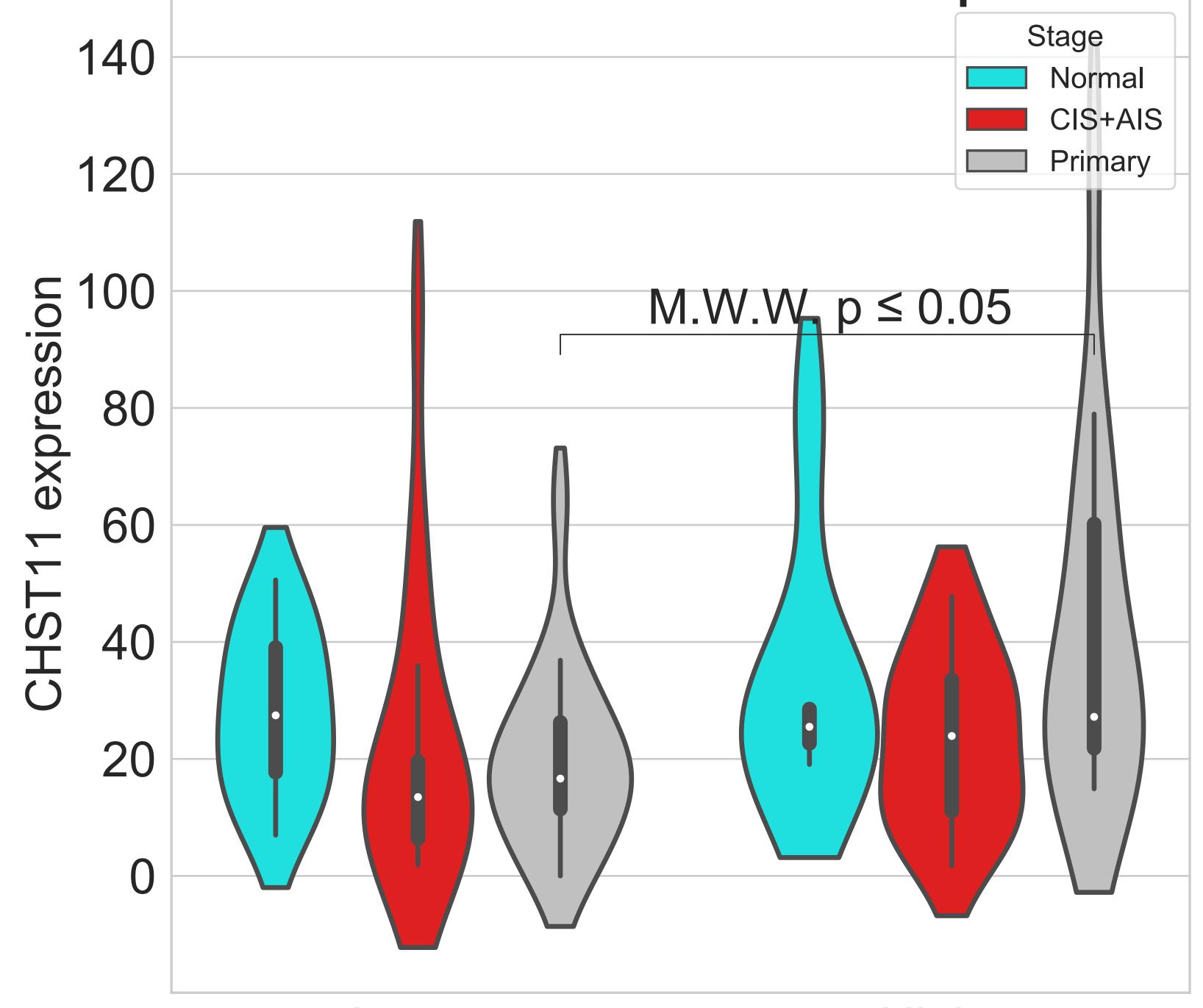
CHST11: Kruskal-Wallis p=0.043



Lower Higher Mutation Shared Proportion (Union & SYN)