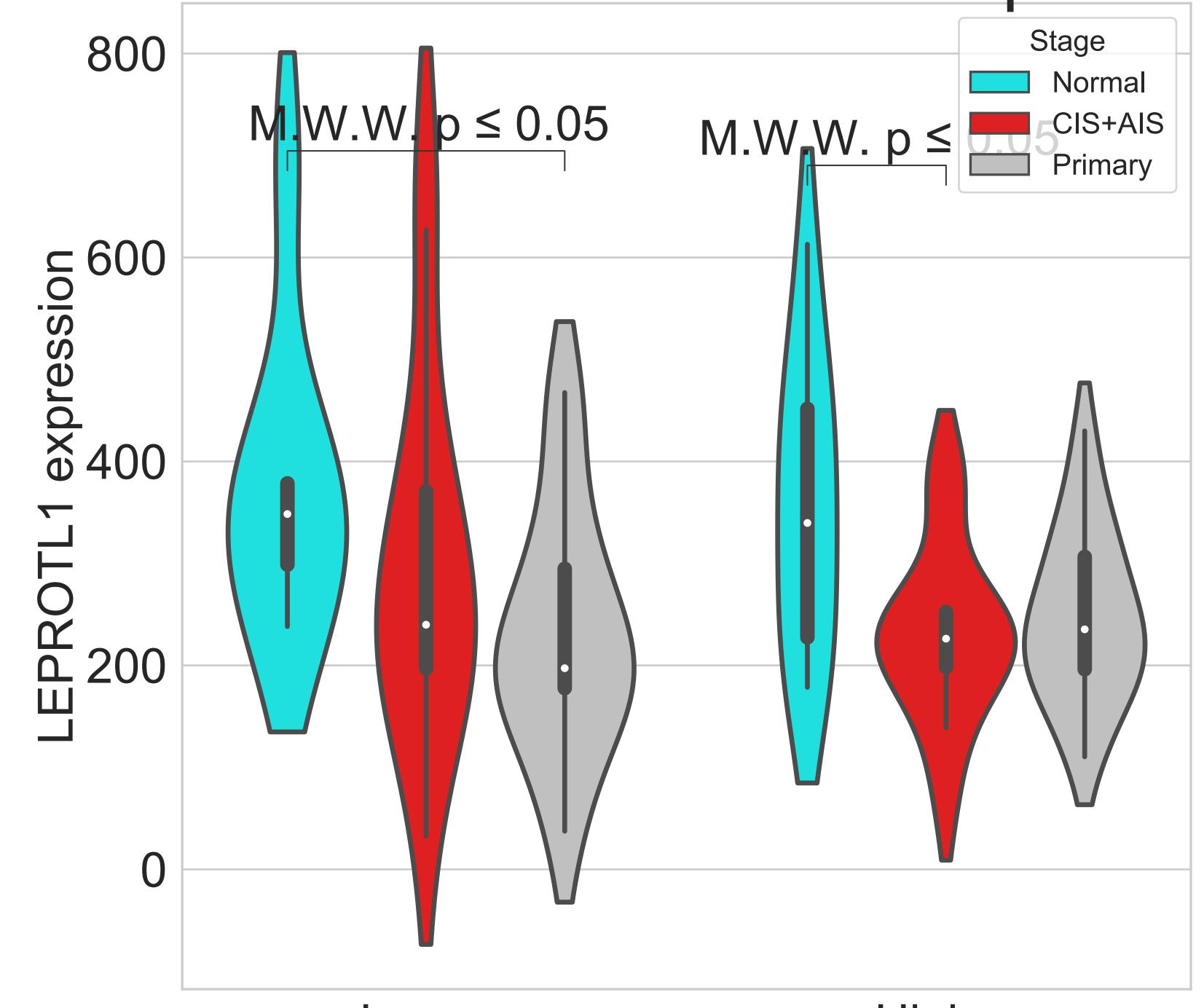
LEPROTL1: Kruskal-Wallis p=0.040



Lower Higher Mutation Shared Count per TMB