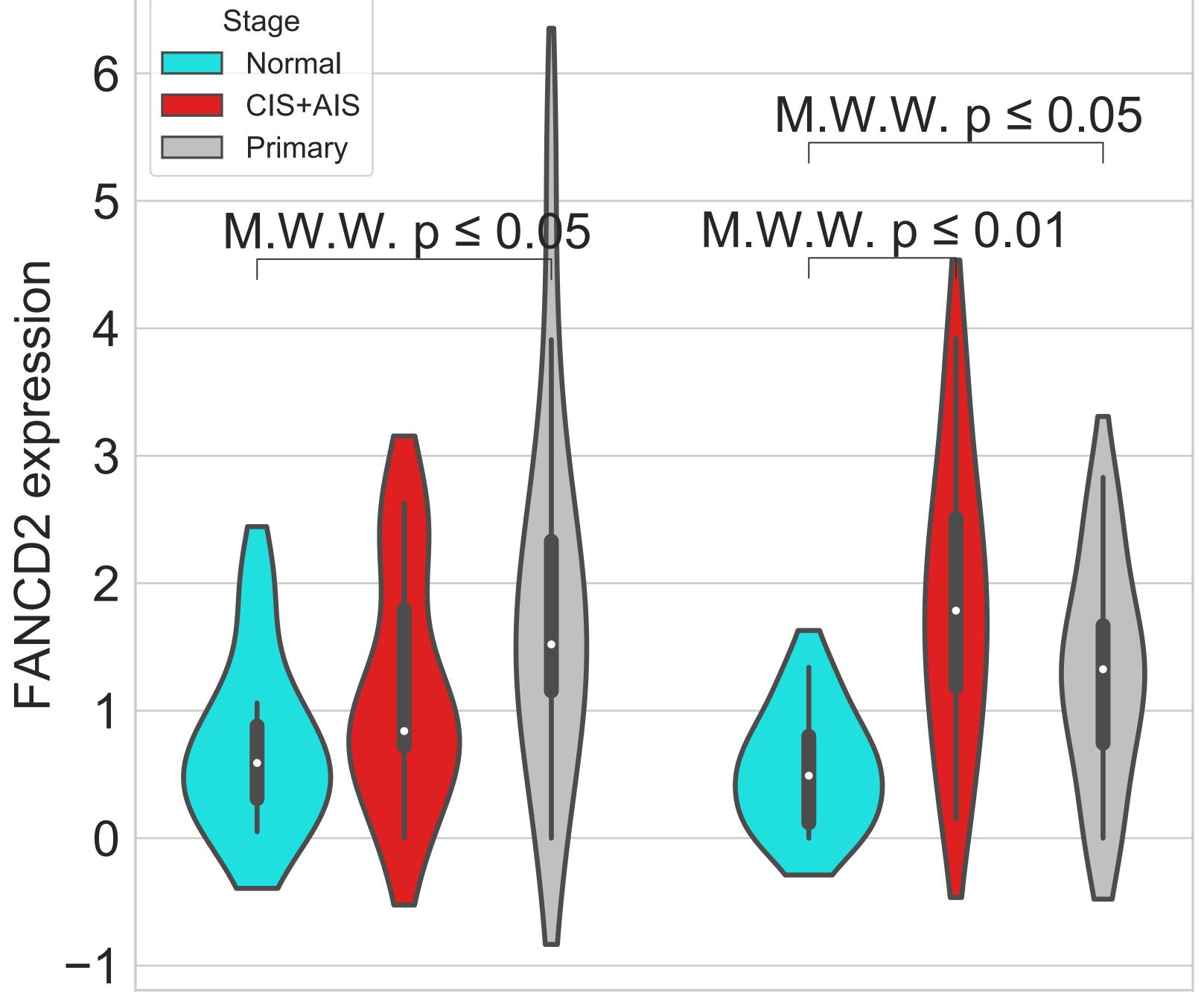
FANCD2: Kruskal-Wallis p=0.007

Stage
Normal



Lower
Mutation Shared Count per TMB