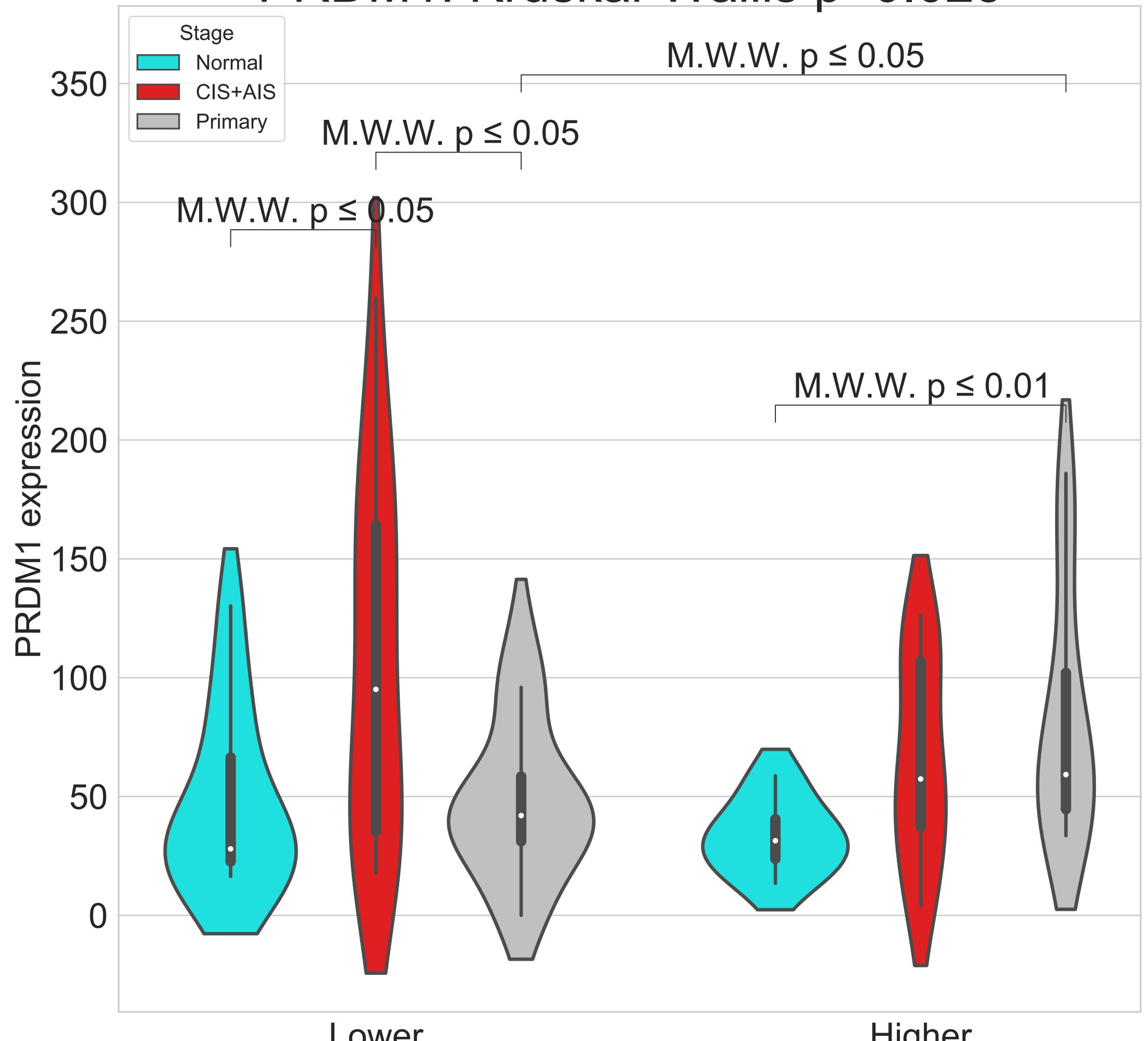
PRDM1: Kruskal-Wallis p=0.025



Lower
Mutation Shared Count per TMB