

0.1 Zipper model

It is a model introduced by Kittel to describe oligomers undergoing *denaturation transition*. Simplest model of DNA thermal denaturation transition (no bubbles). Better model for the denaturation of short oligomers.

The hypothesis are: the binding energy between two bases located at the end of the molecule is smaller than the one for pairs away from the ends. The unbinding starts and develops from the ends as a *zipper*.

Lecture 9.

Friday 8th

November, 2019.

Compiled: Tuesday

10th December,

2019.

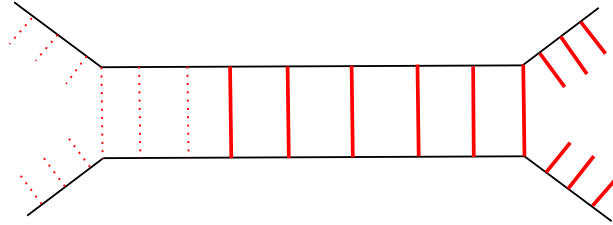


Figure 1: Sequential unzipping from the ends.

In this denaturation transition we do not allow bubbles. Let us consider first the zingle-ended zipper, i.e. a molecular zipper of N parallel links that can be opened only from one end as in Figure 2.

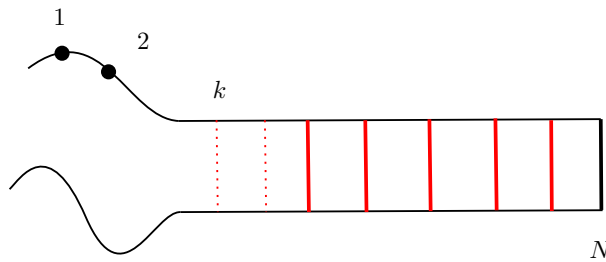


Figure 2

If the first k bonds are open (unbounded pairs) the energy to open the $k+1$ is ε_0 . Note that if at least one of the previous k bond is closed the energy needed to open the $k+1$ band is infinite! One a bond is open it can orient itself in G different ways. In other words there is an entropy

$$S_0 = k_B \log G \quad (1)$$

associated to each open band. If k bands ($i = 1, \dots, k$) are open the associated weight is given by

$$G^k e^{-k\varepsilon_0/k_B T} \quad (2)$$

By summing over the possible values of k , the partition function is

$$Z_N(T, G, \varepsilon_0) = \sum_{k=0}^{N-1} G^k e^{-k\varepsilon_0/k_B T} = \sum_{k=0}^{N-1} e^{k(S_0 T - \varepsilon_0)/k_B T} \quad (3)$$

Let us call

$$\chi \equiv G e^{-\varepsilon_0/k_B T} \quad (4)$$

and simplify the previous expression

$$Z_N = \sum_{k=0}^{N-1} \chi^k = \frac{1 - \chi^N}{1 - \chi} \quad (5)$$

We see immediately there is a single pole singularity. The free energy is

$$F_N = -k_B T \ln Z_N = -k_B T \ln \left[\frac{1 - \chi^N}{1 - \chi} \right] \quad (6)$$

We can now compute some observables of interest.

Calculate average number of open links

The equal of our magnetization in this case is

$$\langle k \rangle_N \equiv \frac{\sum_{k=0}^{N-1} k \chi^k}{\sum_{k=0}^{N-1} \chi^k} = \chi \frac{d}{d\chi} \ln Z_N = \frac{N \chi^N}{\chi^N - 1} - \frac{\chi}{\chi - 1} \quad (7)$$

There is a pole in $\chi = 1$ and the last quantity can be drawn as in Figure 3.

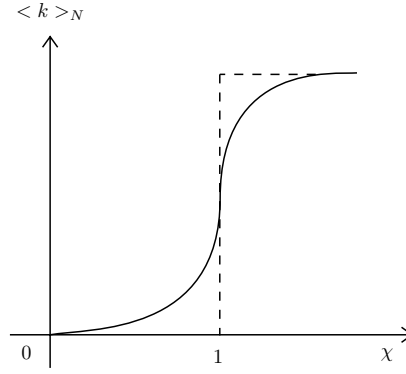


Figure 3: Description.

In order to analyze what happens near 1, we expand $\chi \equiv 1 + \varepsilon$:

$$\begin{aligned} \log Z_N(\chi) &= \log \left[\frac{1 - (1 + \varepsilon)^N}{1 - (1 + \varepsilon)} \right] \\ &= \log \left[\frac{1 - (1 + \varepsilon N + \frac{N(N-1)}{2!} \varepsilon^2 + \frac{N(N-1)(N-2)}{3!} \varepsilon^3 + O(\varepsilon^4))}{\varepsilon} \right] \\ &= \log \left[N + \frac{N(N-1)}{2} \varepsilon + \frac{N(N-1)(N-2)}{6} \varepsilon^2 + \dots \right] \\ &= \log N + \log \left[1 + \frac{N-1}{2} \varepsilon + \frac{N(N-1)}{6} \varepsilon^2 \right] \\ &= \log N + \log \left[1 + \frac{N\varepsilon}{2} + \frac{N^2 \varepsilon^2}{6} + \dots \right] \\ &= \log N + \left(\frac{N\varepsilon}{2} + \frac{N^2 \varepsilon^2}{6} + \dots \right) + \frac{1}{2} \left(\frac{N\varepsilon}{2} + \frac{N^2 \varepsilon^2}{6} + \dots \right)^2 + \dots \\ &= \log N + \frac{N\varepsilon}{2} + \frac{N^2 \varepsilon^2}{24} + \dots \end{aligned} \quad (8)$$

By doing the same for $\langle k \rangle_N = \frac{N \chi^N}{\chi^N - 1} - \frac{\chi}{\chi - 1}$, one gets

$$\langle k \rangle_N = \frac{N}{2} \left(1 + \frac{N\varepsilon}{6} - \frac{N^3 \varepsilon^3}{360} + \dots \right) \quad (9)$$

this is true for $N \gg 1, \varepsilon \ll 1$. At the transition point $\chi_C = 1$, where $\varepsilon = 0$:

$$\langle k \rangle_N \simeq \frac{N}{2} \quad (10)$$

We can define the variation (slope per site) as a response function (the derivative with respect to the parameter):

$$\frac{1}{N} \frac{d \langle k \rangle}{d \varepsilon} \simeq \frac{N}{12} - \frac{N^3 \varepsilon^3}{240} + \dots \quad (11)$$

It is max at $\varepsilon = 0$ and diverges as $N \rightarrow \infty$ (linearly). The response function diverges linearly to N . This is a good signal that we have a transition. The temperature T_c corresponding to the pole $\chi = 1$ is given

$$G e^{-\varepsilon_0/k_B T_C} = 1 \quad (12)$$

Hence

$$T_C = \frac{\varepsilon_0}{k_B \log G} \quad (13)$$

Note that as $G \rightarrow 1$, $T_c \rightarrow 0$. For $G = 1$ there is no solution and hence the model does not display a phase transition for any finite T ! This is telling you that if $G = 1$ what is important it is the energy, you have no entropy as disorder. At that point everything can happen.

Remark. Despite the model is $1D$, for $G > 1$ there is a phase transition. This is due to two contributions:

1. Existence of forbidden configuration (infinite energy). Necessary condition for a phase transition in $d = 1$ with finite range interactions.
2. Degeneracy of the excited state (G).

0.1.1 Transfer matrix method for the Kittel model

The idea is: we want to map this model to an Ising model. The spin like model consists on associating to each bond a spin such that $S_i = 0$ if the i -esim bond is *closed*, while $S_i = 1, \dots, G$ if the i -esim bond is *open* with G possible orientations.

- Case: $S_i \neq 0$ open. We have two subcases:
 - S_{i-1} open: $S_{i-1} \neq 0 \Rightarrow E(S_i \neq 0 | S_{i-1} \neq 0) = \varepsilon_0$.
 - S_{i-1} closed: $S_{i-1} = 0 \Rightarrow E(S_i \neq 0 | S_{i-1} = 0) = \varepsilon_0 + V_0$
- Case: $S_i = 0$ closed. We have $E(S_i = 0) = 0$ irrespective of S_{i-1} .

Therefore

$$E(S_i, S_{i-1}) = (\varepsilon_0 + V_0 \delta_{S_{i-1}, 0})(1 - \delta_{S_i, 0}) \quad (14)$$

The boundary condition is $S_N = 0$ (always closed). The full Hamiltonian of the model can be written as (it could be also a function of delta, but it is not a problem):

$$\mathcal{H}_N = \varepsilon_0(1 - \delta_{S_1, 0}) + \sum_{i=2}^{N-1} (\varepsilon_0 + V_0 \delta_{S_{i-1}, 0})(1 - \delta_{S_i, 0}) \quad (15)$$

The Kittel's version is obtained by assuming $V_0 = \infty$.

The partition function is

$$Z_N = \sum_{\{S\}} \exp(-\beta \mathcal{H}_N) \quad (16)$$

In order to implement the transfer matrix formalism we rewrite Z_N as follows (DUB-BIIOOOOOO per la prossima equation)

$$Z_N = \sum_{\{S\}} e^{-\beta \varepsilon_0 (1 - \delta_{S_1, 0})} \prod_{i=1}^{N-2} e^{-\beta \varepsilon_0 (1 - \delta_{S_{i+1}, 0})} \left(1 + (e^{-\beta V_0} - 1) \delta_{S_i, 0} (1 - \delta_{S_{i+1}, 0}) \right) \quad (17)$$

Consider the Kittel model, $V_0 = \infty$ which implies $\exp(-\beta V_0) = 0$. We can define the transfer matrix as

$$\mathbb{T} = \{ \langle S | \mathbb{T} | S' \rangle \equiv t_{S, S'} \} \quad (18)$$

where

$$t_{S, S'} = e^{-\beta \varepsilon_0 (1 - \delta_{S', 0})} [1 - \delta_{S, 0} (1 - \delta_{S', 0})] \quad (19)$$

or in matrix form

$$\mathbb{T} = \begin{bmatrix} 1 & 0 & \dots & 0 \\ 1 & a & \dots & a \\ \vdots & \vdots & & \vdots \\ \vdots & \vdots & & \vdots \\ 1 & a & \dots & a \end{bmatrix} \quad (20)$$

where $a \equiv e^{-\beta \varepsilon_0}$.

The first think to notice that the constraint that the bond S_{i+1} cannot be open if bond S_i is closed ($S_i = 0$) yields the null entries in the first row of \mathbb{T} . This violates the hypothesis of the Perron-Frobenius theorem!

The matrix \mathbb{T} has three differen eigenvalues

$$\lambda_1 = Ga, \quad \lambda_2 = 1, \quad \lambda_3 = 0 \quad (21)$$

The partition function can be written as

$$Z_N = (1, a, \dots, a) \mathbb{T}^{N-2} \begin{pmatrix} 1 \\ 1 \\ \vdots \\ 1 \end{pmatrix} \quad (22)$$

$$\lambda_1 \rightarrow \vec{v}_1 = \begin{pmatrix} 0 \\ 1 \\ \vdots \\ 1 \end{pmatrix}, \quad \lambda_2 \rightarrow \vec{v}_2 = \begin{pmatrix} 1 - Ga \\ 1 \\ \vdots \\ 1 \end{pmatrix} \quad (23)$$

We can then write

$$\begin{pmatrix} 1 \\ a \\ \vdots \\ a \end{pmatrix} = \frac{a(1 - Ga) - 1}{1 - Ga} \vec{v}_1 + \frac{1}{1 - Ga} \vec{v}_2 \quad (24a)$$

$$\begin{pmatrix} 1 \\ 1 \\ \vdots \\ 1 \end{pmatrix} = \frac{-Ga}{1 - Ga} \vec{v}_1 + \frac{1}{1 - Ga} \vec{v}_2 \quad (24b)$$

Therefore

$$Z_N = \frac{1 - (Ga)^N}{1 - Ga} = \frac{1 - (Ge^{-\beta \varepsilon_0})^N}{1 - Ge^{-\beta \varepsilon_0}} \quad (25)$$

or

$$Z_N = \frac{1}{1 - Ge^{-\beta\epsilon_0}} (-\lambda_1^N + \lambda_2^N) \quad (26)$$

Since in the thermodynamic limit only the contribution of the largest eigenvalue matters for f_b we have

$$f_b = -k_B T \ln \max(\lambda_1, \lambda_2) \quad (27)$$

Remark. Given that the λ_1 and λ_2 are positive, analytic function of T ($\lambda_1 = Ga, \lambda_2 = 1$). In order to have a phase transition (i.e. non analyticity of f_b) the two eigenvalues must cross for a given value of T . It is true if and only if:

$$Ga_c = 1 \Leftrightarrow Ge^{-\beta_c \epsilon_0} = 1 \Leftrightarrow T_c = \frac{\epsilon_0}{k_B \ln G} \quad (28)$$

It is agree with previous calculation.

0.2 Transfer matrix for 2D Ising

Dimensional redudation from d to $d-1$. You can to the same for the 3D Ising model. In order to go smoothly to the one dimensional to the two dimensional the idea is to solve the Ising model to surfaces.

Consider a square lattice with N rows and M columns, as in Figure 4, with periodic boundary conditions (wrapped around a torus). The spin in a site is identified by $S_{\text{site}} = S_{m,n}$. The Hamiltonian is (if $H \neq 0$ we have also the term with the h):

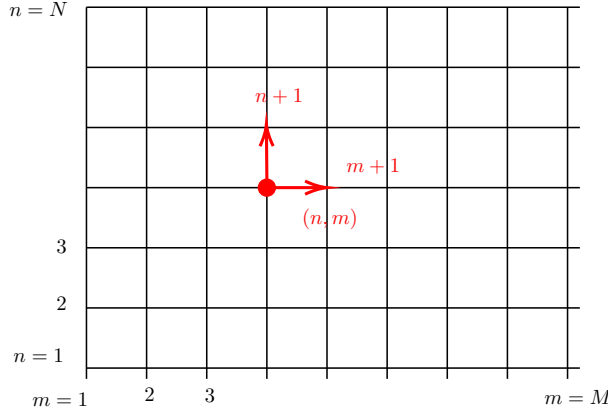


Figure 4: Description.

$$\begin{aligned} -\beta \mathcal{H}_\Omega(\{S\}) &= k \sum_{\langle ij \rangle} S_i S_j + h \sum_i S_i \\ &= k \sum_{n=1}^N \sum_{m=1}^M (S_{m,n} S_{m+1,n} + S_{m,n} S_{m,n+1}) + h \sum_{n=1}^N \sum_{m=1}^M S_{m,n} \end{aligned} \quad (29)$$

in fact

$$\sum_{\langle i,j \rangle} \rightarrow \sum_{i,j \in nn(i)} \Rightarrow S_{m,n} \rightarrow S_{m+1,n} \quad (30)$$

The Hamiltonian can be rewritten as follows:

$$-\beta \mathcal{H}_\Omega(\{S\}) = \sum_{m=1}^M [E[\mu_m, \mu_{m+1}] + E[\mu_m]] \quad (31)$$

where the first term is the interaction between columns (two body interaction), the second term is the one body interaction of one column. Moreover, the μ is a m dimensional vector, in particular each μ_m represents the set of N spins along column m :

$$\mu_m = \{S_{m,1}, S_{m,2}, \dots, S_{m,N}\} \quad (32)$$

We have:

$$E[\mu_m, h] = k \sum_{n=1}^N S_{m,n} S_{m,n+1} + h \sum_{n=1}^N S_{m,n} \quad (33a)$$

$$E[\mu_m, \mu_{m+1}, h] = k \sum_{n=1}^N S_{m,n} S_{m+1,n} \quad (33b)$$

where the first equation is the one body interaction, while the second equation represents the interaction between neirest neighbours columns.

We can write a transfer matrix between these new variables. The transfer matrix permit to transfer along the m . To make it simpler suppose $h = 0$ (so the energy does not depend on h):

$$\langle \mu_m | \mathbb{T} | \mu_{m+1} \rangle = \exp[k(E[\mu_m, \mu_{m+1}] + E[\mu_m])] \quad (34)$$

Now we have to diagonalize. In the 2x2 tranfer matrix in the two dimensional we have 2 possible values. Now we have to do the same in principle, but we have to do for all of the (32). \mathbb{T} is a matrix of dimension $2^N \times 2^N$. In the thermodynamic limit is an infinite matrix (violation of Perron-Frobenius). According to the formalism

$$Z_N(k, h) = \text{Tr}(\mathbb{T}^N) \quad (35)$$

To find the eigenvalues of \mathbb{T} given by (34) is highly non trivial. The big problem it is that in the thermodynamic limit is that the dimension of the transfer matrix goes to infinity, then it is difficult to be diagonalized. This was first achieved by Onsanger in 1944 for the case $H = 0$ and in the $N \rightarrow \infty$ limit. The result is given by

$$f_b(T) = -k_B T \log(2 \cosh(2\beta J)) - \frac{k_B T}{2\pi} \int_0^{2\pi} \log \left[\frac{1}{2} \left(1 + \sqrt{1 - g^2 \sin^2(\Phi)} \right) \right] d\Phi \quad (36)$$

that is the density bulk free energy, where

$$g = \frac{2}{\cosh(2\beta J) \coth(2\beta J)} \quad (37)$$

The equation for the critical T is

$$2 \tanh^2 \left(\frac{2J}{k_B T_c} \right) = 1 \Rightarrow T_c \simeq 2,264 J / k_B \neq 0! \quad (38)$$

One can also show that

$$c \propto A \left[-\ln \left(1 - \frac{T}{T_c} \right) + B \right] \quad (39)$$

The specific heat displays at the transition a logarithmic divergence (no power law!). Therefore in $d = 2$ the Ising model as $\alpha_{\text{Ising } 2D} = 0$. You do Montecarlo simulation.