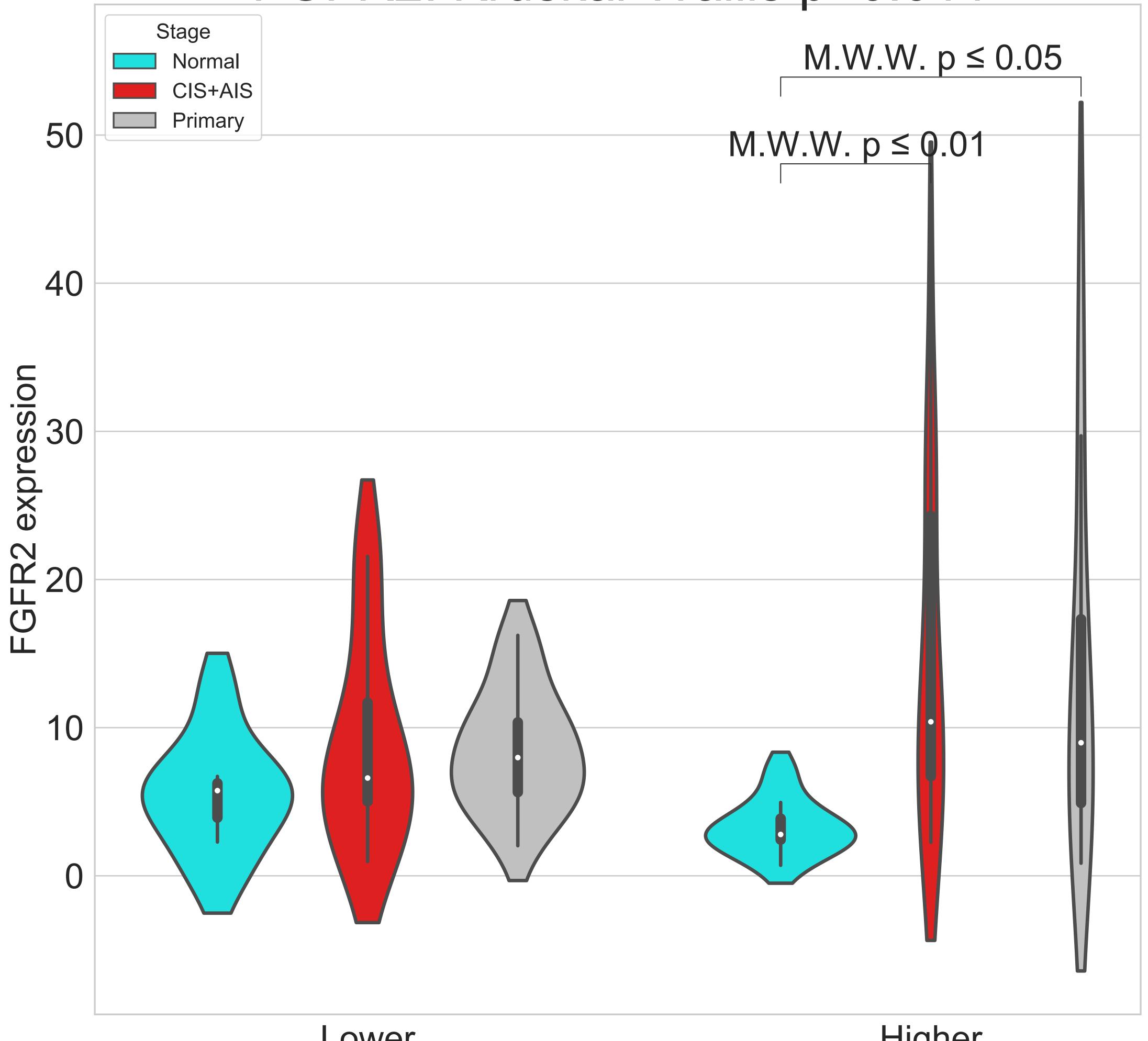
FGFR2: Kruskal-Wallis p=0.011



Lower
Mutation Shared Count per TMB