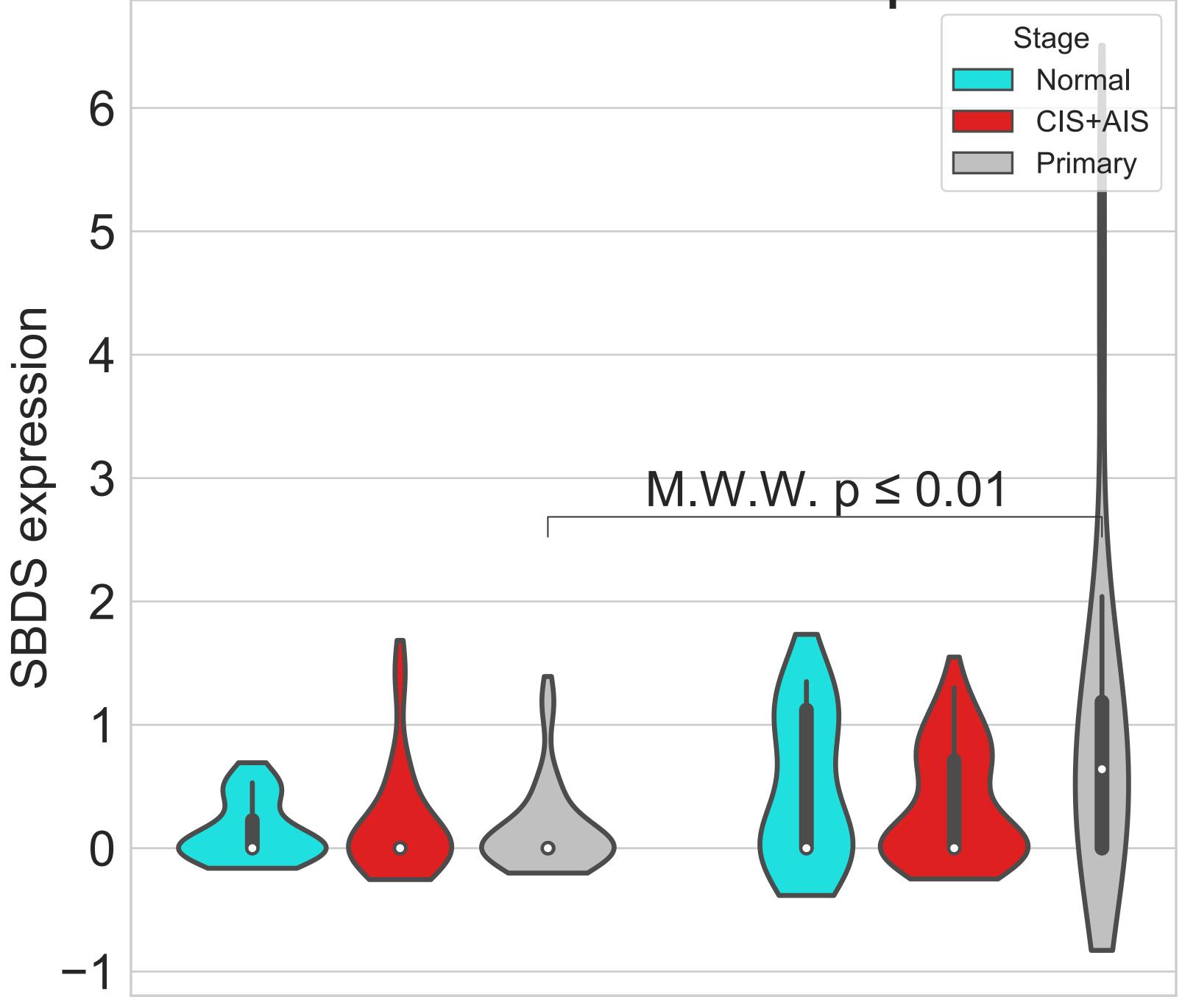
SBDS: Kruskal-Wallis p=0.038



Lower
Mutation Shared Count (SYN) per TMB