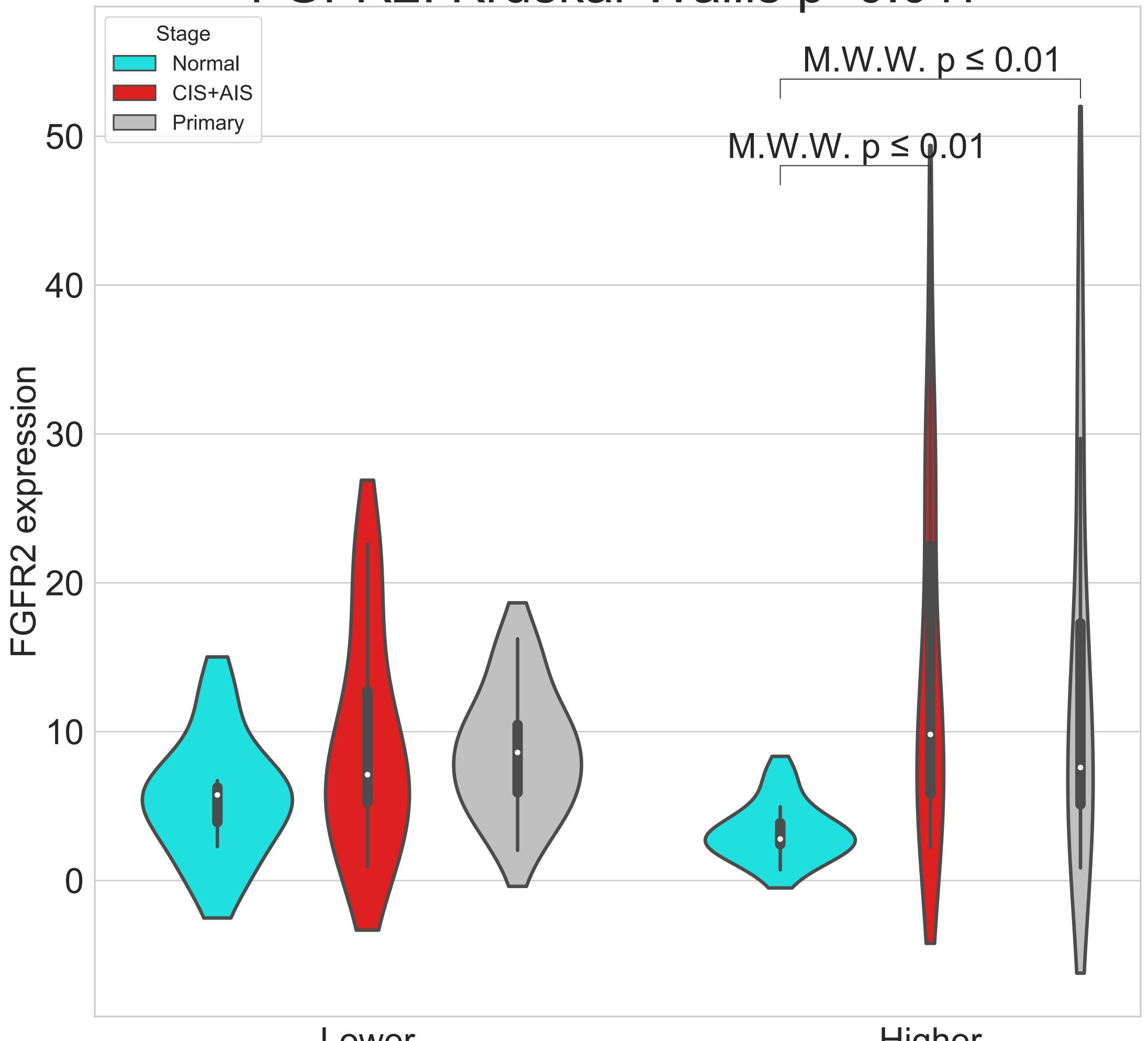
FGFR2: Kruskal-Wallis p=0.017



Lower
Mutation Shared Count (SYN) per TMB