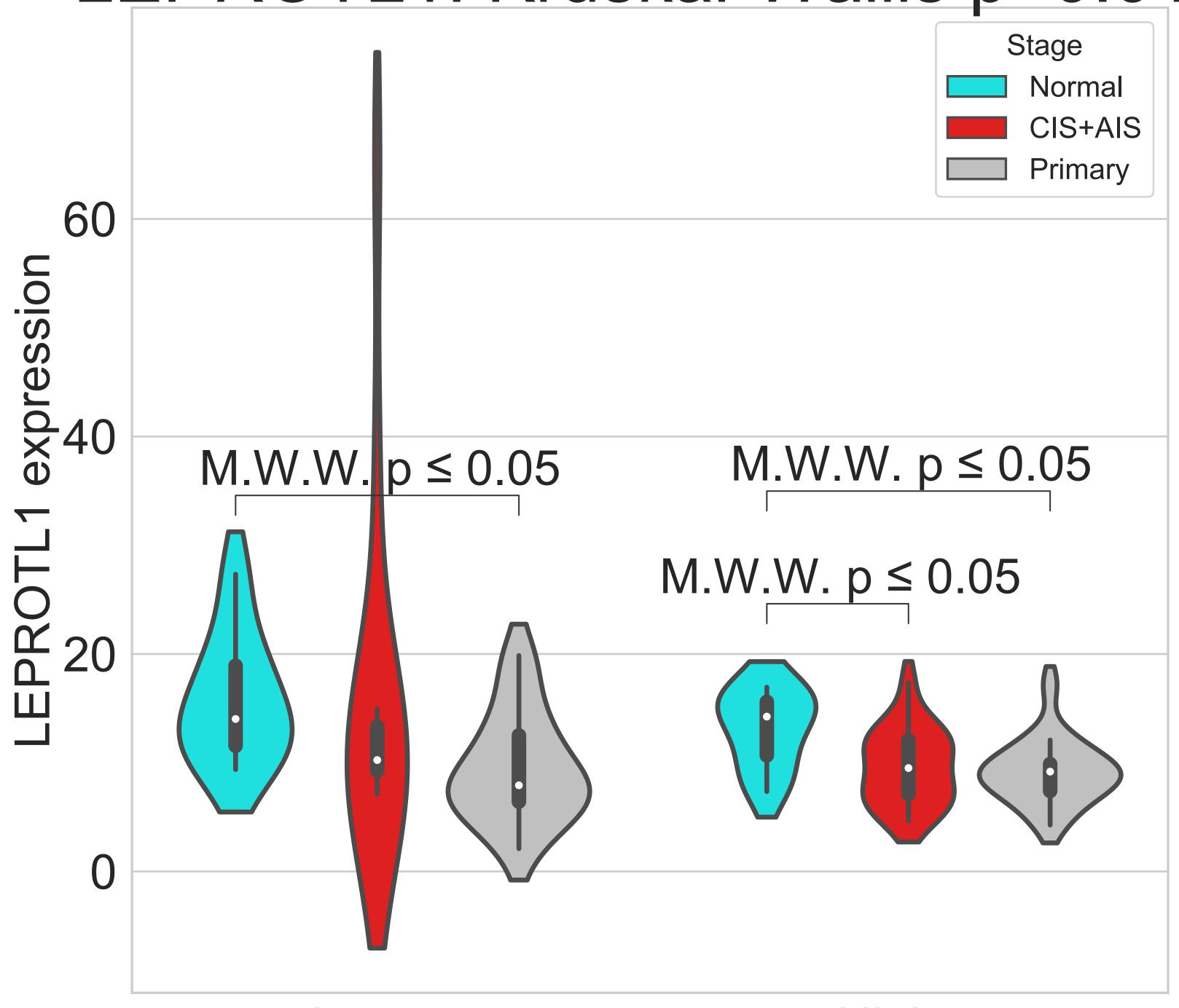
LEPROTL1: Kruskal-Wallis p=0.011



Lower Higher Mutation Shared Proportion (Union & SYN)