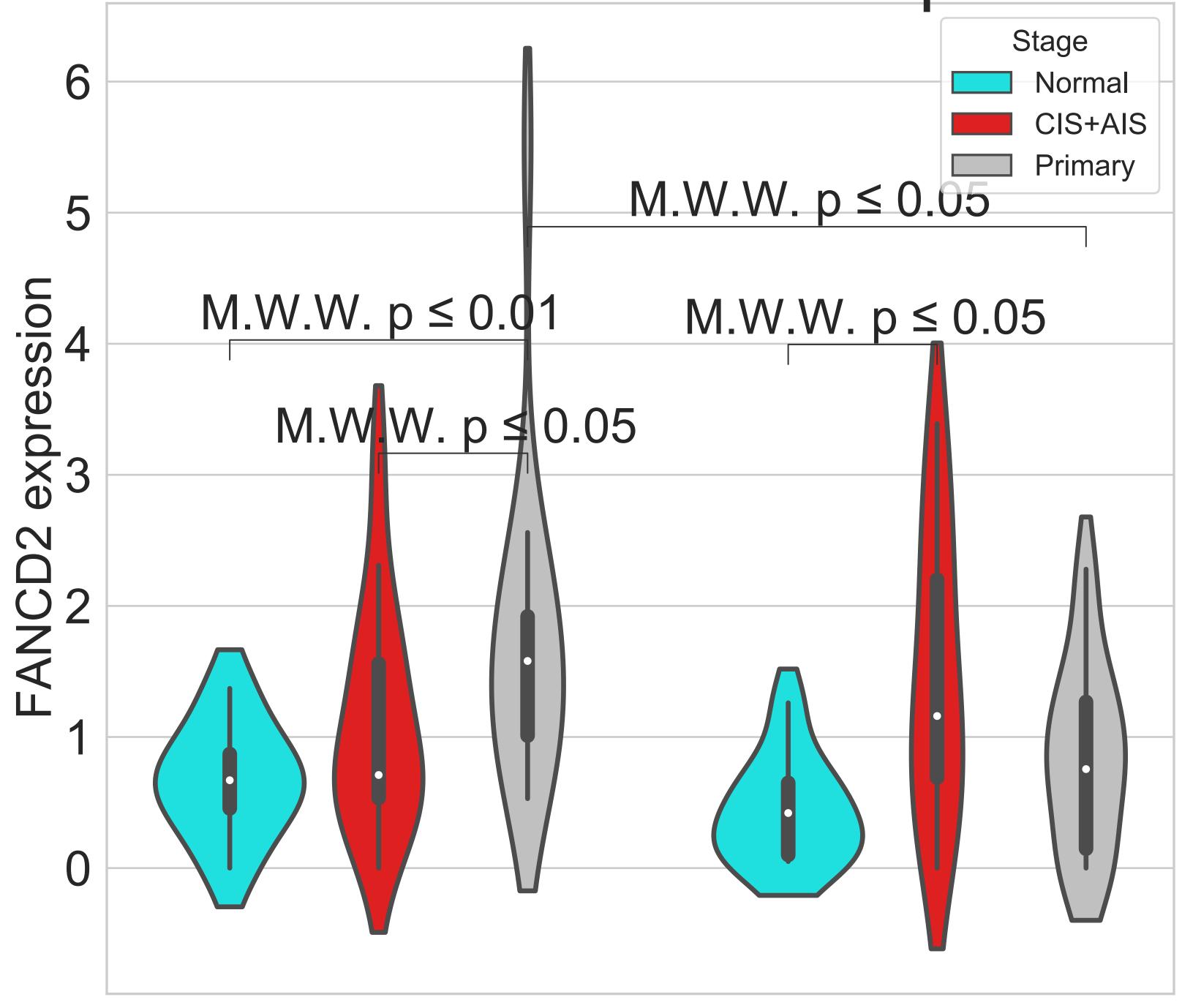
FANCD2: Kruskal-Wallis p=0.005



Lower Higher Mutation Shared Count