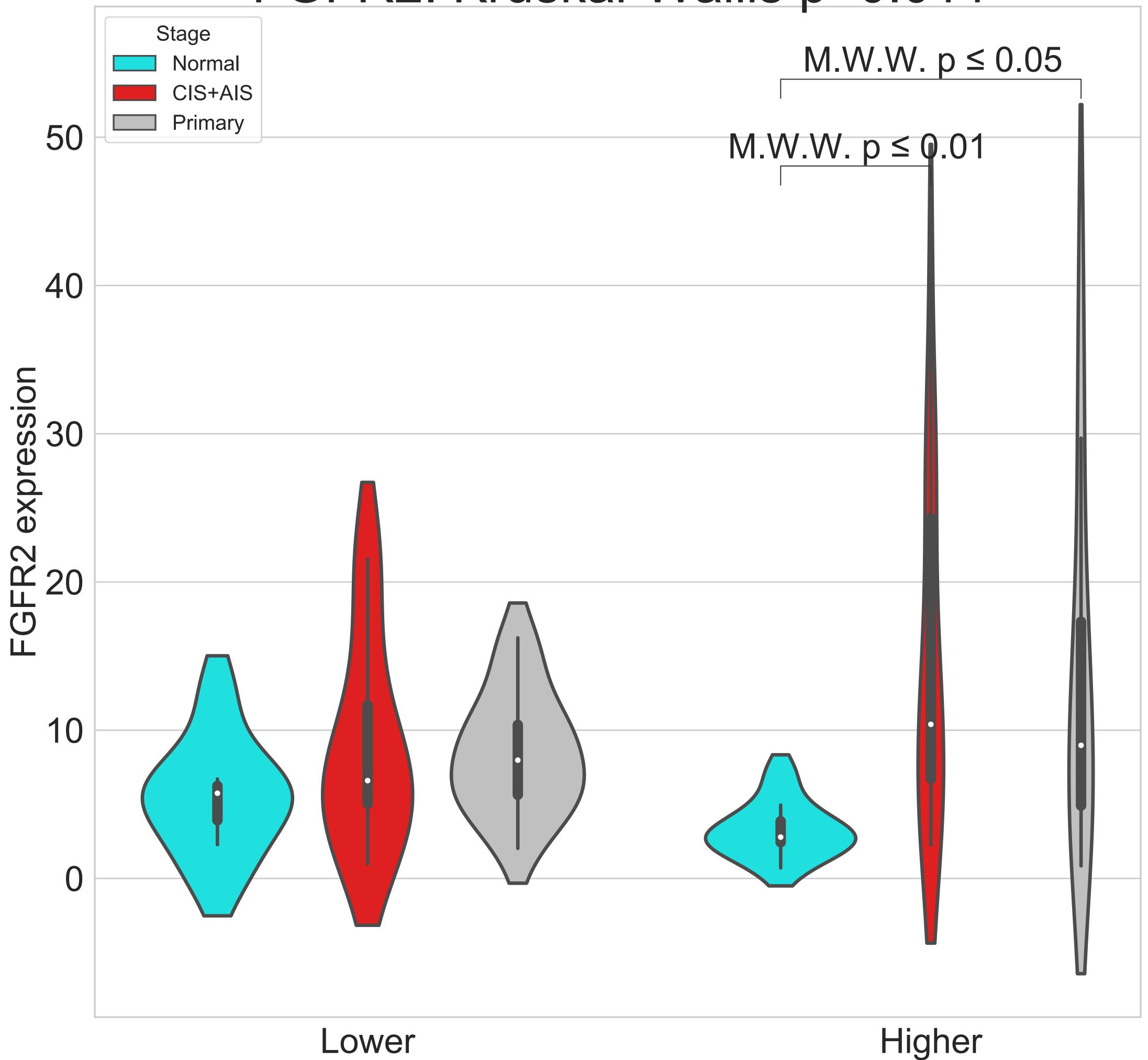
FGFR2: Kruskal-Wallis p=0.011



Mutation Shared Proportion