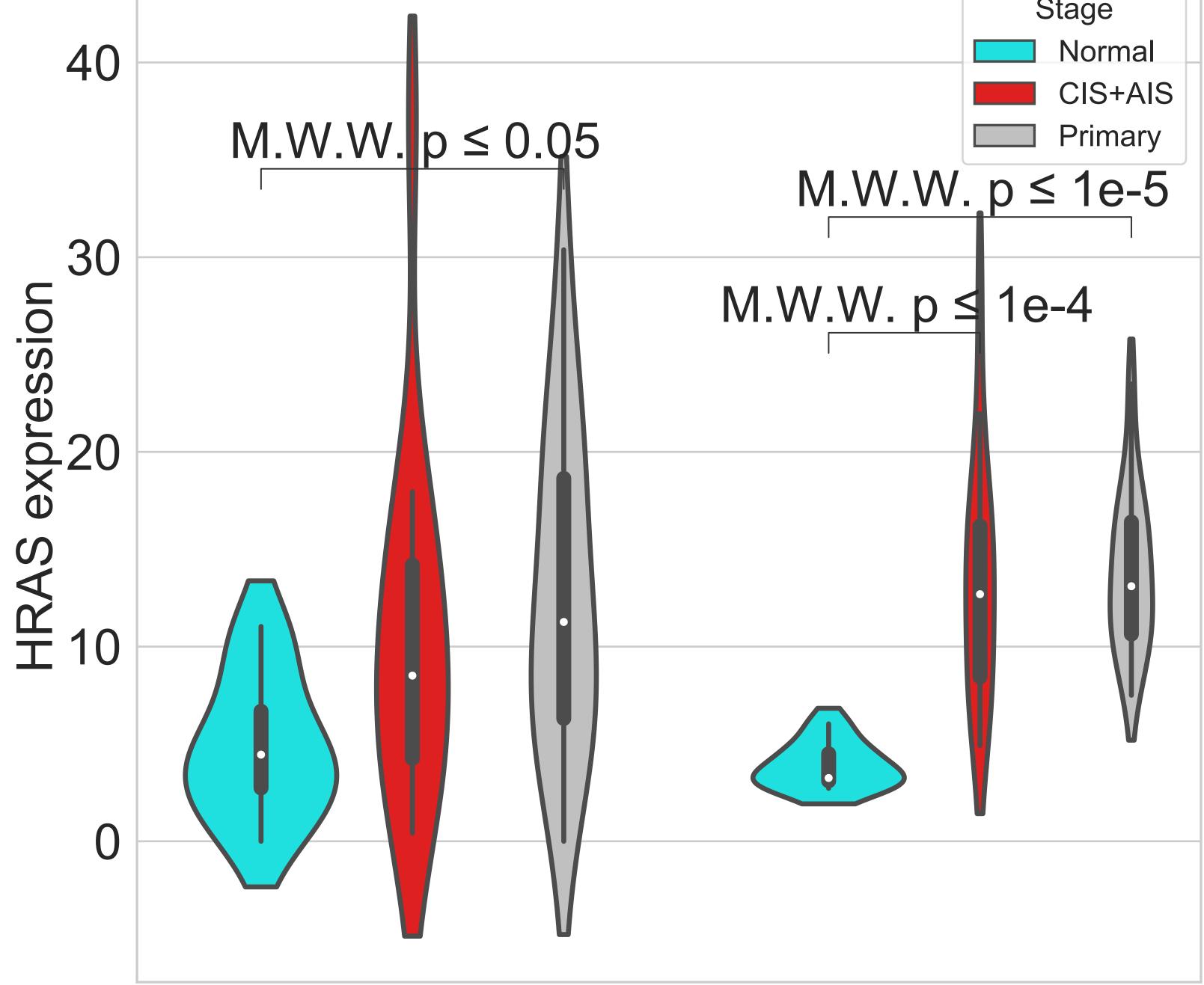
HRAS: Kruskal-Wallis p=0.000

Stage
Normal
CIS+AIS



Lower
Mutation Shared Count (SYN)