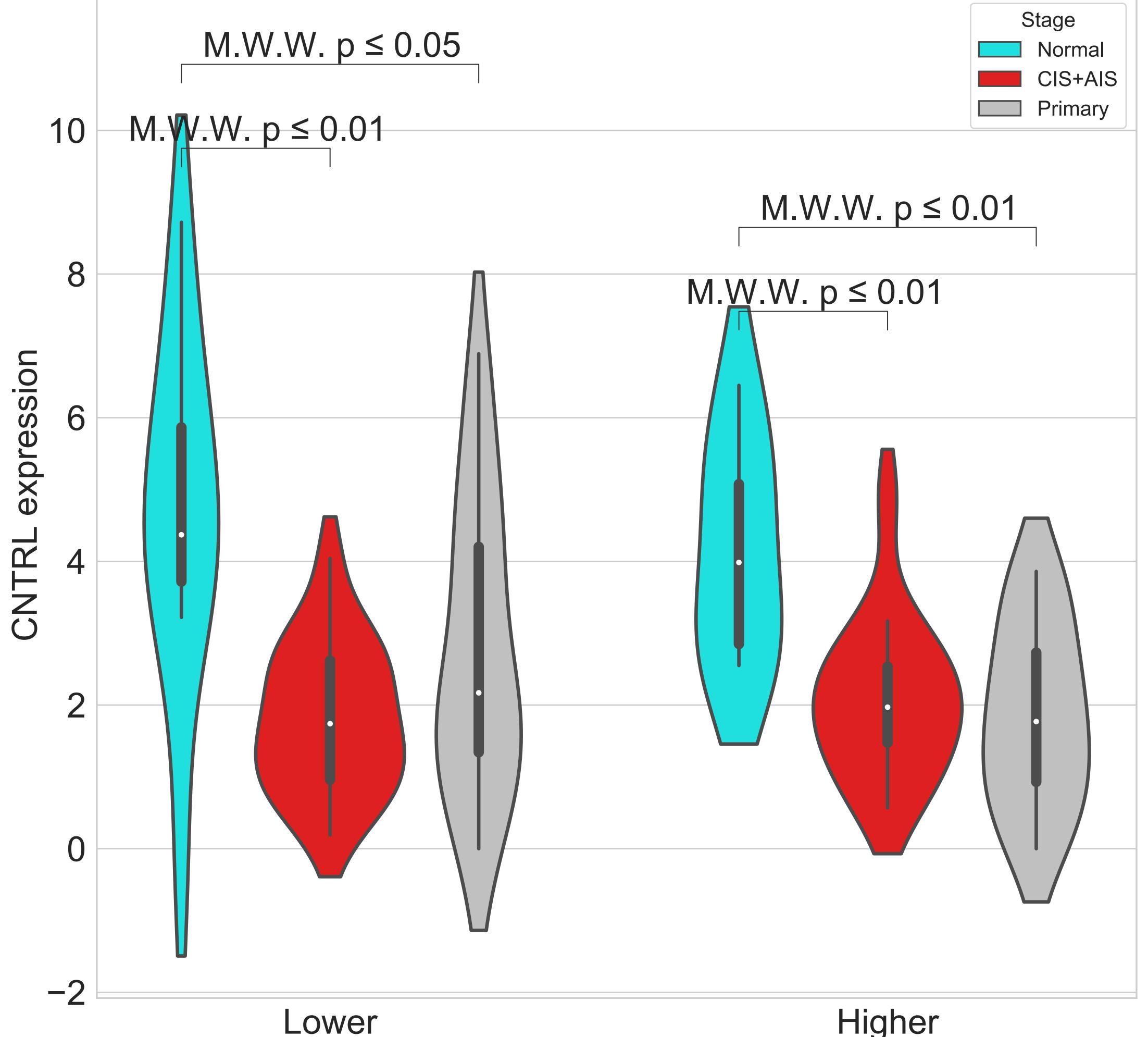
CNTRL: Kruskal-Wallis p=0.001
.W.W. p ≤ 0.05



Lower
Mutation Shared Count (SYN) per TMB