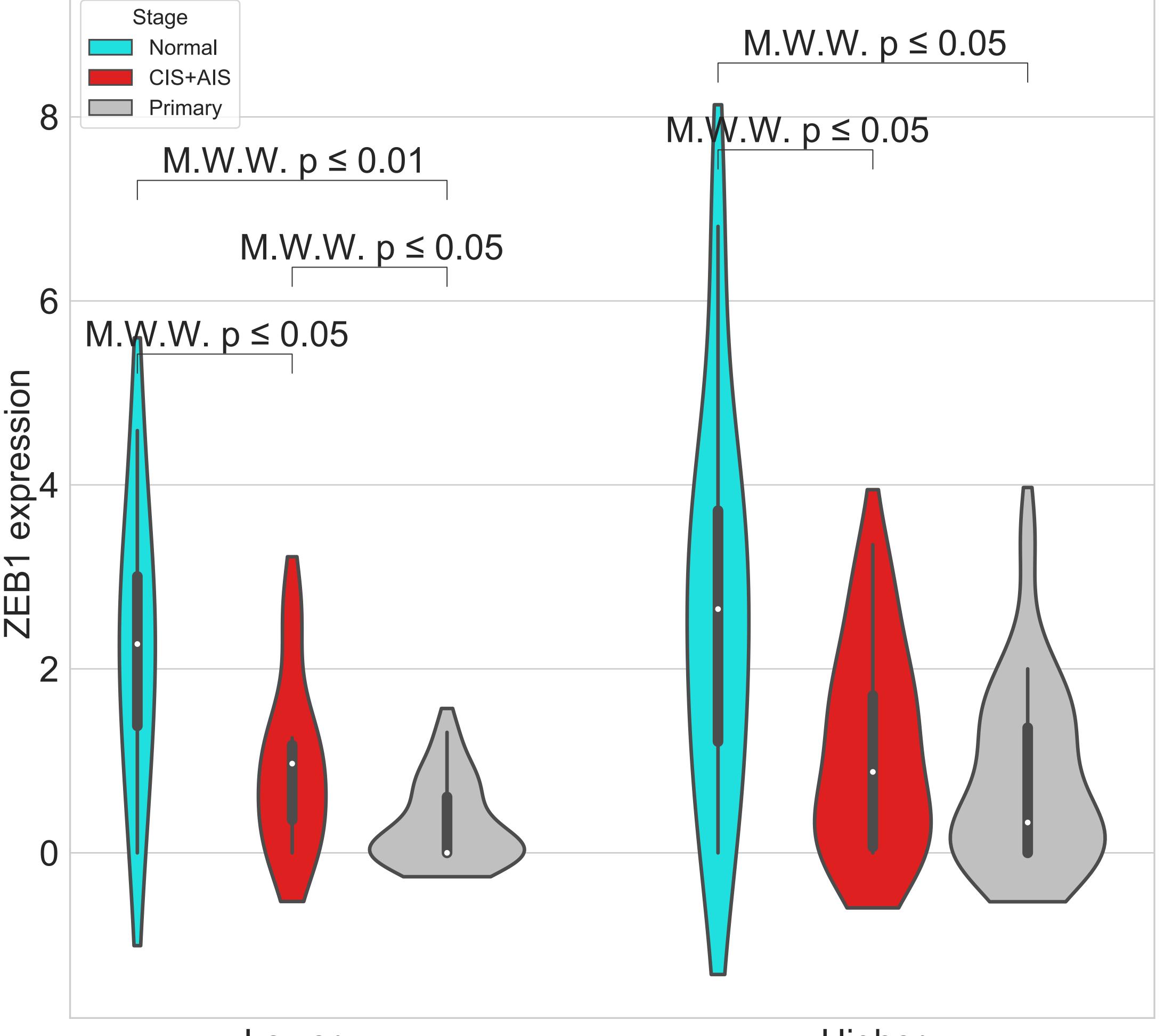
ZEB1: Kruskal-Wallis p=0.002



Higher Lower Mutation Shared Proportion (SYN)