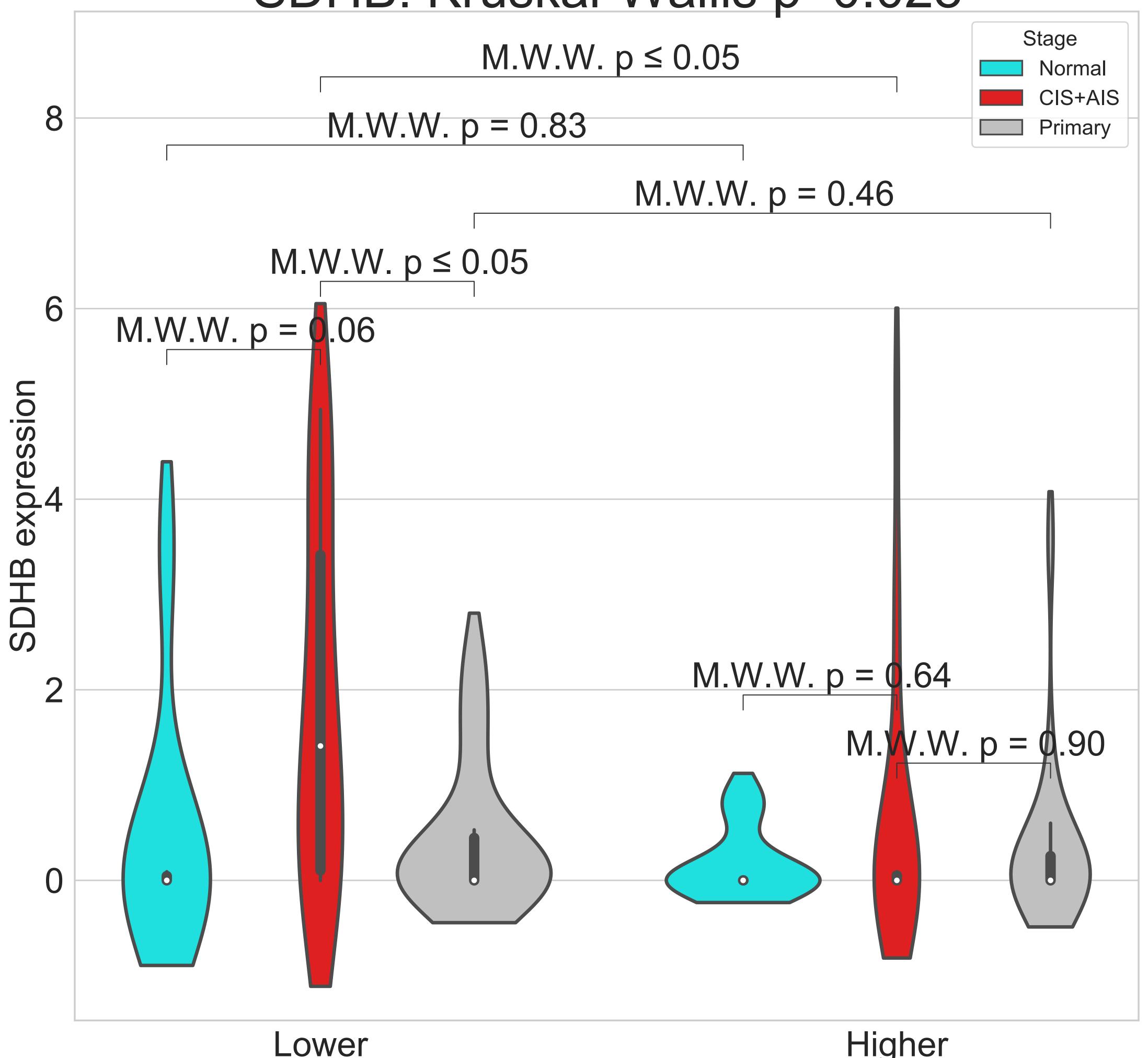
SDHB: Kruskal-Wallis p=0.028



wer
Mutation Shared Count per TMB