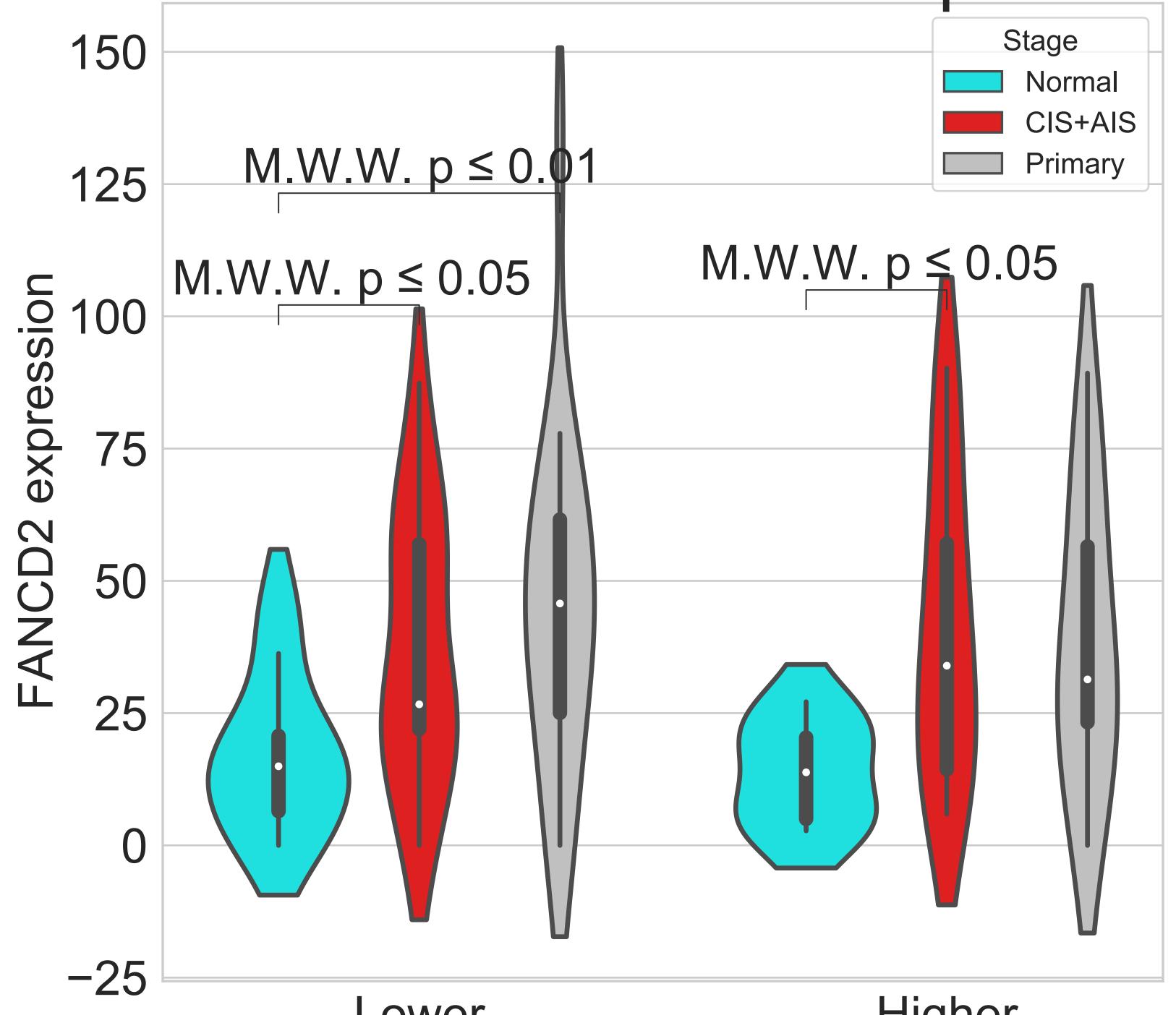
FANCD2: Kruskal-Wallis p=0.031



Lower
Mutation Shared Proportion (Union)