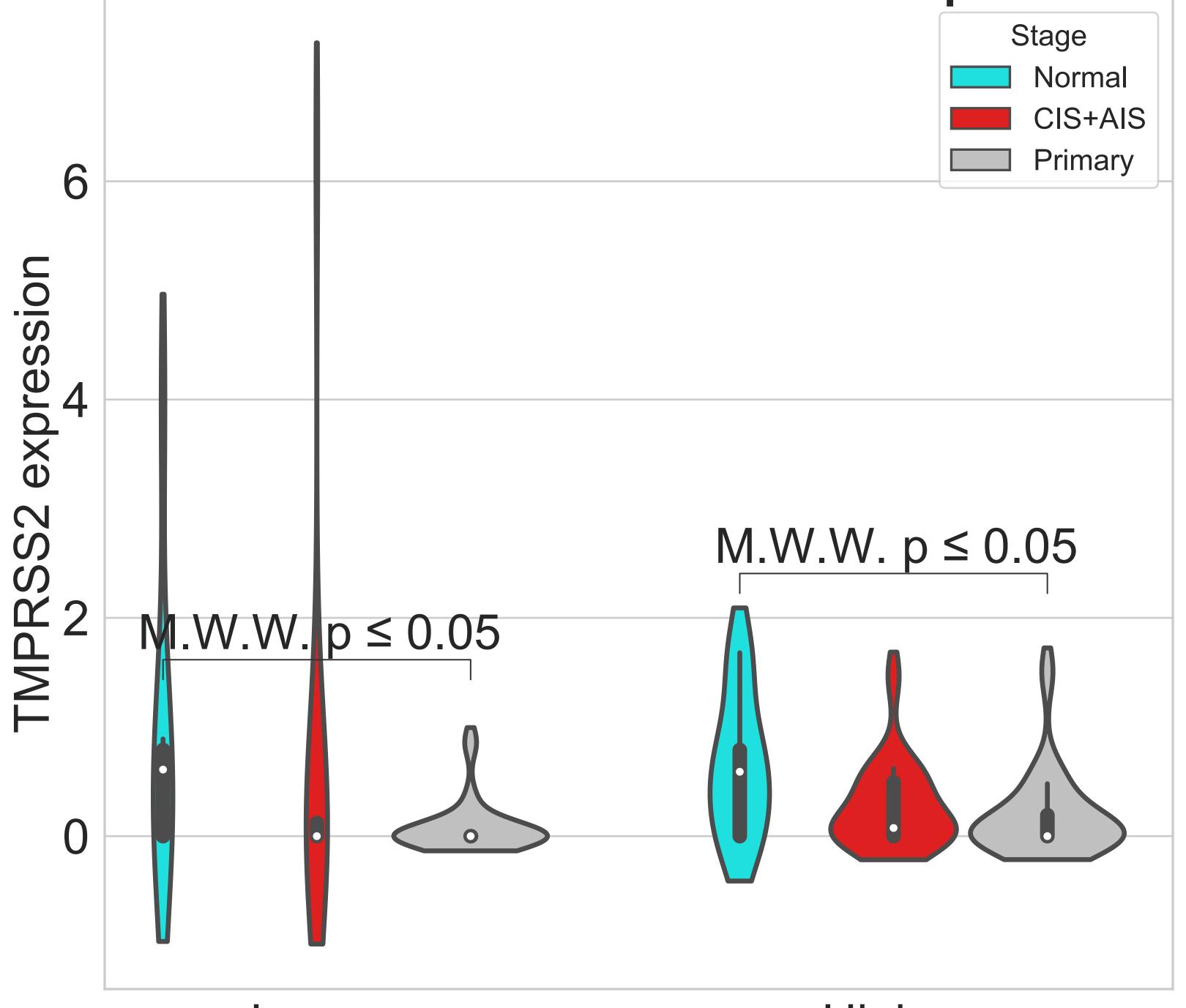
TMPRSS2: Kruskal-Wallis p=0.036



Lower
Mutation Shared Count per TMB