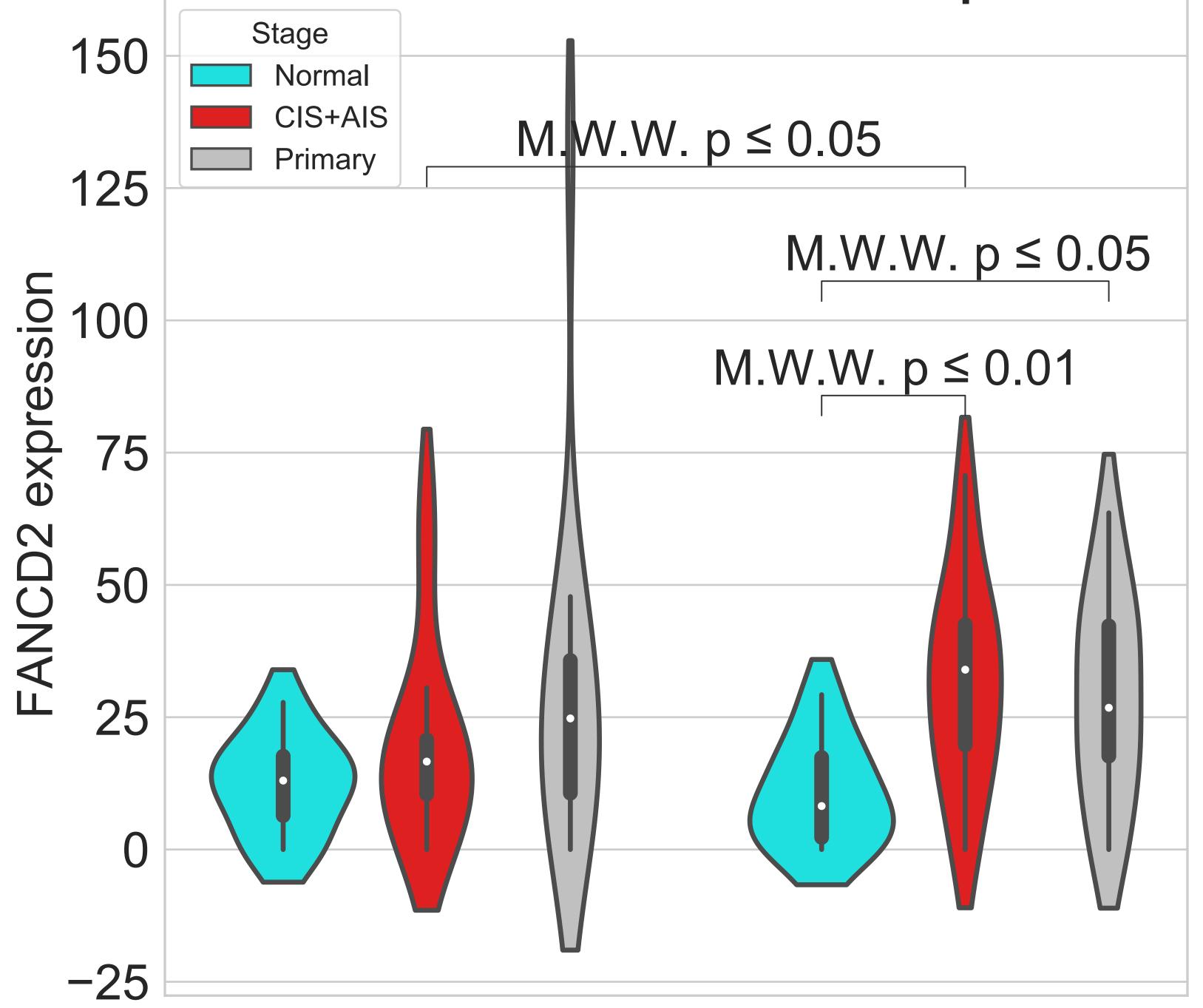
FANCD2: Kruskal-Wallis p=0.020



Lower Higher Mutation Shared Proportion (Union & SYN)