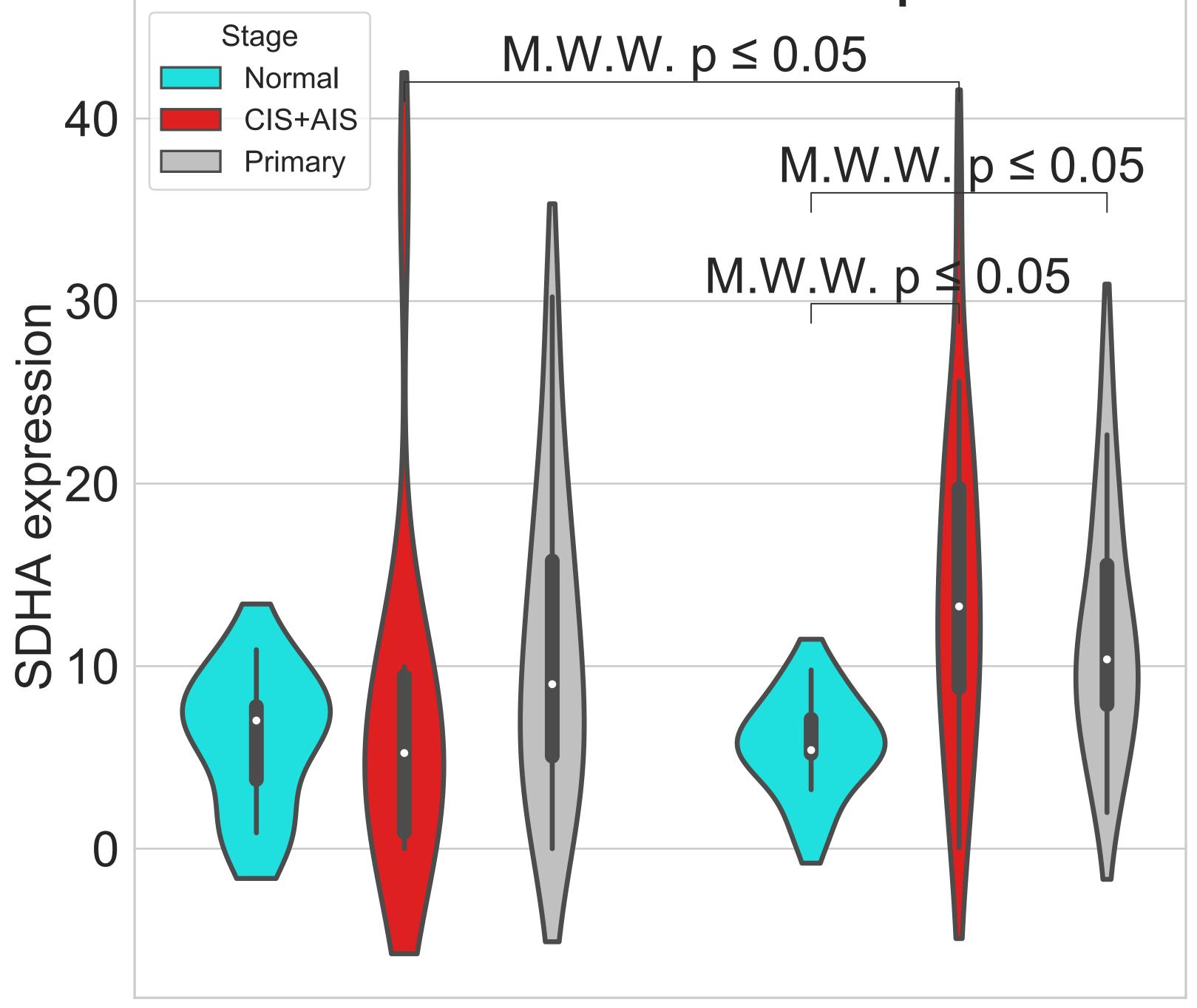
SDHA: Kruskal-Wallis p=0.008



Lower
Mutation Shared Count (SYN) per TMB