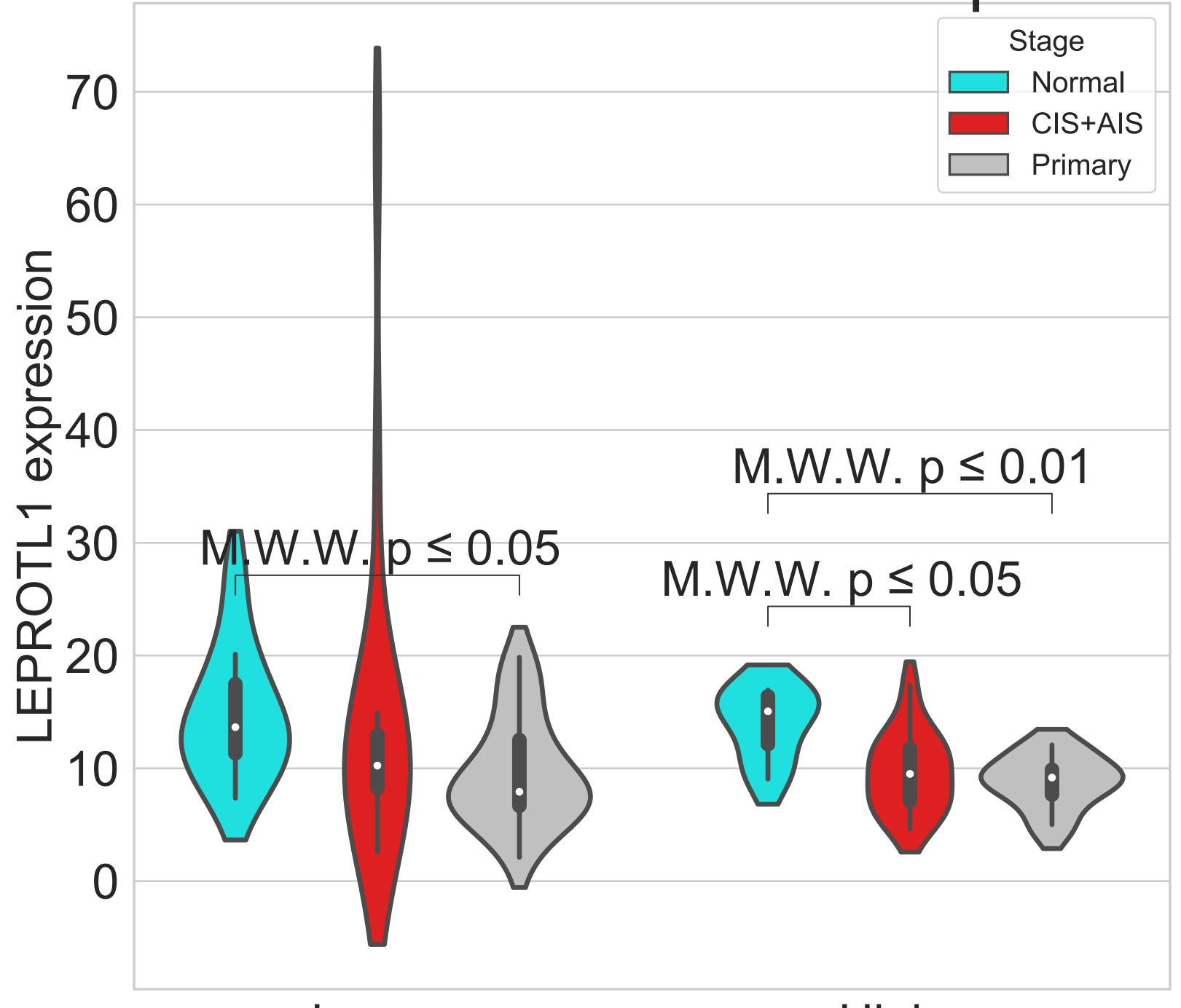
LEPROTL1: Kruskal-Wallis p=0.016



Lower
Higher
Mutation Shared Proportion