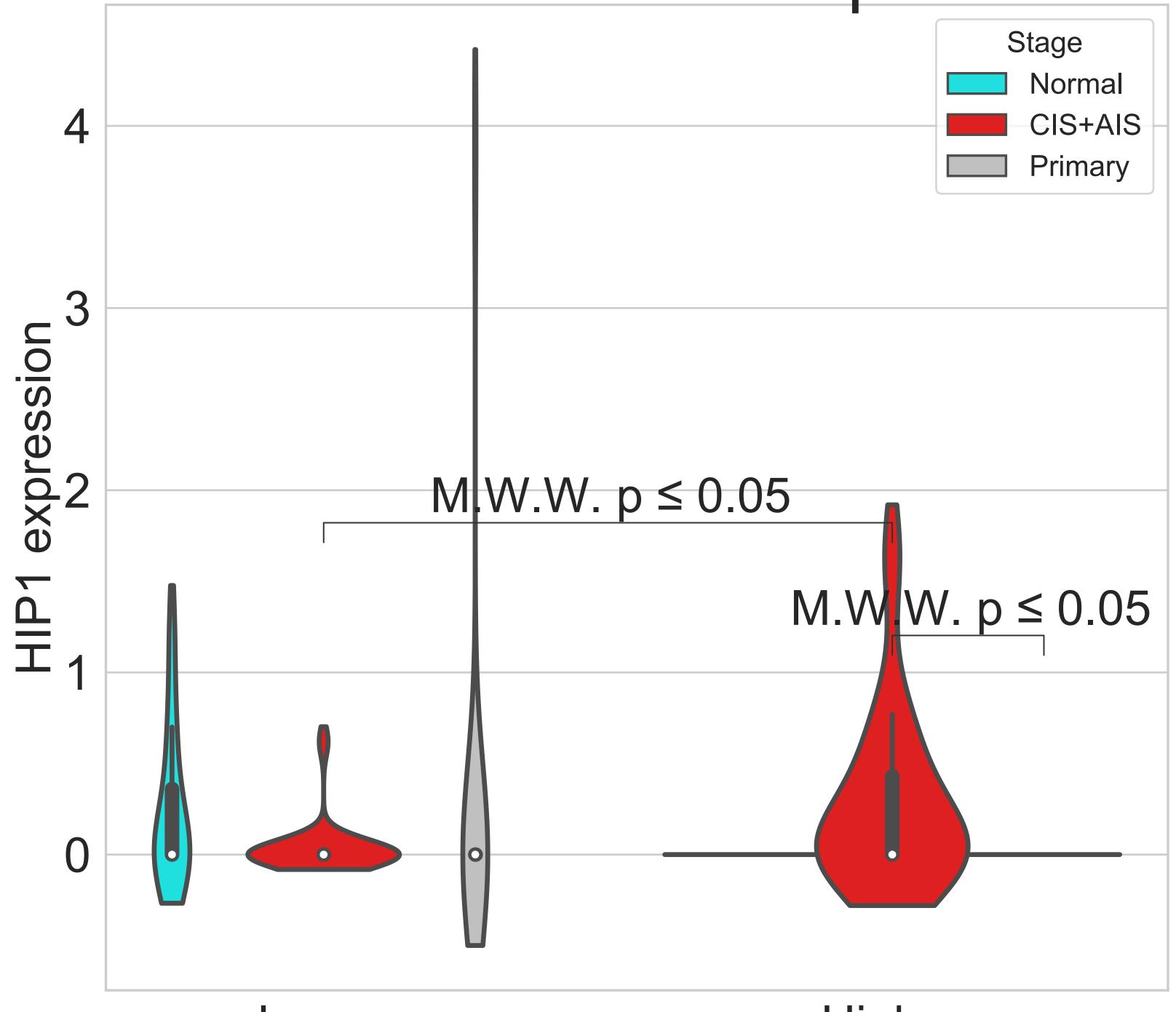
HIP1: Kruskal-Wallis p=0.018



Lower Higher Mutation Shared Count per TMB