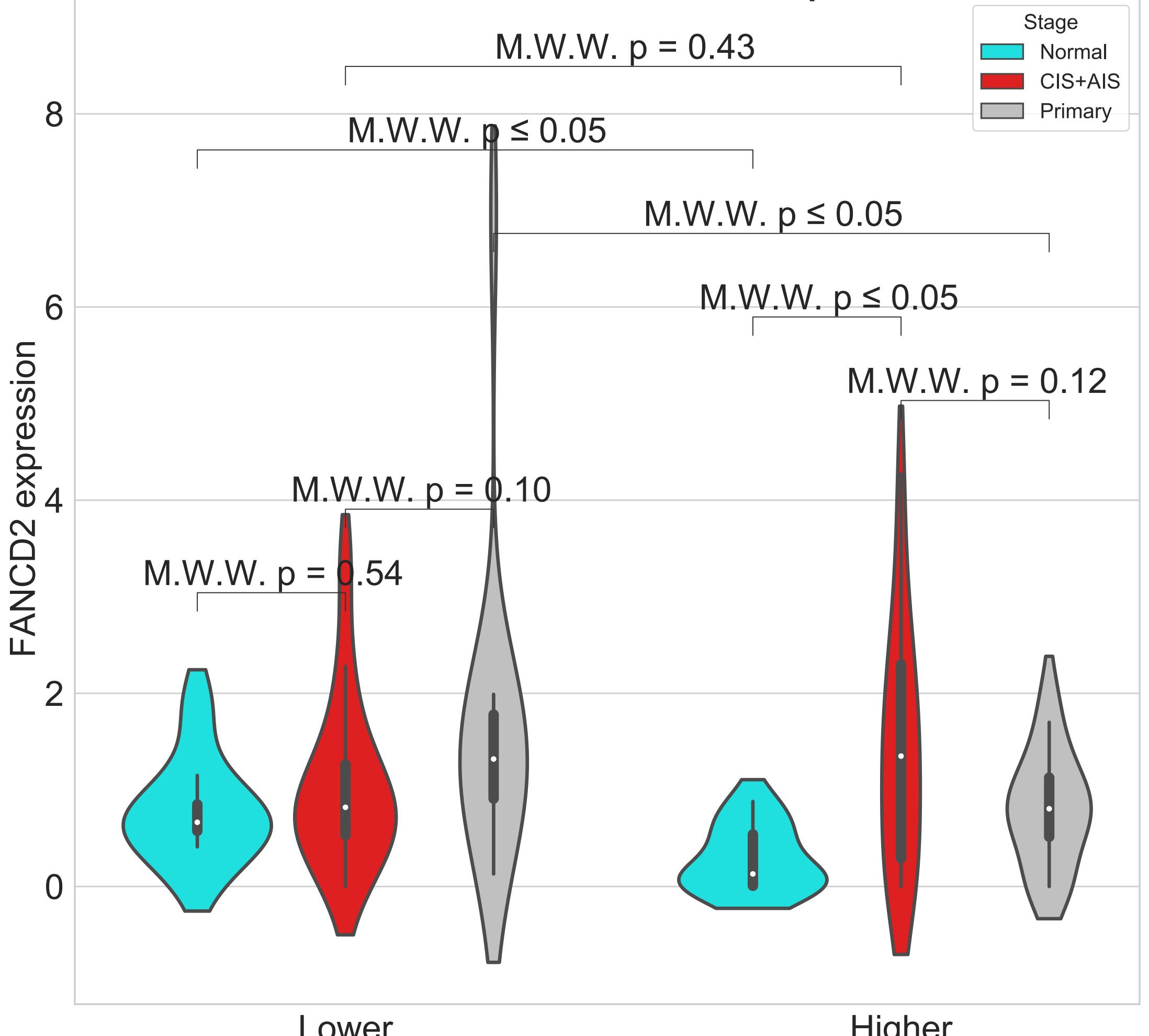
FANCD2: Kruskal-Wallis p=0.010 M.W.W. p = 0.43



Higher Lower Mutation Shared Count (SYN)