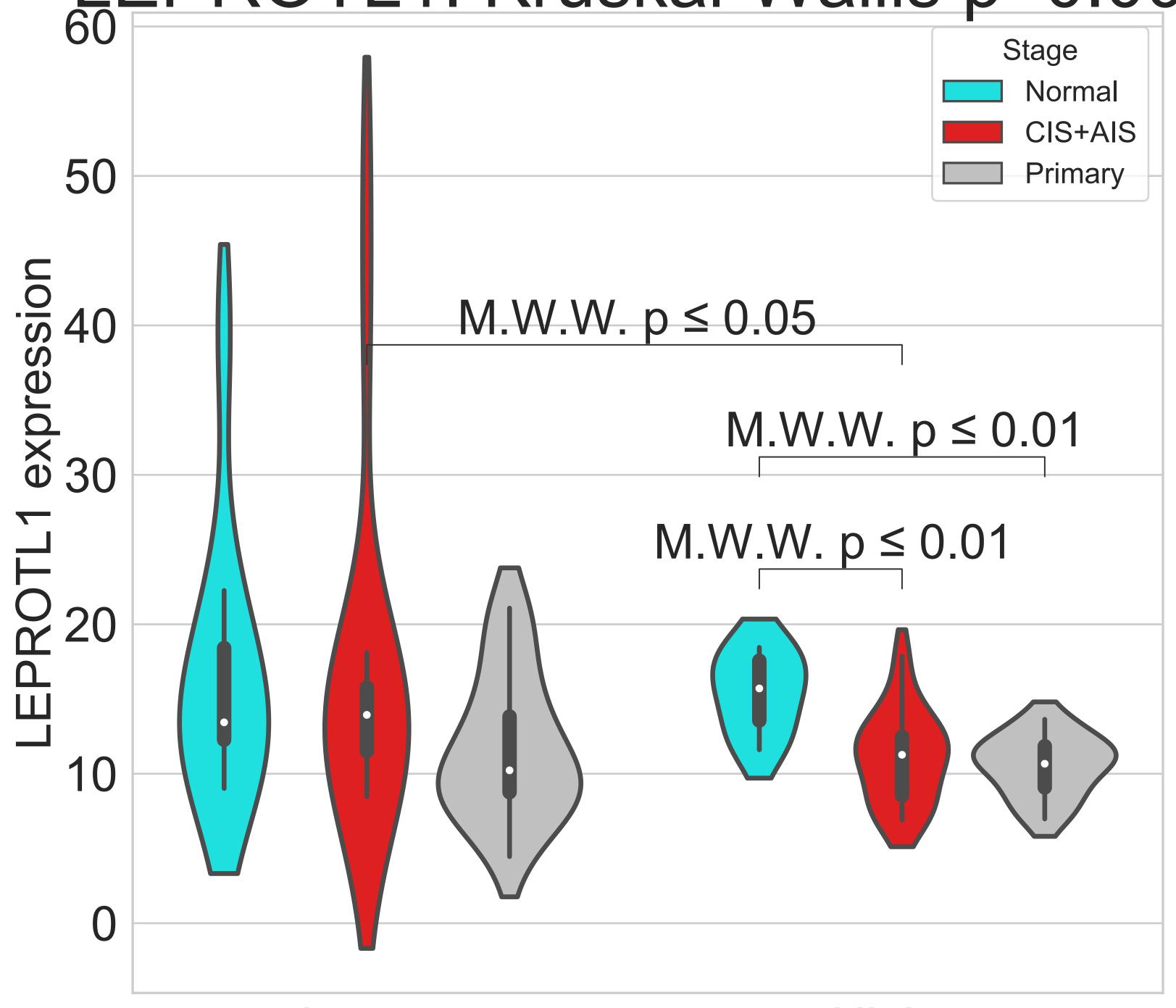
LEPROTL1: Kruskal-Wallis p=0.006



Lower Higher Mutation Shared Proportion (SYN)