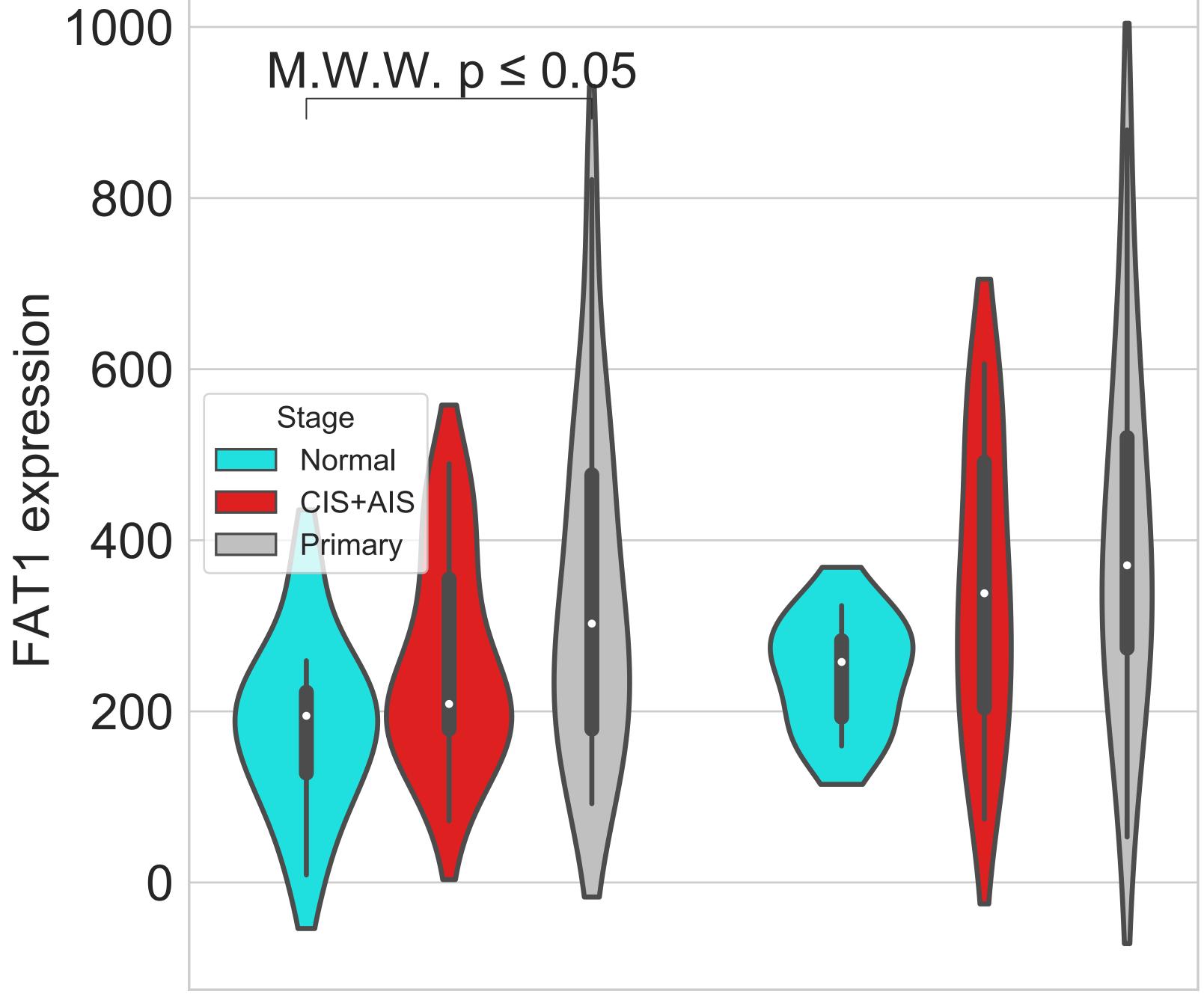
FAT1: Kruskal-Wallis p=0.034 M.W.W. p ≤ 0.05



Lower Higher Mutation Shared Count (SYN) per TMB