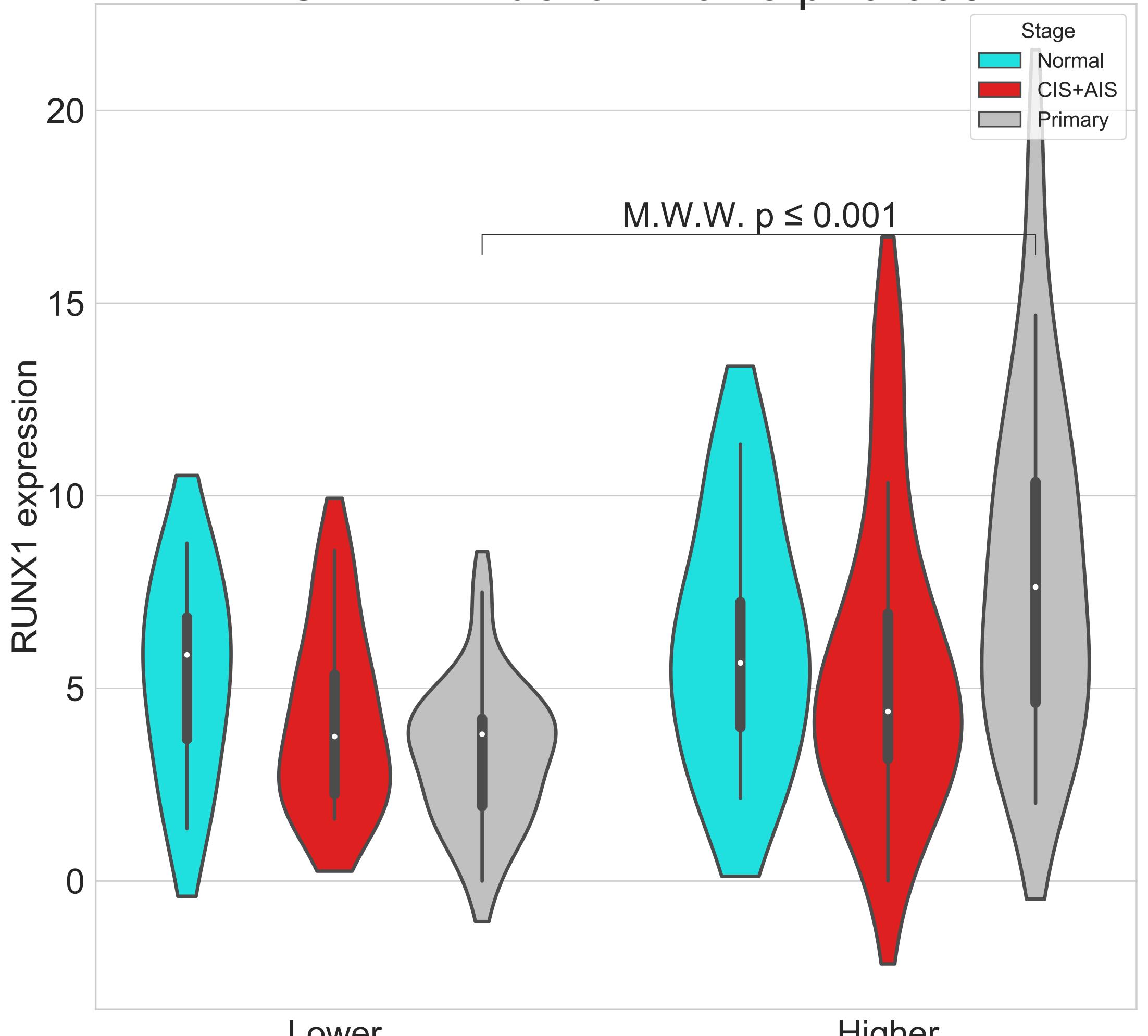
RUNX1: Kruskal-Wallis p=0.003



Lower
Mutation Shared Count per TMB