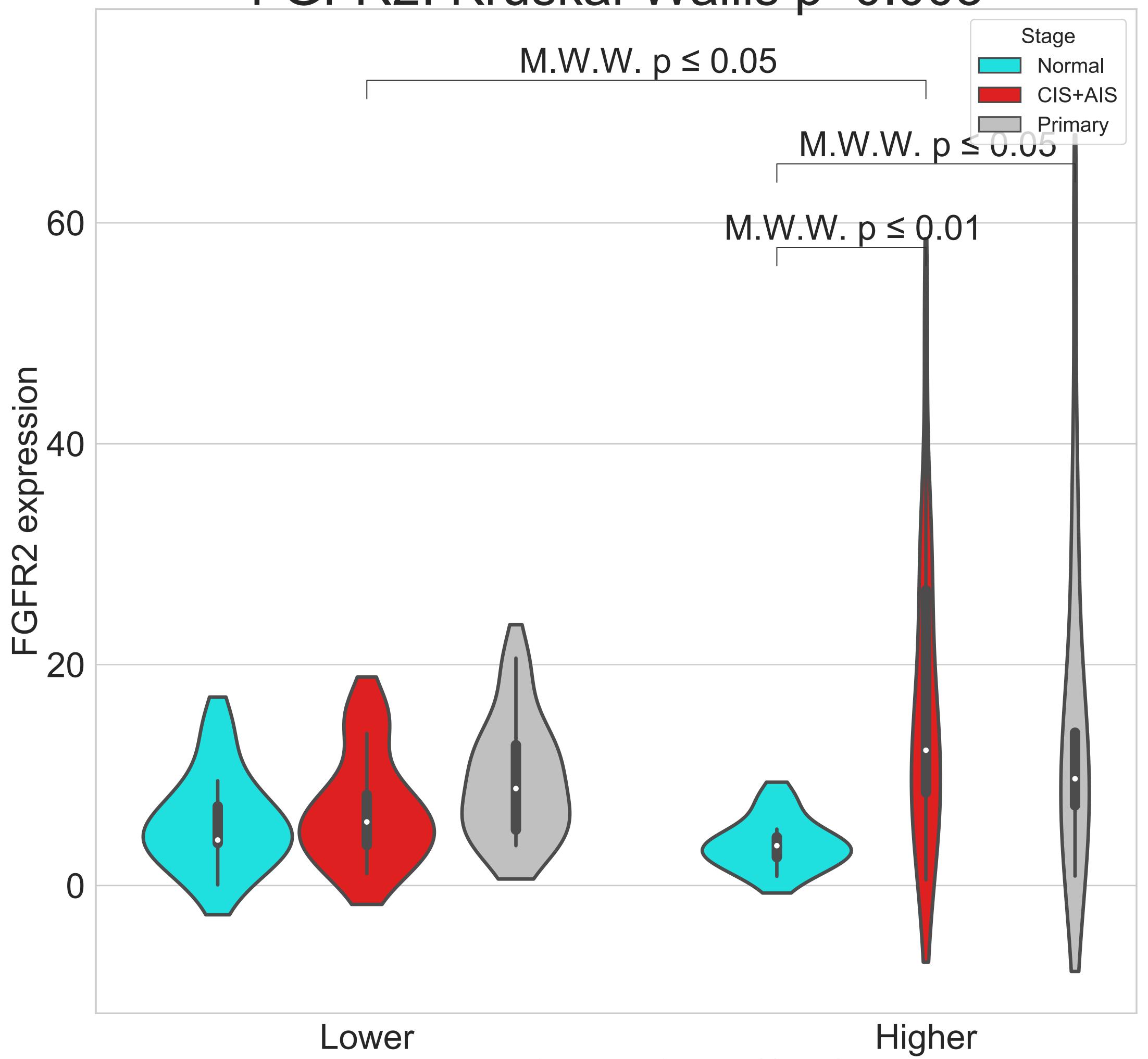
FGFR2: Kruskal-Wallis p=0.003



Mutation Shared Count (SYN)