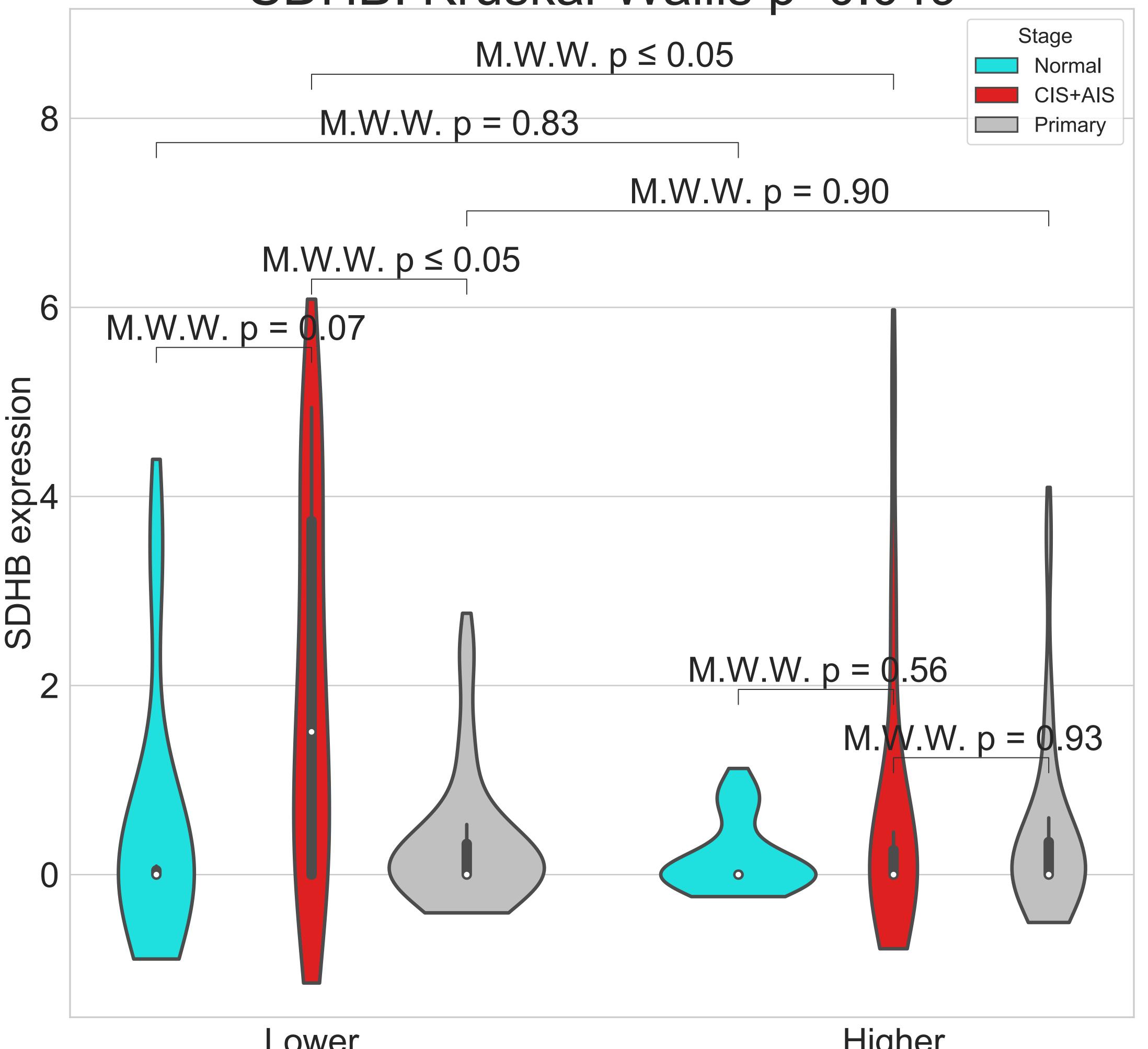
SDHB: Kruskal-Wallis p=0.049



Lower
Mutation Shared Count (SYN) per TMB