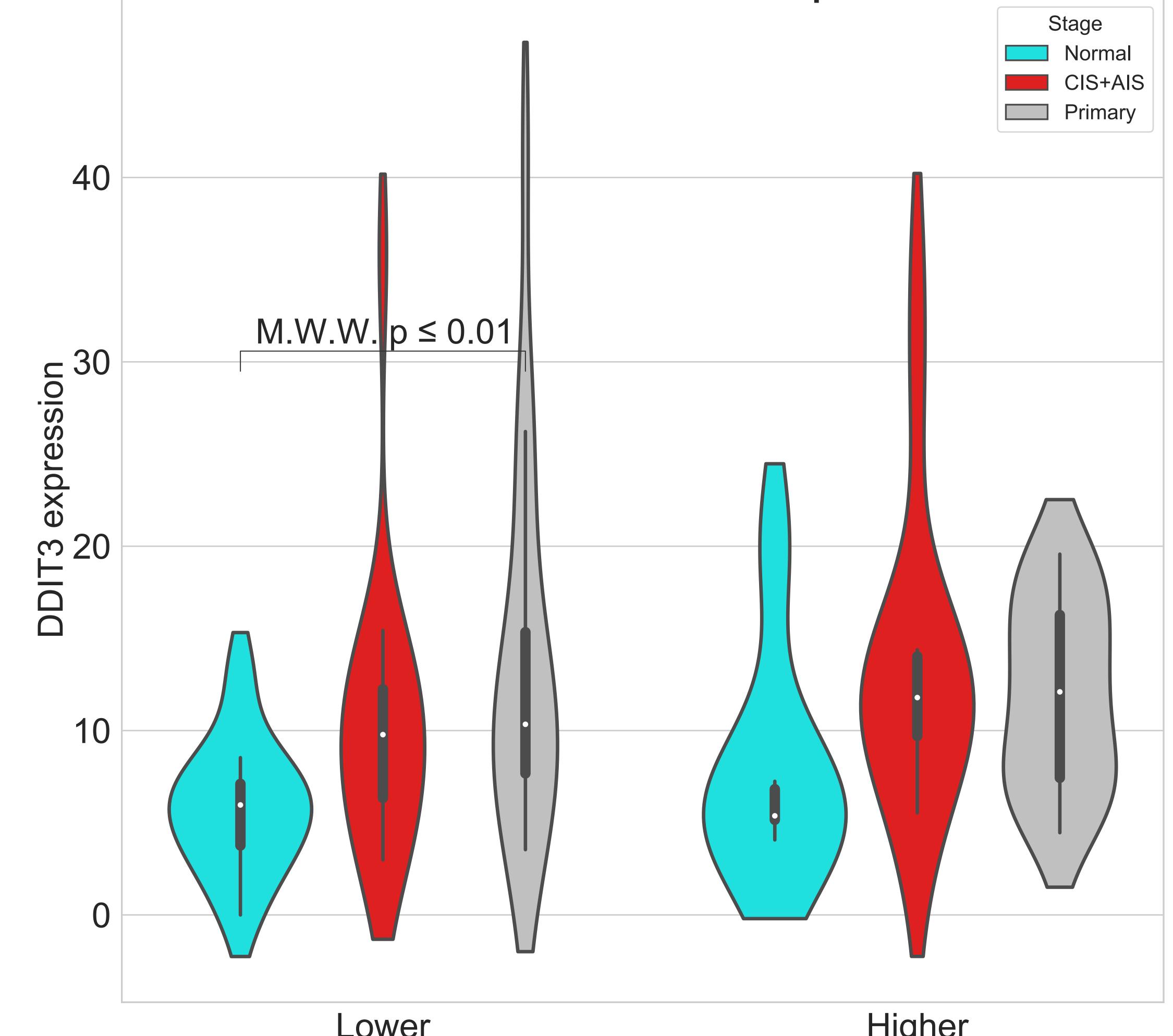
DDIT3: Kruskal-Wallis p=0.014



Lower Mutation Shared Count (SYN) per TMB