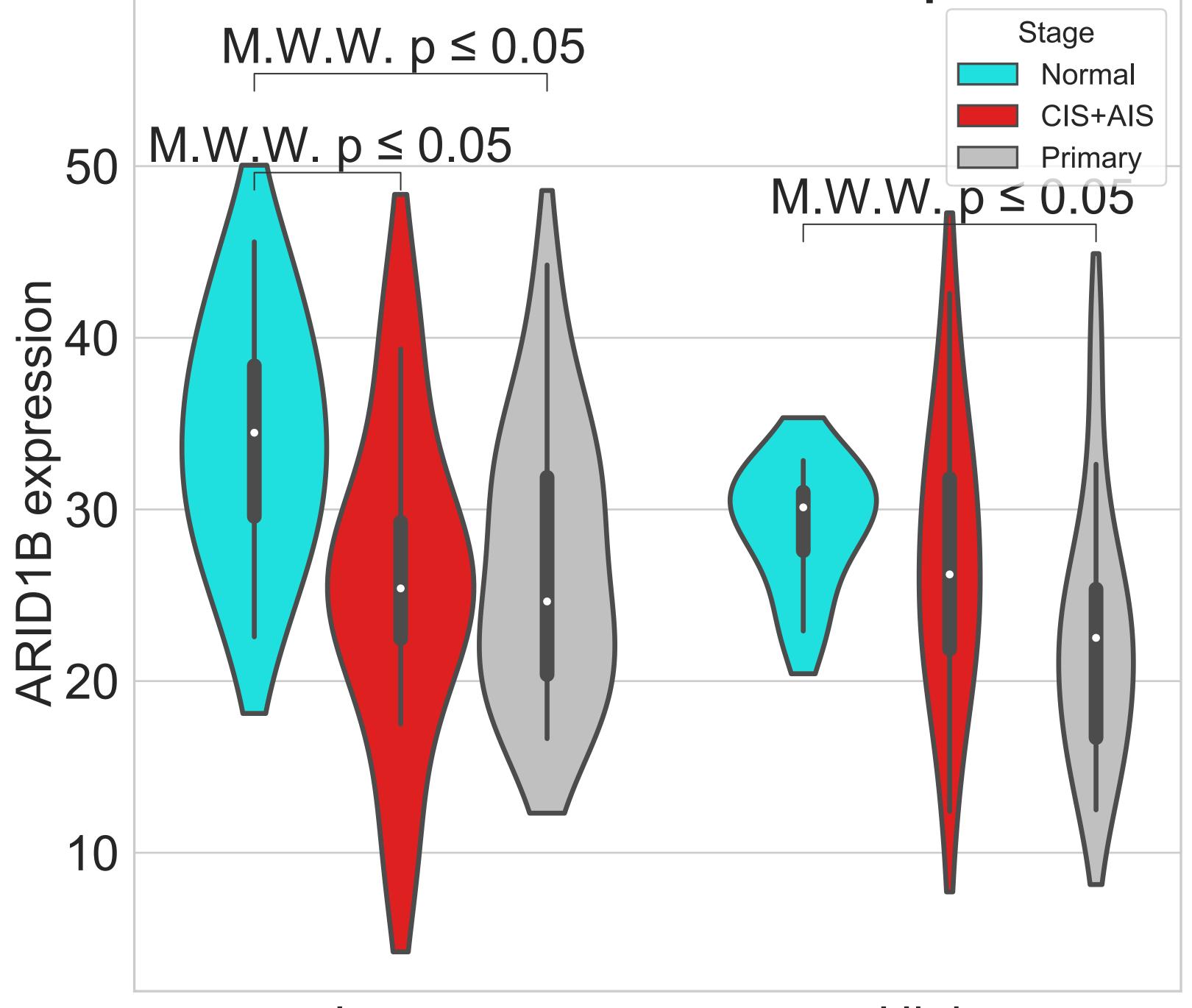
ARID1B: Kruskal-Wallis p=0.024



Lower
Mutation Shared Count per TMB