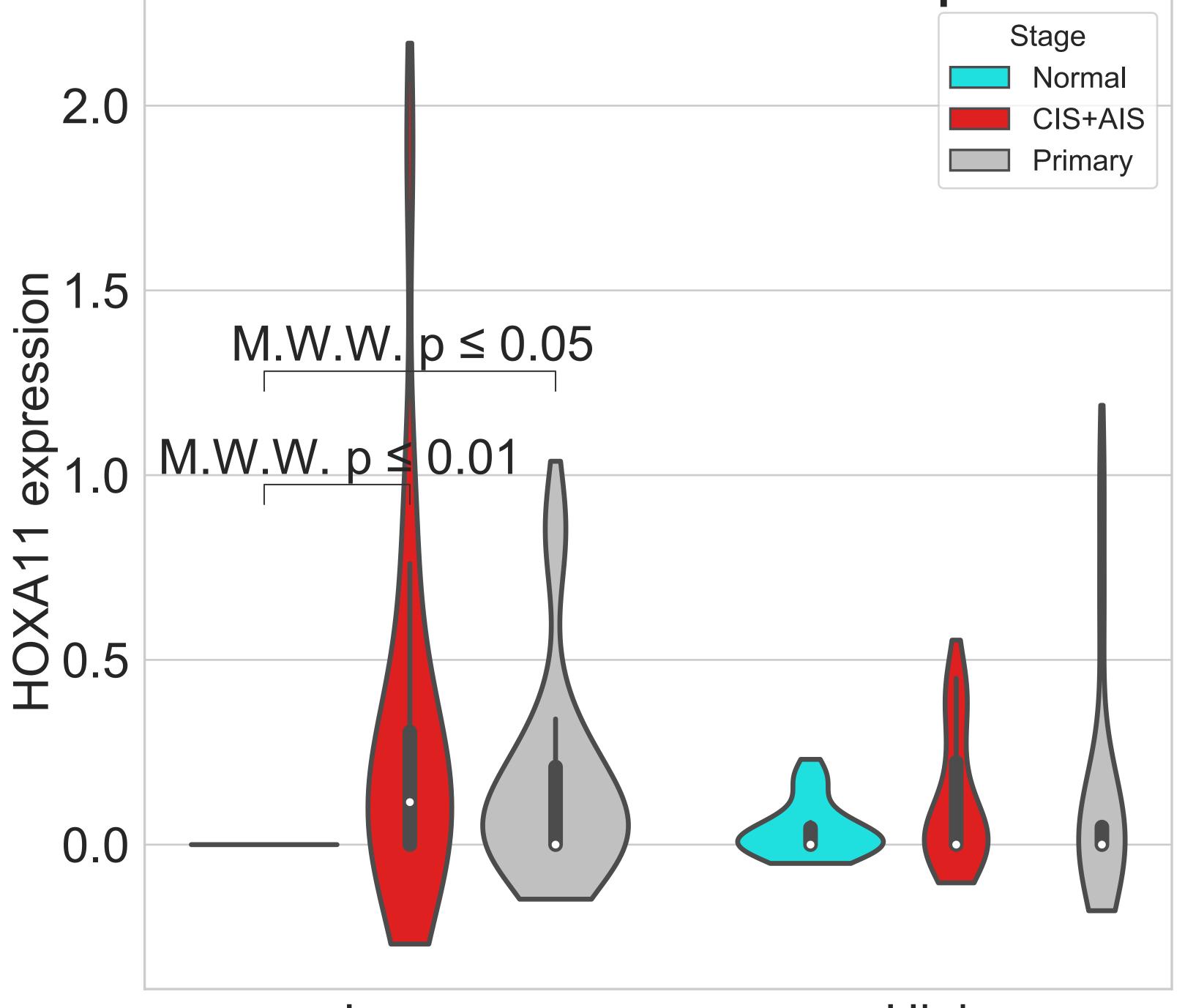
HOXA11: Kruskal-Wallis p=0.033



Lower
Mutation Shared Count per TMB