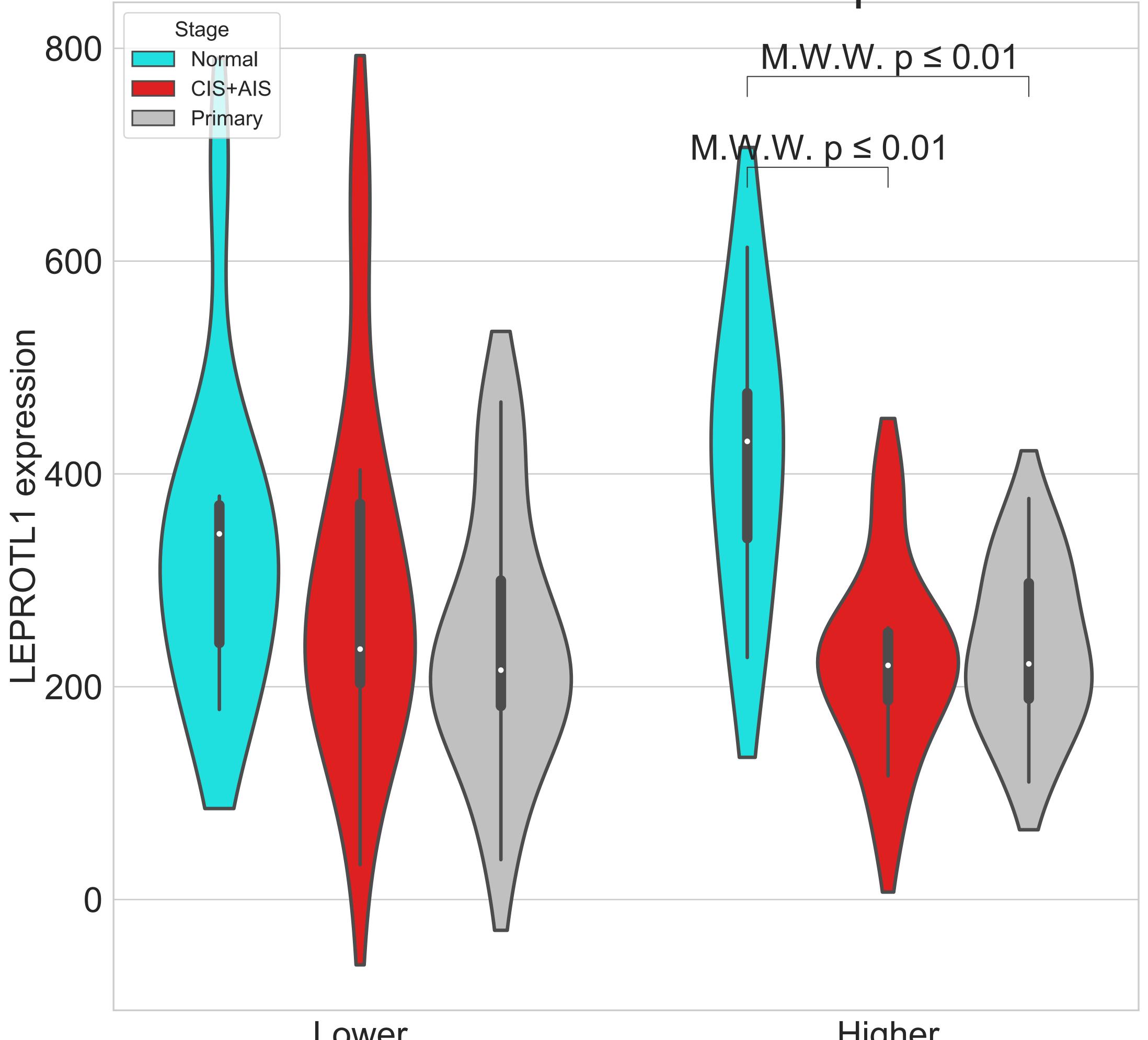
LEPROTL1: Kruskal-Wallis p=0.028



Lower
Mutation Shared Proportion (SYN)