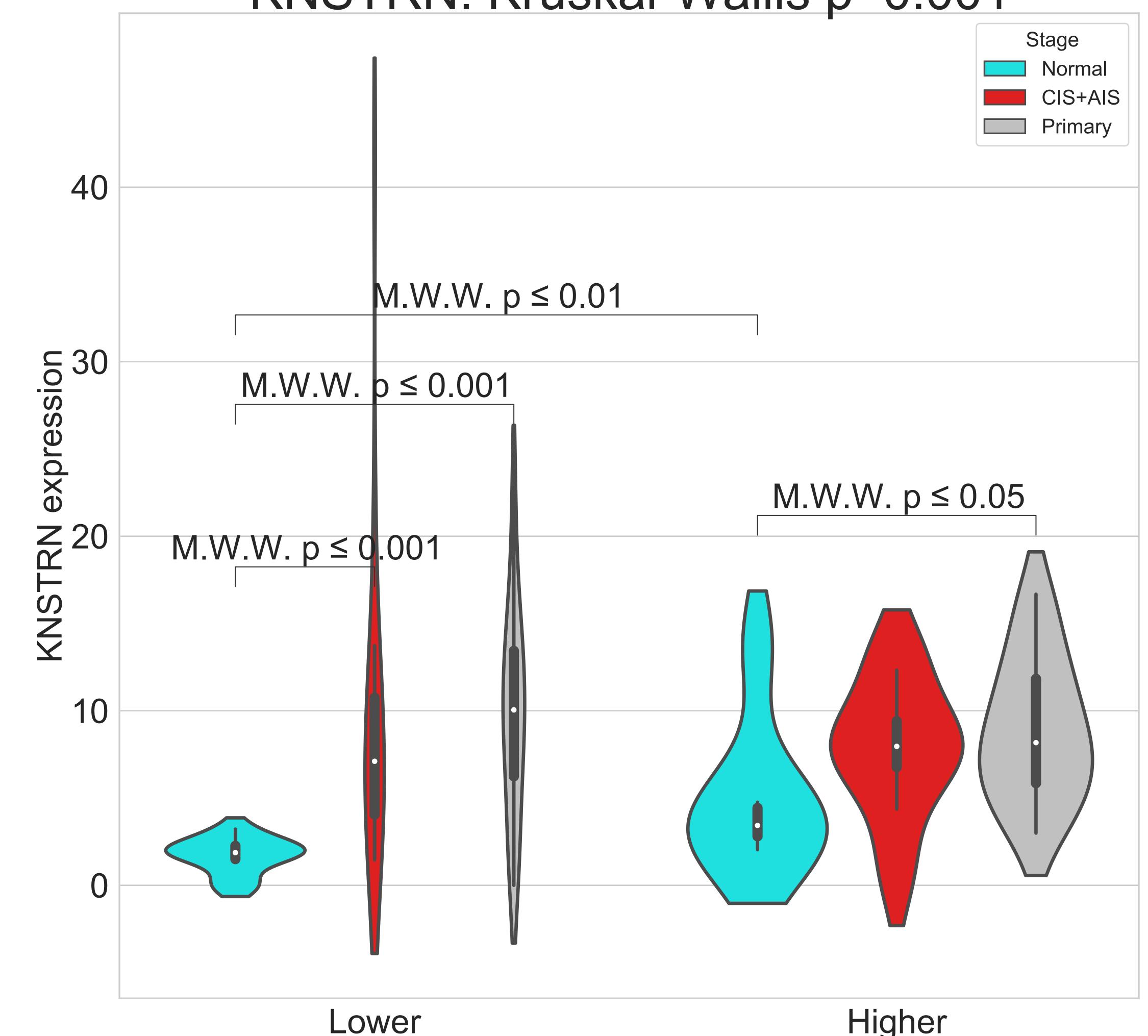
KNSTRN: Kruskal-Wallis p=0.001



wer
Mutation Shared Count per TMB