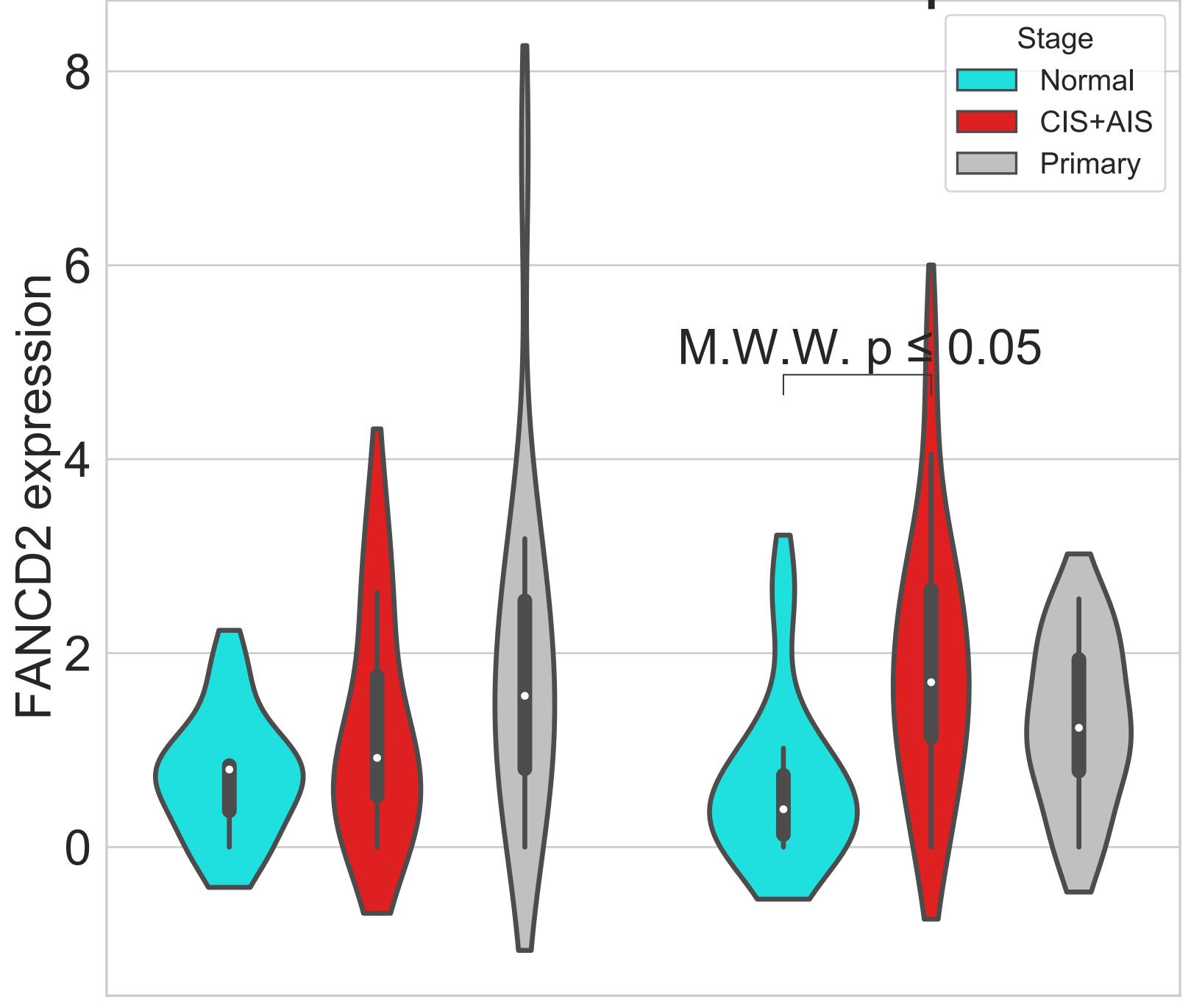
FANCD2: Kruskal-Wallis p=0.033



Lower
Mutation Shared Proportion (SYN)