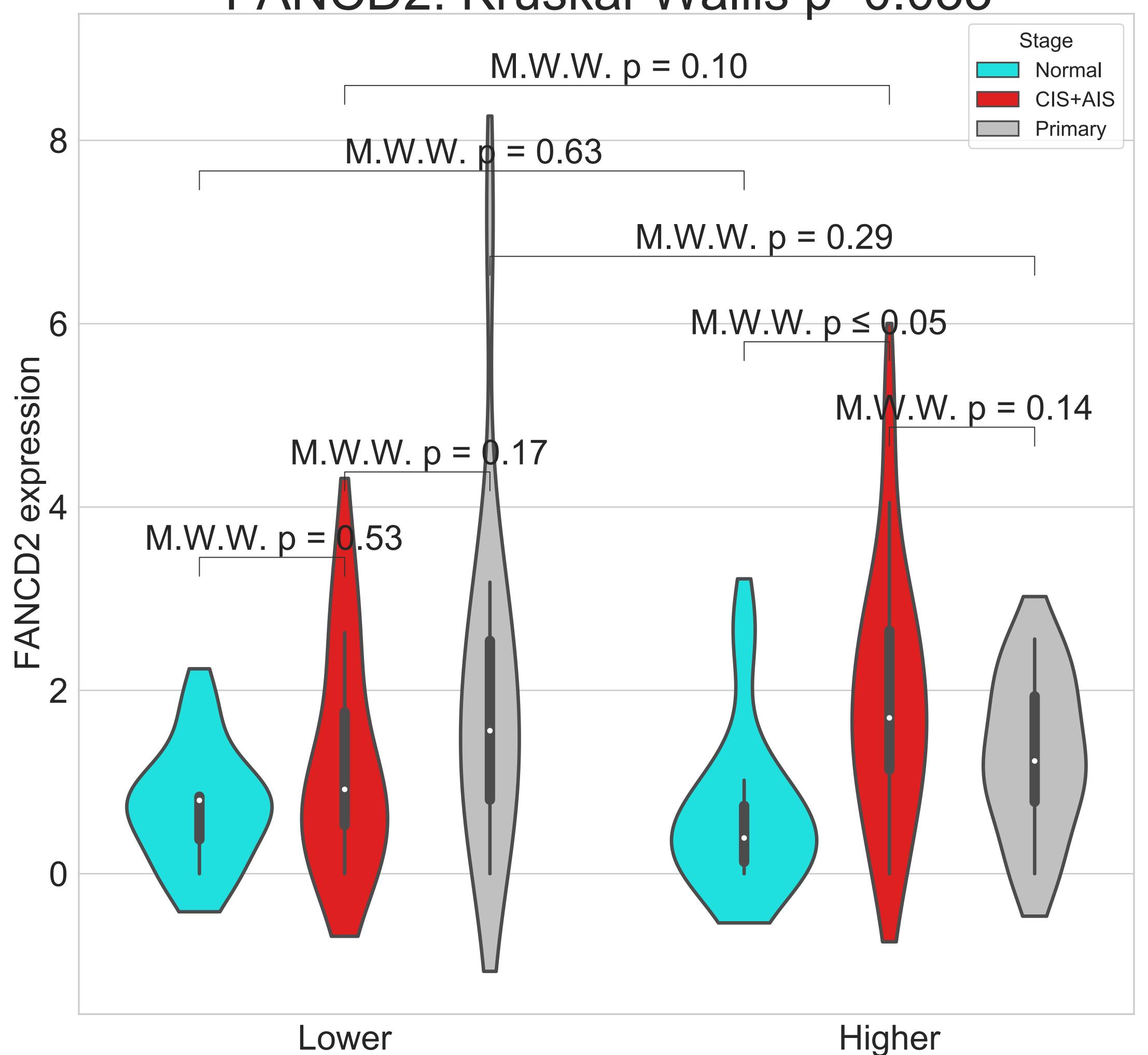
FANCD2: Kruskal-Wallis p=0.033



wer Mutation Shared Proportion (SYN)