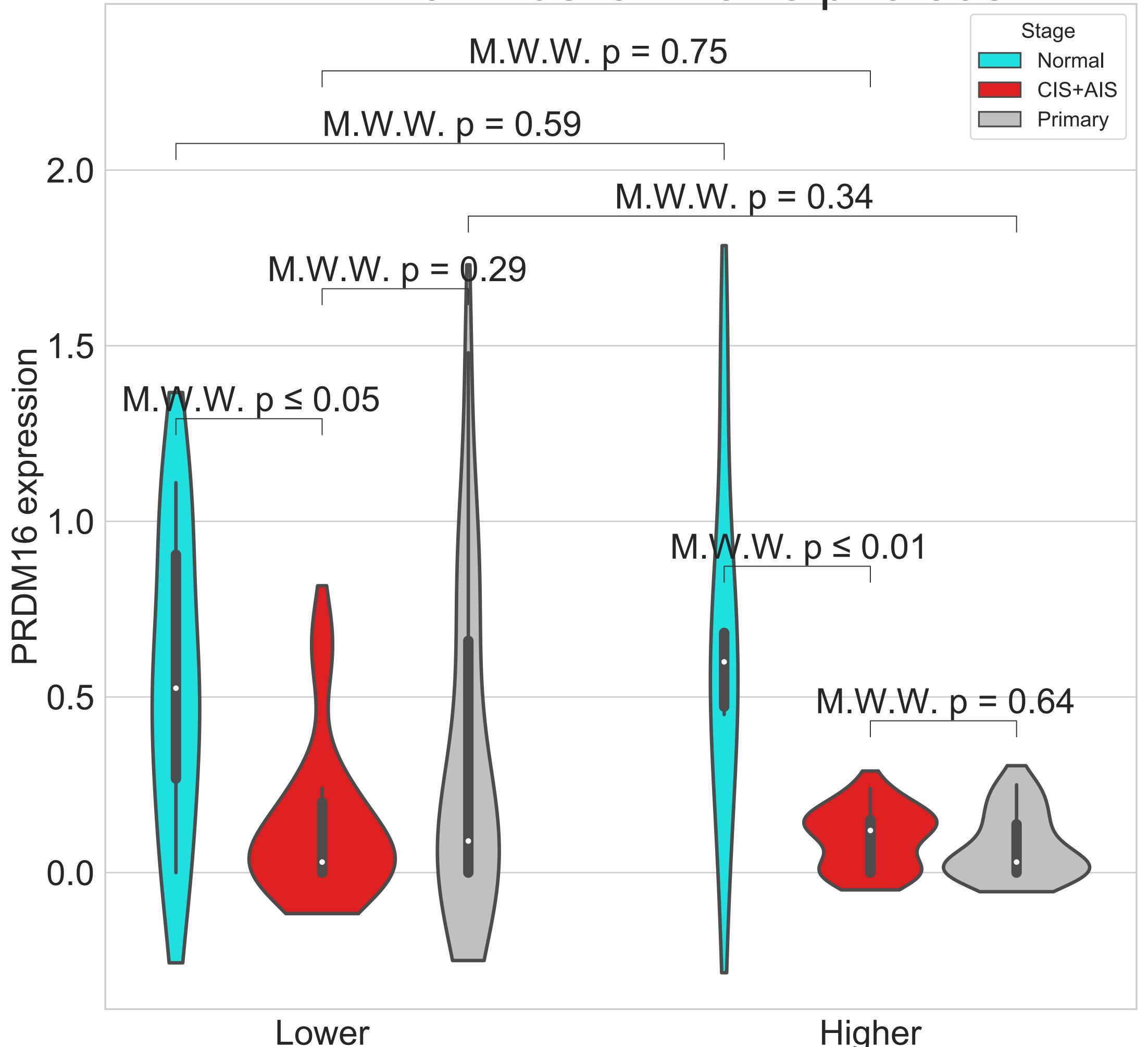
PRDM16: Kruskal-Wallis p=0.008



Higher
Mutation Shared Count per TMB