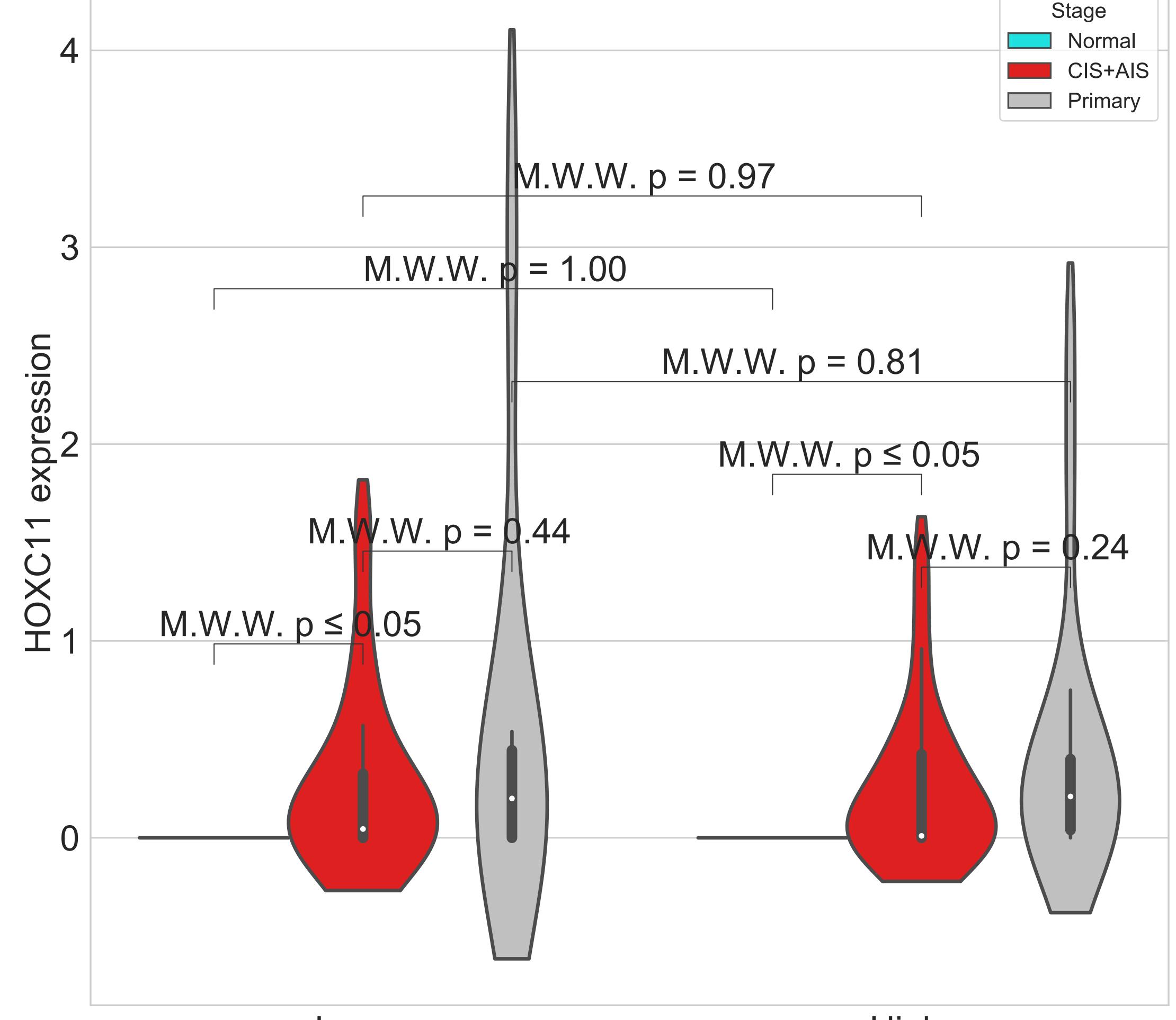
HOXC11: Kruskal-Wallis p=0.002



Higher Lower Mutation Shared Count per TMB