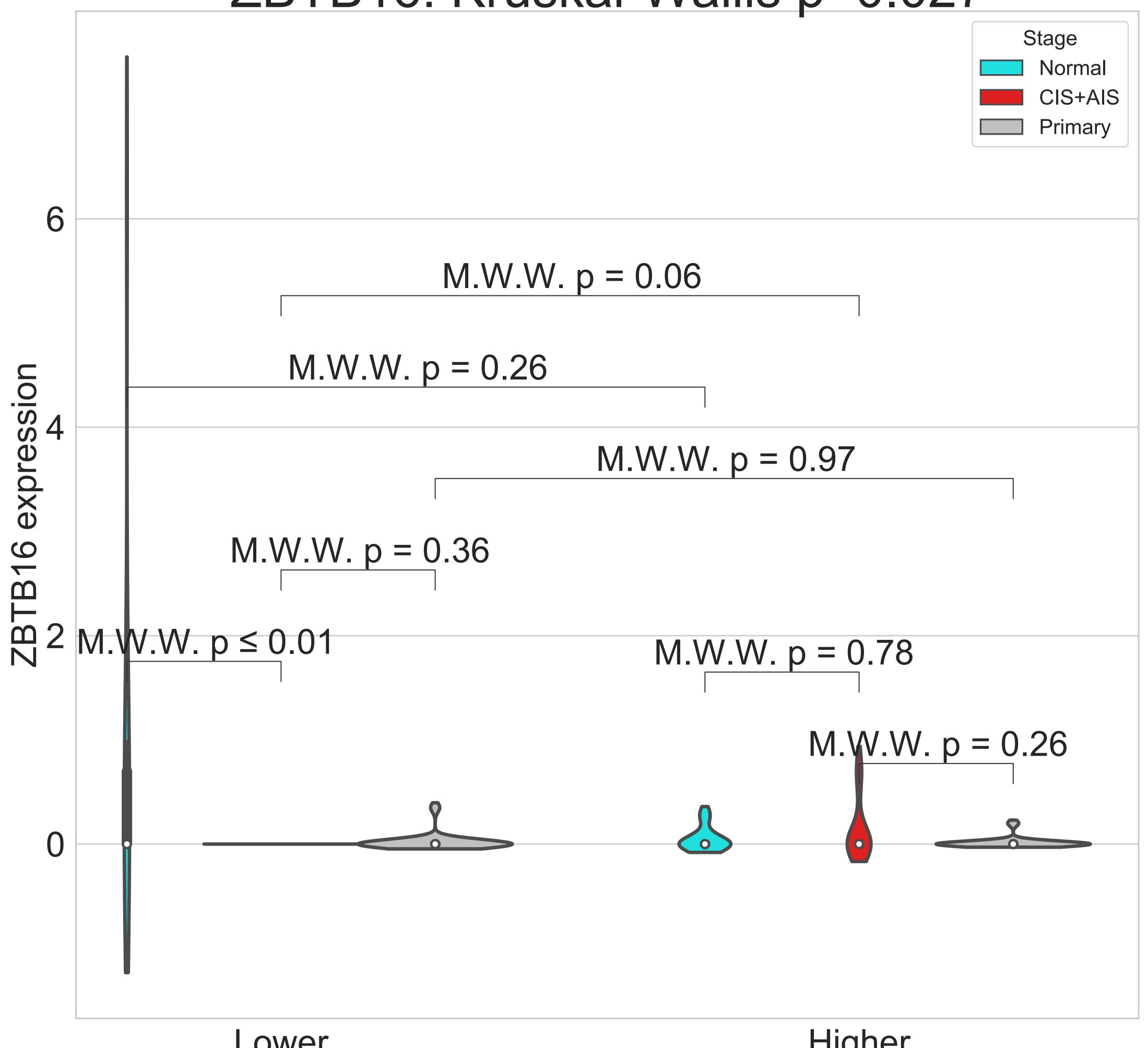
ZBTB16: Kruskal-Wallis p=0.027



Lower Higher Mutation Shared Proportion (SYN)