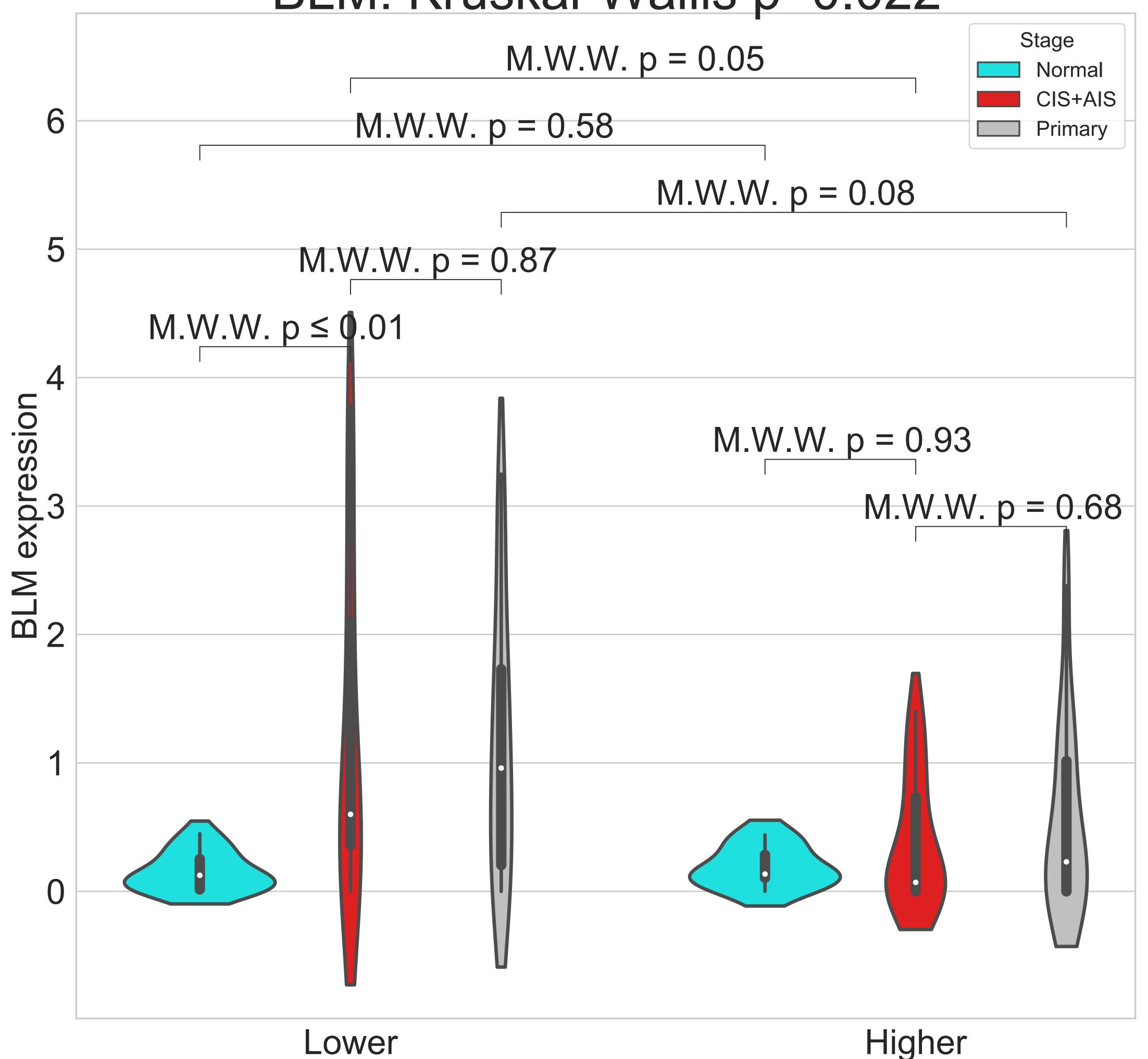
BLM: Kruskal-Wallis p=0.022



ower.
Mutation Shared Count per TMB