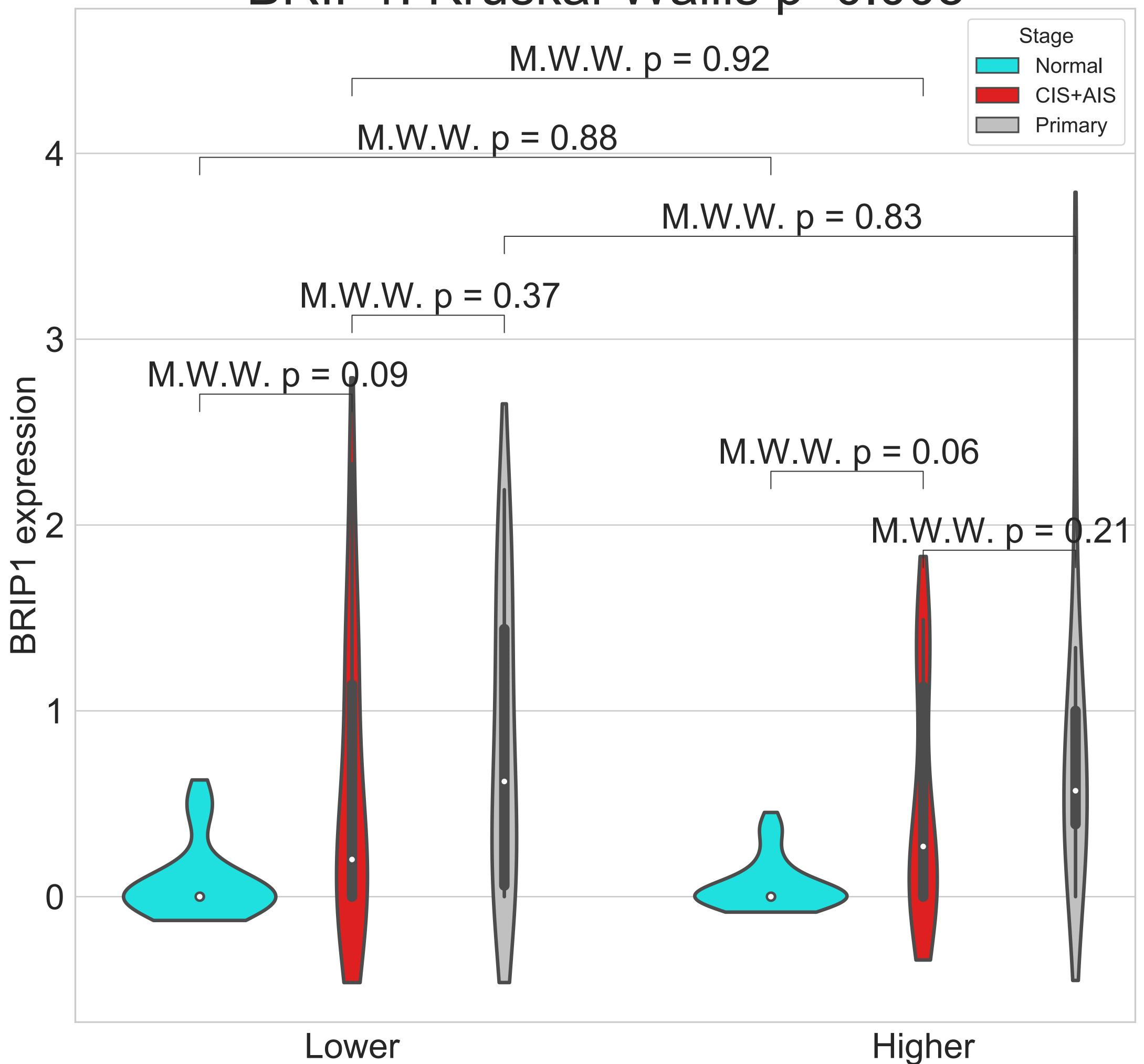
BRIP1: Kruskal-Wallis p=0.005



Mutation Shared Count (SYN) per TMB