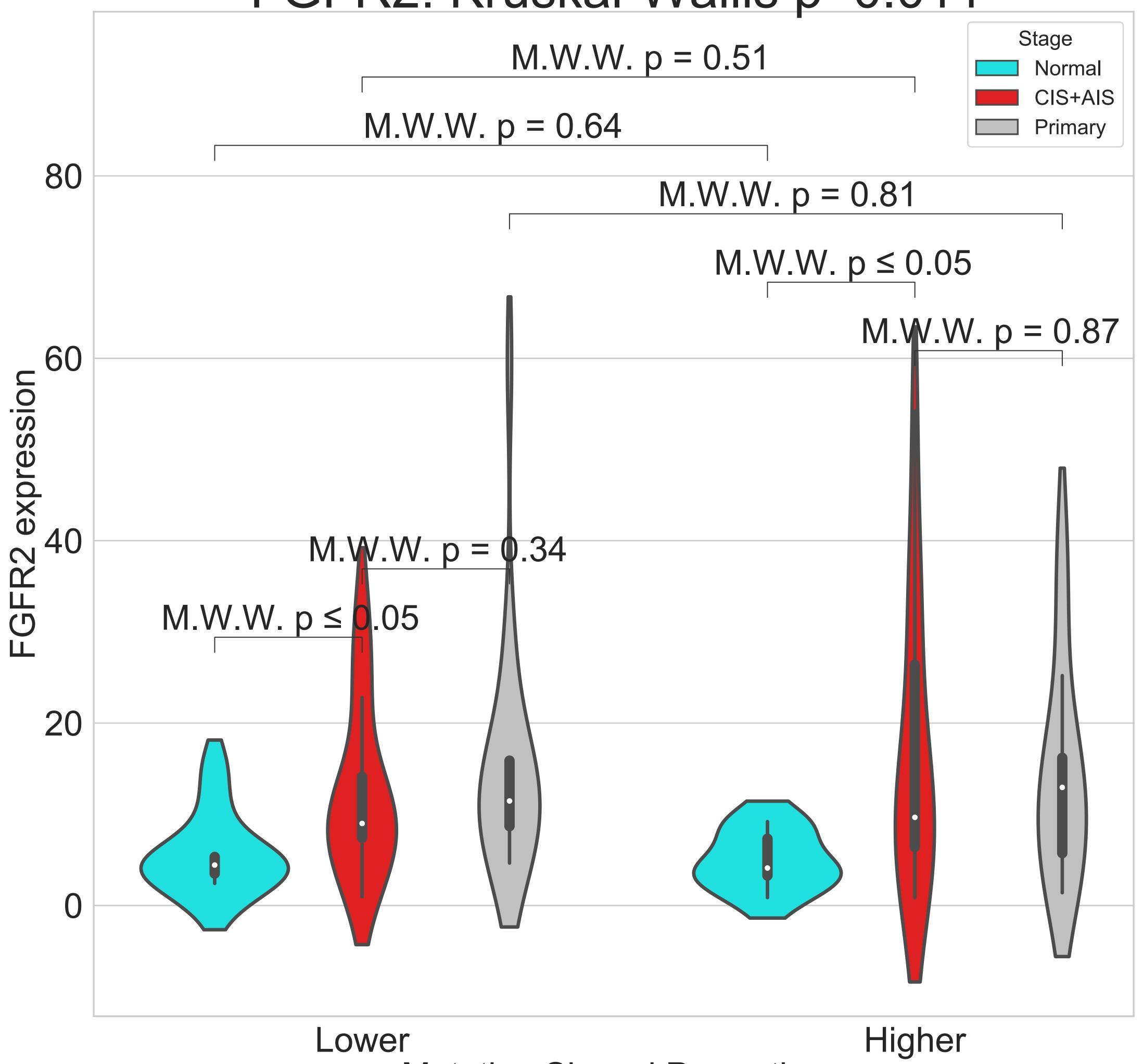
FGFR2: Kruskal-Wallis p=0.011



**Mutation Shared Proportion**