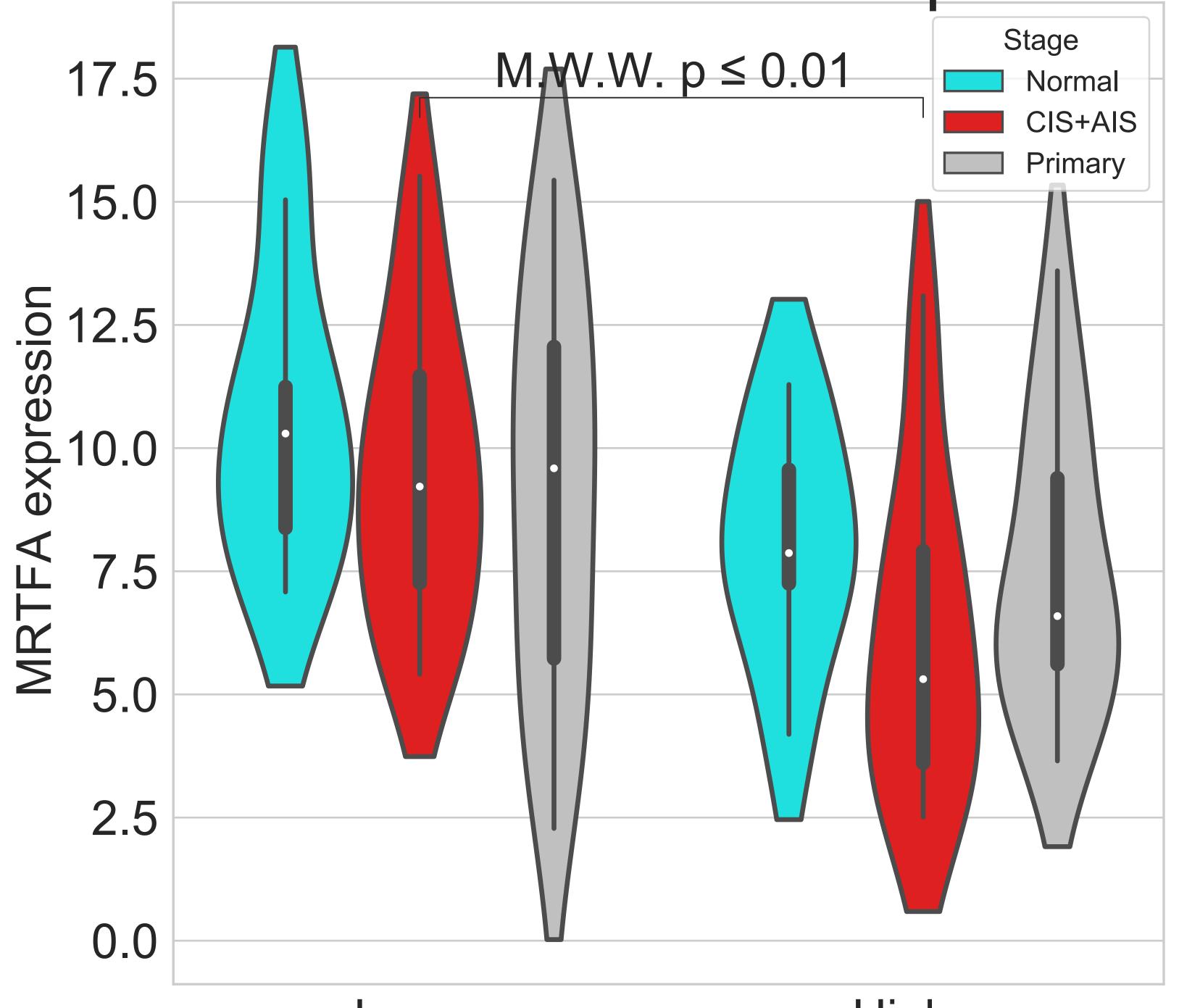
MRTFA: Kruskal-Wallis p=0.021



Lower Higher Mutation Shared Count per TMB