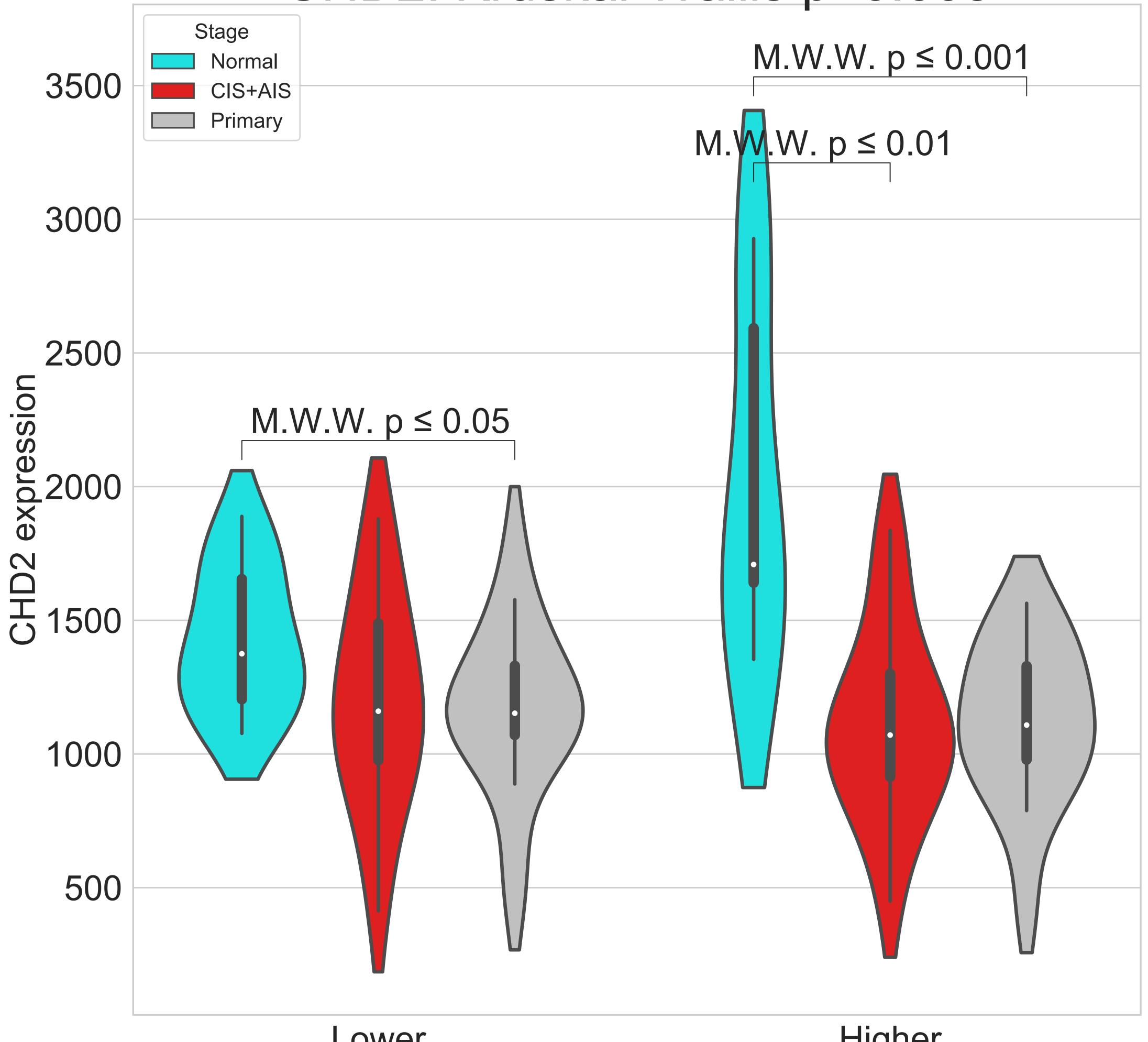
CHD2: Kruskal-Wallis p=0.003



Lower Mutation Shared Count (SYN) per TMB