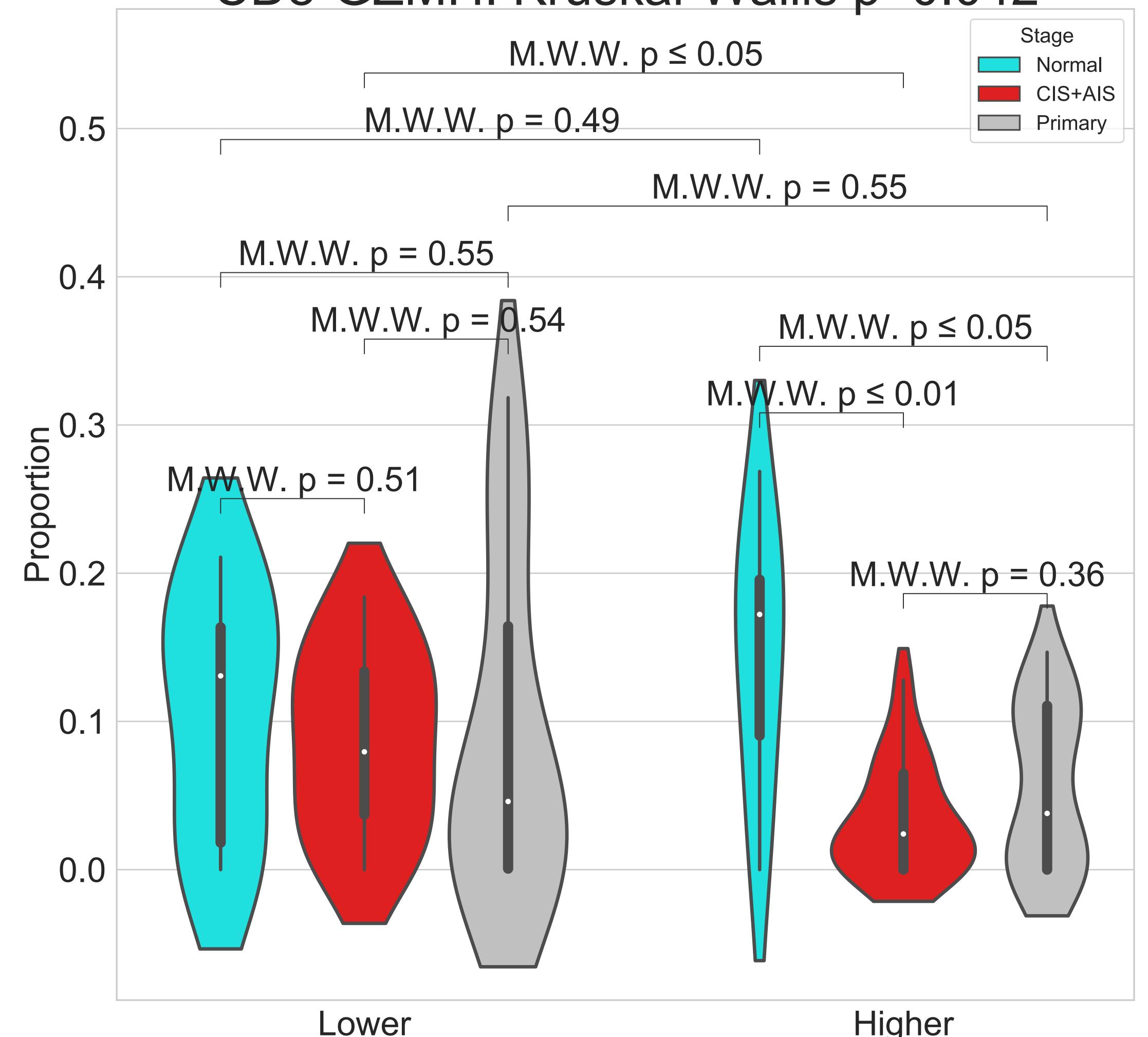
CD8-GZMH: Kruskal-Wallis p=0.042



wer
Mutation Shared Count (SYN)