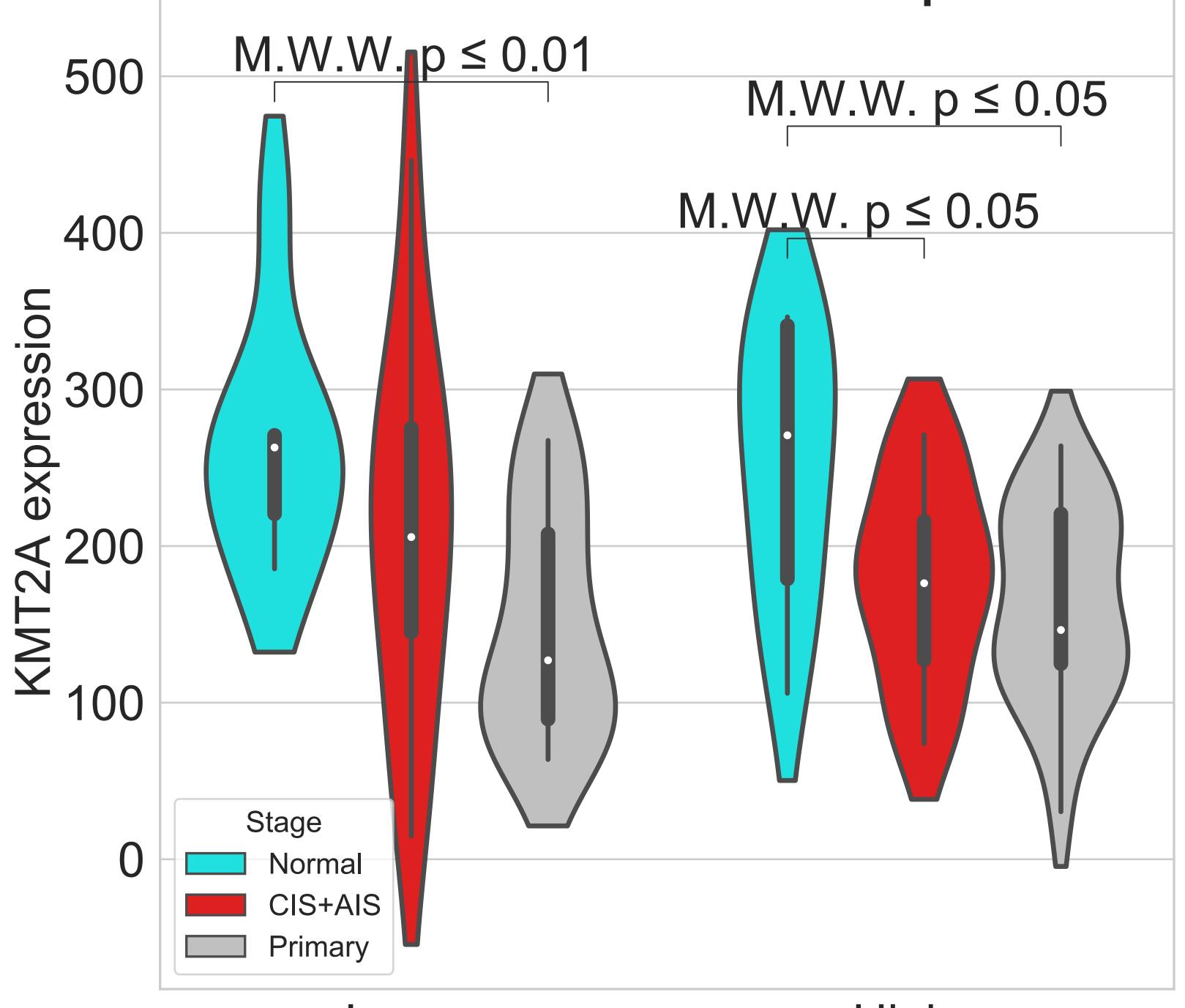
KMT2A: Kruskal-Wallis p=0.012



Lower Higher Mutation Shared Count per TMB