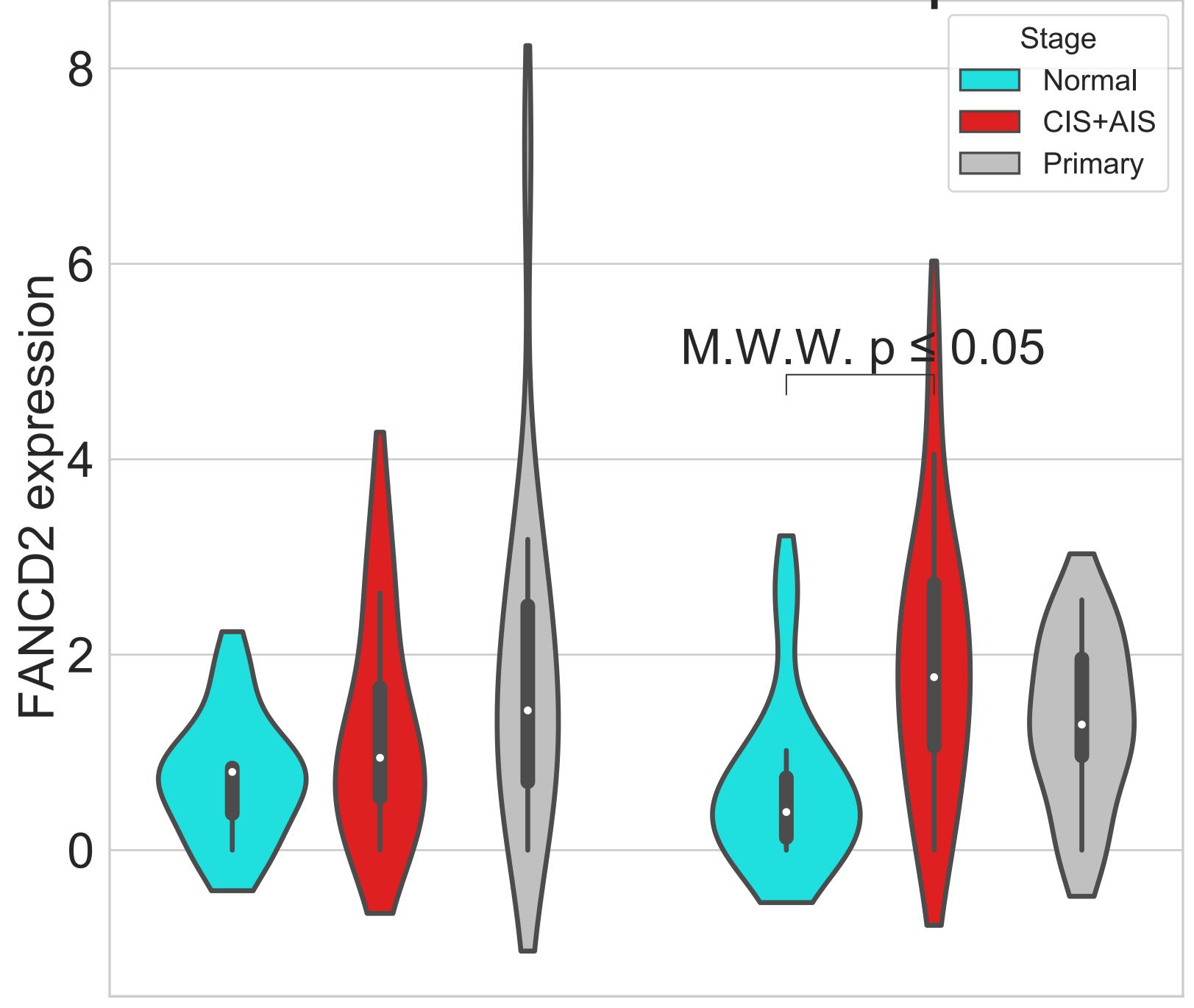
FANCD2: Kruskal-Wallis p=0.043



Lower Higher Mutation Shared Count per TMB