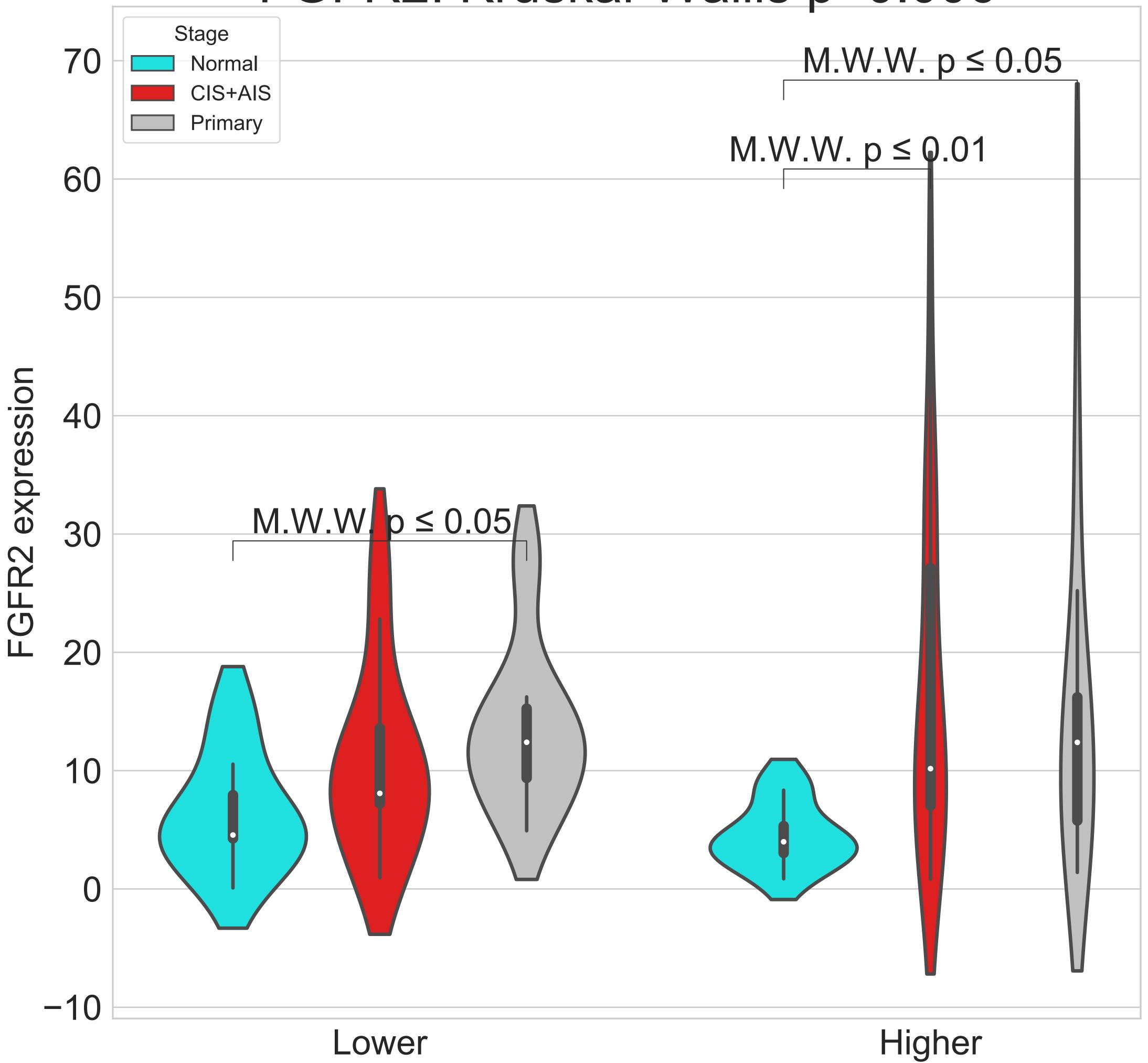
FGFR2: Kruskal-Wallis p=0.006



Mutation Shared Count (SYN) per TMB