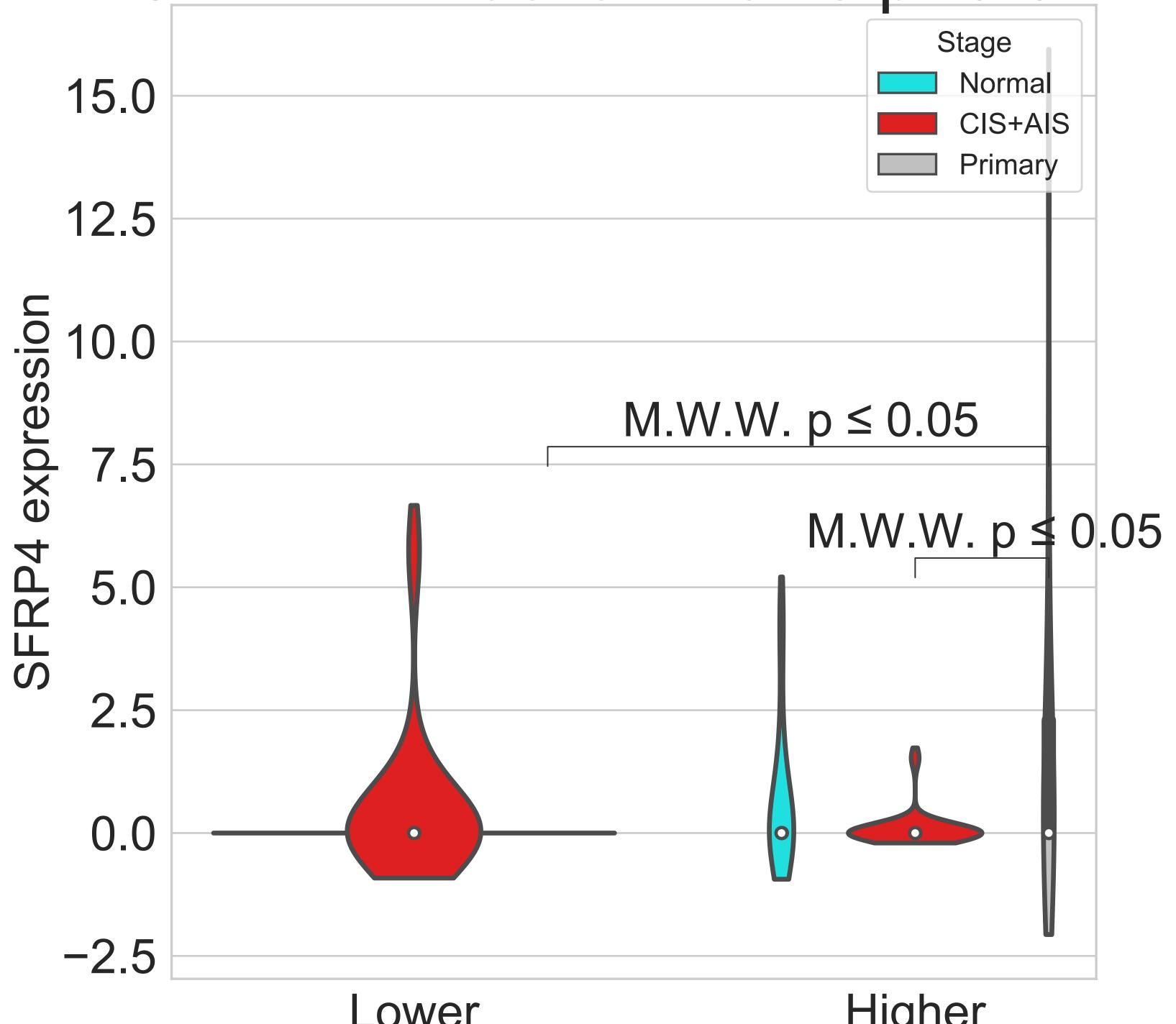
SFRP4: Kruskal-Wallis p=0.047



Lower
Mutation Shared Count per TMB