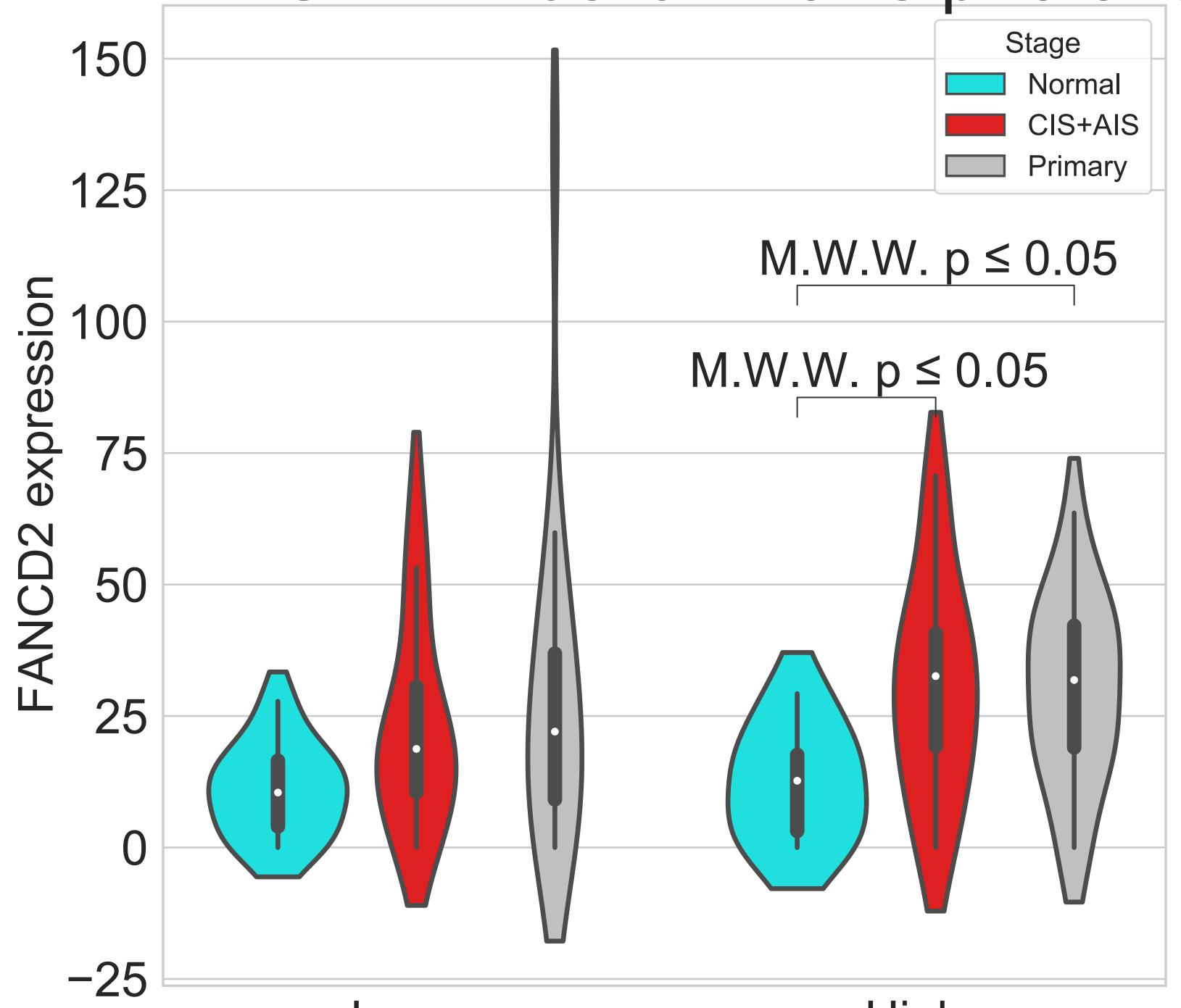
FANCD2: Kruskal-Wallis p=0.040



Lower
Mutation Shared Count (SYN) per TMB