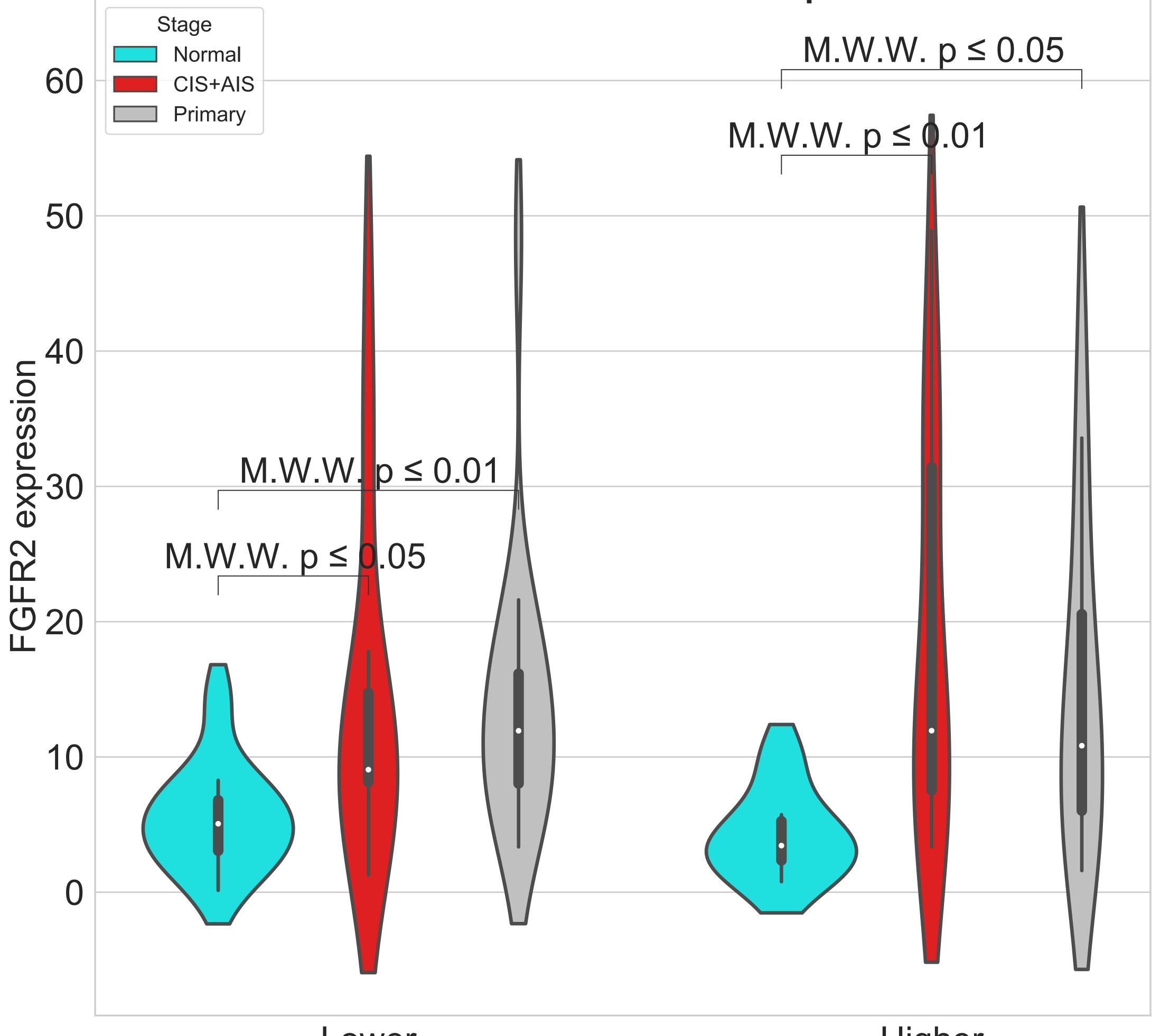
FGFR2: Kruskal-Wallis p=0.007

M.W.W. p ≤ 0



Lower
Mutation Shared Count (SYN) per TMB