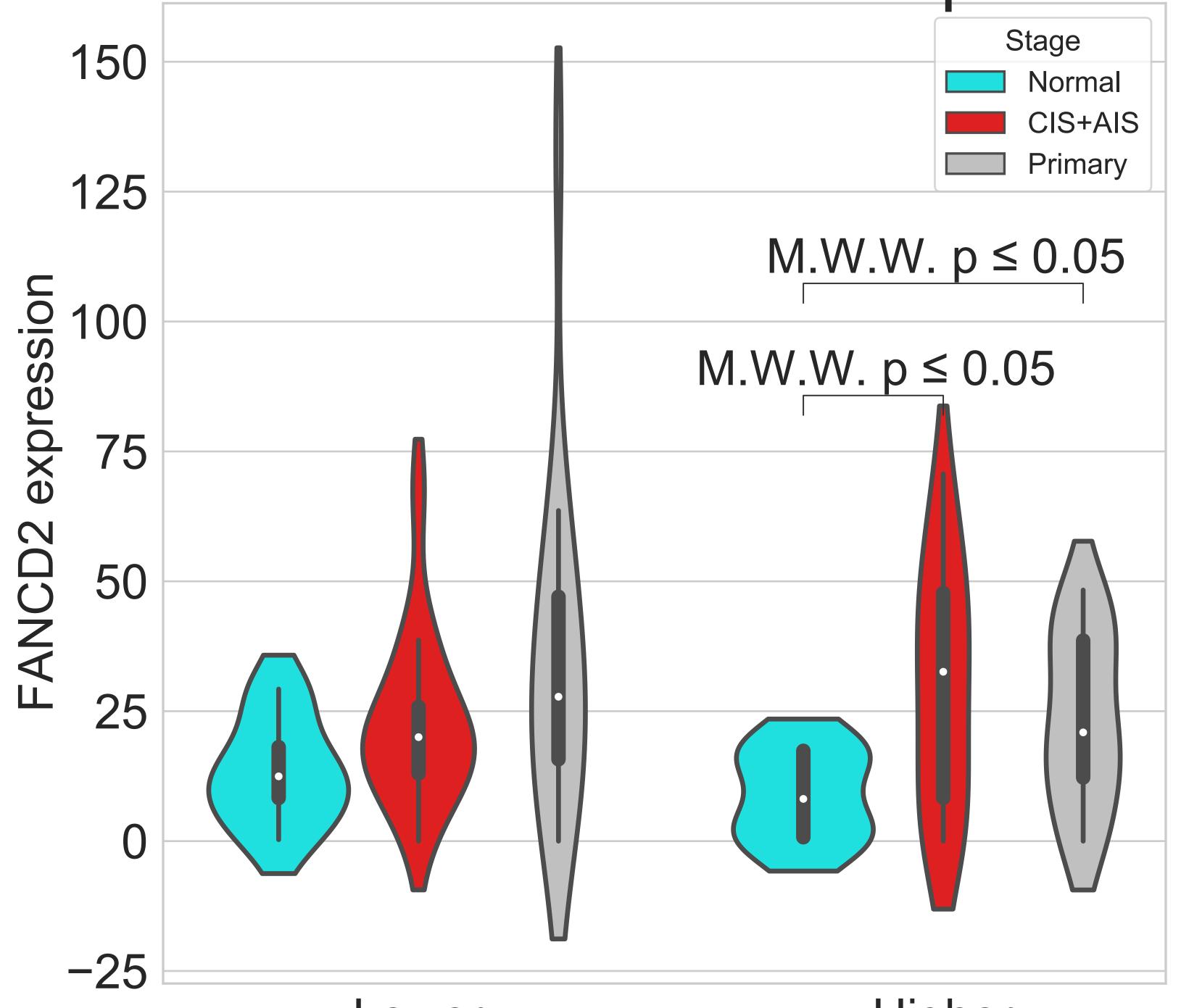
FANCD2: Kruskal-Wallis p=0.046



Lower
Mutation Shared Count (SYN)