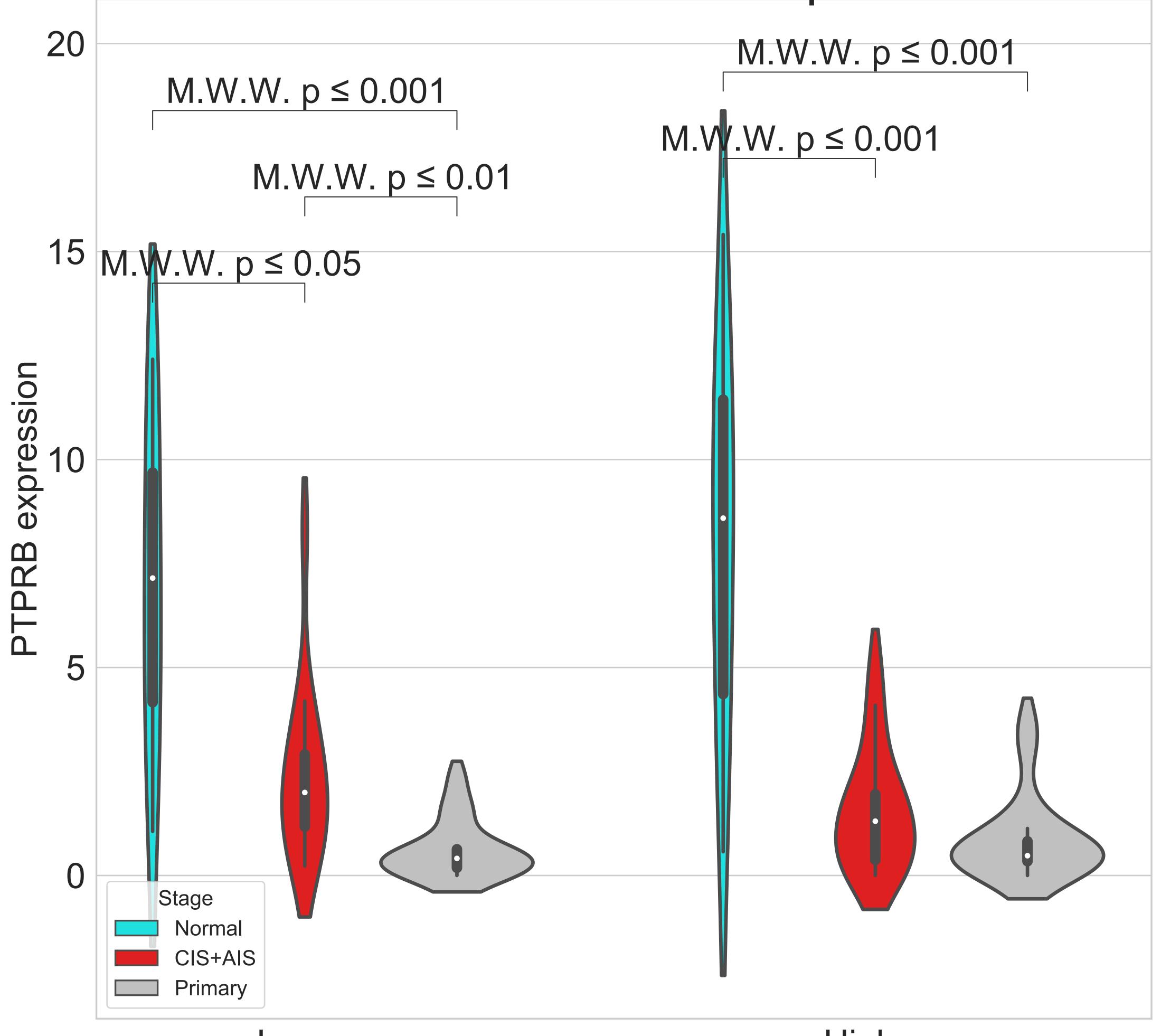
PTPRB: Kruskal-Wallis p=0.000



Higher Lower Mutation Shared Count per TMB