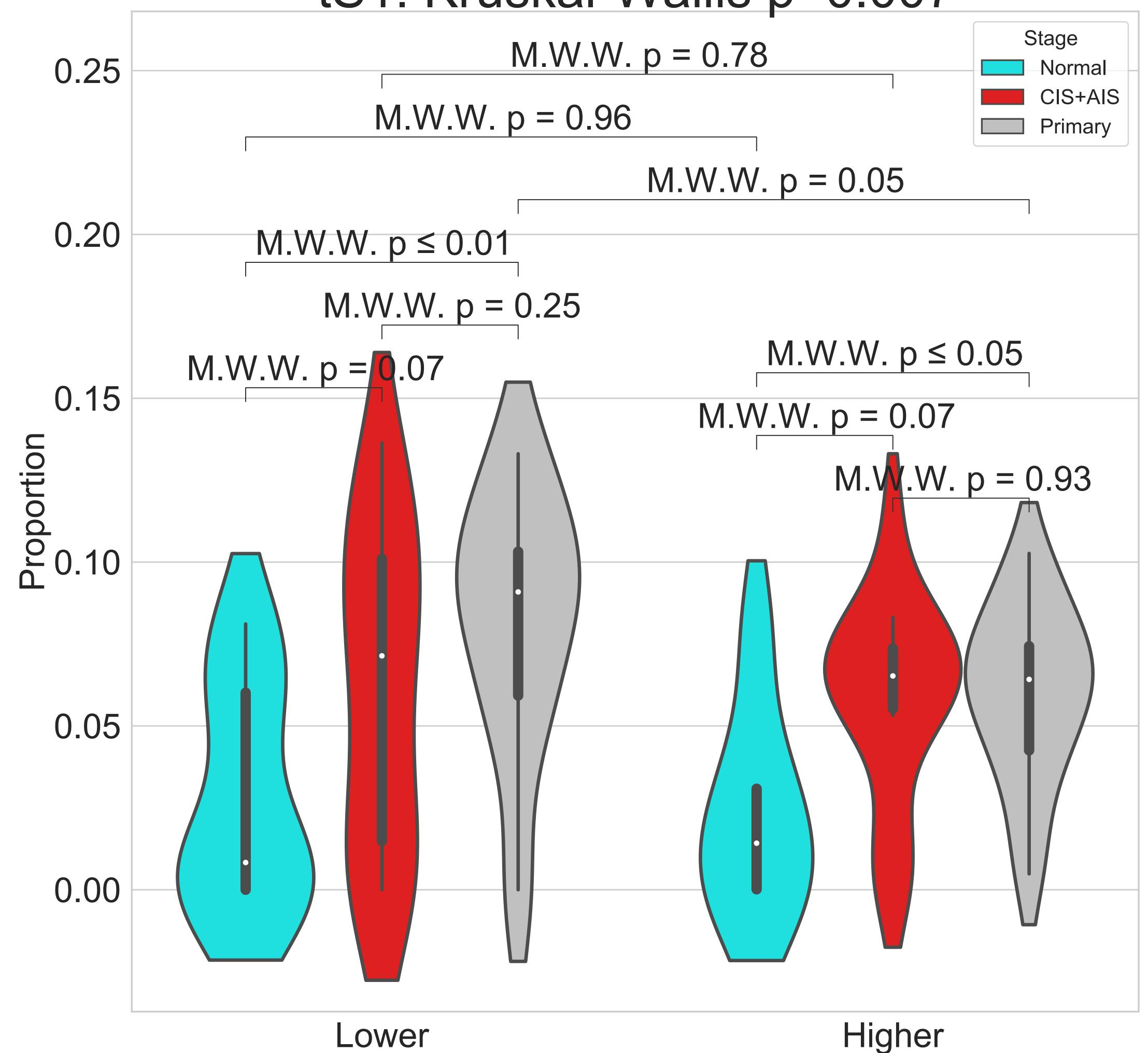
tS1: Kruskal-Wallis p=0.007



Mutation Shared Count (SYN) per TMB