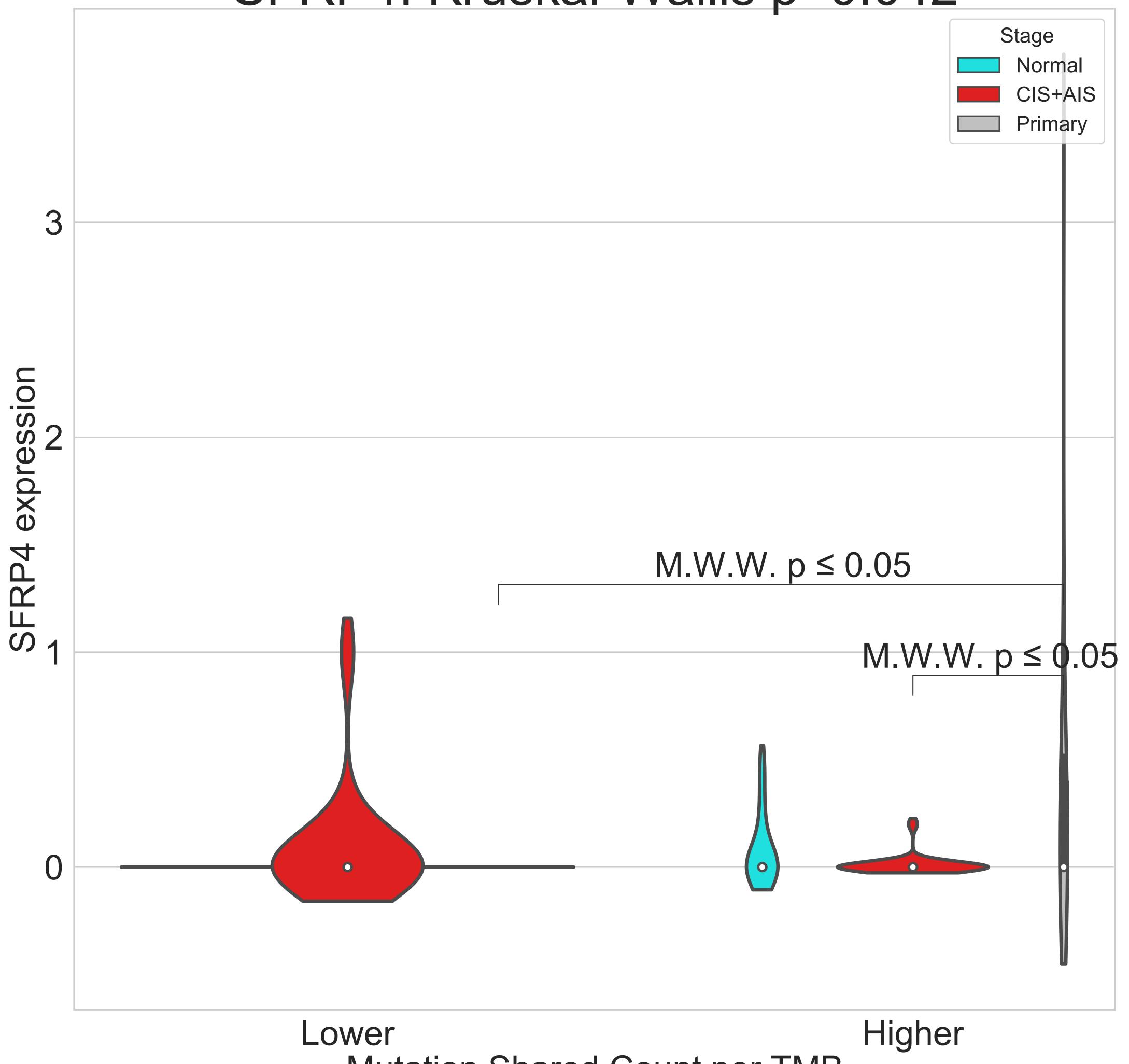
SFRP4: Kruskal-Wallis p=0.042



Mutation Shared Count per TMB