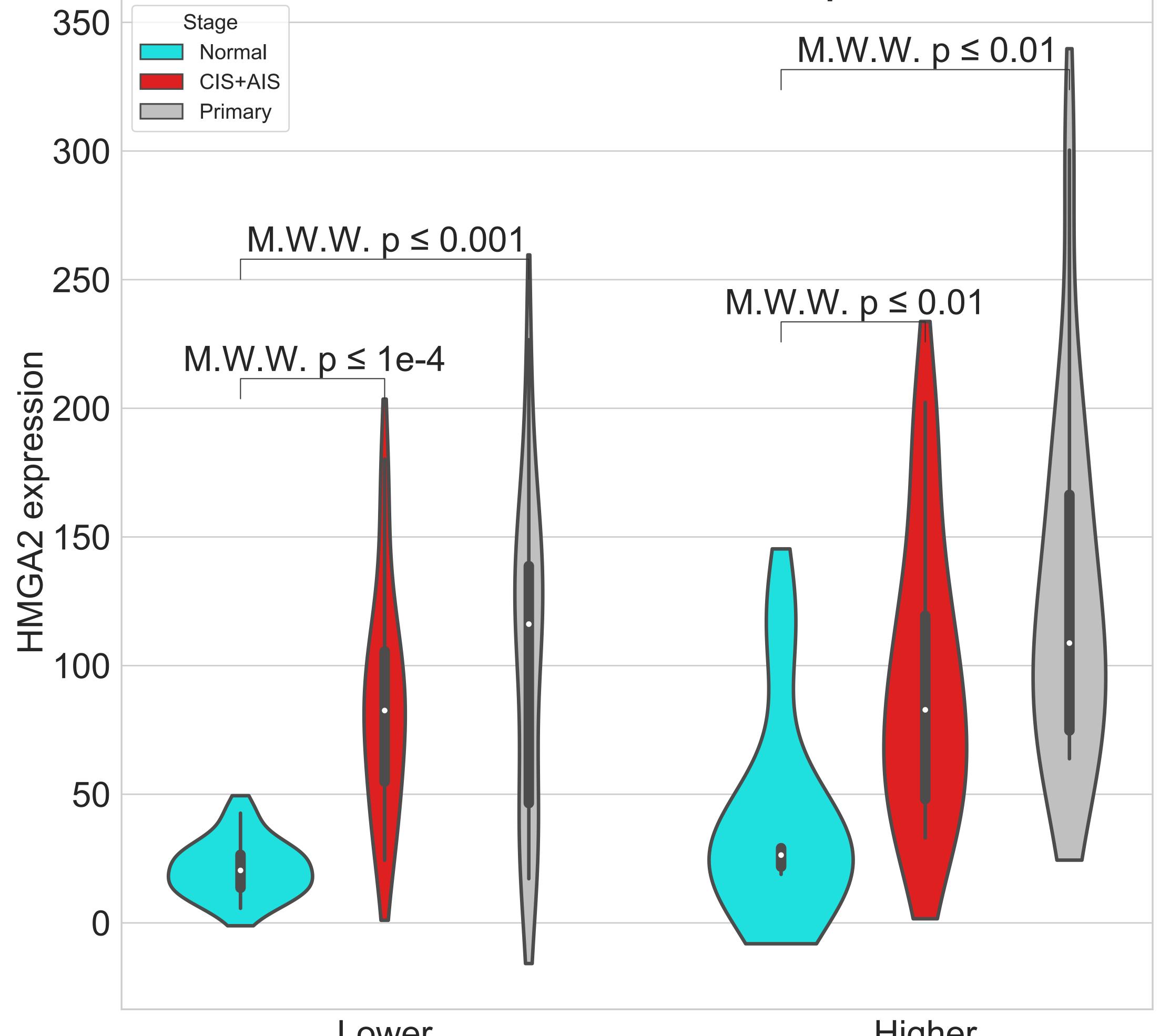
HMGA2: Kruskal-Wallis p=0.000



Higher Lower **Mutation Shared Proportion**