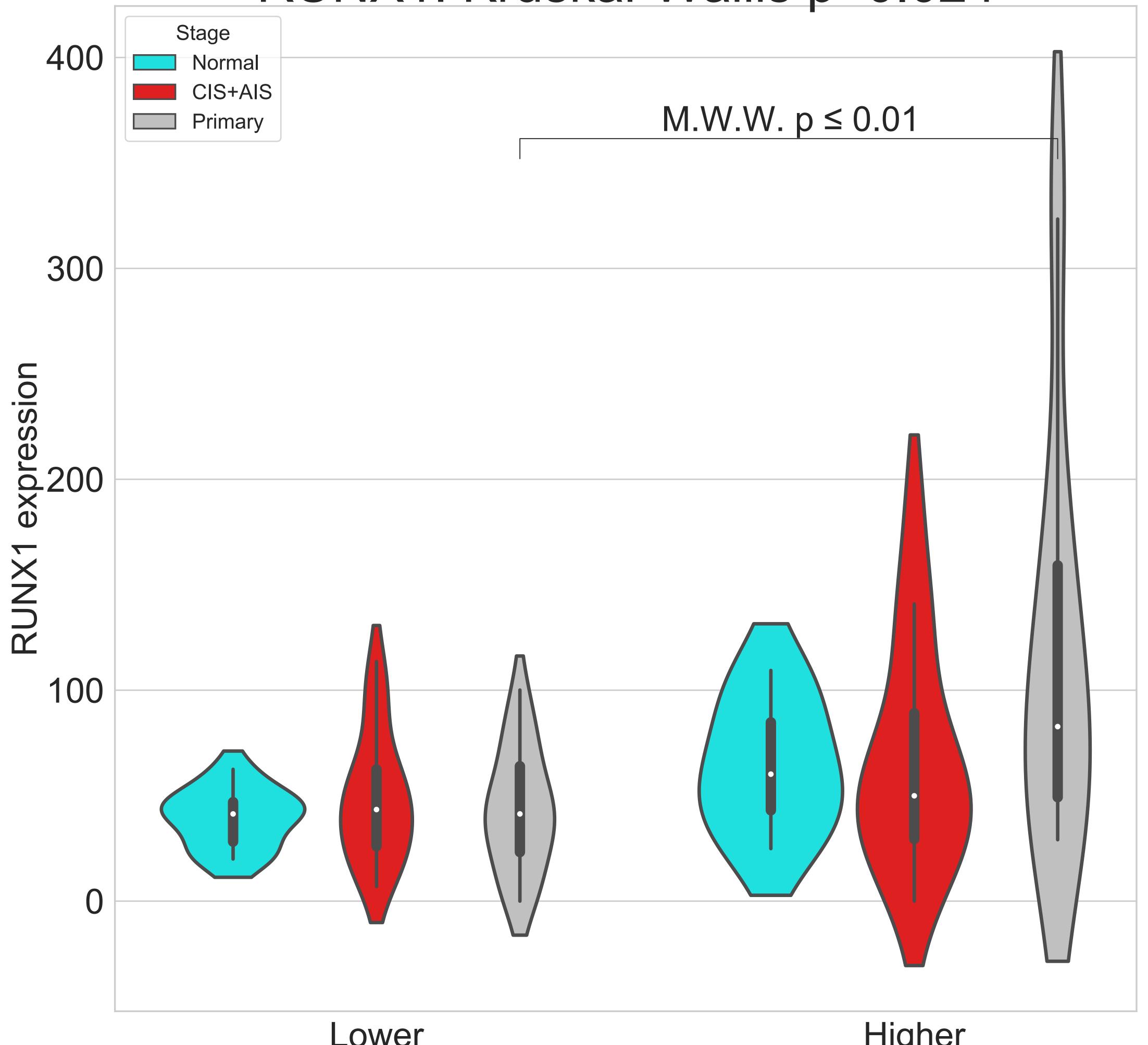
RUNX1: Kruskal-Wallis p=0.024



Lower
Mutation Shared Count (SYN) per TMB