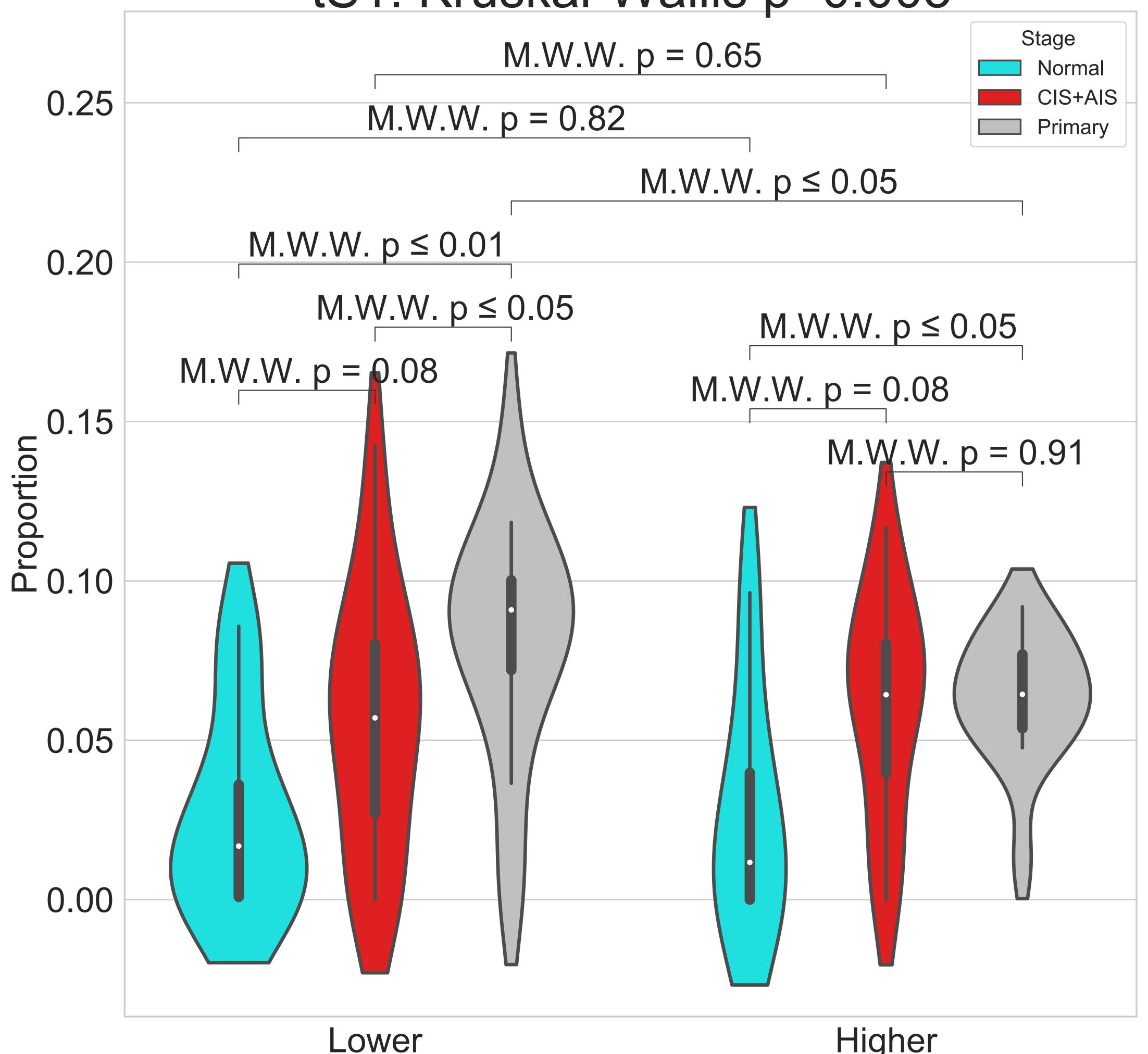
tS1: Kruskal-Wallis p=0.003



wer
Mutation Shared Count per TMB