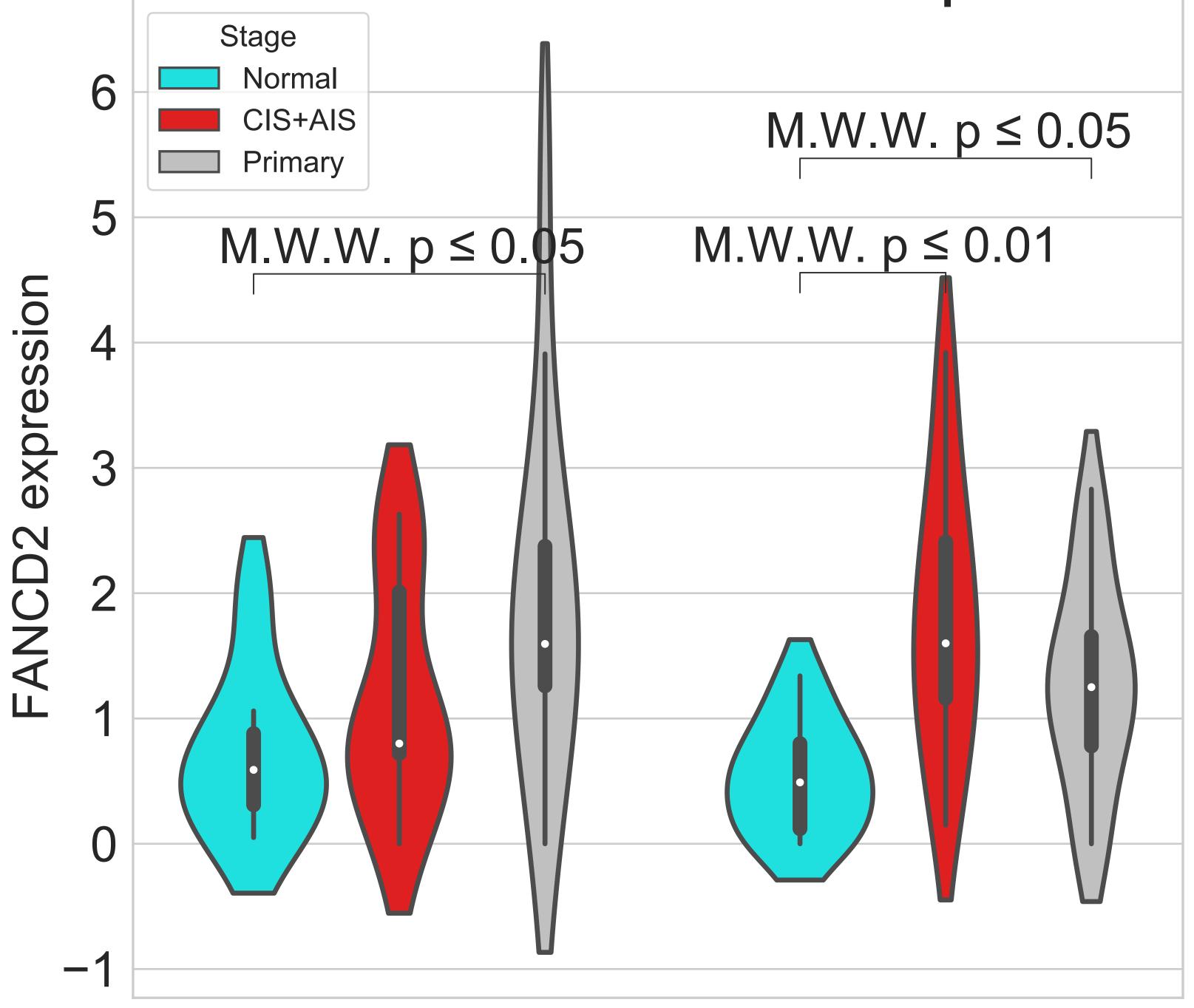
FANCD2: Kruskal-Wallis p=0.007



Lower
Mutation Shared Proportion (SYN)