Notes on window strat

Windower Class

- · Looks like Windower objects control what part of a given sequence energy calculations are made for
 - The runtime of energy calcs is not clear right now but guessing it is probably linear-ish for the length of the windowed sequence

```
In [13]:
          seq = 'ATGCCCCAGGGT'
          start, stop, min_size = 0, 1, 2
          c = 0
          while True:
              if start == stop - min_size+1 and stop == len(seq) -1:
                  break
              print(seq[start:stop+1])
              c += 1
              if stop < len(seq) - 1:</pre>
                  stop += 1
              else:
                  start += 1
                  stop = start + min_size -1
          print("Sequence of len {} produced {} structures".format(len(seq), c))
         ΑТ
         ATG
         ATGC
         ATGCC
         ATGCCC
         ATGCCCC
         ATGCCCCA
         ATGCCCCAG
         ATGCCCCAGG
         ATGCCCCAGGG
         ATGCCCCAGGGT
         TGC
         TGCC
         TGCCC
         TGCCCC
         TGCCCCA
         TGCCCCAG
          TGCCCCAGG
         TGCCCCAGGG
         TGCCCCAGGGT
         GC
         GCC
         GCCC
         GCCCC
         GCCCCA
         GCCCCAG
         GCCCCAGG
         GCCCCAGGG
         GCCCCAGGGT
         CC
         CCC
         CCCC
         CCCCA
         CCCCAG
         CCCCAGG
         CCCCAGGG
         CCCCAGGGT
         CC
         CCC
         CCCA
```

CCCAG **CCCAGG** CCCAGGG CCCAGGGT CC CCA CCAG CCAGG **CCAGGG** CCAGGGT CA CAG CAGG CAGGG CAGGGT AG AGG AGGG AGGGT GG GGG **GGGT** GG

Energy Calcs

What is happening to each of the structures produced by the algo above?

- Called in model.compute_structure and pushed back onto the rloop_structures vector
 - for each rloop structure we then call compute residuals and pass in the Model we are using
 - compute_residuals computes the residual linking difference and the residual twist and stores both of these in the structure instance that was passed in

Residual linking difference

$$(rac{4C(\pi^2)}{4C(\pi^2)+(K\cdot L)})lpha+L*A$$

Where

- \bullet C = tortional stiffness of ssDNA winding. (Could be 3.6 for ds or 1.8 for ss winding)
- A = turns/bp
- ullet L = length of the sequence
- α = linking difference: topological parameter
- $K = \text{Hooke's law coefficient: } (2200 \textit{ideal_gas_constant in kcal/mol} \text{absolute_temp_in_kelvin})/N$

Above should actually be a fast calculation though $\theta(1)$. The big time loss is the number of structures that have to be computed for and the base-pairing energy over the each structure

Base-pairing energy

 $Two \ things \ are \ computed \ here \ in \ a \ for \ loop \ in \ the \ Rloop_equilibrium_model::compute_structure \ method$

- 1. Free Energy
 - Take adjacent bases call step_forward_bps which calls compute_bps_interval which returns a cached energy value based on the identity of the first and then second base
 - $\theta(1)$
- 2. Boltzmann Factor
 - ullet Call compute_boltzman_factor and pass in the free energy value we just calculated for the last two adjacent bases and constant T

Base-pairing energy improvements

It might make more sense to calculate free energy and boltzman while we are generating each struct instead of calculating all structures and then for each structure do the energy calc again. We end up repeating a **TON** of calculations it seems like

Example

```
In [2]:
         # create class to represent rloop structures (windows)
         class Struct:
             seq = None
             def __init__(self, start, stop):
                 self.start = start
                 self.stop = stop
                 self.boltz = 0
                 self.free_energy = 0
             @property
             def sequence(self):
                 return self.seq[self.start:self.stop]
In [3]:
         # set class variable seq to the seq we want to use for calcs
         Struct.seq = 'ATGCCCCAGGGT'
In [4]:
         T = 0.64 # stand in for T constant
         calc free = lambda a, b: ord(a) + ord(b) # not actual calc just take ascii values for now as stand in
         calc_boltz = lambda e: e + T
```

Dynamic programming approach

If we are at the same start point and just building the sequence out we (I think) don't need to recalculate all the free energy calcs done on the preceding bases, just the new base (n) along with the n - 1 base.

```
In [5]:
         def EH structs():
             import time
             start, stop, min_size = 0, 1, 2
             structures = []
             cur_boltz, cur_energy = 0, 0
             s = time.time()
             while True:
                  if start == stop - min_size+1 and stop == len(Struct.seq) -1:
                 if stop < len(Struct.seq) - 1:</pre>
                     new_struct = Struct(start, stop+1)
                     cur_energy += calc_free(Struct.seq[new_struct.stop -2], Struct.seq[new_struct.stop-1])
                     cur_boltz += calc_boltz(cur_energy)
                     new_struct.boltz = cur_boltz
                     new_struct.free_energy = cur_energy
                     structures.append(new_struct)
                     stop += 1
                  else:
                     start += 1
                     stop = start + min_size - 1
                     cur boltz = 0
                     cur_energy = 0 # reset as we have moved to new start point
             total_time = time.time() - s
             print("Total Time: {} seconds".format(time.time() - s))
             print("Free Energy | Boltzman | Seq")
             #for s in structures:
                  print(s.free_energy, round(s.boltz, 2), Struct.seq[s.start:s.stop])
             return total_time
```

Same thing as above but with current method

```
In [6]:
         def 0G_structs():
              import time
              start, stop, min_size = 0, 1, 2
              structures = []
              s = time.time()
              while True:
                  if start == stop - min_size+1 and stop == len(Struct.seq) -1:
                      break
                  if stop < len(Struct.seq) - 1:</pre>
                      new_struct = Struct(start, stop+1)
                      for i in range(len(new_struct.sequence)-1):
                          a, b = new_struct.seq[i], new_struct.seq[i+1]
                          new_struct.free_energy += calc_free(a, b)
                      new_struct.boltz += calc_boltz(new_struct.free_energy)
                      structures.append(new_struct)
                      stop += 1
                  else:
                      start += 1
                      stop = start + min_size - 1
              total_time = time.time() - s
              print("Total Time: {} seconds".format(time.time() - s))
              print("Free Energy | Boltzman | Seq")
              #for s in structures:
              # print(s.free_energy, round(s.boltz, 2), Struct.seq[s.start:s.stop])
              return total time
In [11]:
         EH_time = []
          0G_{time} = []
          l = []
          Struct.seq = 'ATGCCCCAGGGT'
          for i in range(0, 7):
    Struct.seq = Struct.seq * 2
              l.append(len(Struct.seq))
              print('Current length:', l[-1])
              EH_time.append(EH_structs())
              OG_time.append(OG_structs())
         Current length: 24
         Total Time: 0.0006544589996337891 seconds
         Free Energy | Boltzman | Seq
         Total Time: 0.0015773773193359375 seconds
         Free Energy | Boltzman | Seq
         Current length: 48
         Total Time: 0.0027348995208740234 seconds
         Free Energy | Boltzman | Seq
         Total Time: 0.012103557586669922 seconds
         Free Energy | Boltzman | Seq
         Current length: 96
         Total Time: 0.007555484771728516 seconds
         Free Energy | Boltzman | Seq
         Total Time: 0.046959638595581055 seconds
         Free Energy | Boltzman | Seq
         Current length: 192
         Total Time: 0.016002178192138672 seconds
         Free Energy | Boltzman | Seq
         Total Time: 0.2863423824310303 seconds
         Free Energy | Boltzman | Seq
         Current length: 384
         Total Time: 0.10921955108642578 seconds
         Free Energy | Boltzman | Seq
         Total Time: 2.176204204559326 seconds
         Free Energy | Boltzman | Seq
         Current length: 768
         Total Time: 0.39378976821899414 seconds
         Free Energy | Boltzman | Seq
         Total Time: 17.390355348587036 seconds
```

```
Free Energy | Boltzman | Seq
Current length: 1536
Total Time: 1.4360833168029785 seconds
Free Energy | Boltzman | Seq
Total Time: 142.8125376701355 seconds
```

In [12]: import matplotlib.pyplot as plt
plt.plot(l, EH_time)
plt.plot(l, OG_time)
plt.show()

Free Energy | Boltzman | Seg

