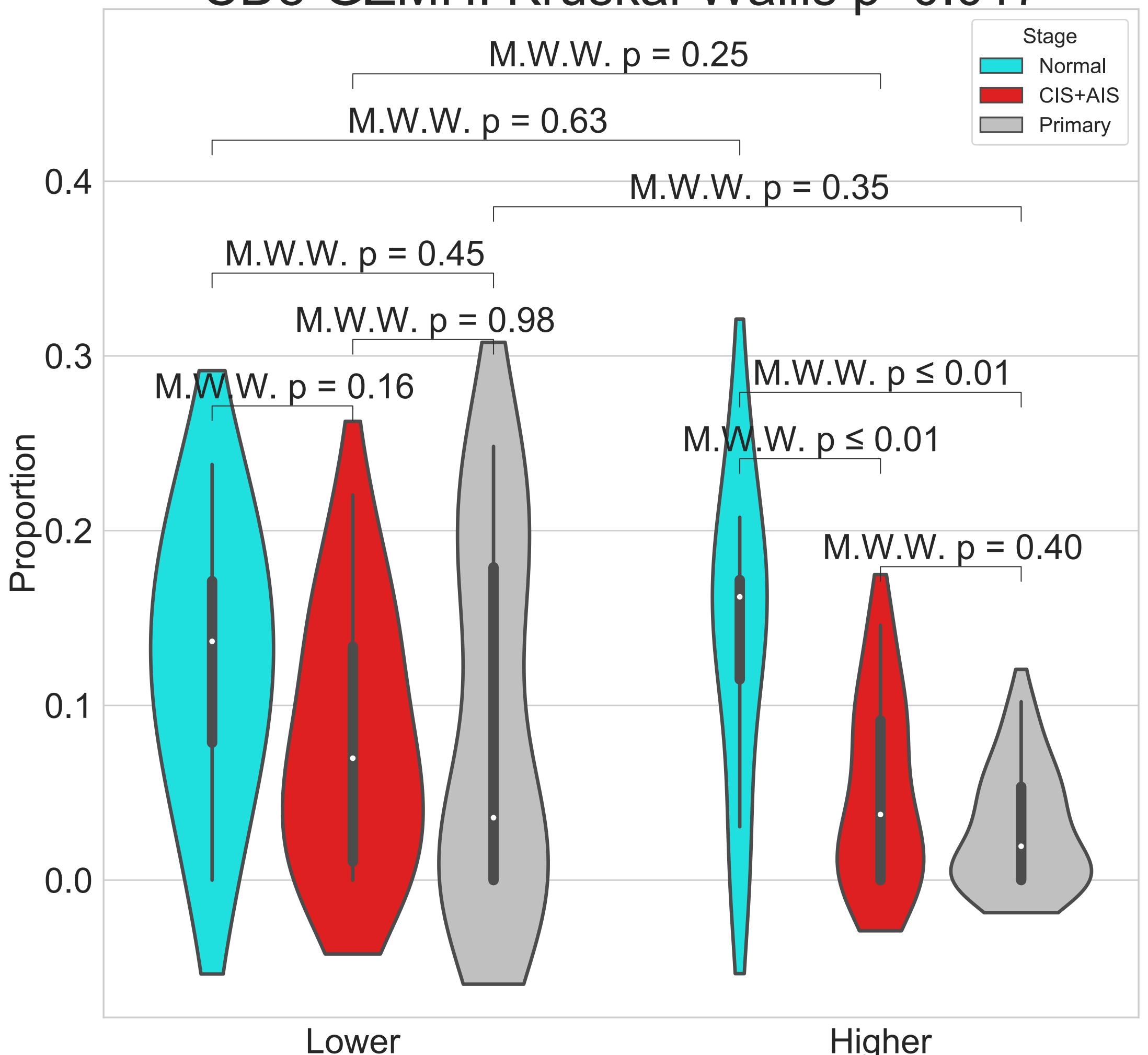
CD8-GZMH: Kruskal-Wallis p=0.017



wer
Mutation Shared Count per TMB