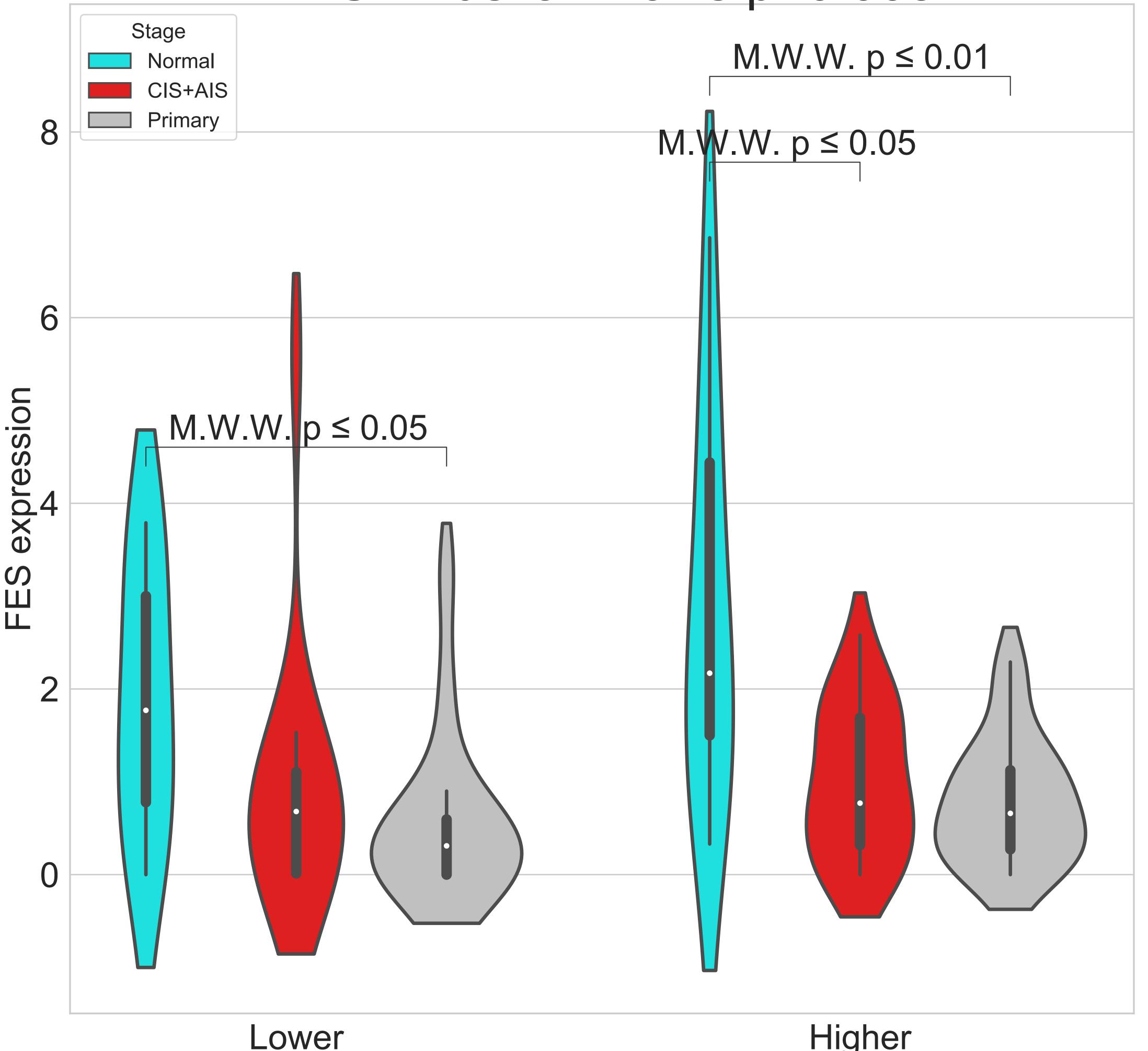
FES: Kruskal-Wallis p=0.005



ver
Mutation Shared Count per TMB