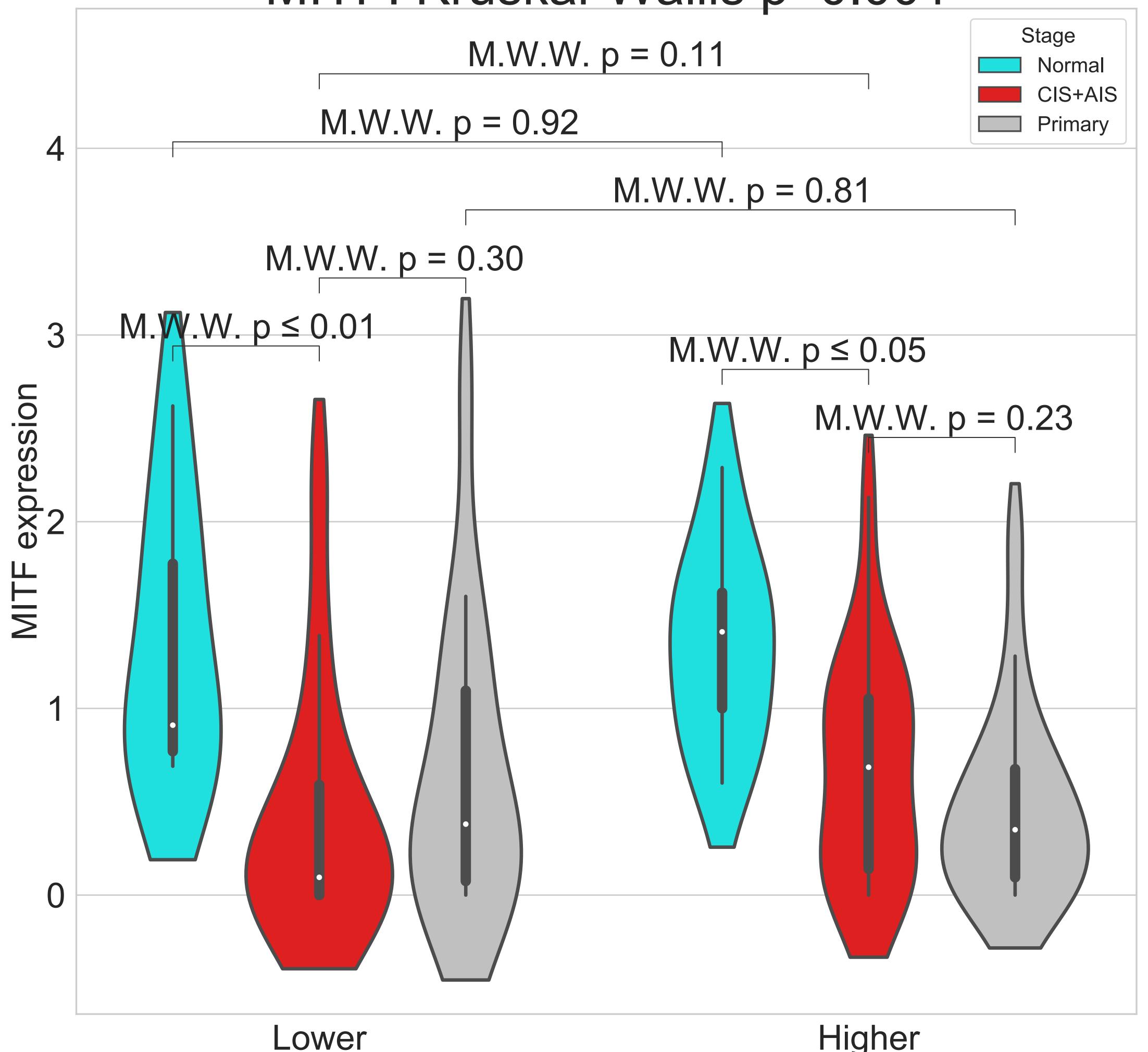
MITF: Kruskal-Wallis p=0.001



Higher Mutation Shared Proportion