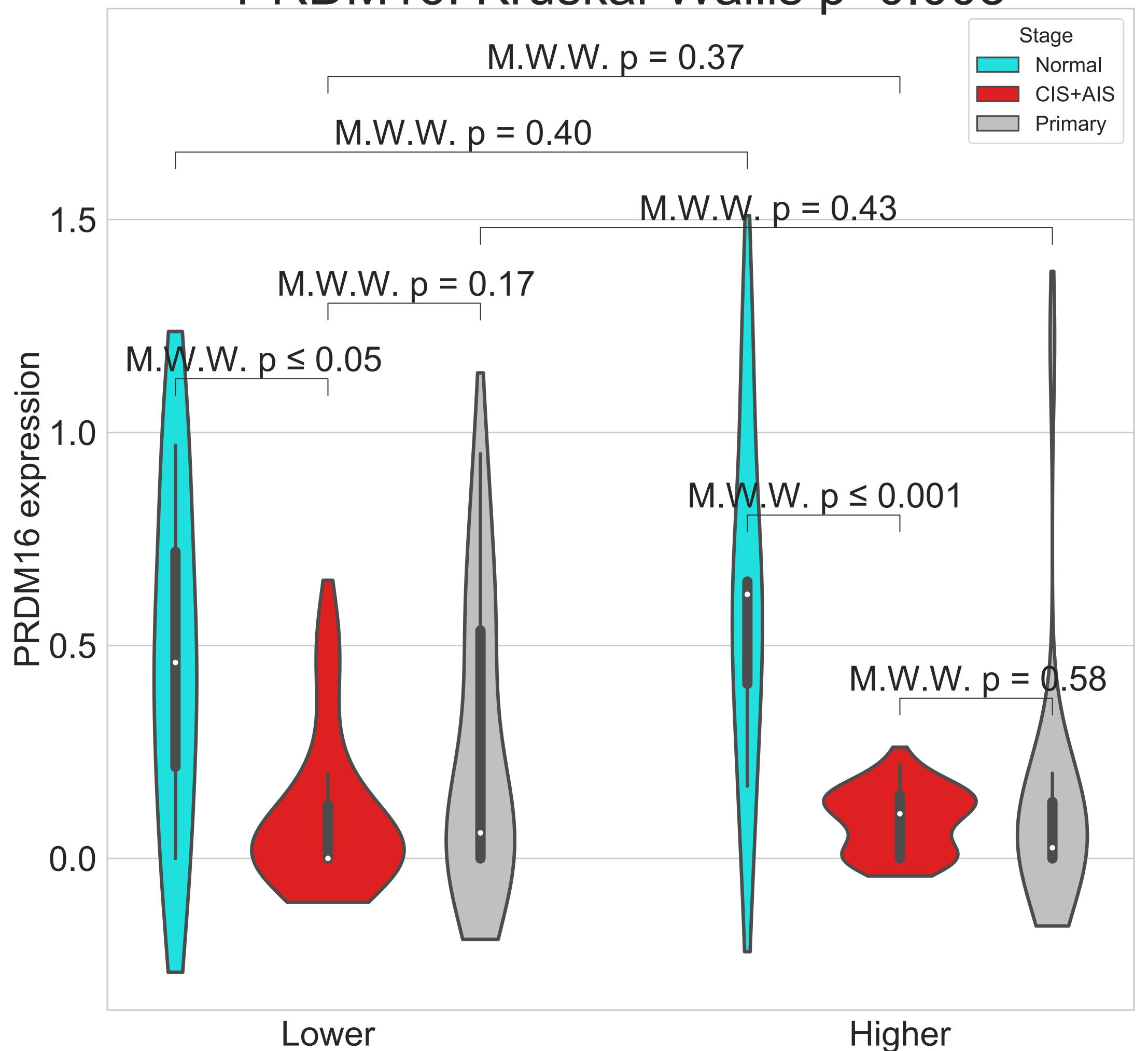
PRDM16: Kruskal-Wallis p=0.003



Mutation Shared Count per TMB