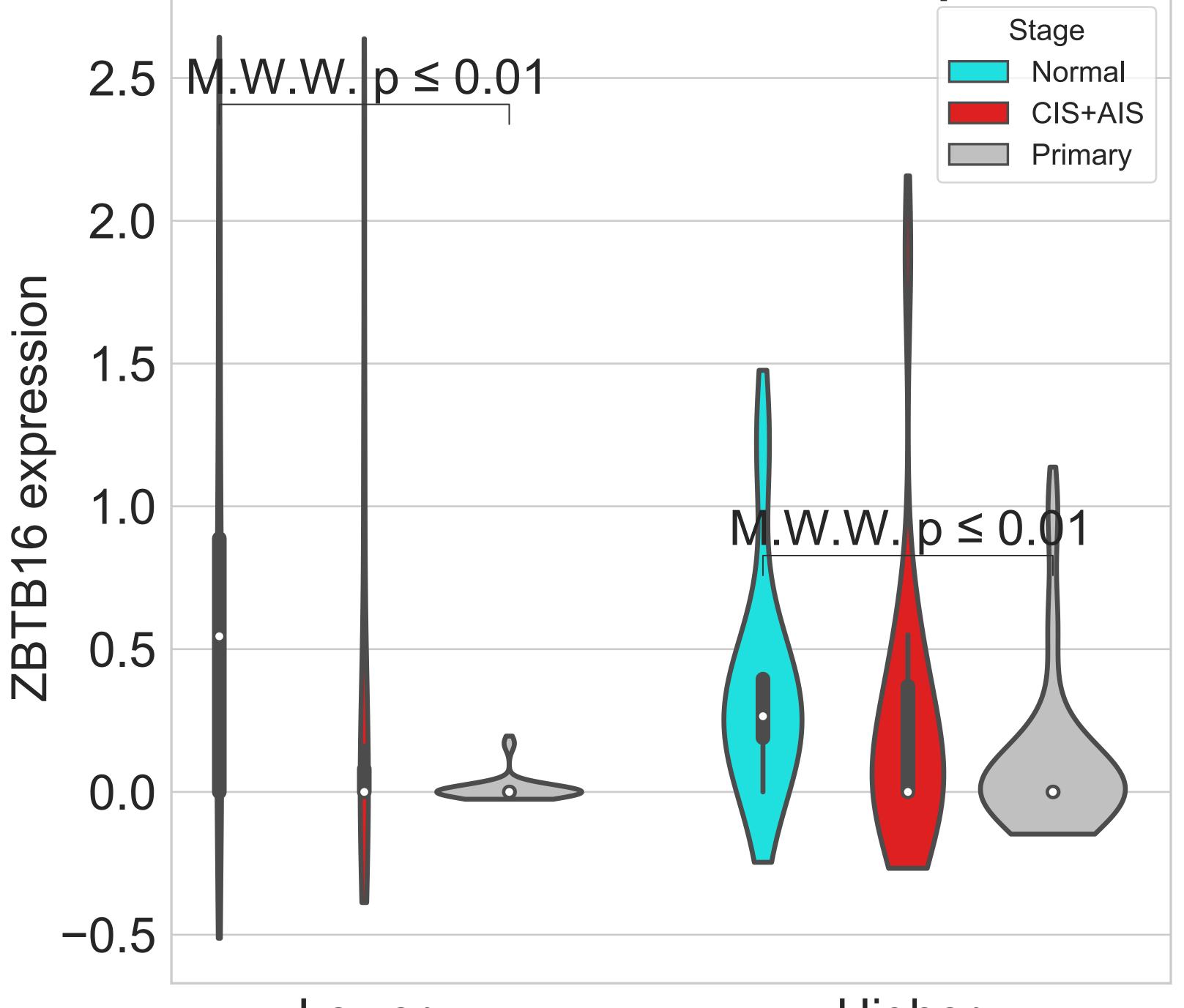
ZBTB16: Kruskal-Wallis p=0.002



Lower

Higher

Mutation Shared Count (SYN)