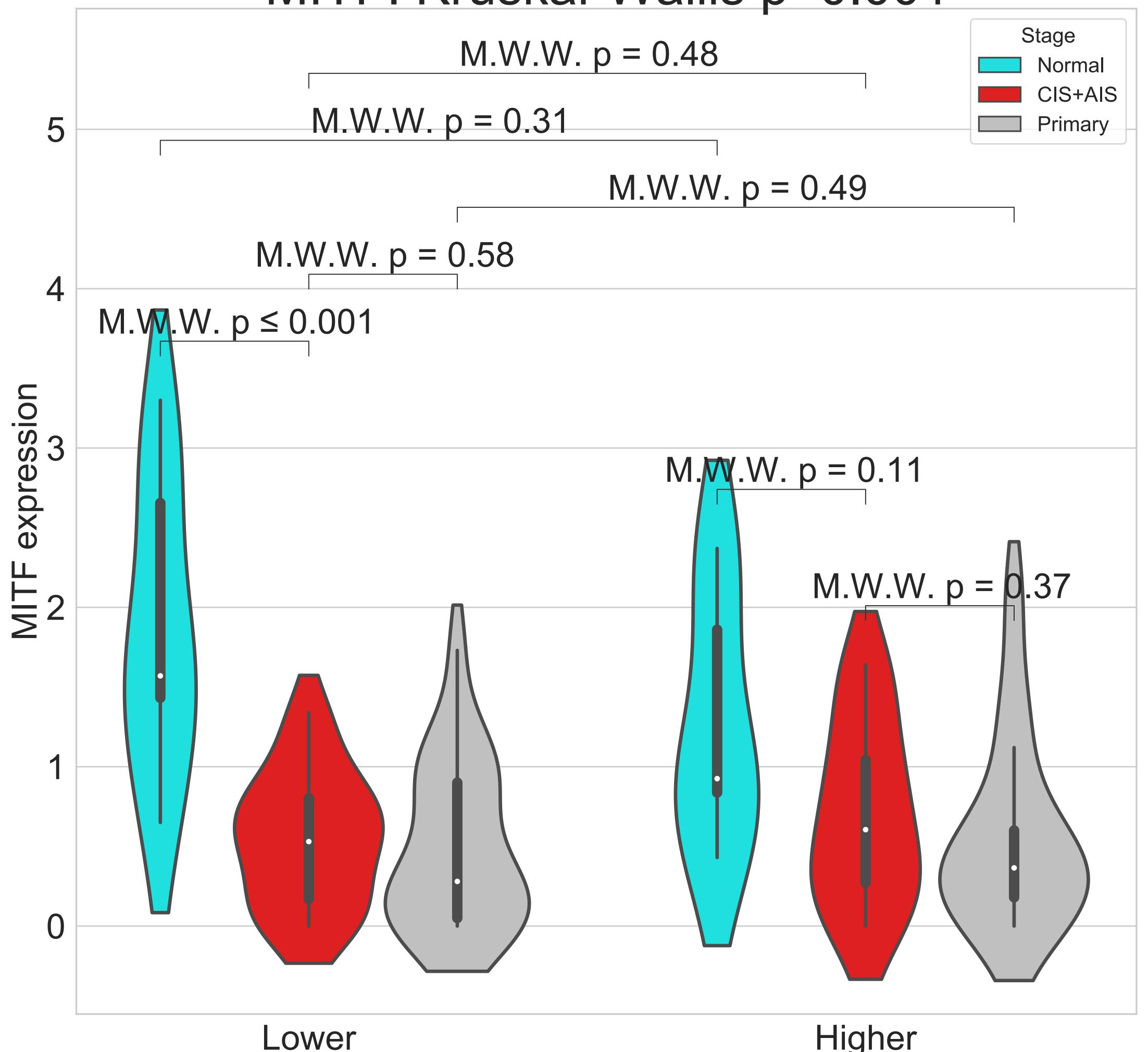
MITF: Kruskal-Wallis p=0.001



wer

Mutation Shared Count per TMB