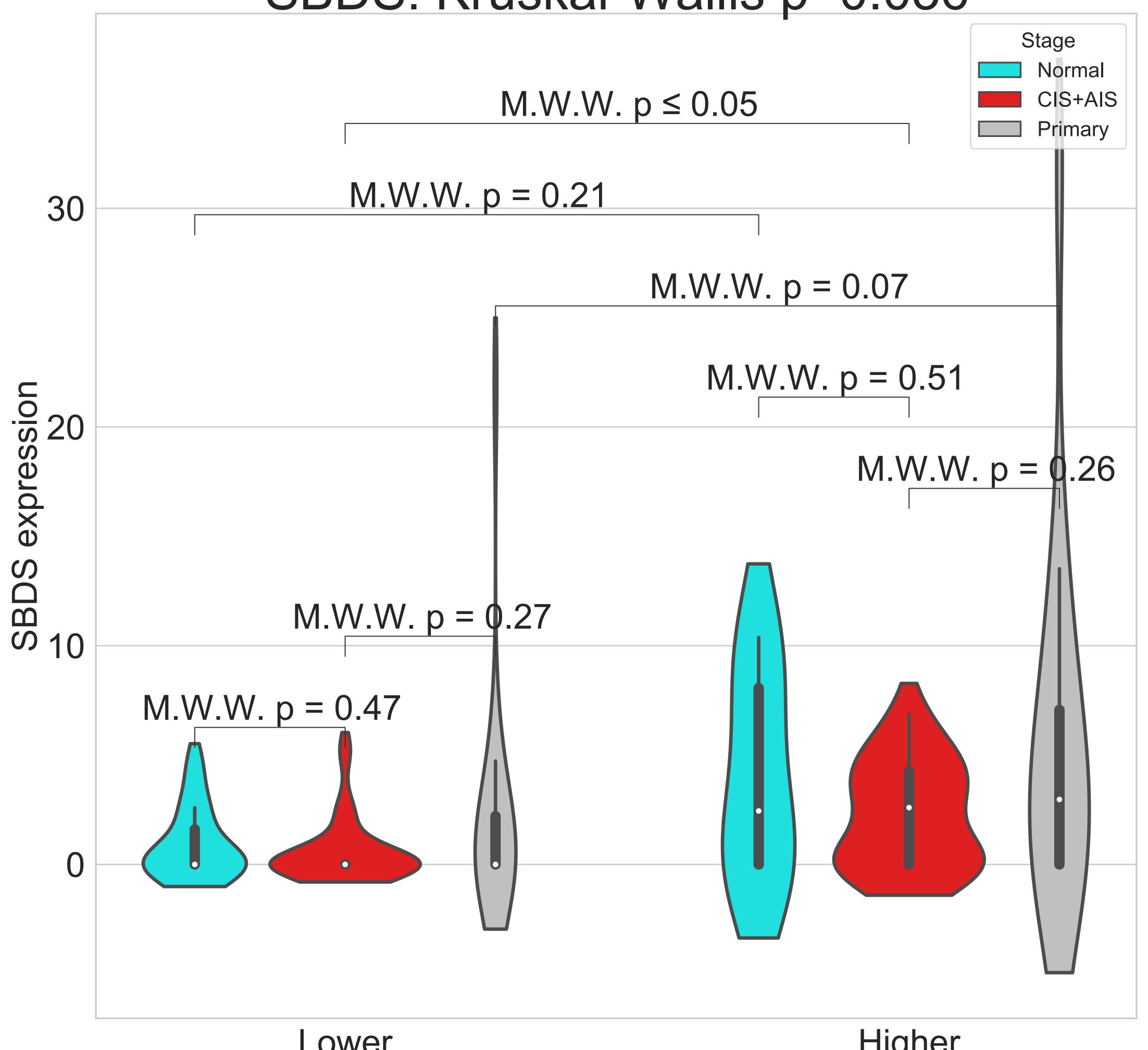
SBDS: Kruskal-Wallis p=0.036



Lower
Mutation Shared Count (SYN) per TMB