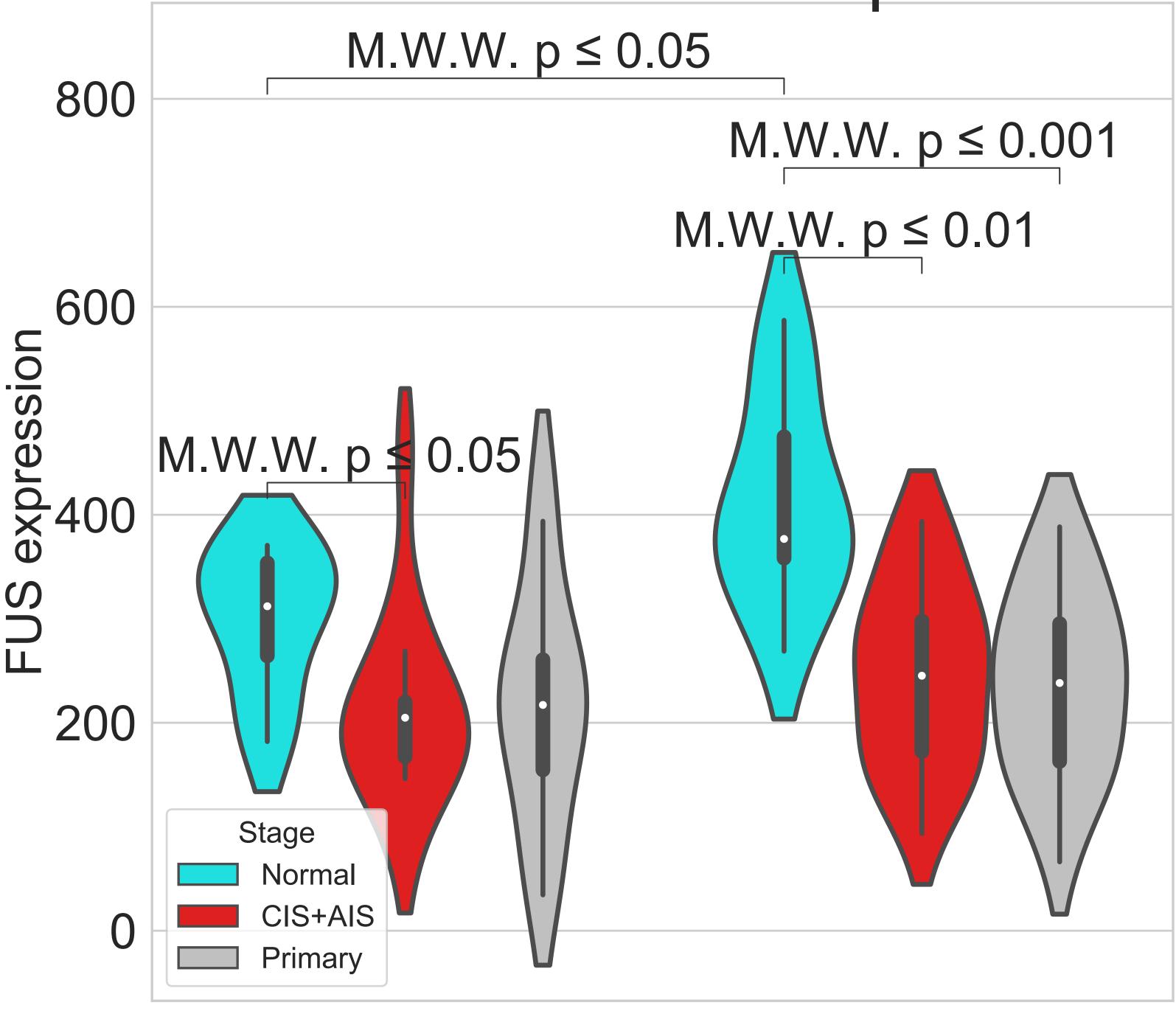
FUS: Kruskal-Wallis p=0.001



Lower Higher Mutation Shared Count (SYN) per TMB