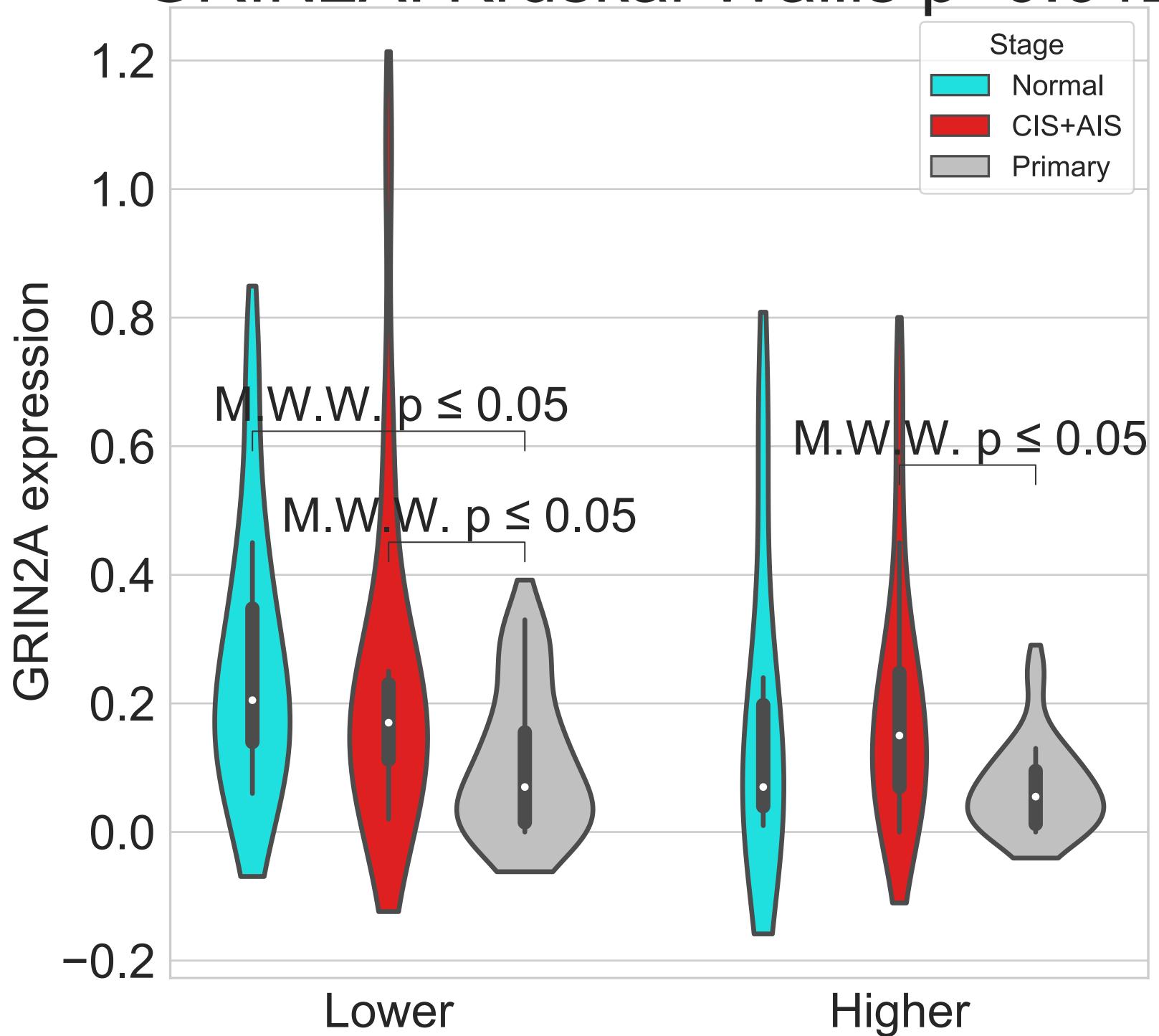
GRIN2A: Kruskal-Wallis p=0.012



Mutation Shared Count per TMB