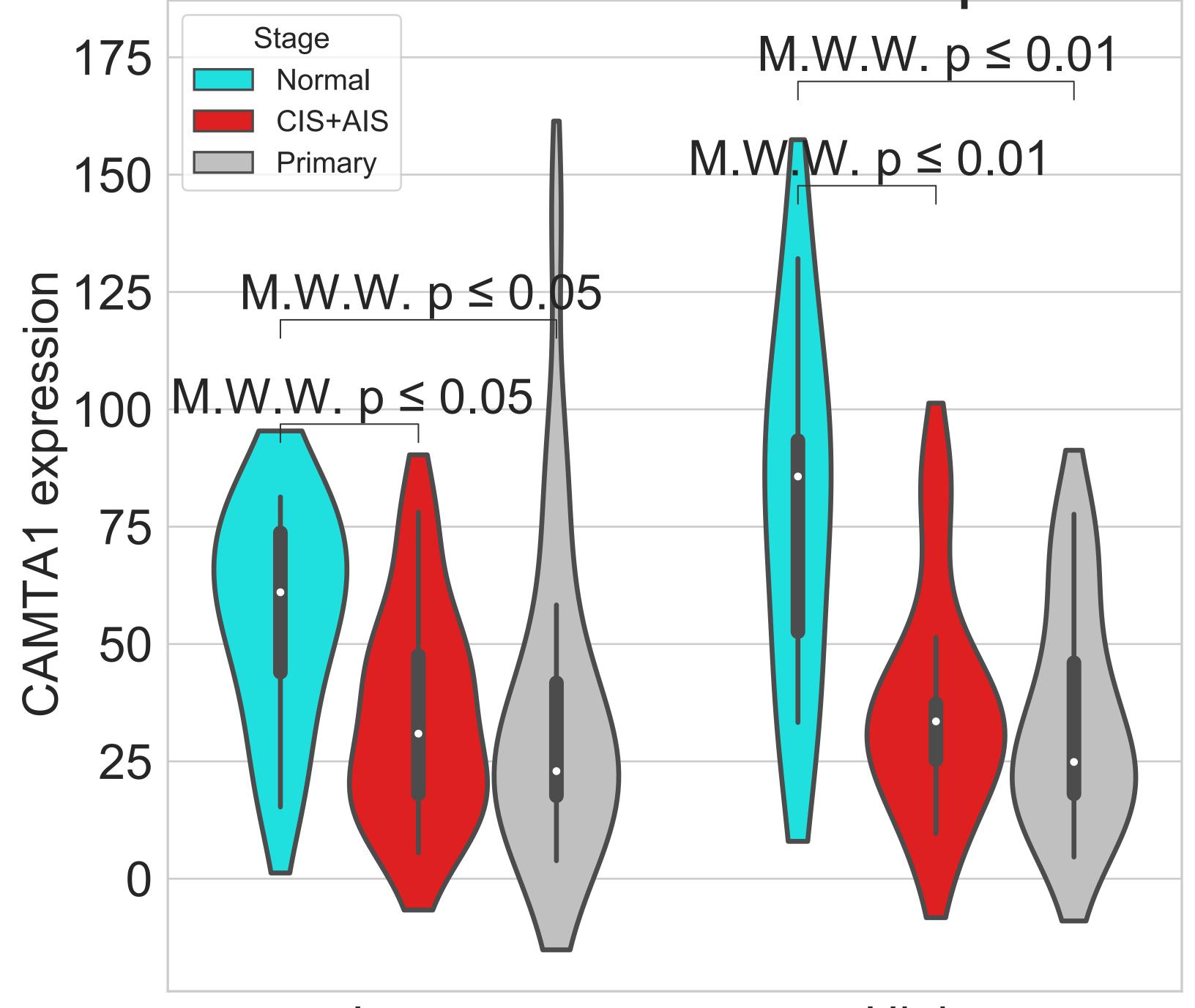
CAMTA1: Kruskal-Wallis p=0.014



Lower Higher Mutation Shared Count per TMB