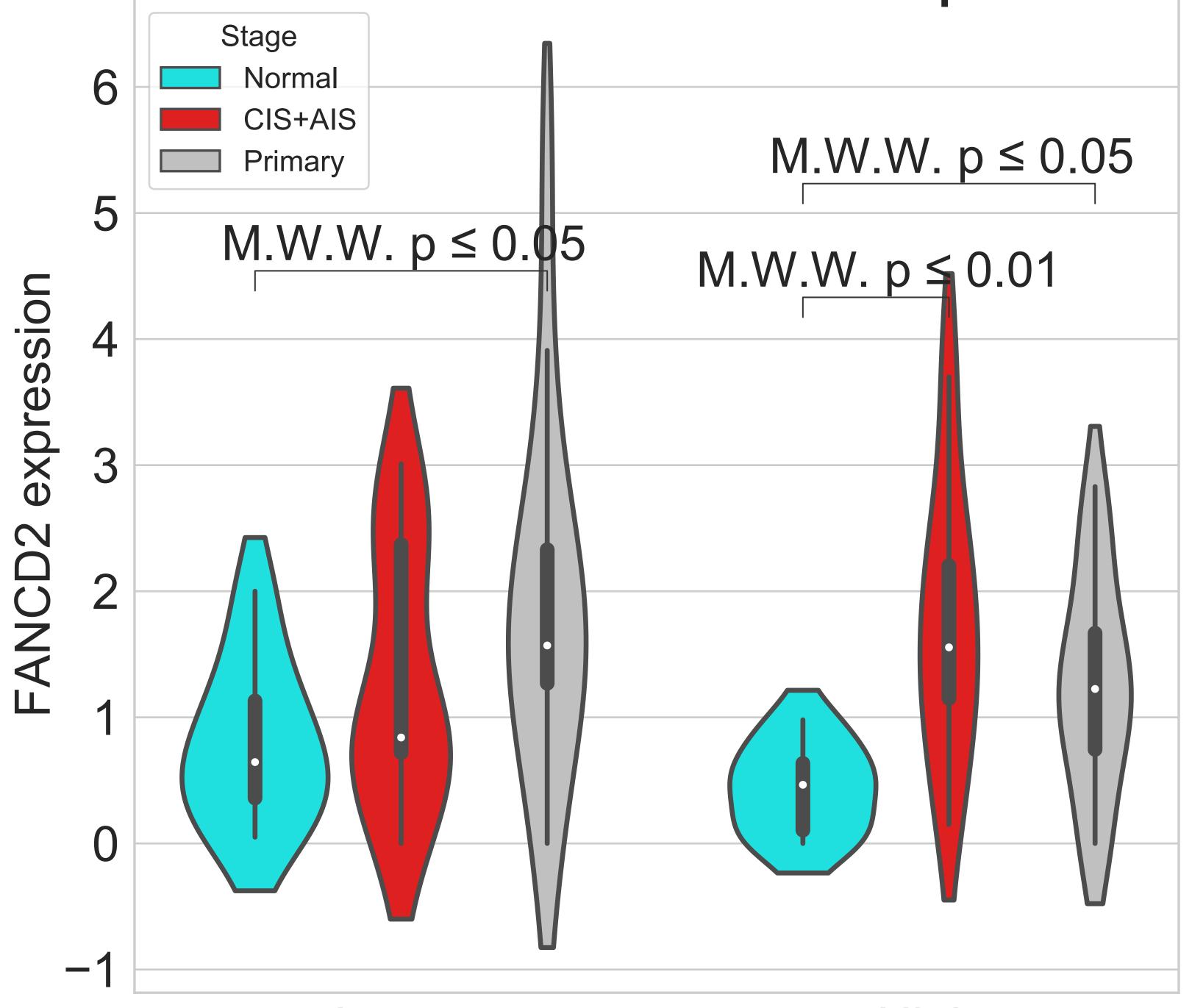
FANCD2: Kruskal-Wallis p=0.009



Lower

Mutation Shared Proportion (Union & SYN)