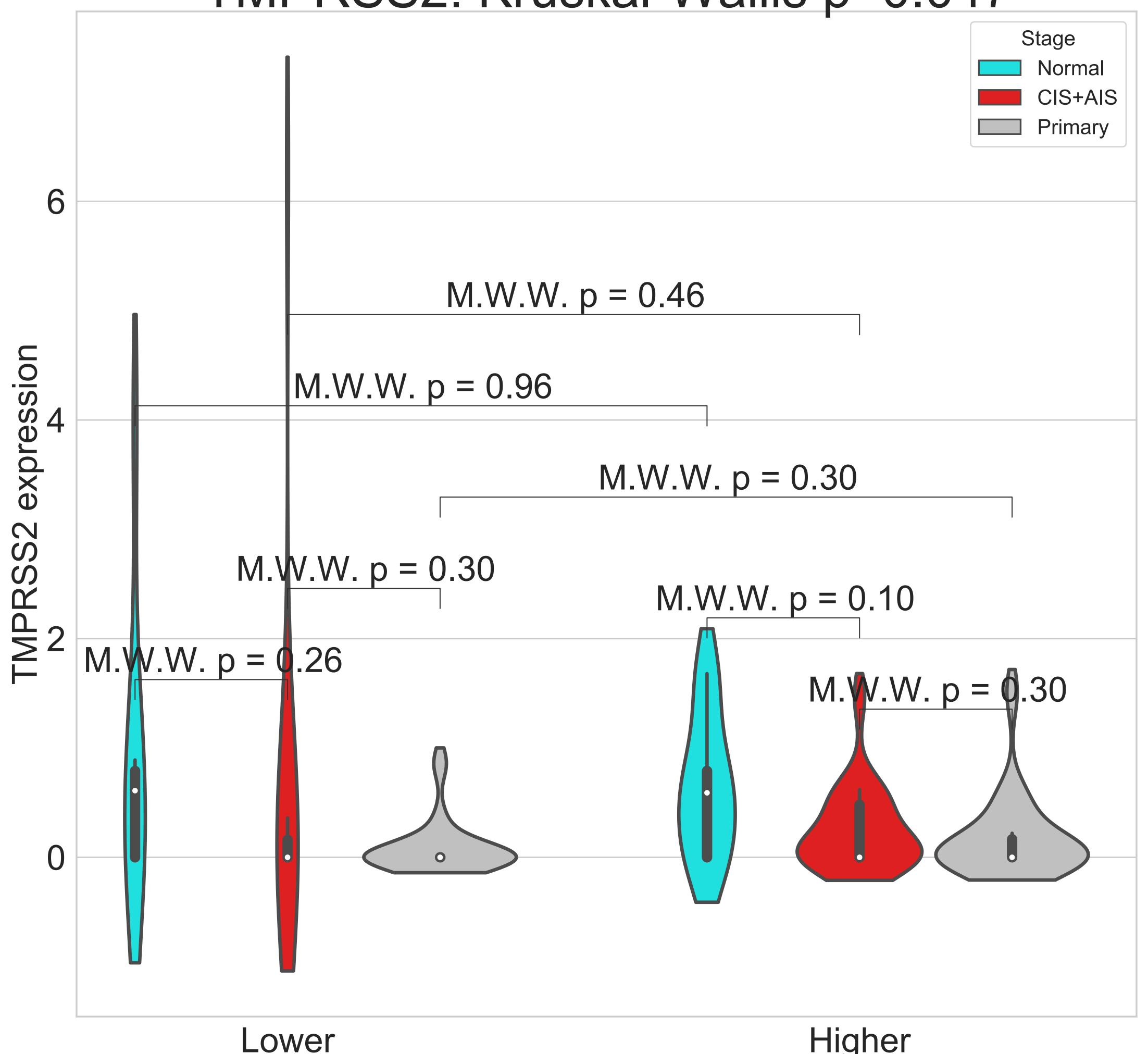
TMPRSS2: Kruskal-Wallis p=0.047



wer
Mutation Shared Count (SYN) per TMB