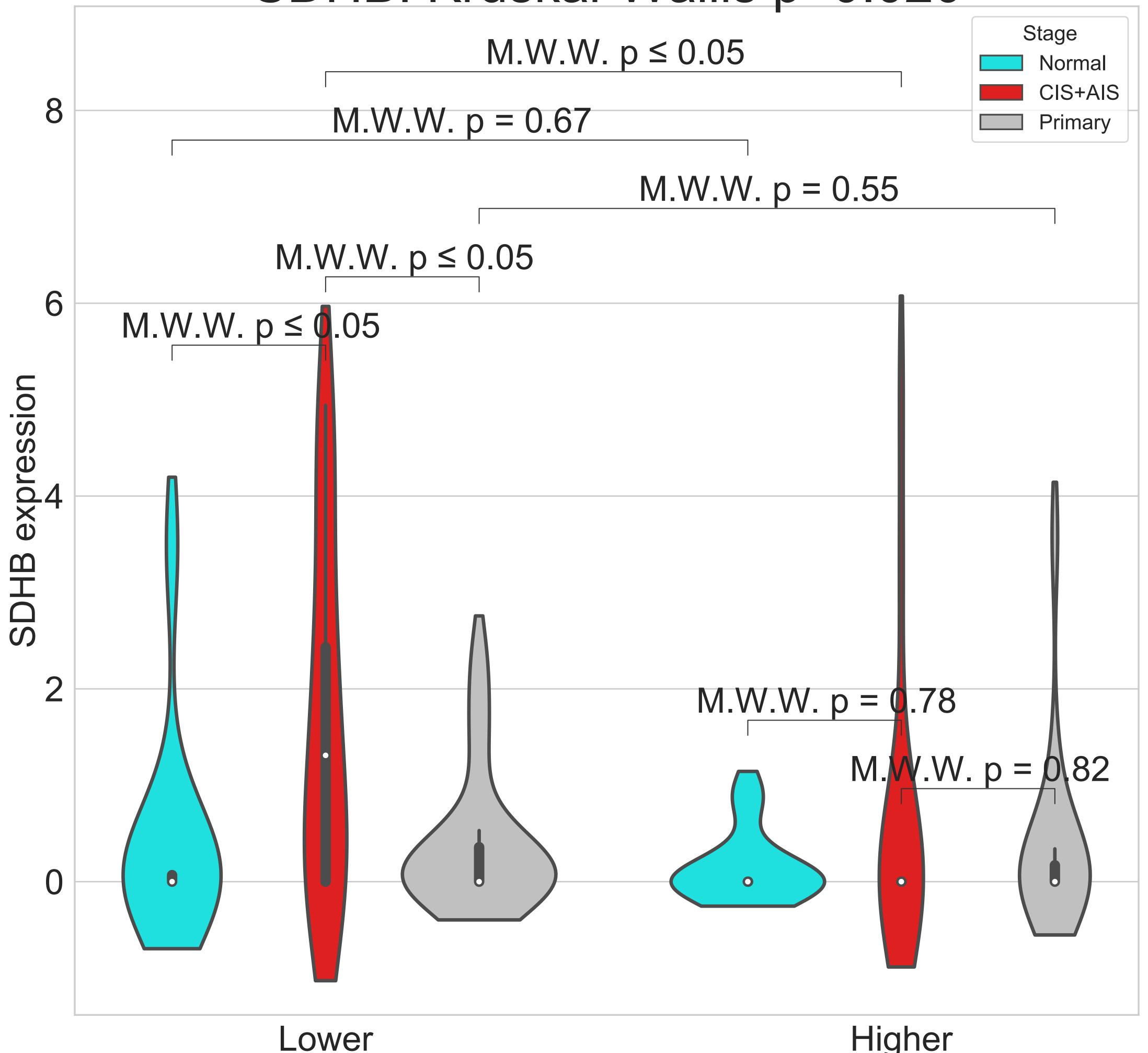
SDHB: Kruskal-Wallis p=0.020



wer
Mutation Shared Proportion (SYN)