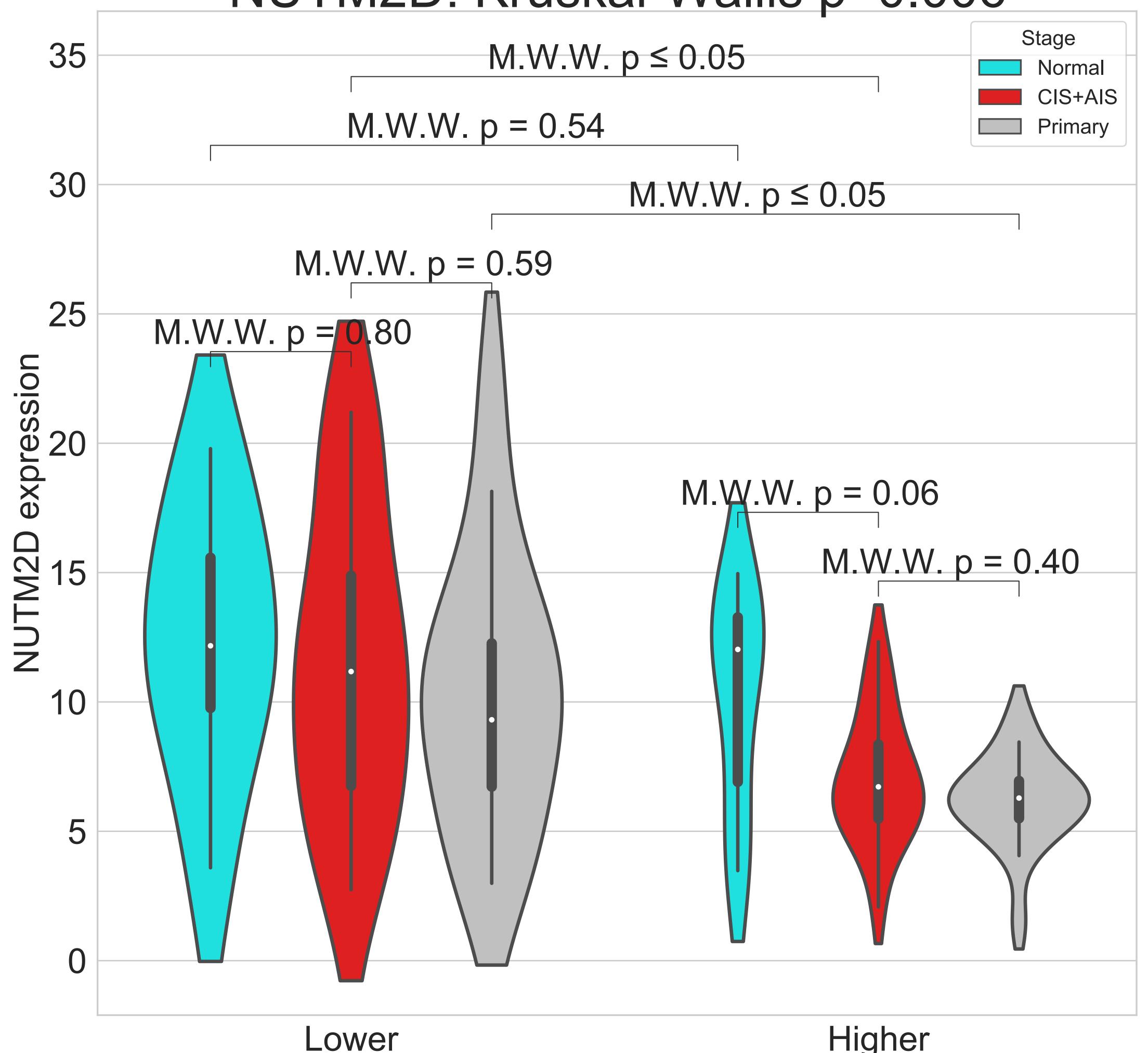
NUTM2D: Kruskal-Wallis p=0.006



wer
Mutation Shared Count per TMB