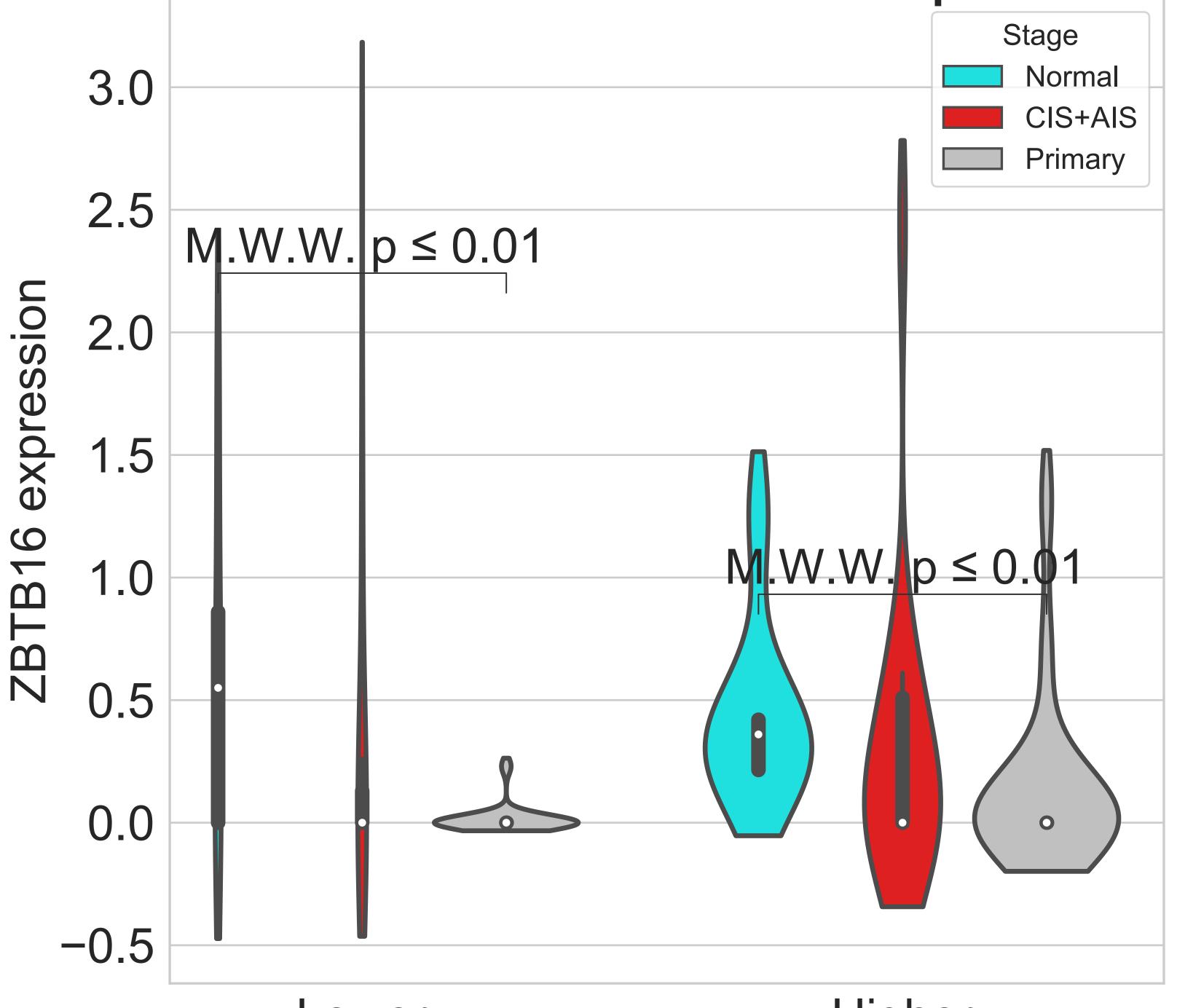
ZBTB16: Kruskal-Wallis p=0.003



Lower

Higher

Mutation Shared Count (SYN)