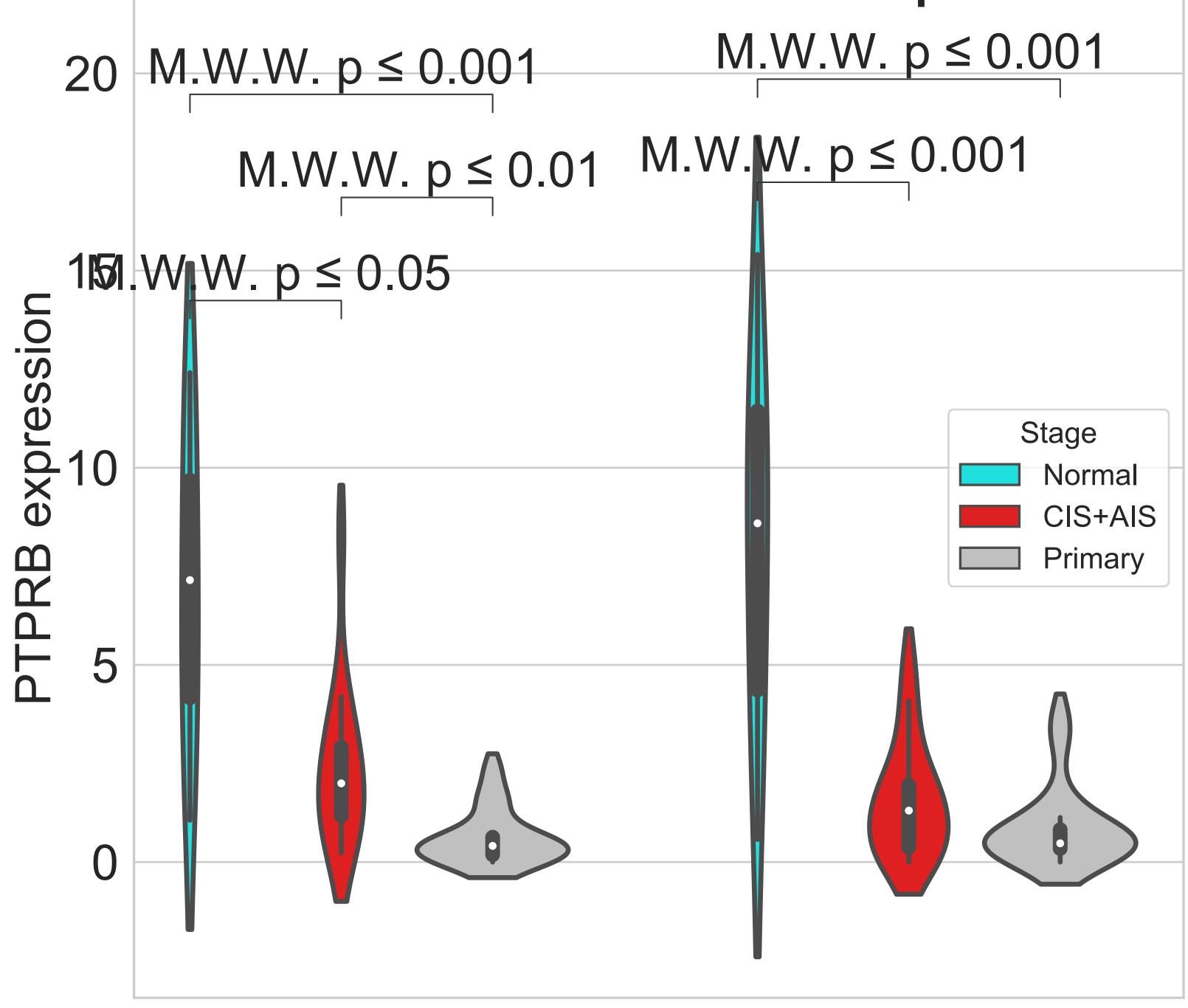
PTPRB: Kruskal-Wallis p=0.000



Lower
Higher
Mutation Shared Count per TMB