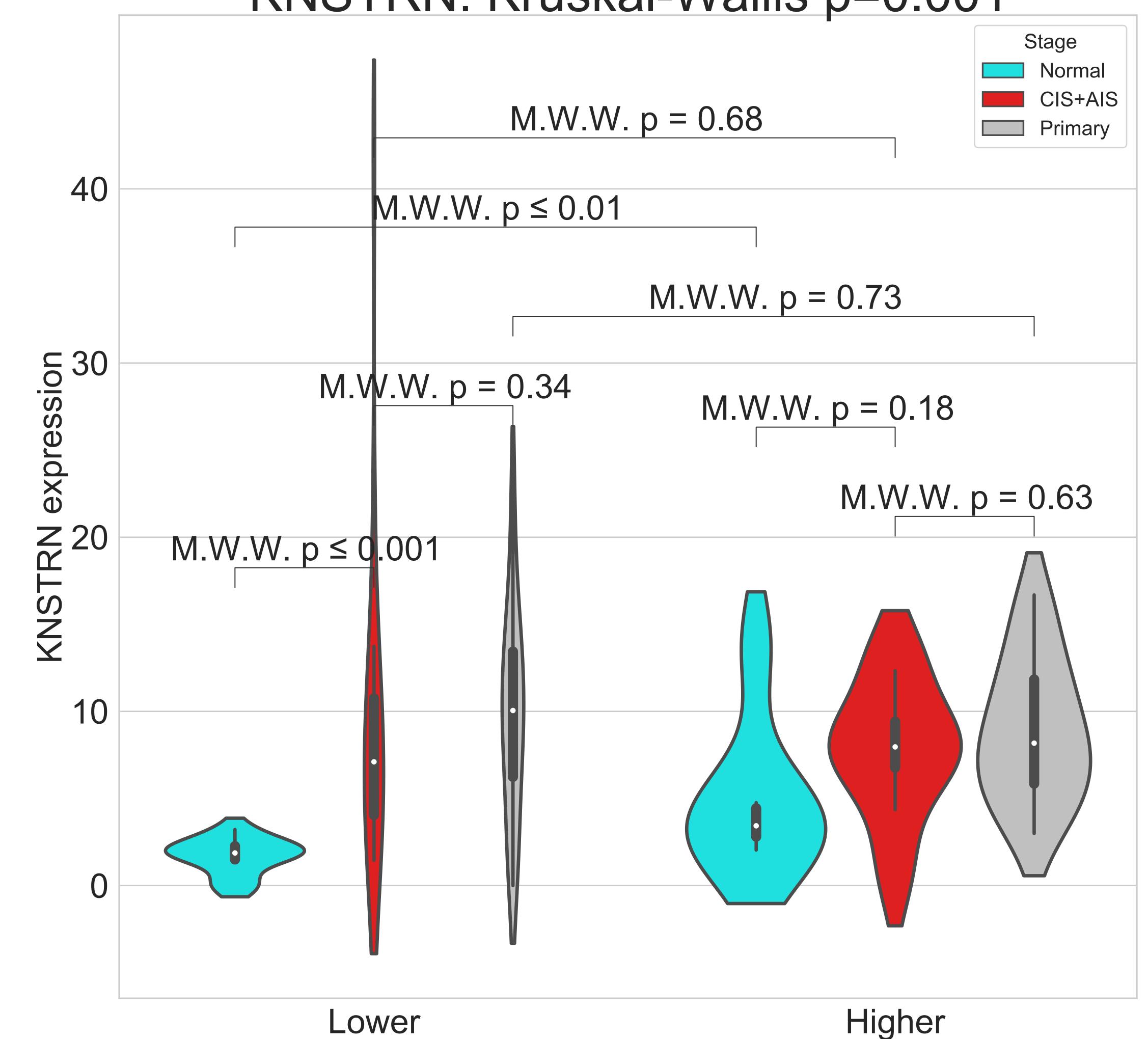
KNSTRN: Kruskal-Wallis p=0.001



wer Mutation Shared Count per TMB