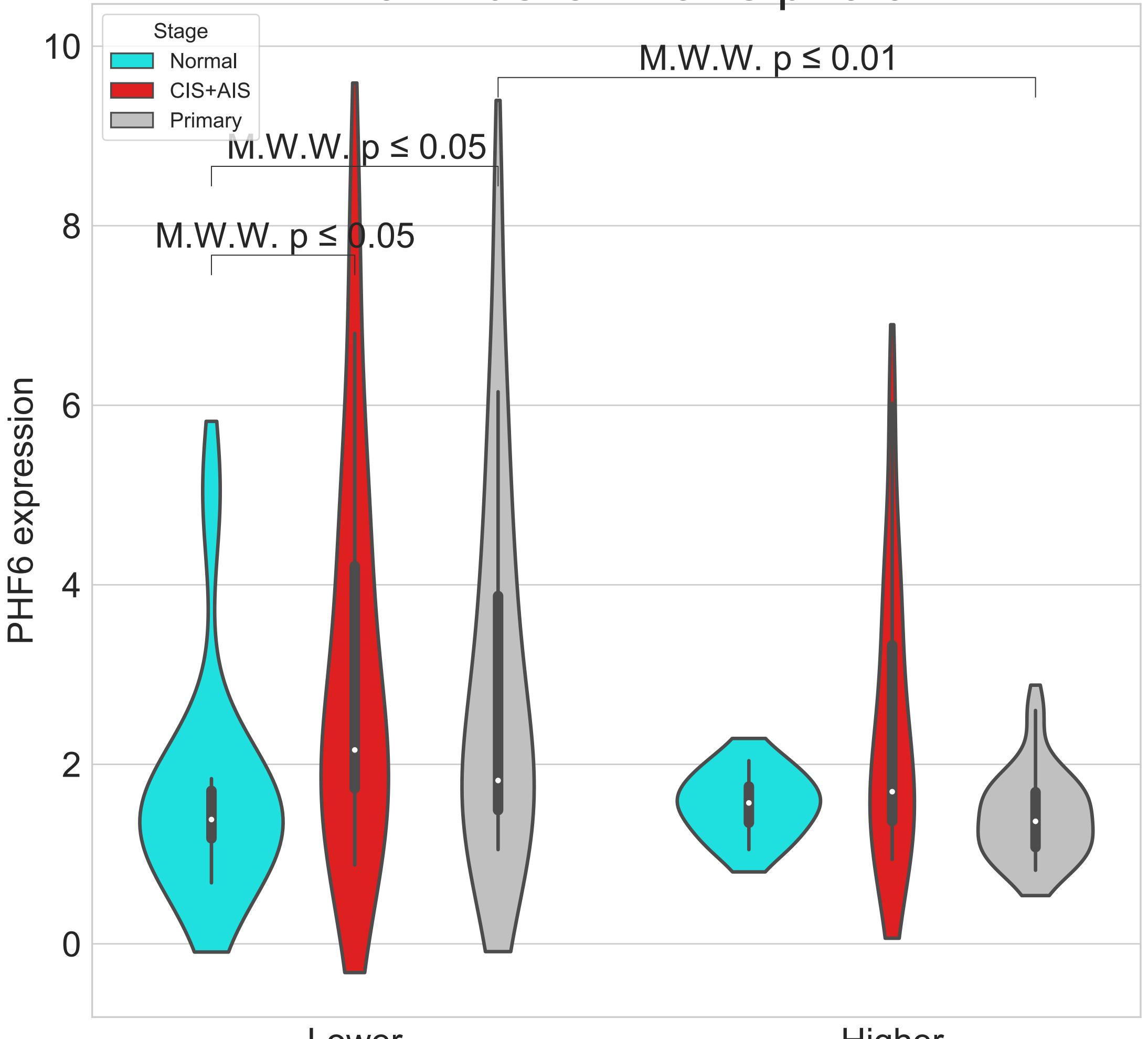
PHF6: Kruskal-Wallis p=0.017



Lower
Mutation Shared Count per TMB