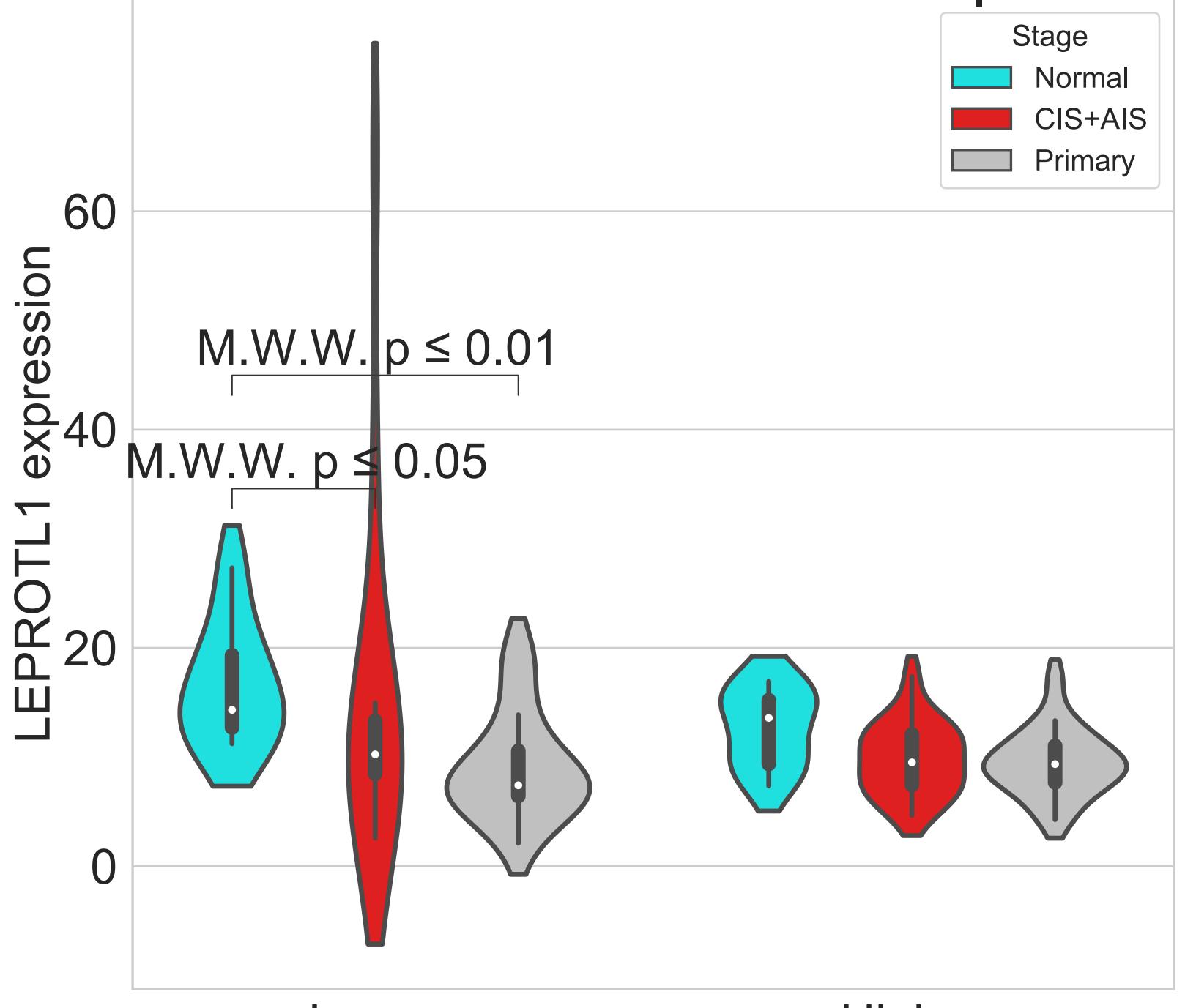
LEPROTL1: Kruskal-Wallis p=0.009



Lower Higher Mutation Shared Count per TMB