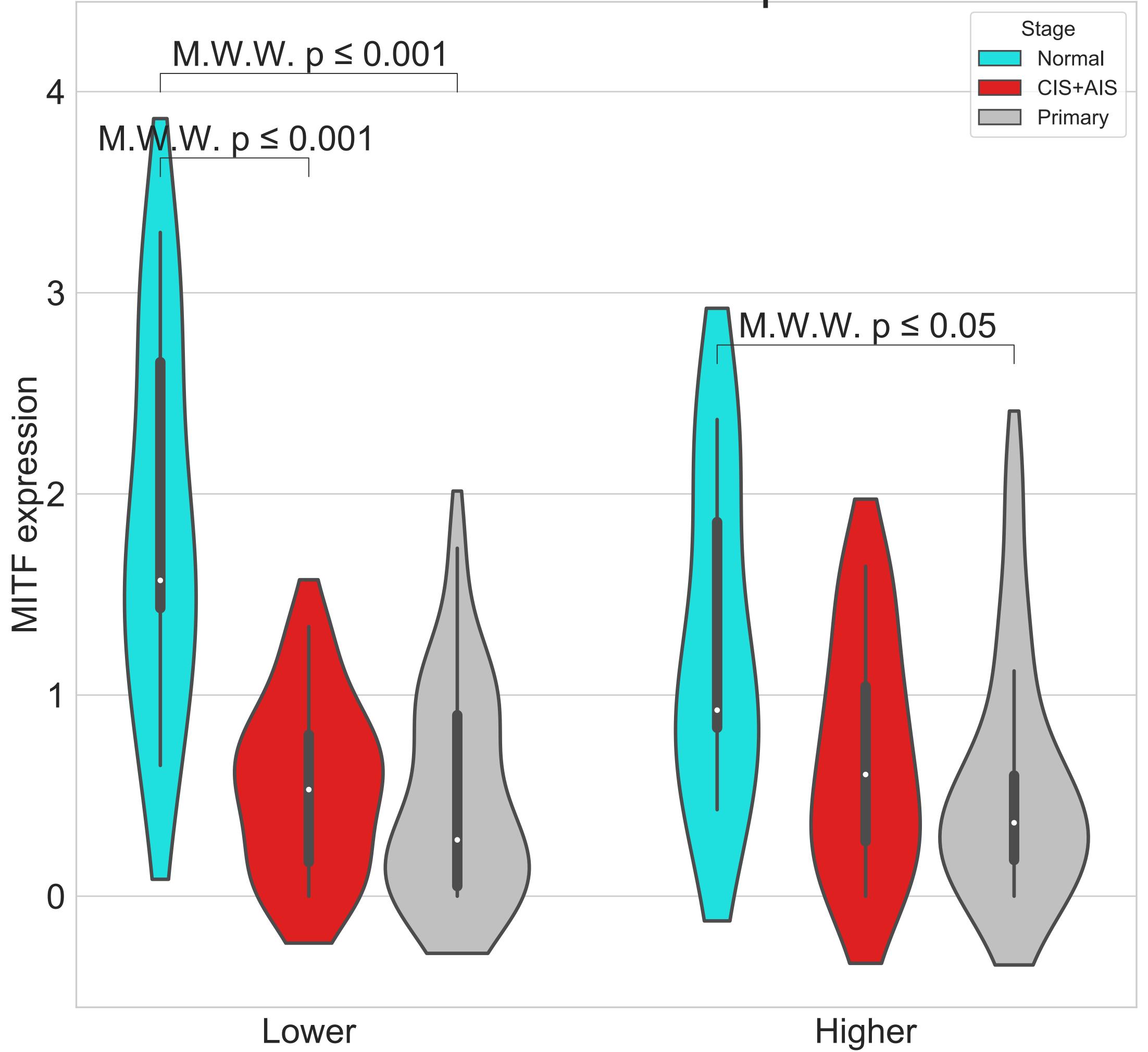
MITF: Kruskal-Wallis p=0.001



Mutation Shared Count per TMB