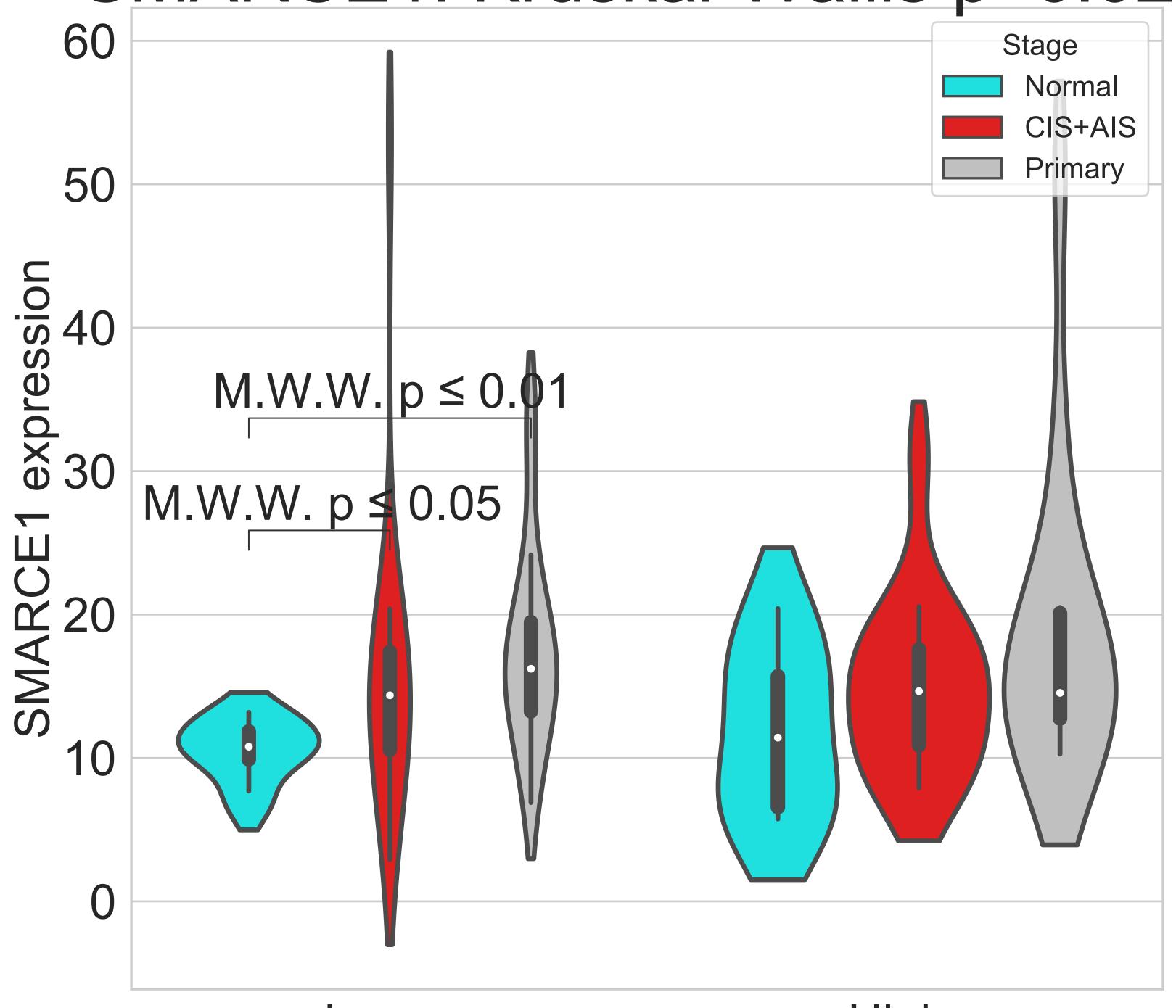
SMARCE1: Kruskal-Wallis p=0.026



Lower
Mutation Shared Count per TMB