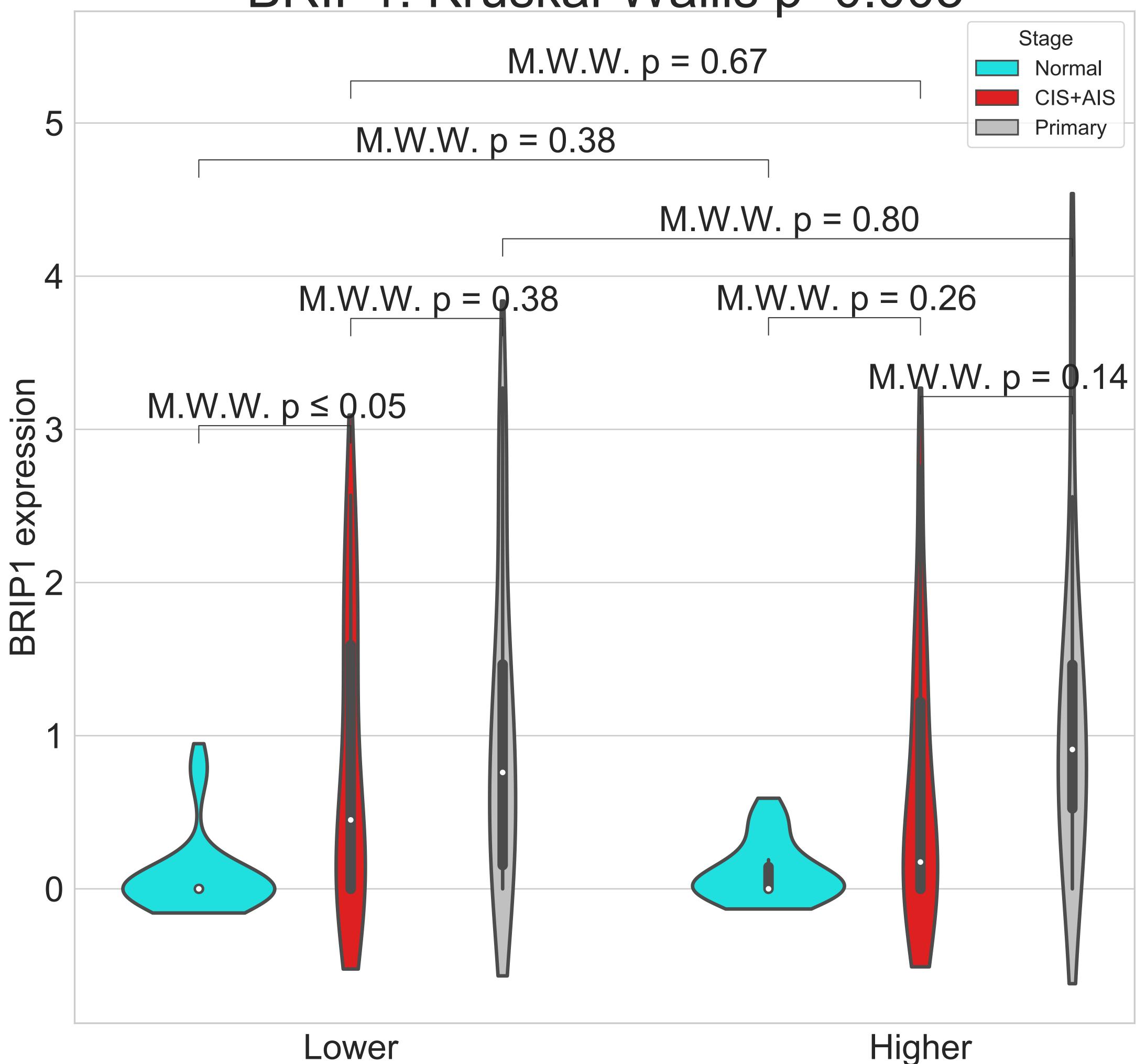
BRIP1: Kruskal-Wallis p=0.005



wer
Mutation Shared Count per TMB