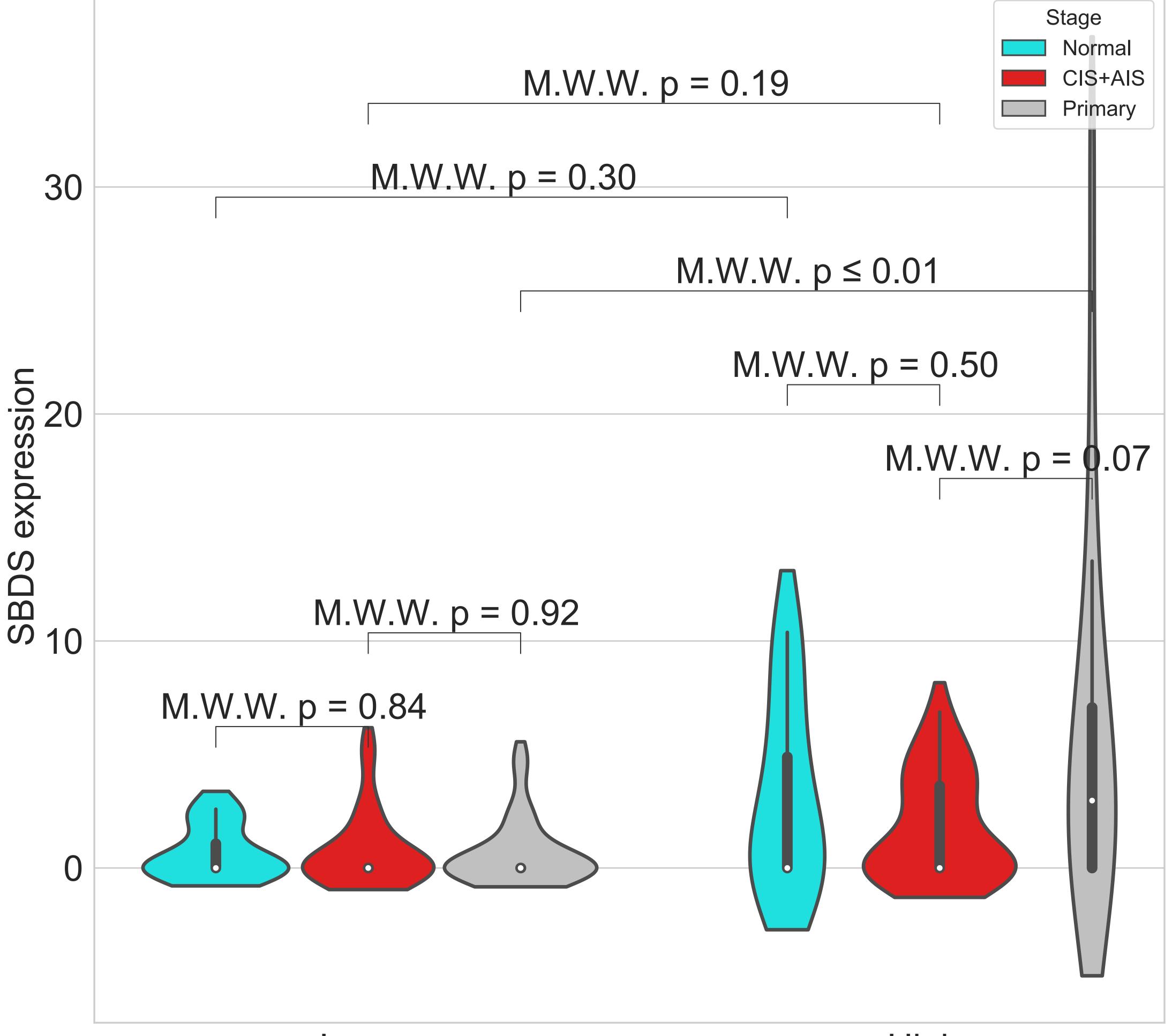
SBDS: Kruskal-Wallis p=0.020 M.W.W. p = 0.19



Higher Lower Mutation Shared Count (SYN) per TMB