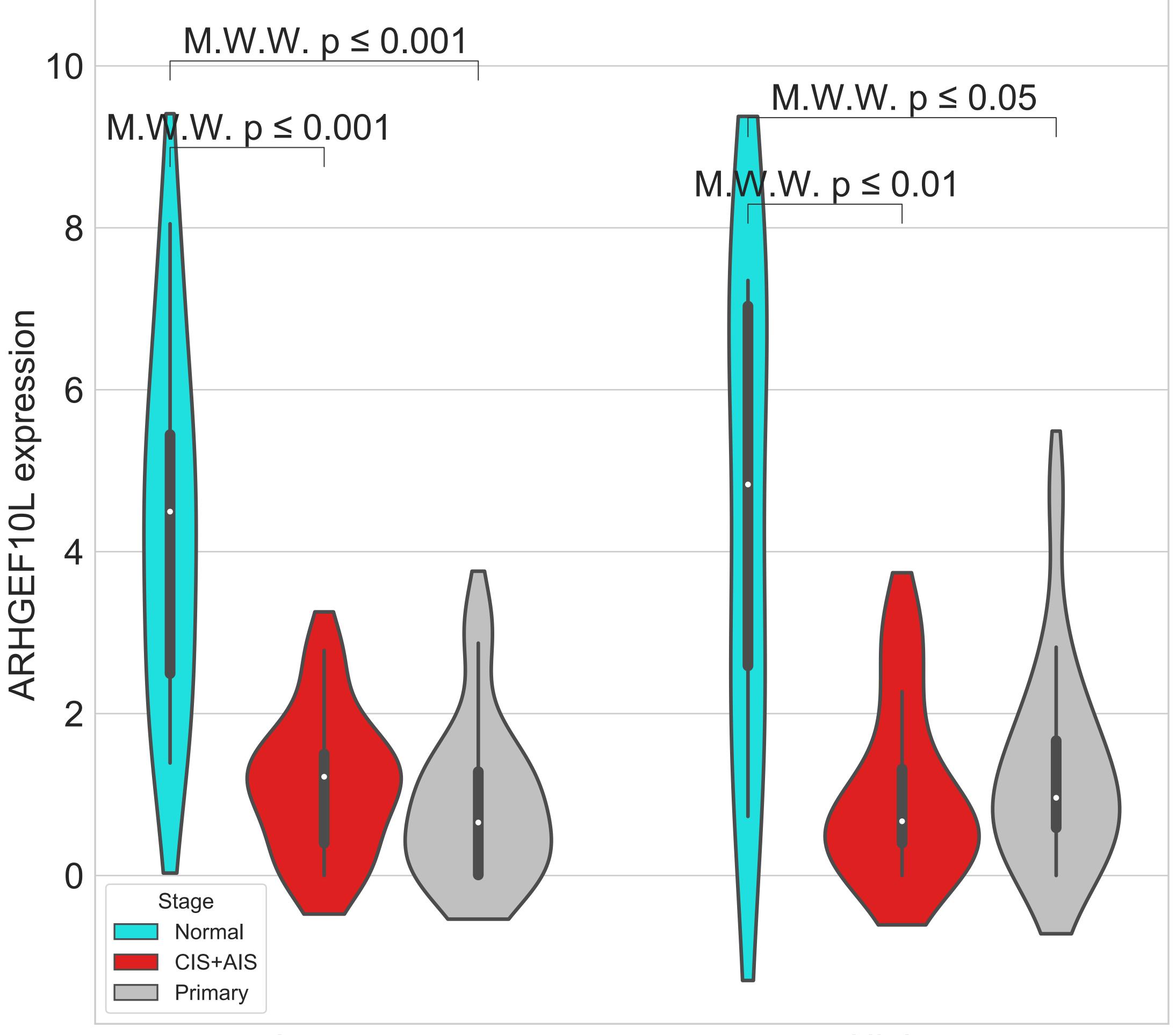
ARHGEF10L: Kruskal-Wallis p=0.000

M.W.W. p ≤ 0.001



Lower
Mutation Shared Count (SYN) per TMB