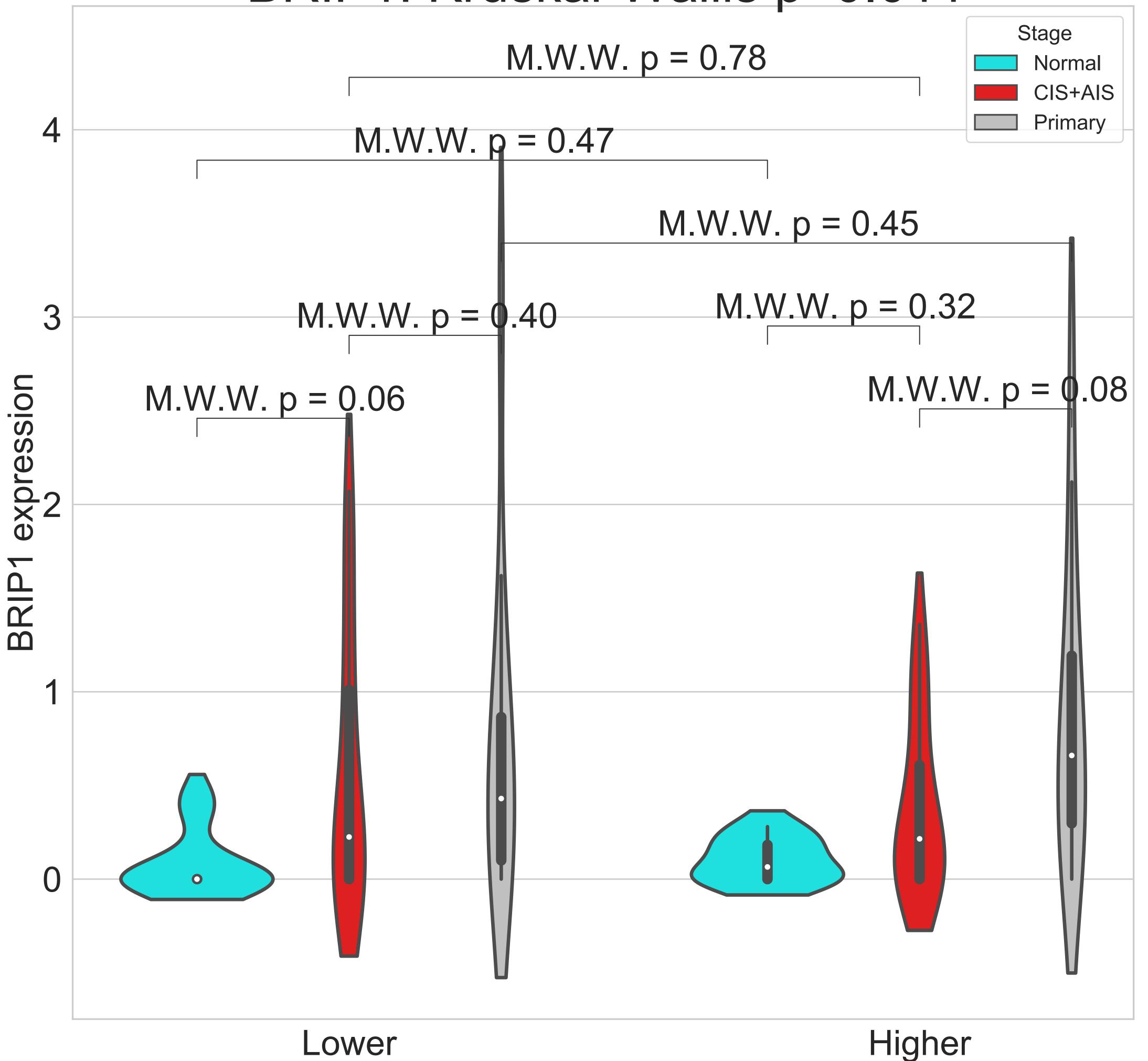
BRIP1: Kruskal-Wallis p=0.011



wer Mutation Shared Count per TMB