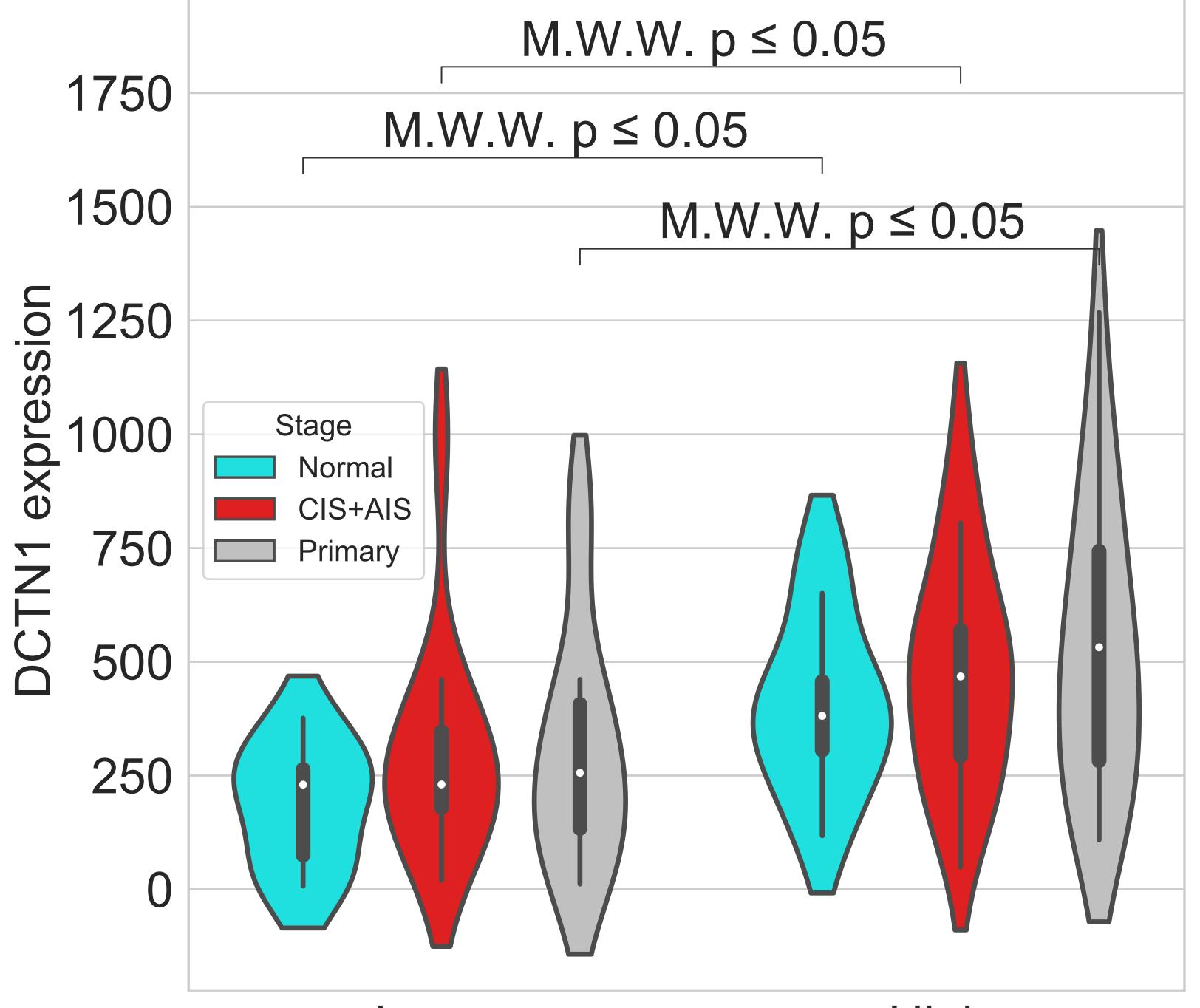
DCTN1: Kruskal-Wallis p=0.010



Lower Higher Mutation Shared Count (SYN) per TMB