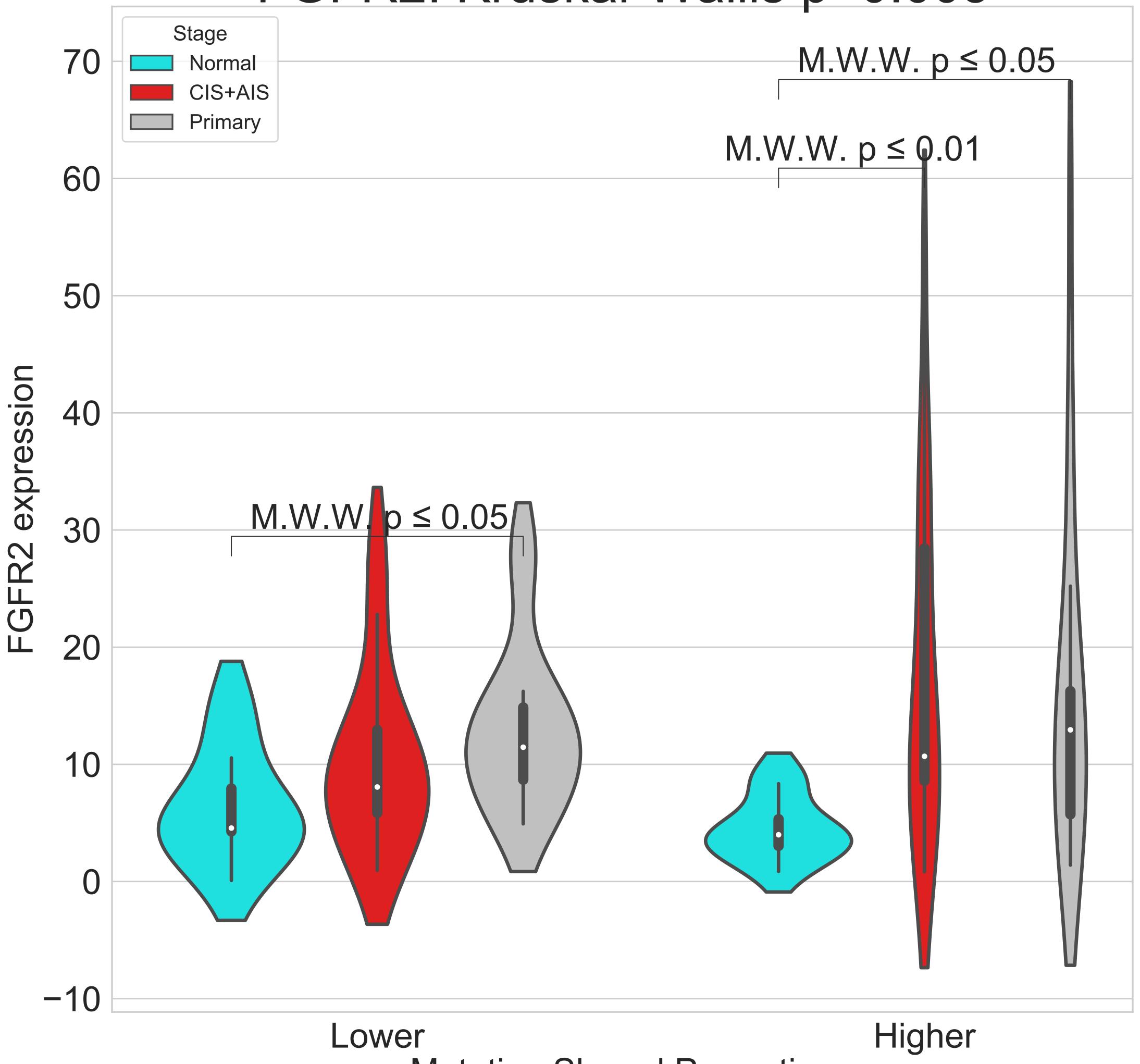
FGFR2: Kruskal-Wallis p=0.005



**Mutation Shared Proportion**