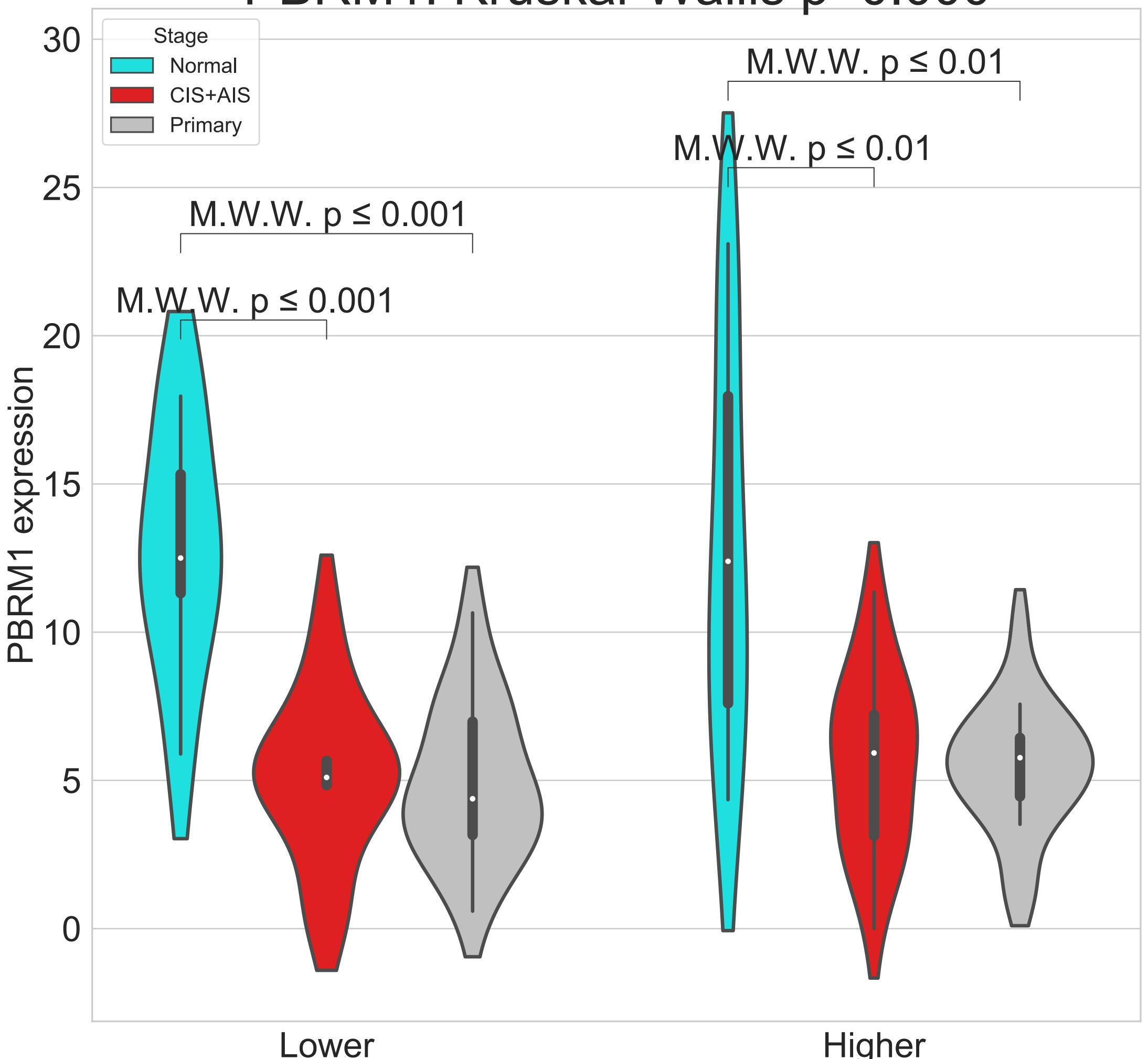
PBRM1: Kruskal-Wallis p=0.000



ver

Mutation Shared Count per TMB