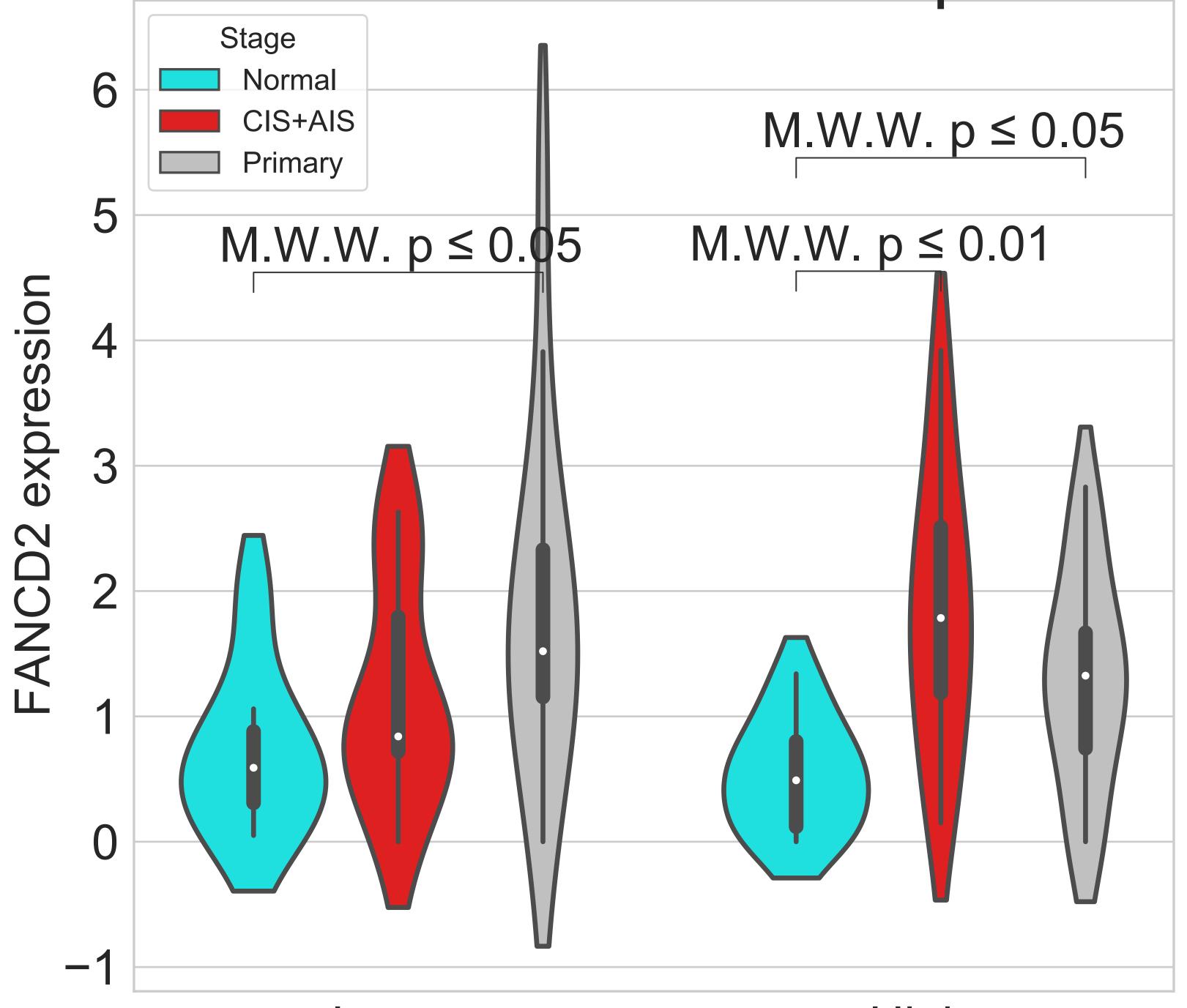
FANCD2: Kruskal-Wallis p=0.007



Lower
Higher
Mutation Shared Proportion