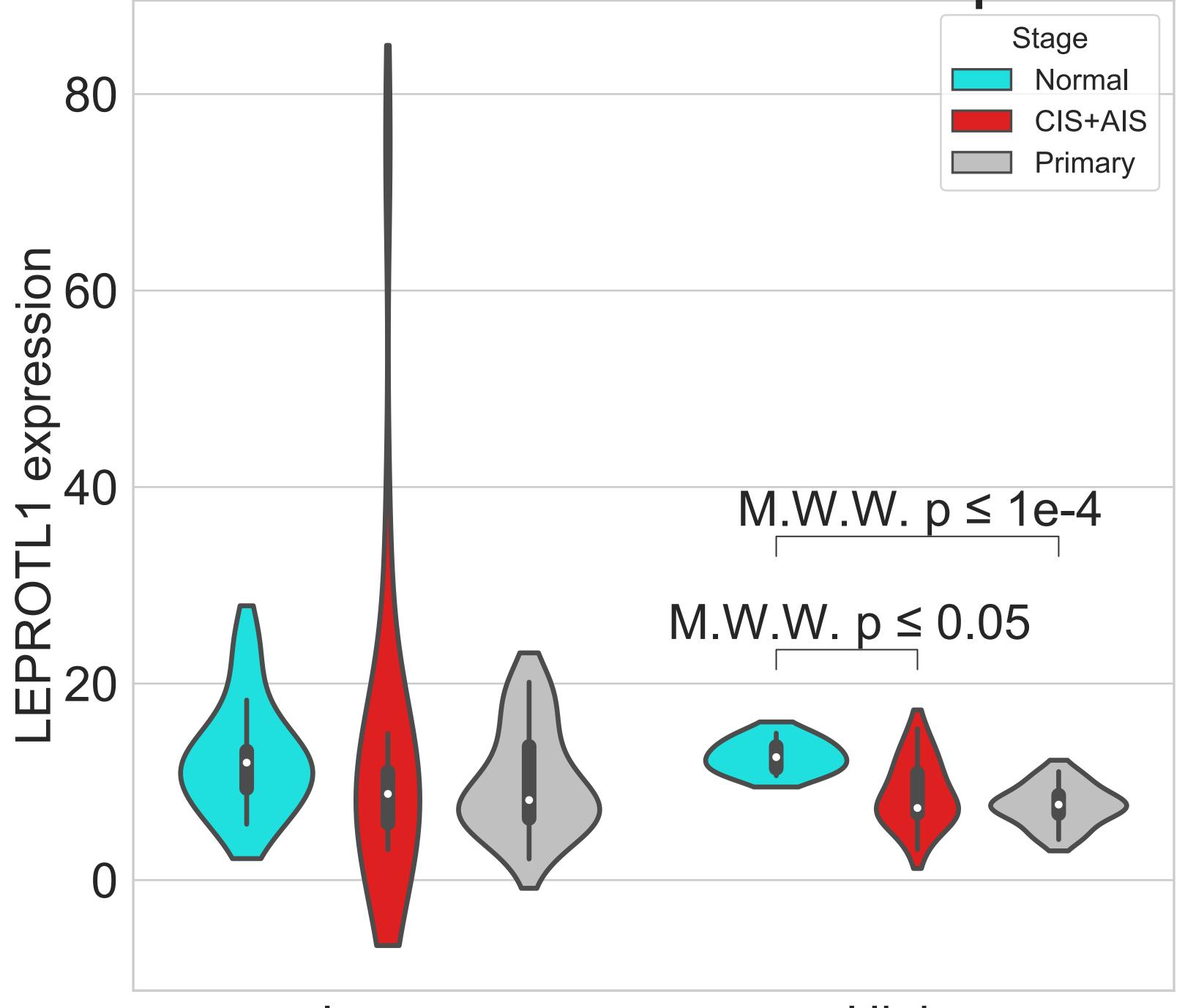
LEPROTL1: Kruskal-Wallis p=0.037



Lower
Mutation Shared Count (SYN) per TMB