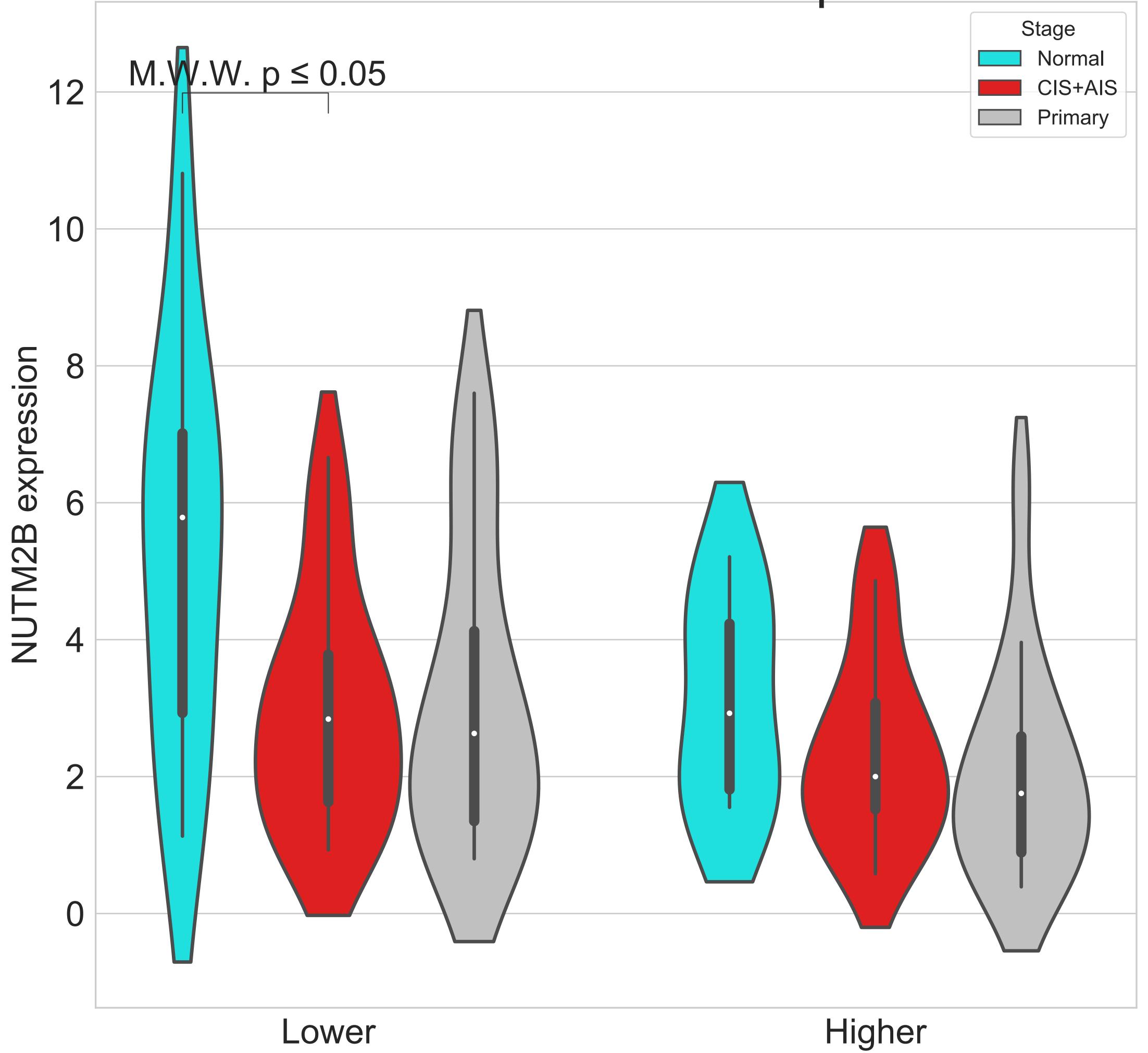
NUTM2B: Kruskal-Wallis p=0.034



Mutation Shared Count per TMB