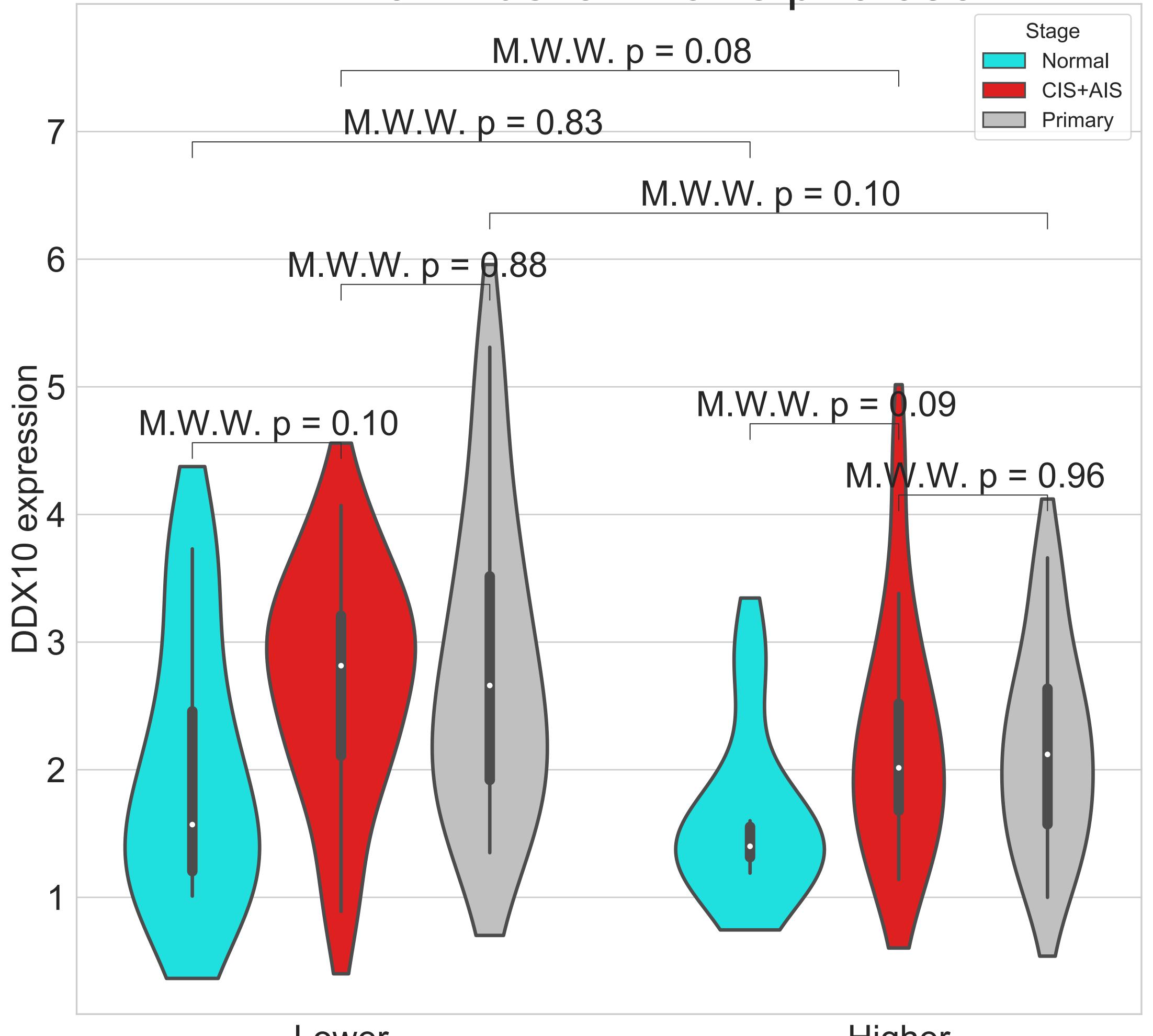
DDX10: Kruskal-Wallis p=0.036



Lower
Mutation Shared Count per TMB