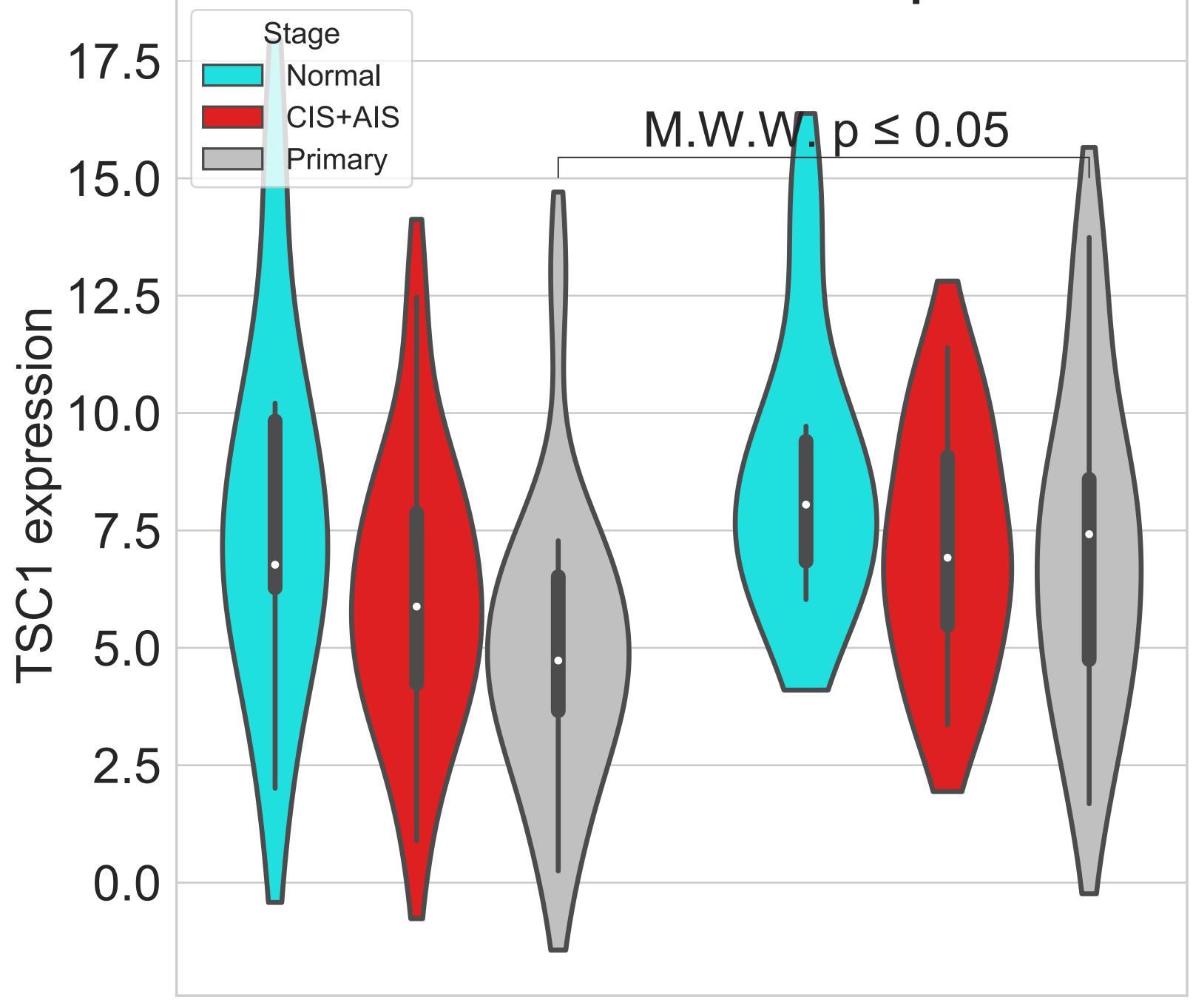
TSC1: Kruskal-Wallis p=0.048



Lower
Mutation Shared Count (SYN)