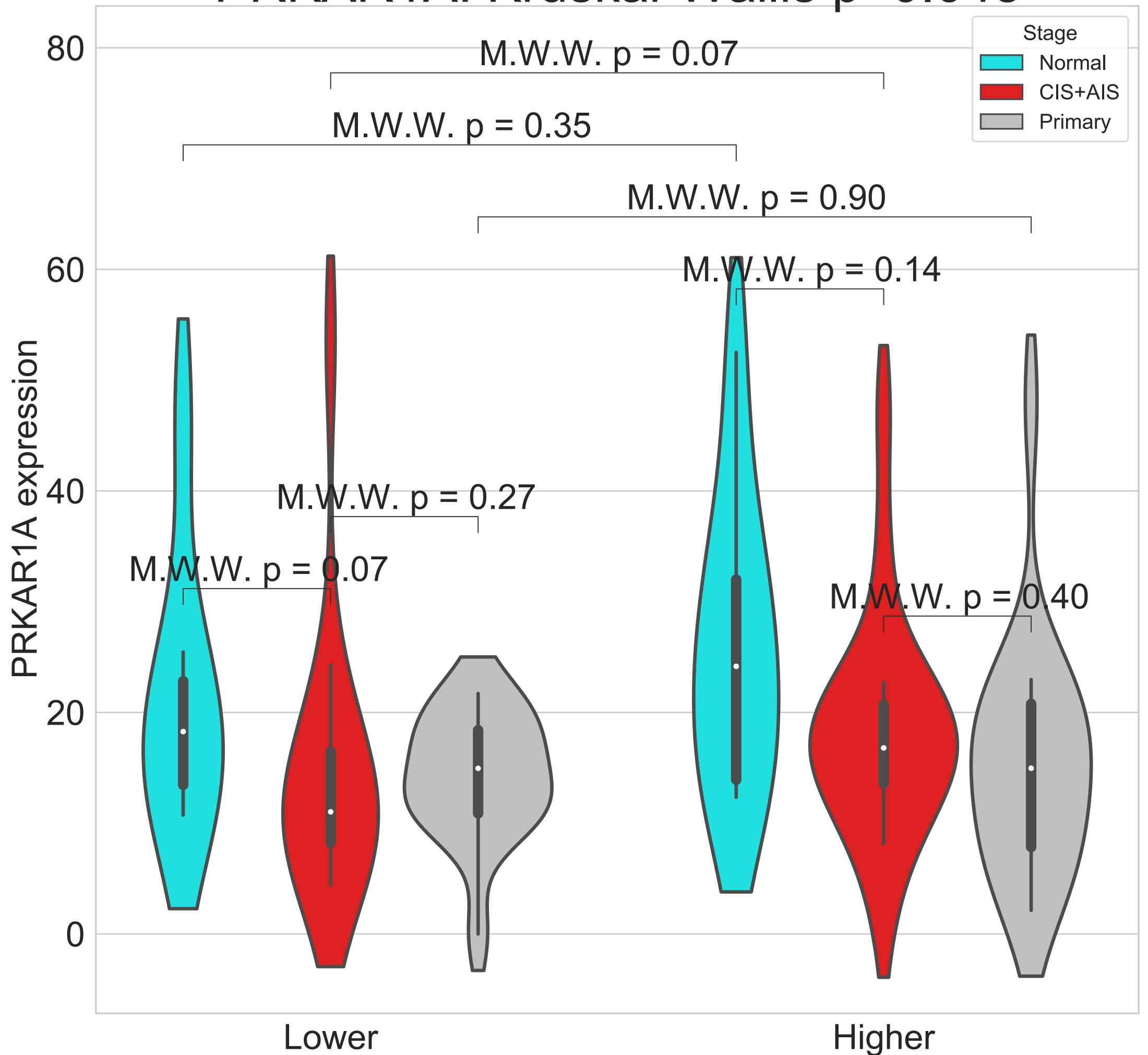
PRKAR1A: Kruskal-Wallis p=0.048



wer
Mutation Shared Count per TMB