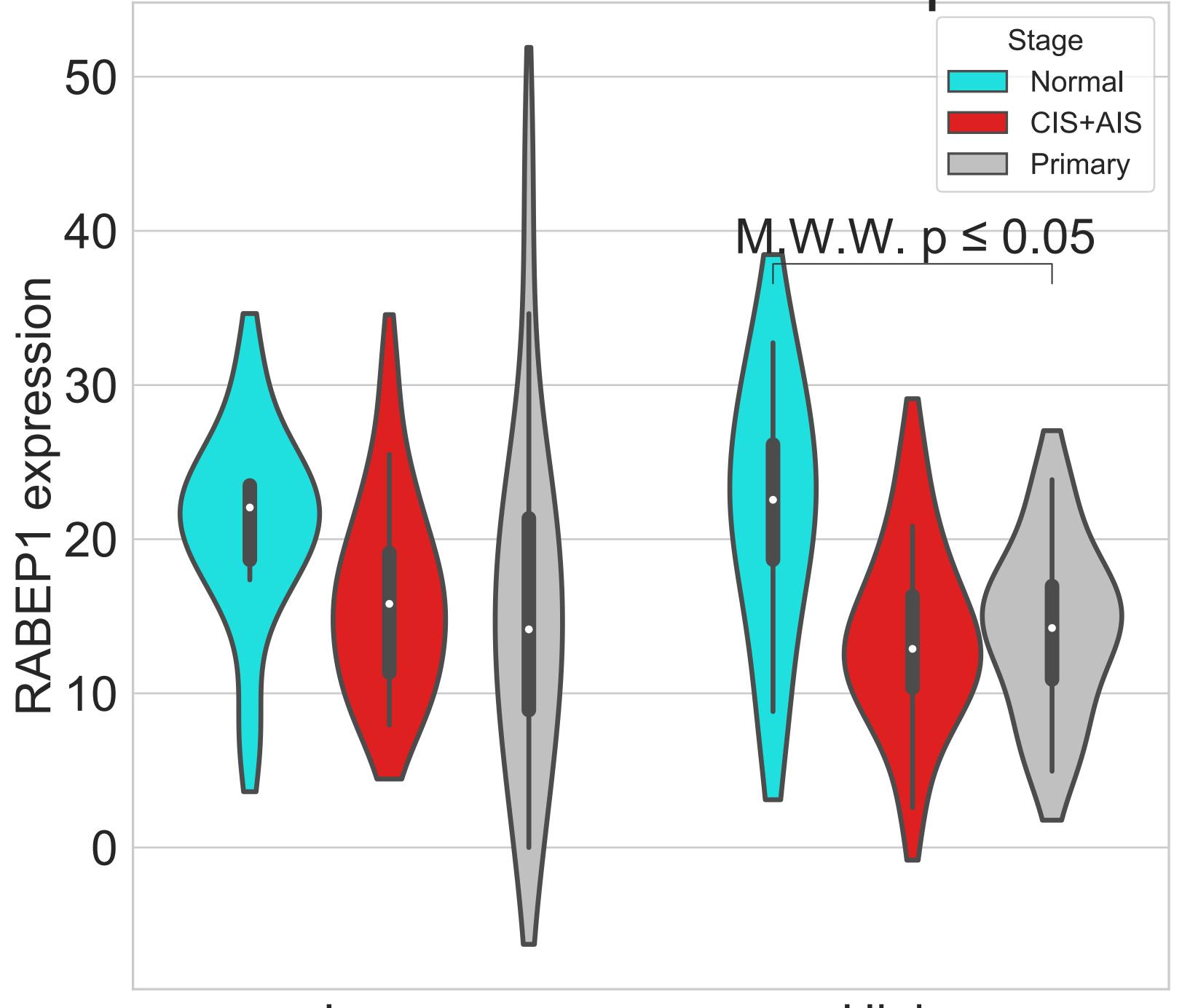
RABEP1: Kruskal-Wallis p=0.047



Lower Higher Mutation Shared Count per TMB