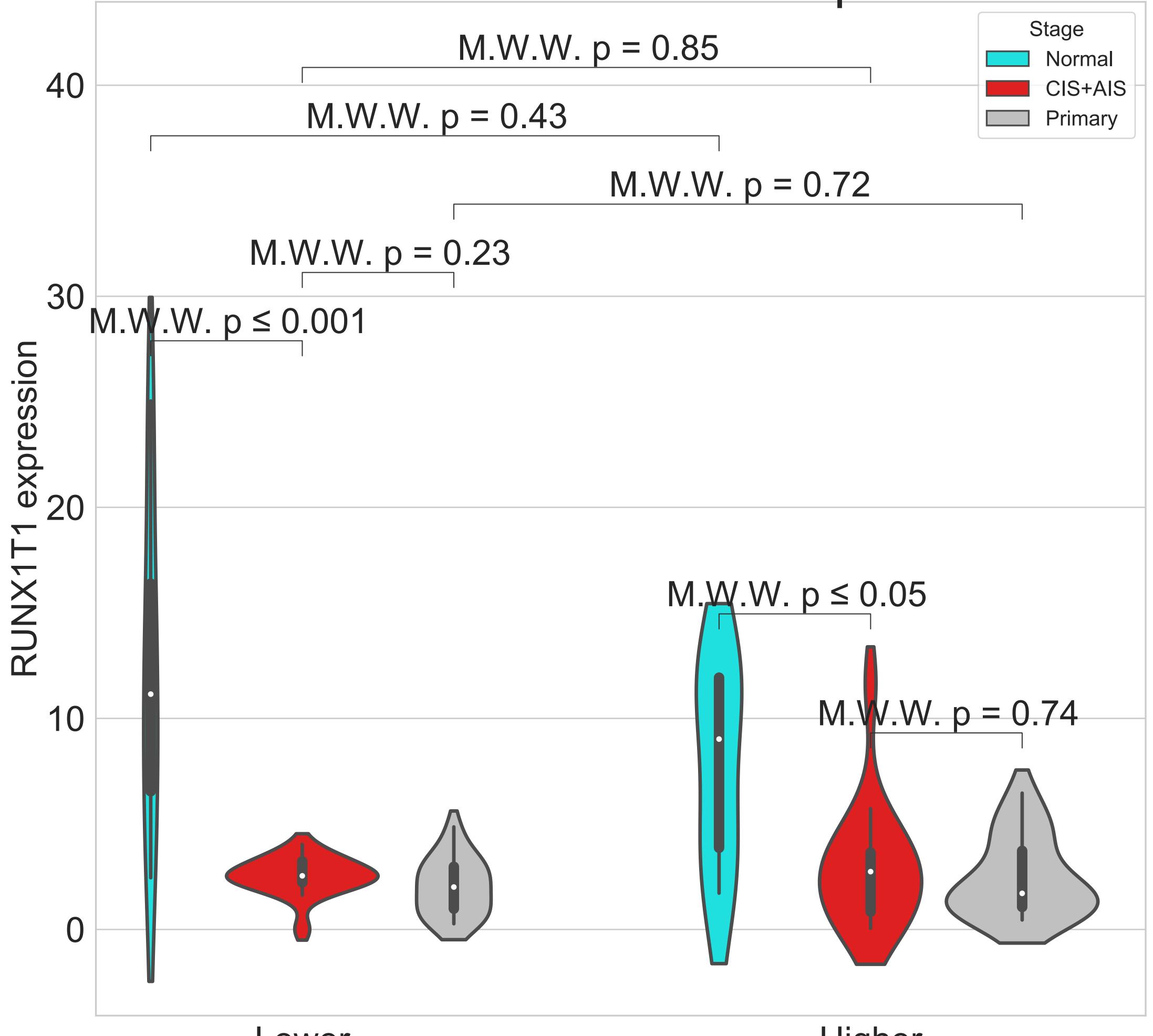
RUNX1T1: Kruskal-Wallis p=0.000 M.W.W. p = 0.85



Higher Lower Mutation Shared Count (SYN) per TMB