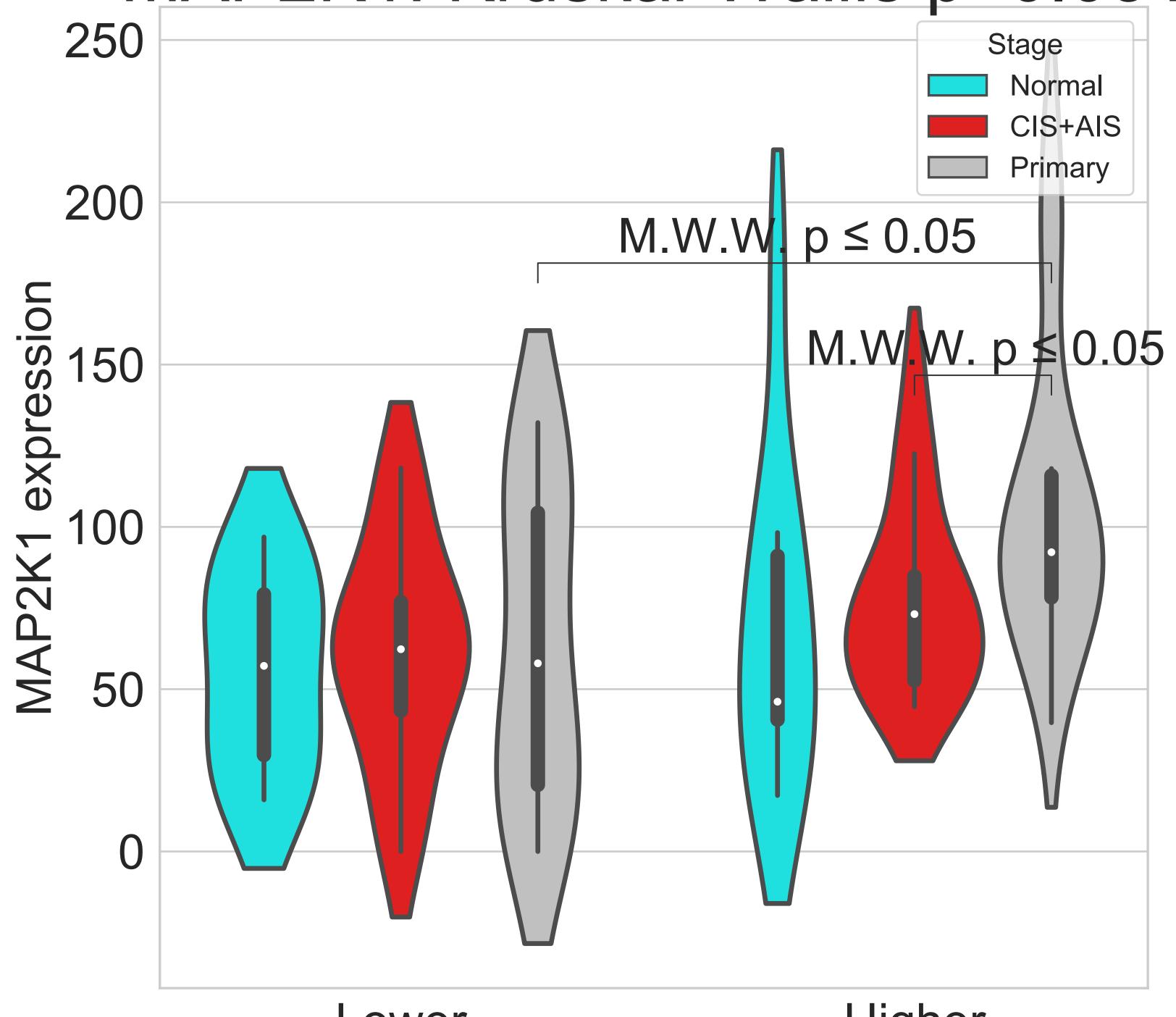
MAP2K1: Kruskal-Wallis p=0.031



Lower
Mutation Shared Count per TMB