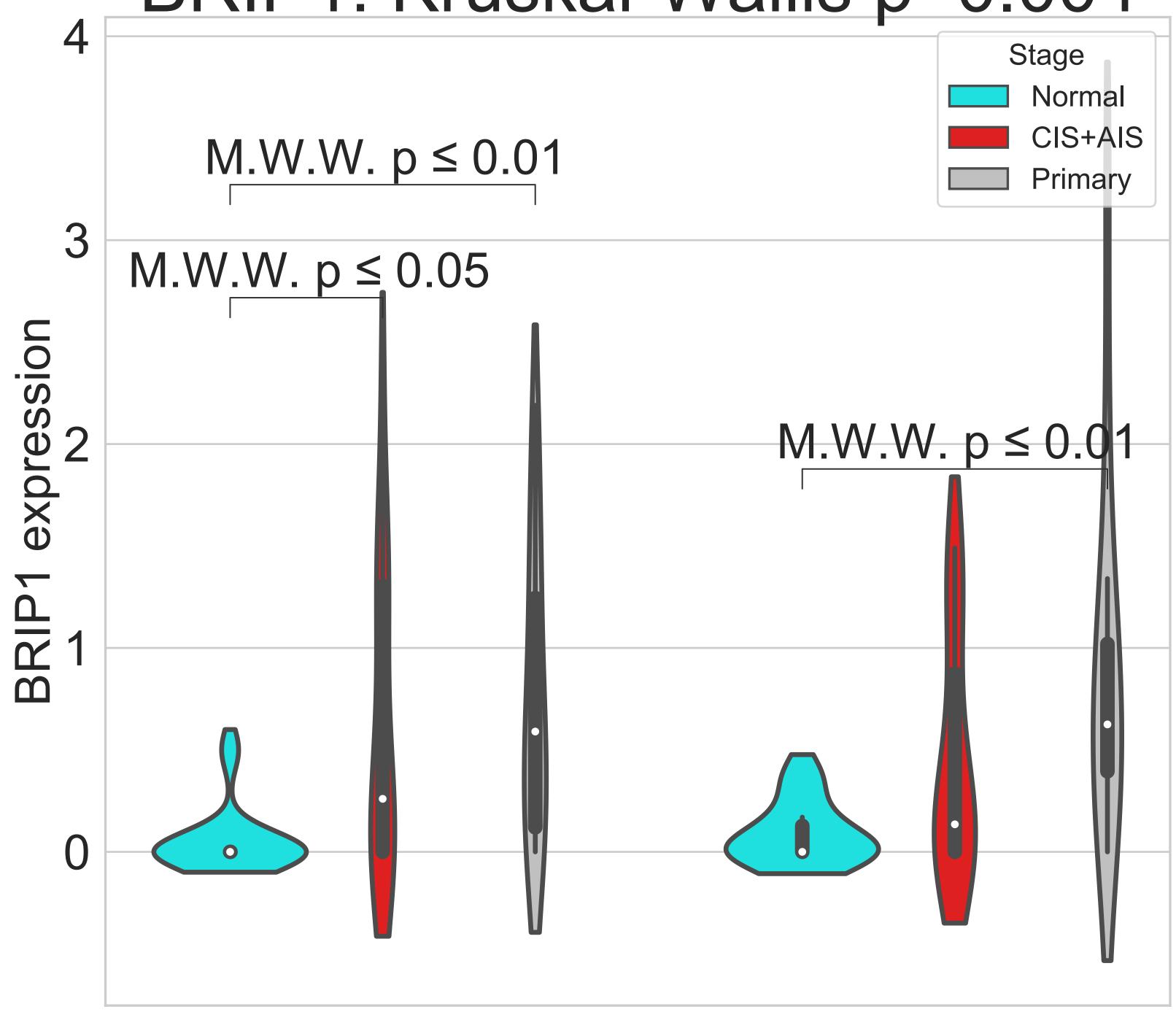
BRIP1: Kruskal-Wallis p=0.004



Lower
Mutation Shared Count per TMB