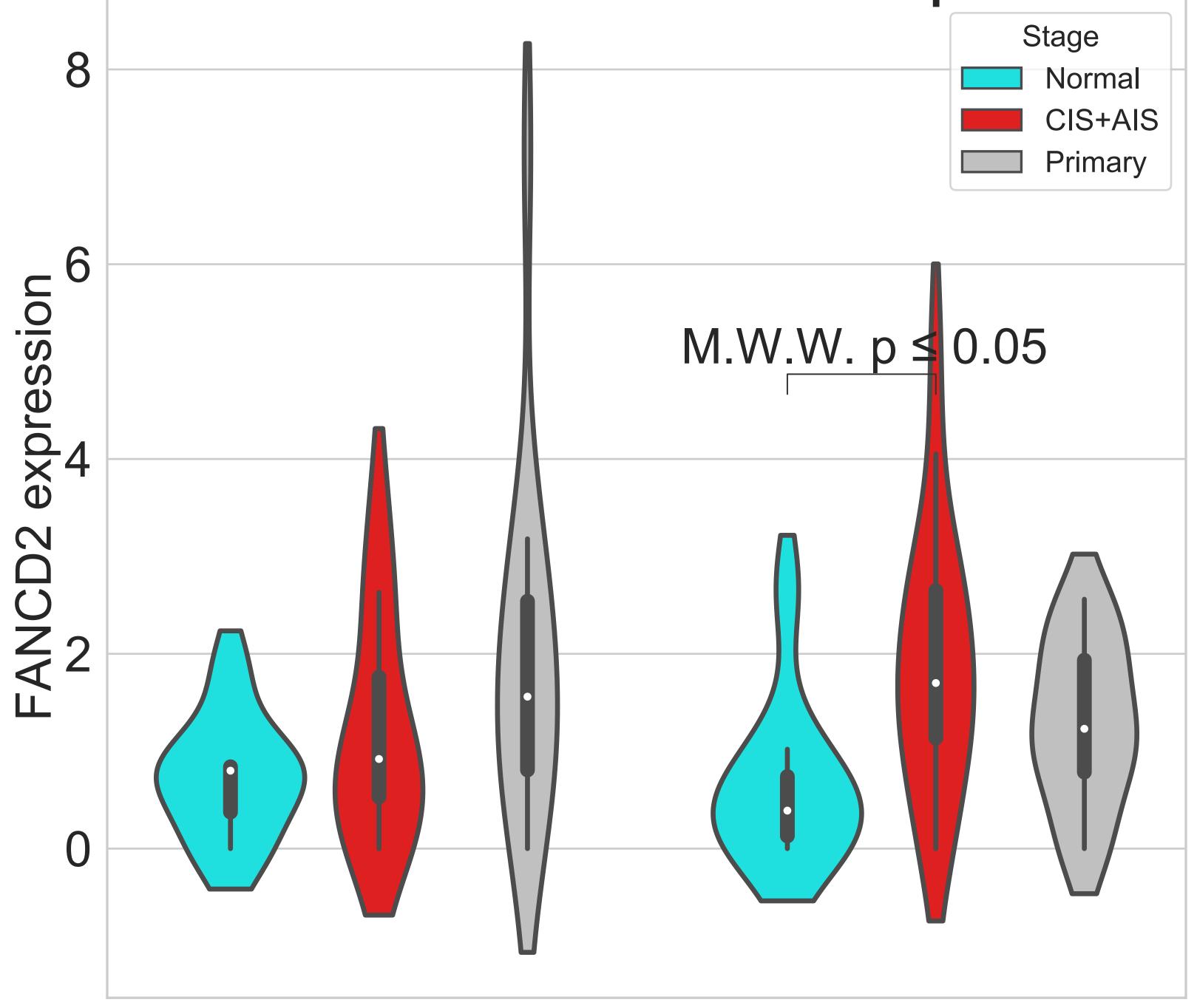
FANCD2: Kruskal-Wallis p=0.033



Lower
Mutation Shared Count (SYN) per TMB