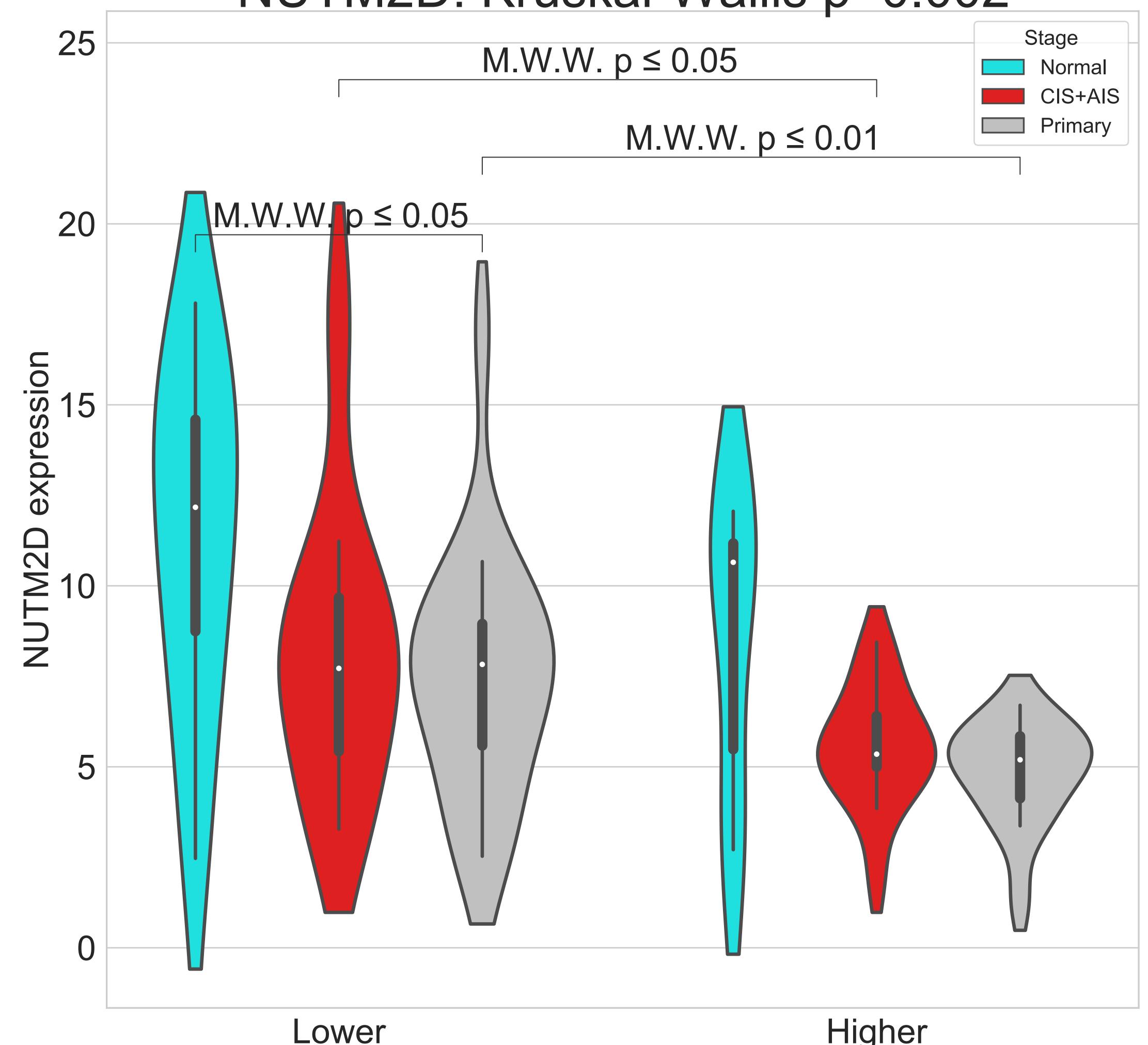
NUTM2D: Kruskal-Wallis p=0.002



wer
Mutation Shared Count per TMB