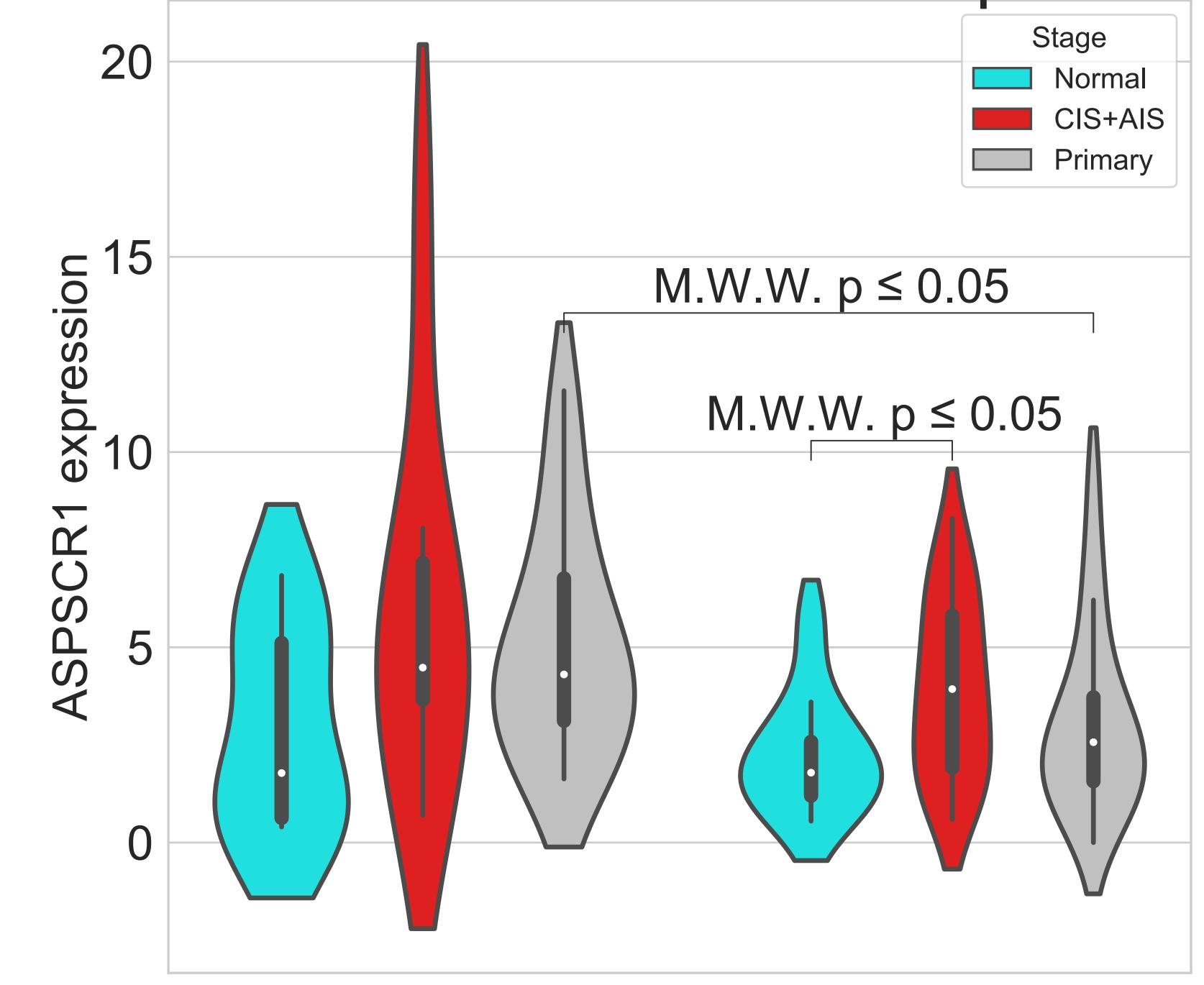
ASPSCR1: Kruskal-Wallis p=0.048



Lower Higher Mutation Shared Proportion (SYN)