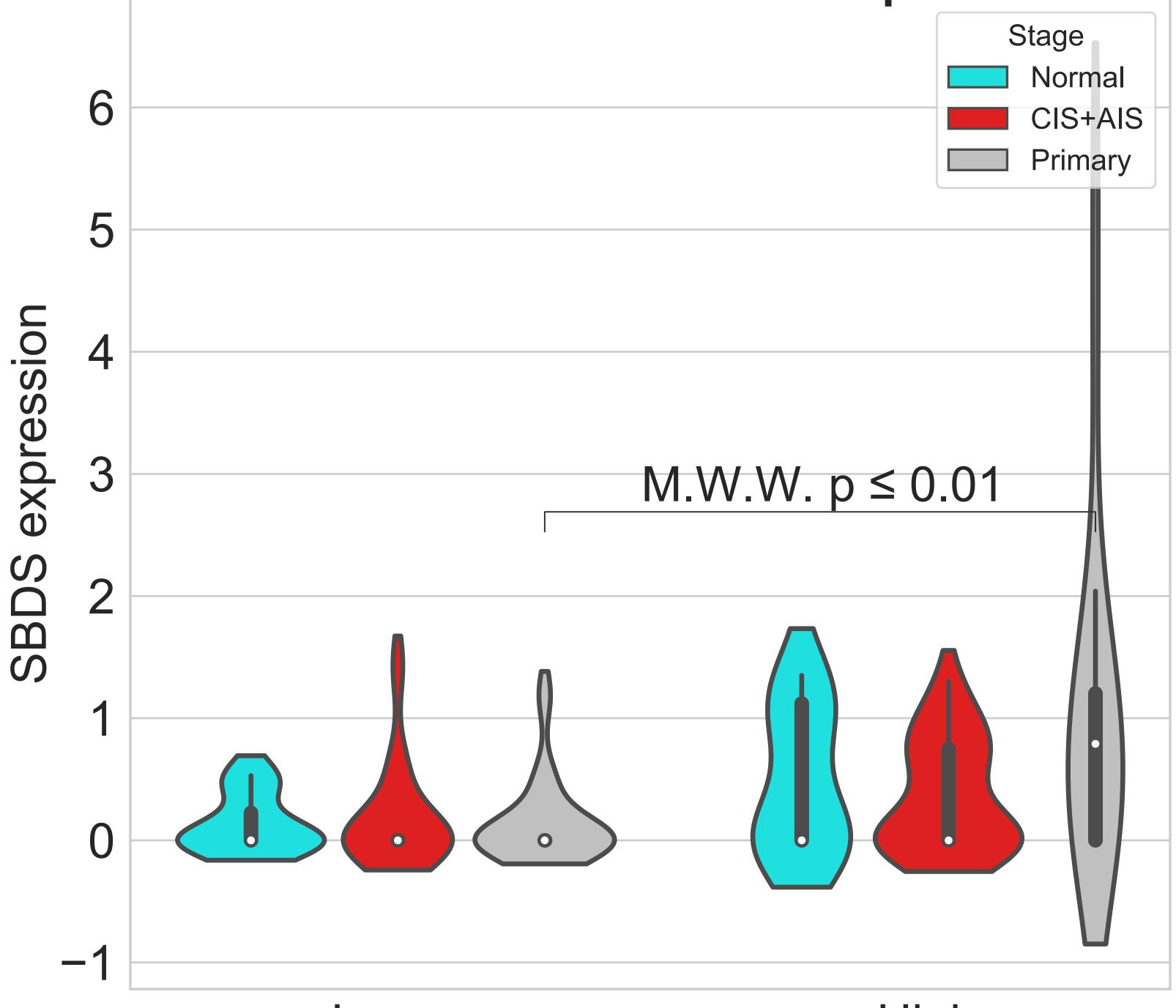
SBDS: Kruskal-Wallis p=0.015



Lower
Mutation Shared Count per TMB