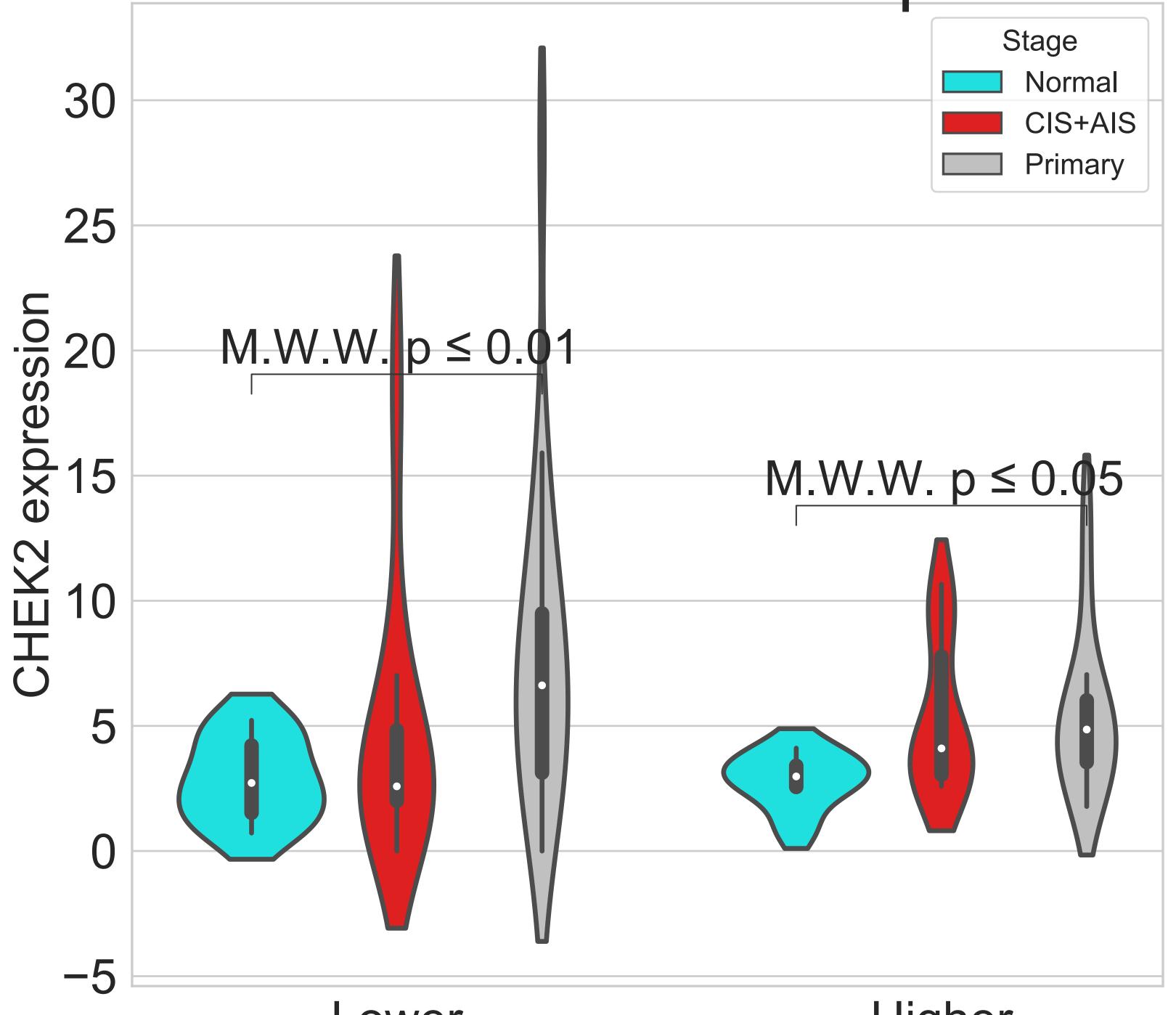
CHEK2: Kruskal-Wallis p=0.018



Lower
Mutation Shared Count per TMB