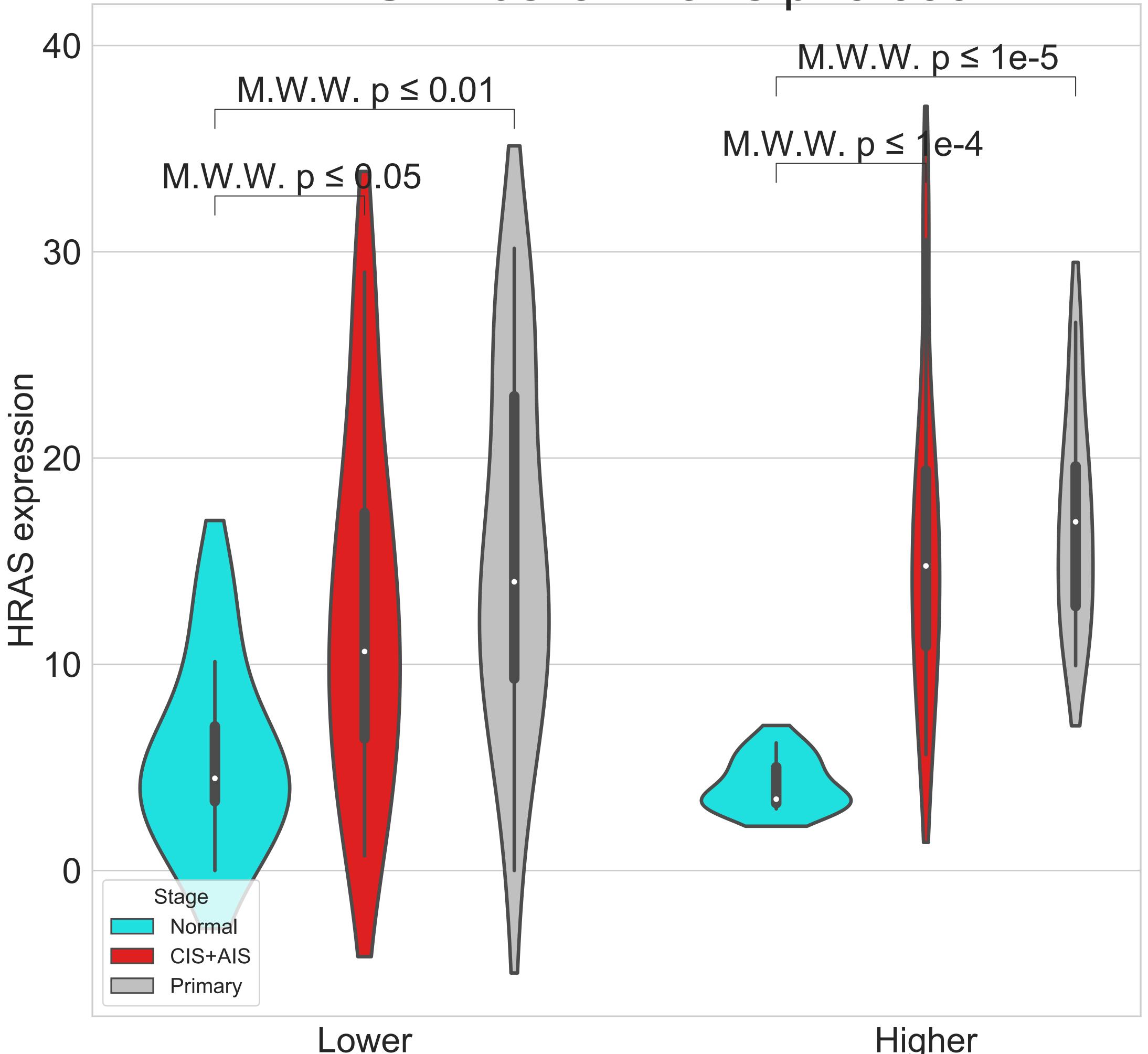
HRAS: Kruskal-Wallis p=0.000



wer
Mutation Shared Count (SYN)