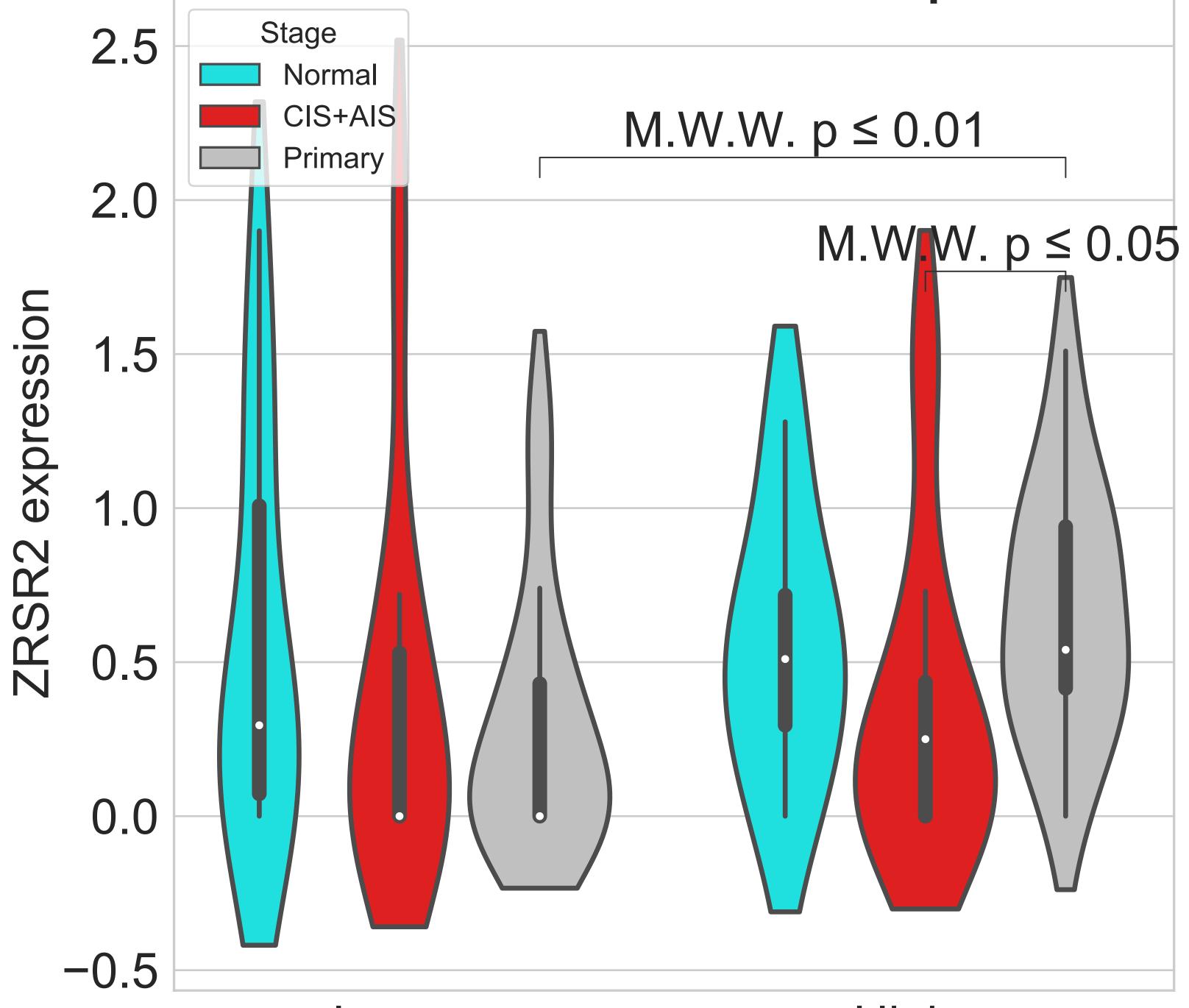
ZRSR2: Kruskal-Wallis p=0.048



Lower Higher Mutation Shared Count (SYN) per TMB