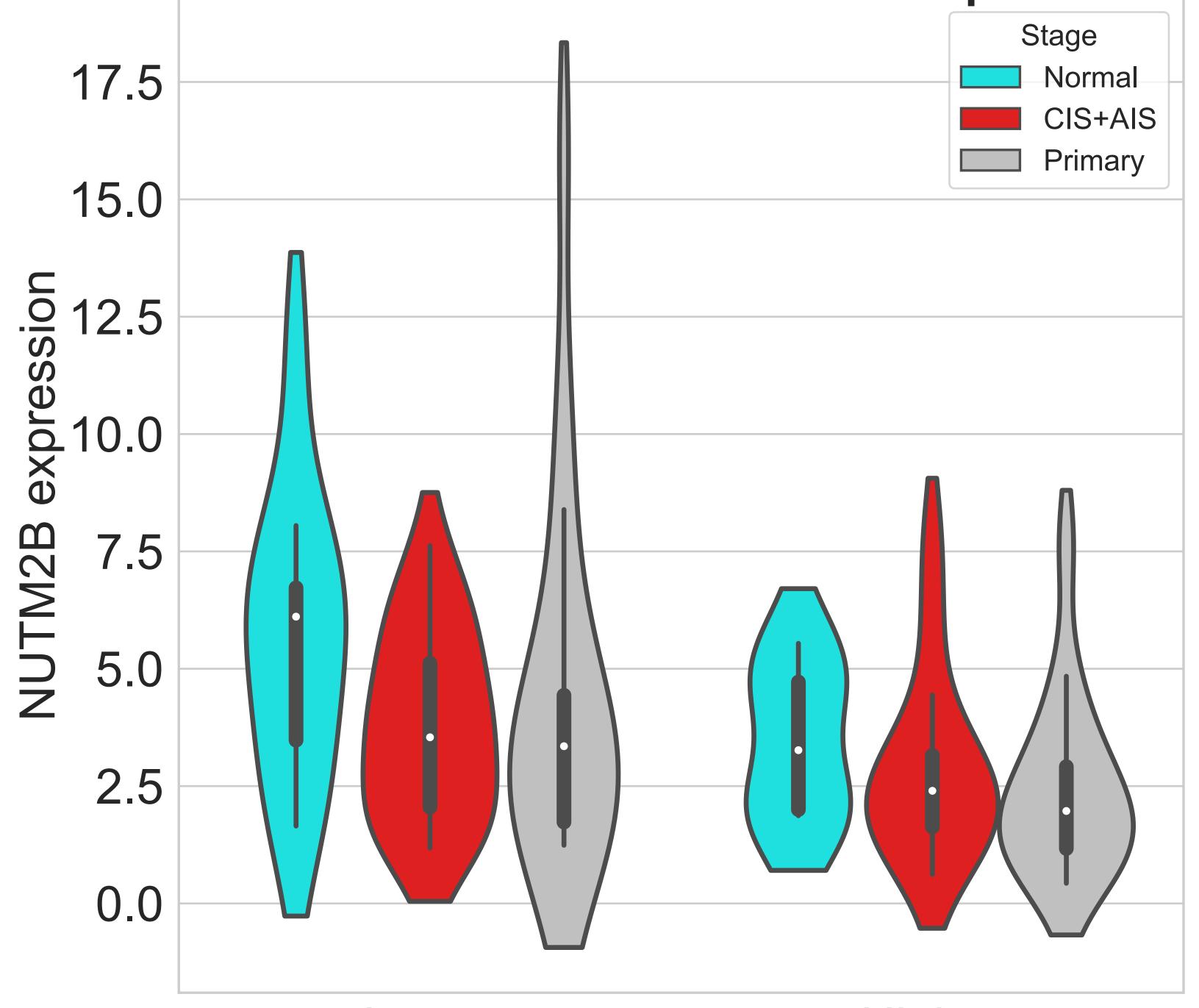
NUTM2B: Kruskal-Wallis p=0.031



Lower Higher Mutation Shared Count (SYN) per TMB