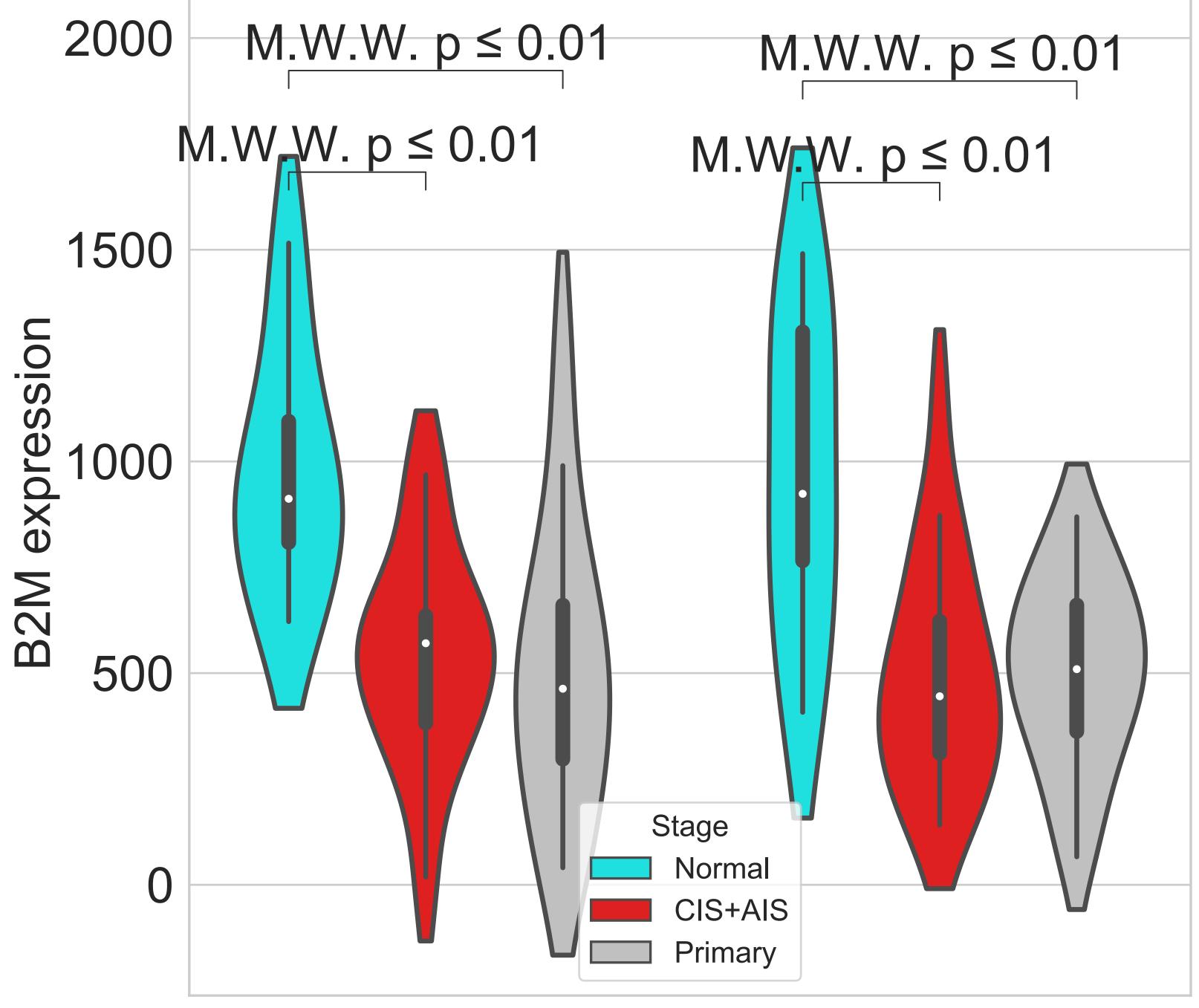
B2M: Kruskal-Wallis p=0.001

M.W. p ≤ 0.01

M.W. w p < 0.01



Lower Higher Mutation Shared Count (SYN) per TMB