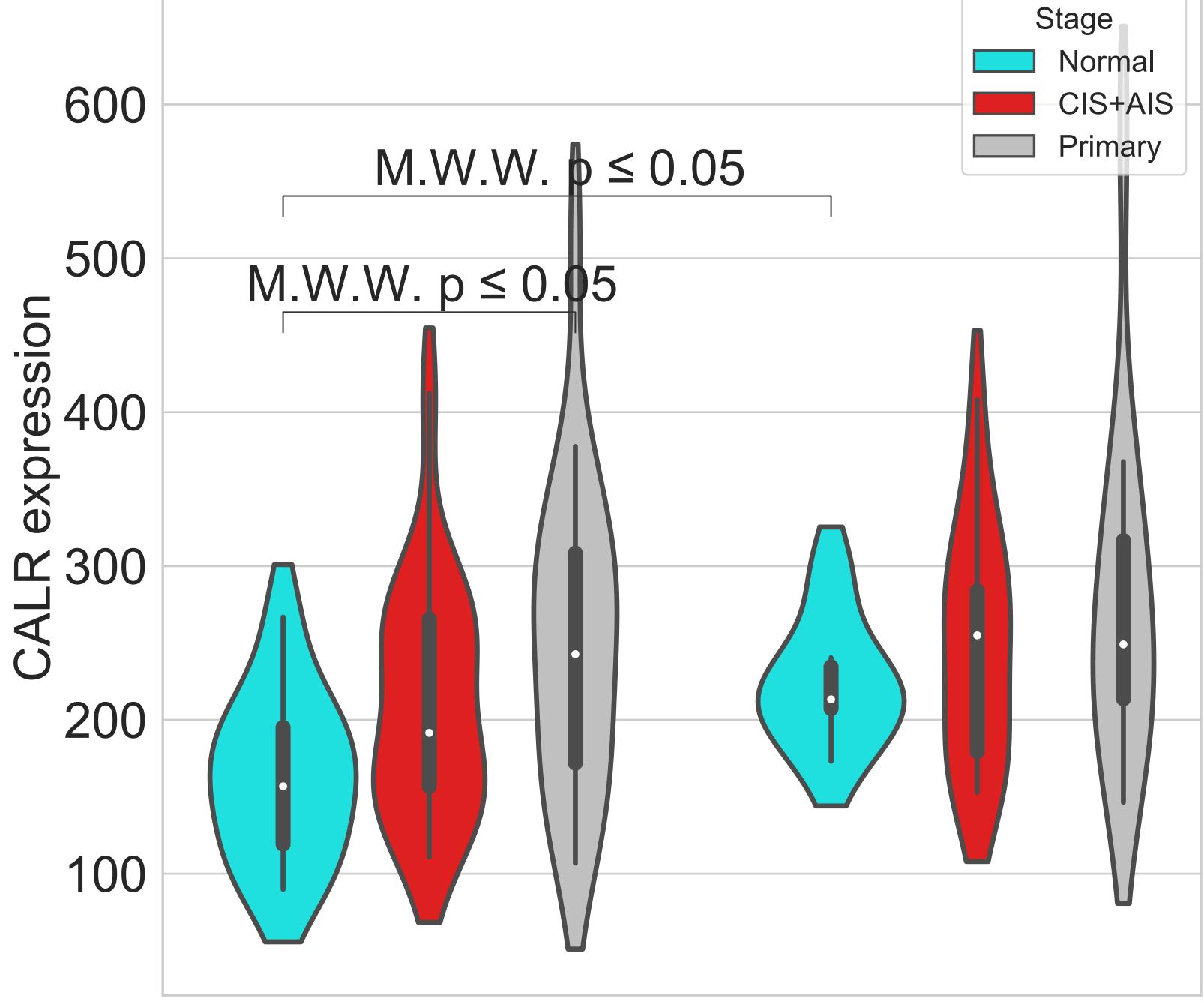
CALR: Kruskal-Wallis p=0.024

Stage
Normal
CIS+AIS



Lower Higher Mutation Shared Count per TMB