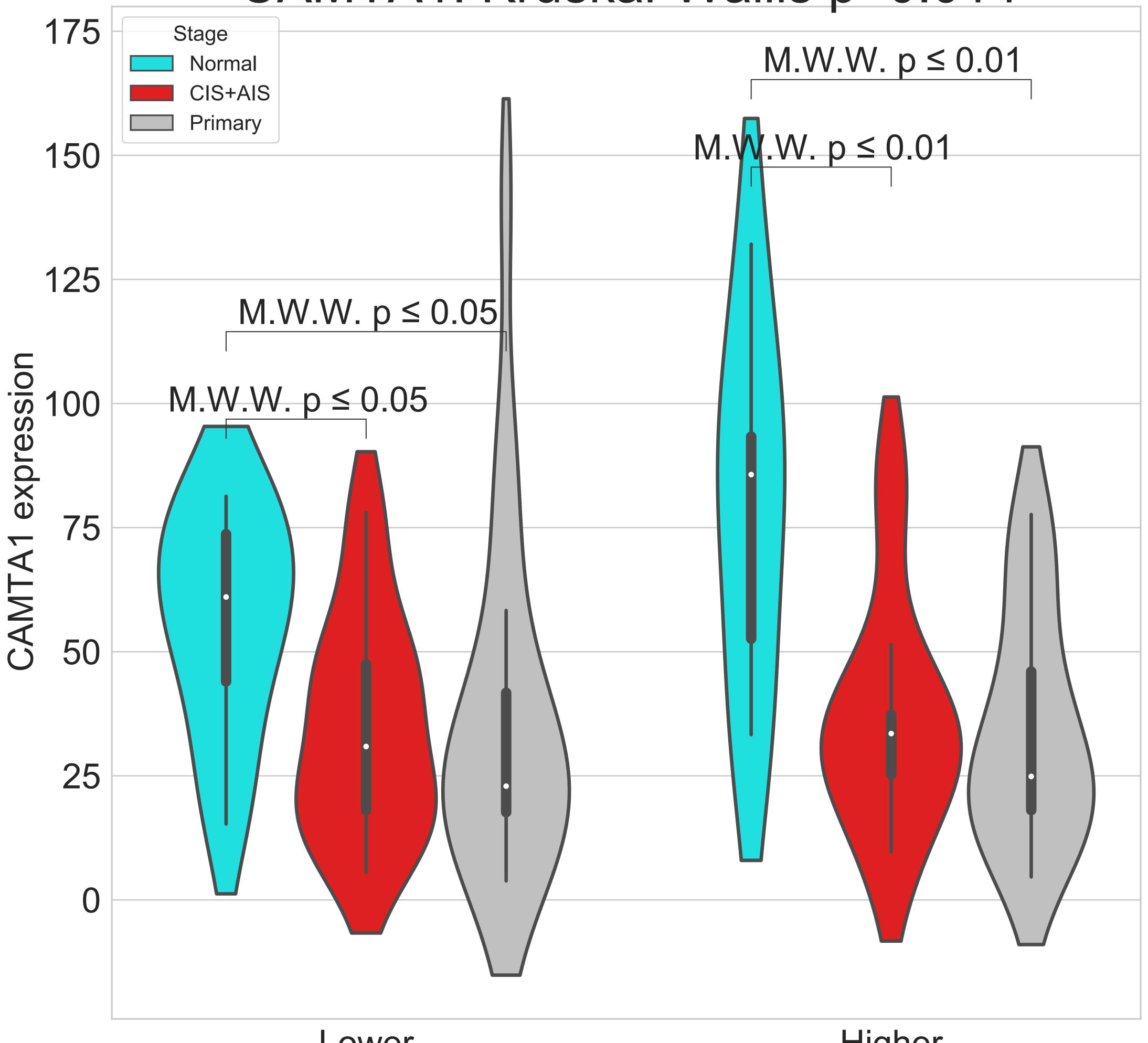
CAMTA1: Kruskal-Wallis p=0.014



Lower
Mutation Shared Count per TMB