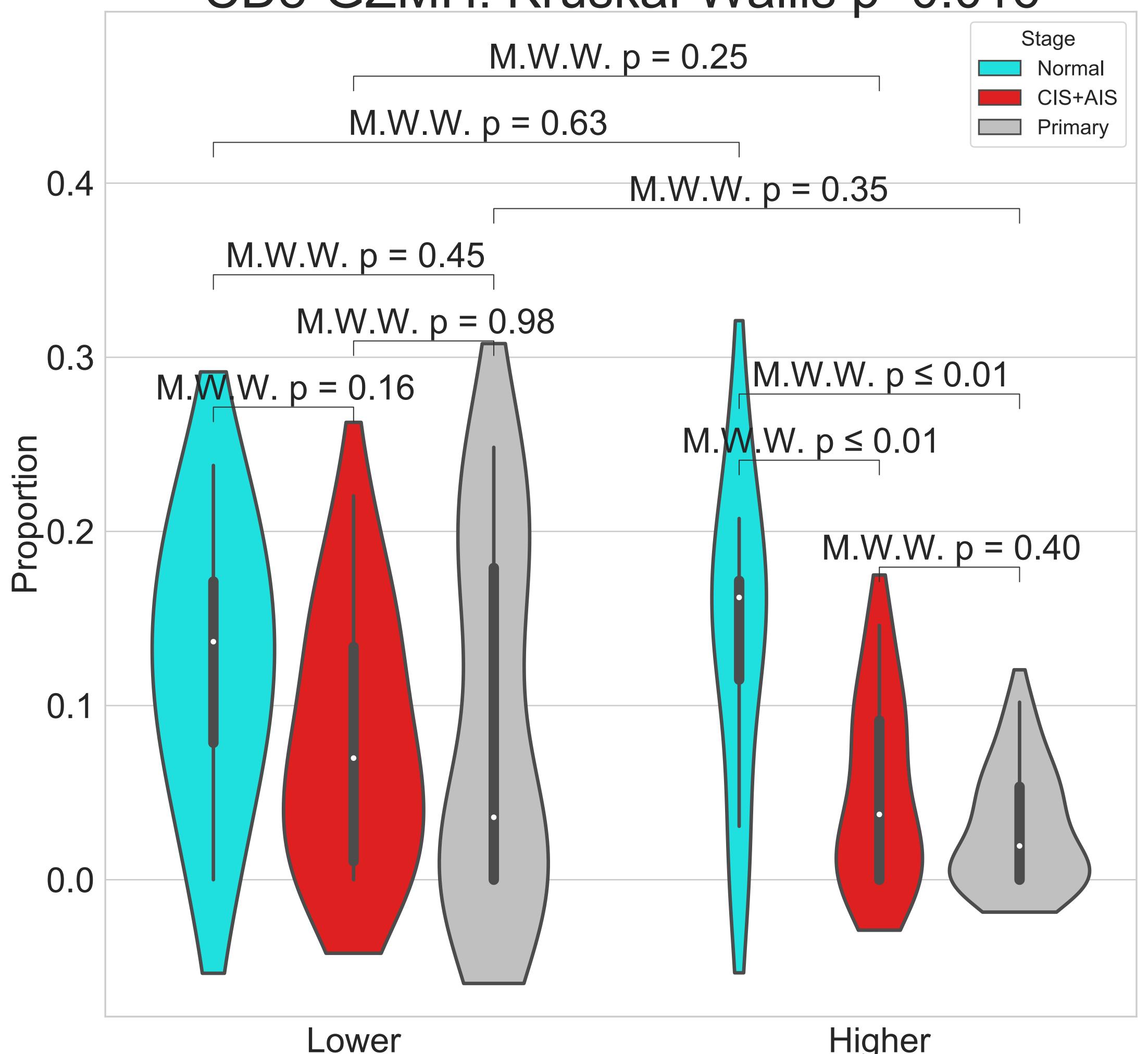
CD8-GZMH: Kruskal-Wallis p=0.016



wer
Mutation Shared Count per TMB