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DNA can be modeled as two parallel polymer strands with links between the strands called *base pairs*. Each base pair can be in a closed state with energy 0 or in an open state with energy ε .

Consider a DNA molecule with *N* base pairs in thermal equilibrium at temperature *T*, as shown below. Thermal fluctuations can cause each base pair to open, leading to separation of the two strands. The molecule can open only from the left end, and only in *sequential order* (i.e. base pair *s* can open only if 1, 2...s-1 to the left of it are already open).



(a) Show that the partition function Z for this system is given by the following expression:

a.
$$Z = \frac{1 - e^{-(N+1)\varepsilon/k_B T}}{1 - e^{-\varepsilon/k_B T}}$$

(b) In the limit that $N \to \infty$, determine the mean number $\langle n \rangle$ of open base pairs. Approximate your expression separately in the limits that $k_B T \gg \varepsilon$, and that $k_B T \ll \varepsilon$.

Next, consider the same DNA molecule now surrounded by a protein *p* at concentration *c*. Protein *p* can bind to the DNA *only at a site that is open*, as shown below. Assume each protein *p* can occupy no more than one base pair. The chemical potential for *p* is $\mu = \Delta + k_B T \ln(c/c_0)$, where c_0 and Δ are constants ($c_0 > 0$).



(d) Using the Grand canonical ensemble, in the limit that N → ∞, determine the mean number of DNA-binding proteins (p) per DNA at a temperature T for a given concentration c.