

Report HW 4

Calling `geneStr.count('ata')` would take `geneStr` and convert it into an array of individual characters like such:

`['a','a','t','g',...,'c']`

You would never find `'ata'` or any length three sub-string in this array. By using a dictionary, you can assign a key to every codon to be able to easily tally them as they appear.