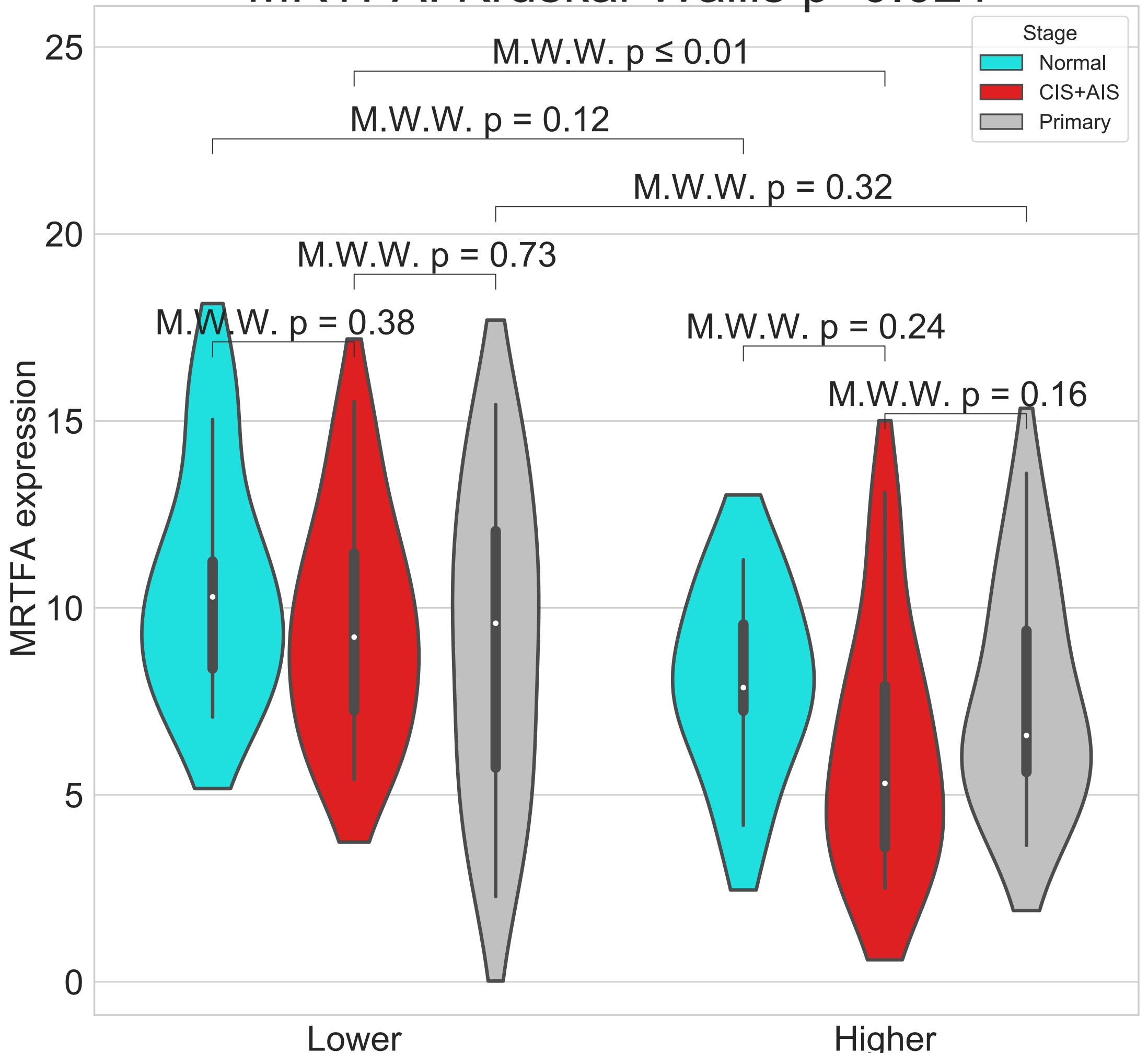
MRTFA: Kruskal-Wallis p=0.021



wer
Mutation Shared Count per TMB