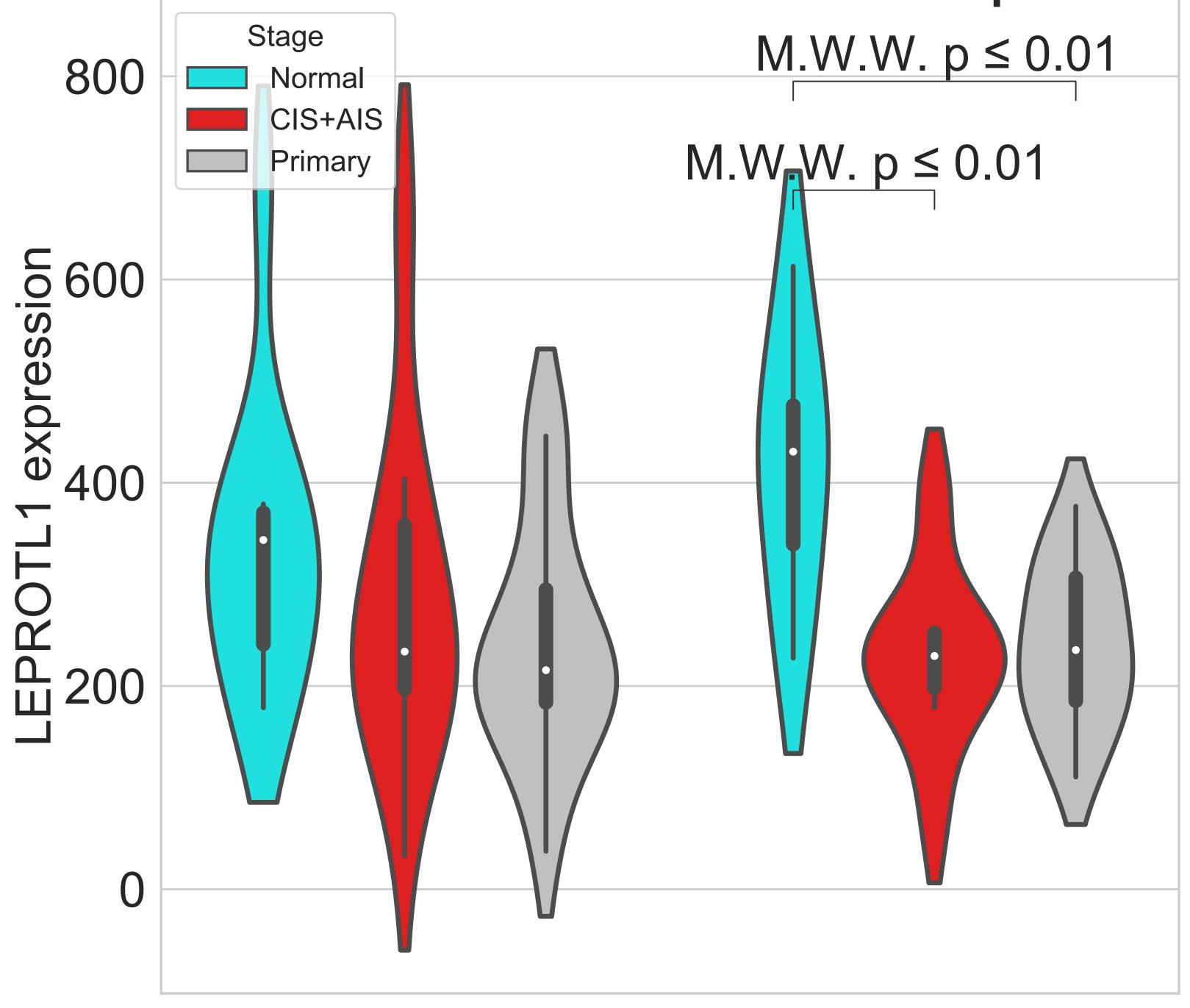
LEPROTL1: Kruskal-Wallis p=0.038



Lower
Mutation Shared Count per TMB