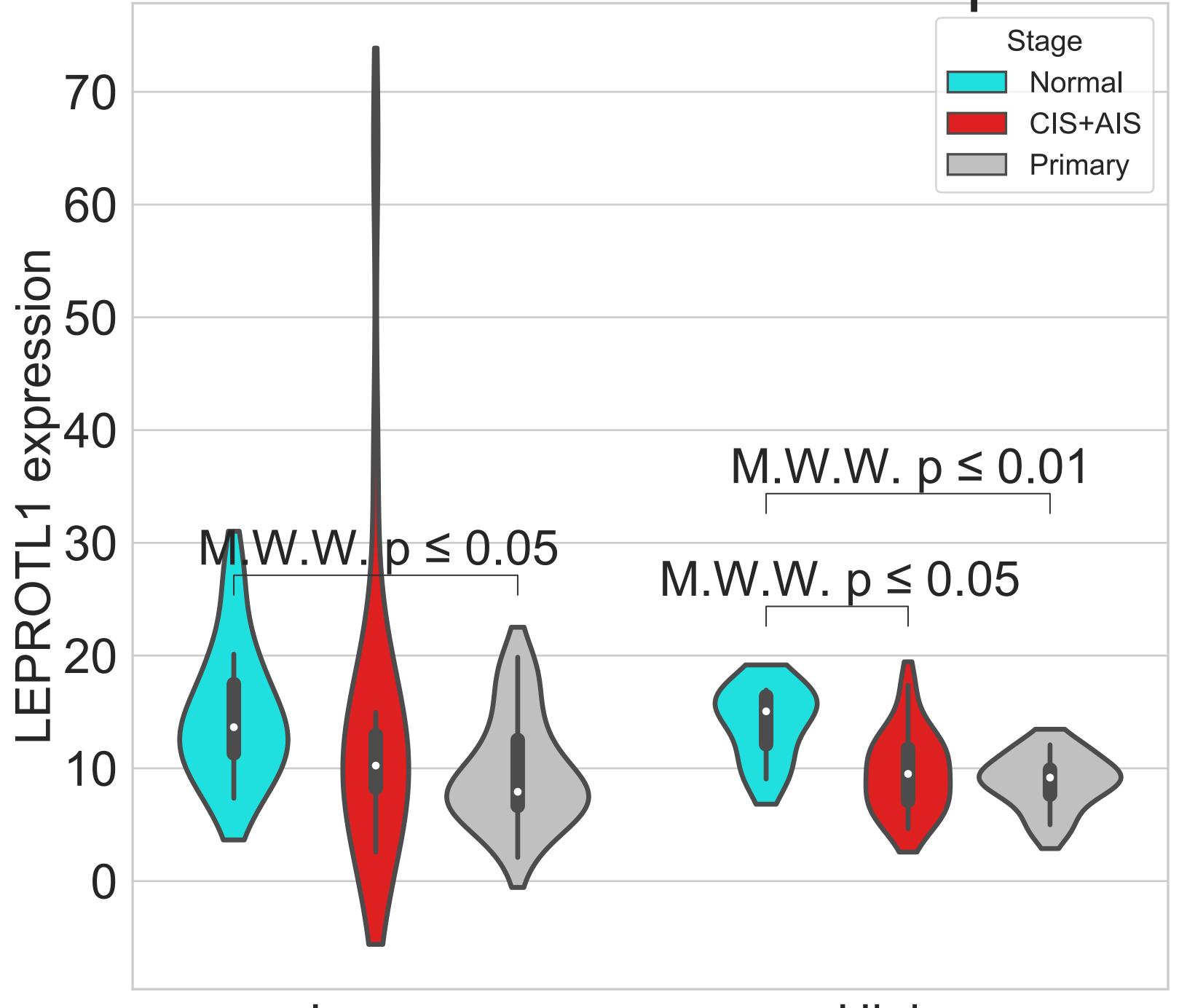
LEPROTL1: Kruskal-Wallis p=0.016



Lower Higher Mutation Shared Count per TMB