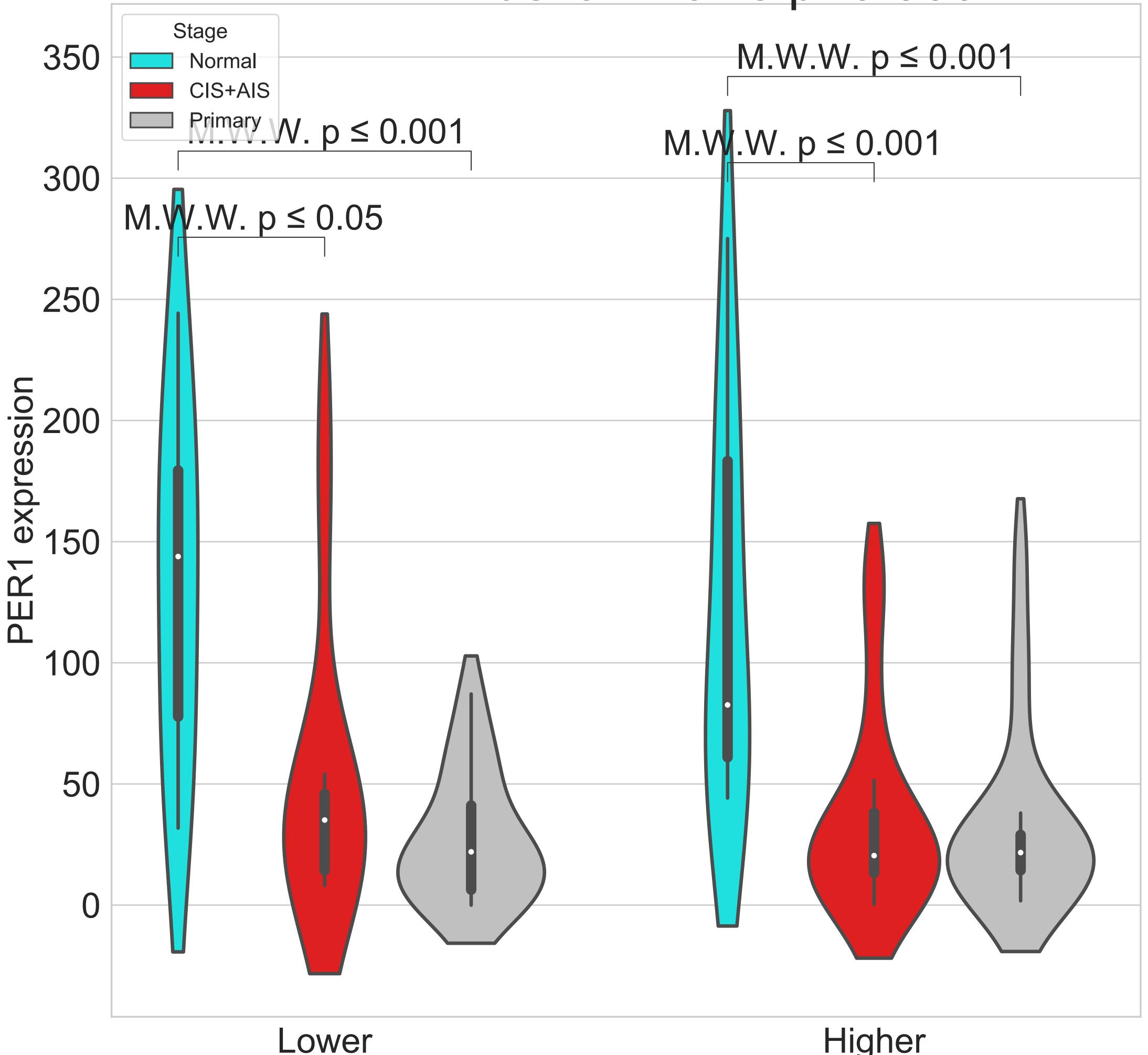
PER1: Kruskal-Wallis p=0.000



wer Mutation Shared Count (SYN) per TMB