- Hamming Distance hamming_distance(p, g): mismatch = 0 for i in range (o, len(p)): if p[i] != q,[i]: mis match t = 1 return misnatch GC G - 3 GCGCG -5 def approxfind (pattern, text, d): indices = I] for i in range (0, len(text) - 1en(paten) +1): if hamming (text[i:i+len(pattern)] pattern) <d: append i to indices

beturn indices.

Vet Sicon Southarn Pro PCR, C10 This approach will only work if pattern appears in the string det frequent_words_ nith_mismatches (text, k,d): K_mers = 33 for i in range (len(text)-k+1): K_mer = text[i:i+k] Count = approxfind(text[i:i+K], text, d) K_men [K_mer] = Count max (k_mers. redusci) maximum = L=[Key for Key in K_mers if K_mes[key] == maximum]

return L

AGICAT CCACT AAAA

CAGA

can be grouped ving

group anagrams method

Pattern might not appear in the string.

Therefore, will have to generate all K_merc g

length K as specified in the problem statement and store their "occurrences" in a dictionary and other the maximum or most frequent K_mer.

bases = ["A" T", "G", "C"]

def generate_ Kmers (K, bases):

if K = = 1:

return bases

Small_|cmen = generate_rmens(K-1, bases)

K_mers = []
for Knor in small_1cmero:
for b in bases
append (K_mer+b) to K-mer
return kmers
4-length 12 mer: -x-x-=4.4.4.4
7 = 44 = 250
4 of tions (Cne
Since boses = ['A','G','T','C']
LK I/
= 9 Kmers - x - x - x - x - = 4 = 1024
AAT
AAT
ATT
K, KC, Ck

- Frequent	Words with Mis Matches and Reverse Complements.
def	foo(text, K,d):
	bours = ['A', T', 'G', 'C']
	s = " " Visited = Set ()
	K_mers = generate_Kmers (K bases)
	K_{mers} Count = $\frac{23}{}$
	for Kmer in K_mers:
Have to	if Kmer not in visited:
make sure if	8_C_ Concr = revose_complement(cmar)
Kner out	Count 1 = len(approx_find(kmer, text, &)[
Mr. M. S. Carce.	Count 2 = len (approx_firel()[i]
Chrispmy aggresse.	[C_men_count [Kmer] = som Cennts,
	Courti
	# seturn Know with most frequery.