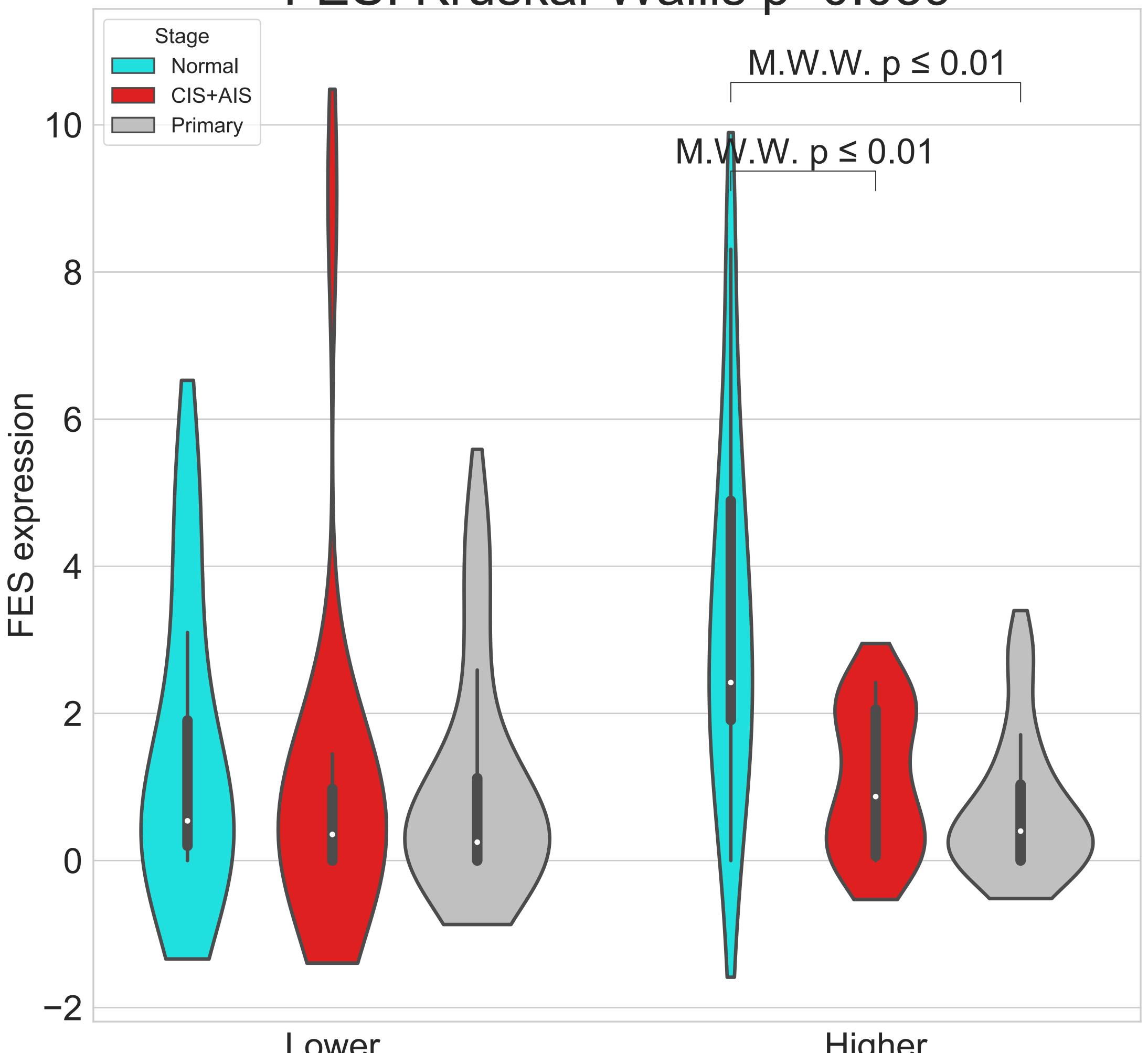
FES: Kruskal-Wallis p=0.039



Lower
Mutation Shared Count per TMB