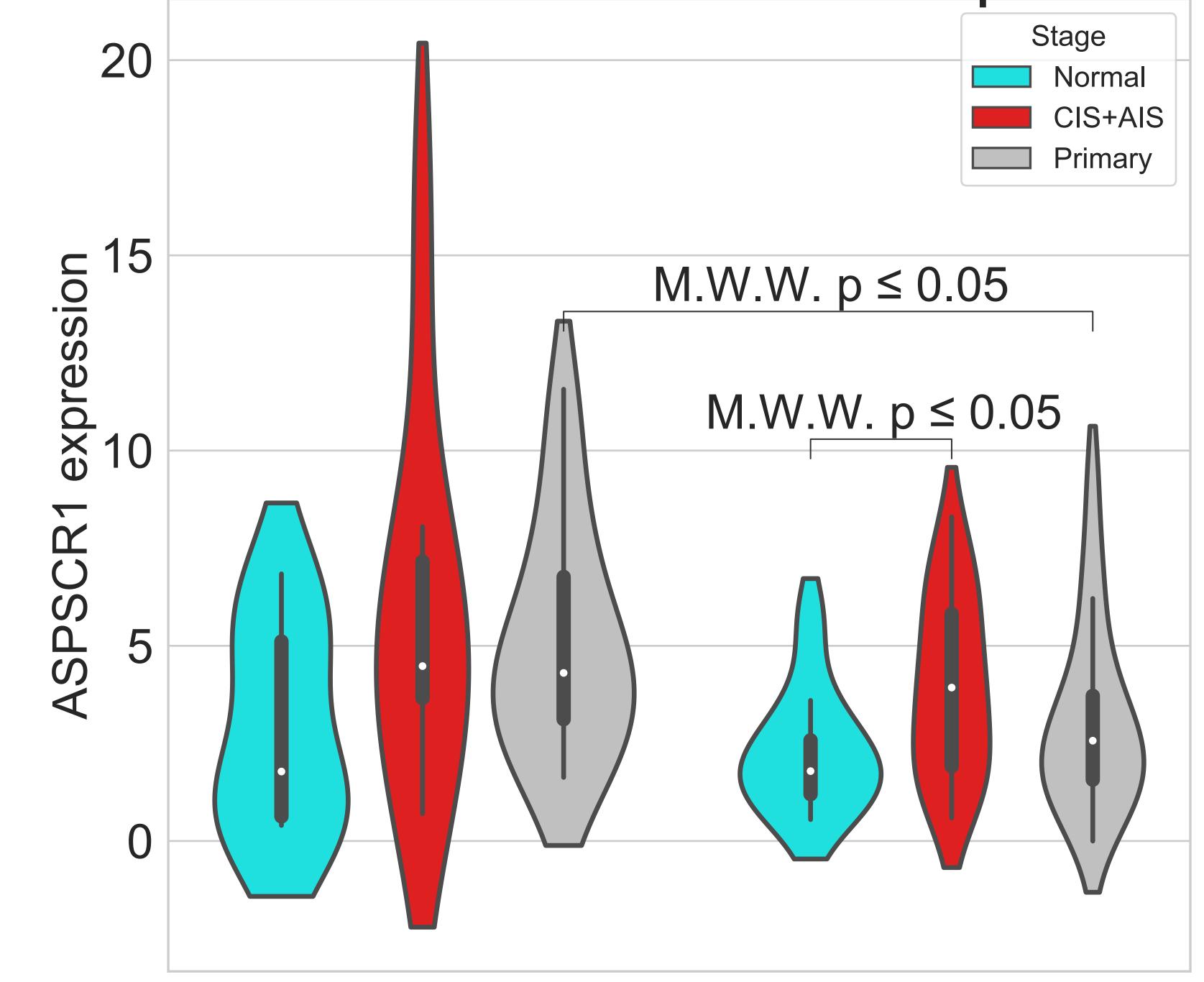
ASPSCR1: Kruskal-Wallis p=0.048



Lower
Mutation Shared Count (SYN) per TMB