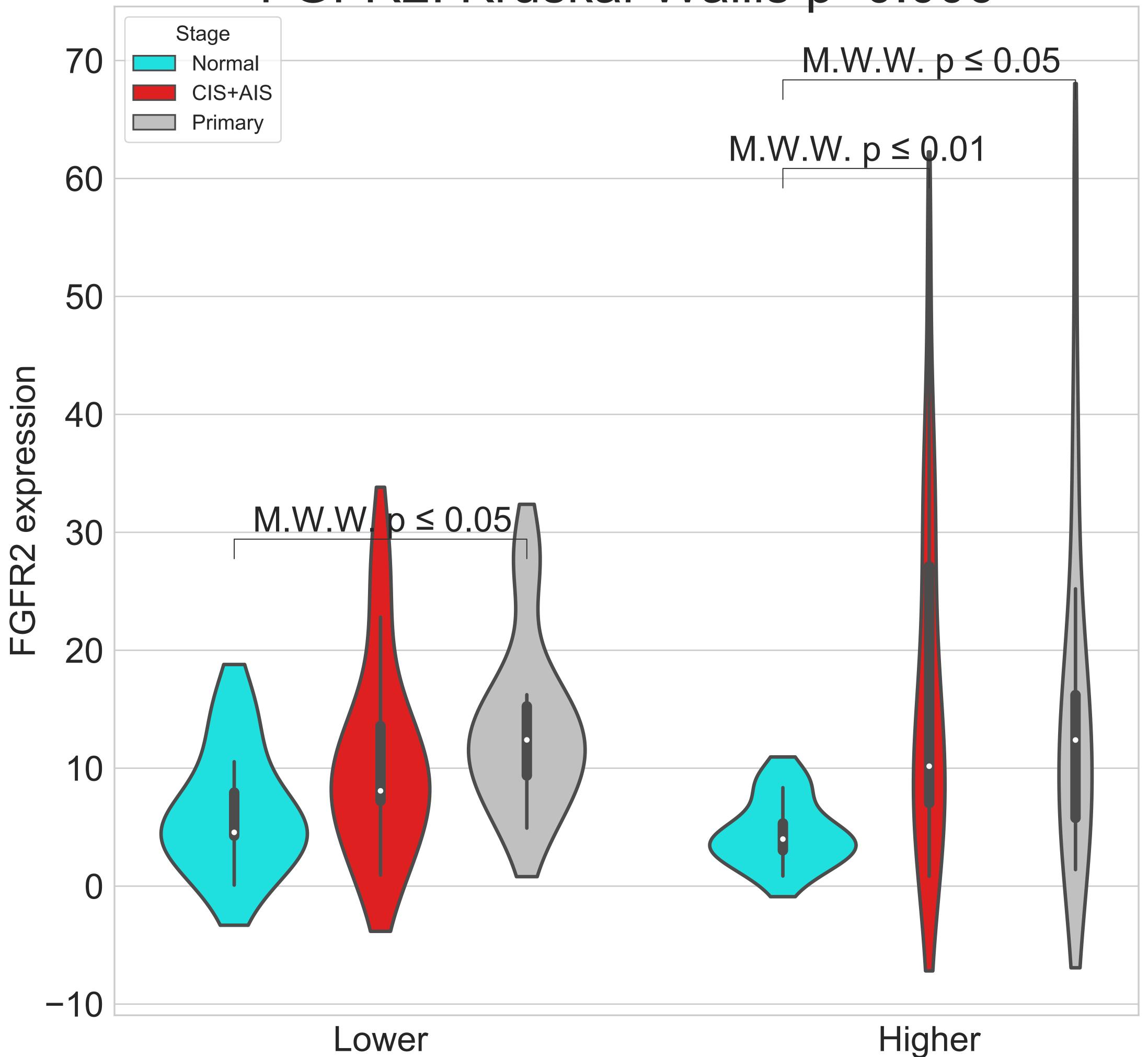
FGFR2: Kruskal-Wallis p=0.006



Mutation Shared Proportion (SYN)