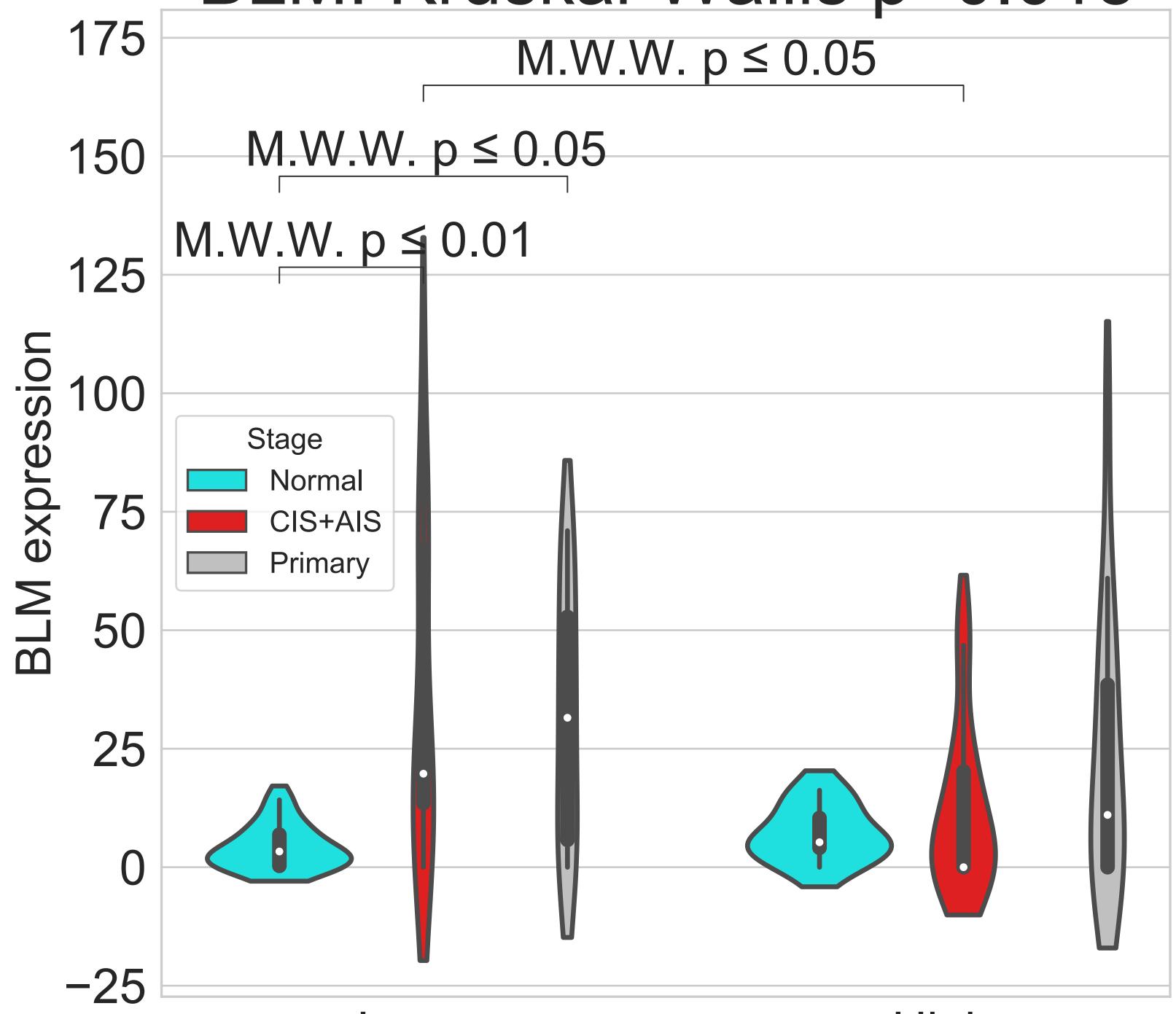
BLM: Kruskal-Wallis p=0.018



Lower
Mutation Shared Count (SYN) per TMB