

Progress Report

Modeling the mechanism of topoisomerase

Aukkawut Ammartayakun

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Worcester Polytechnic Institute

1. Goals
2. Literature Review
3. Initial Ideas

Goals



Overall goal

- Understand the interaction of *TOP2* and the topology of DNA.

This week goal

- Understand the dynamics and fluid-coupled system of DNA.
- (Hopefully) Start to model.

Literature Review

Linking number

$$Lk(C_1, C_2) = \frac{1}{4\pi} \oint_{C_1} \oint_{C_2} \frac{\mathbf{x}'_1(s_1) \times \mathbf{x}'_2(s_2) \cdot [\mathbf{x}_1(s_1) - \mathbf{x}_2(s_2)]}{|\mathbf{x}_1(s_1) - \mathbf{x}_2(s_2)|^3} ds_2 ds_1$$

Writhe

$$Wr(C) = \frac{1}{4\pi} \oint_C \oint_C \frac{\mathbf{x}'(s_1) \times \mathbf{x}'(s_2) \cdot [\mathbf{x}(s_1) - \mathbf{x}(s_2)]}{|\mathbf{x}(s_1) - \mathbf{x}(s_2)|^3} ds_2 ds_1$$

Twist

$$Tw(C_2, C_1) = Lk(C_1, C_2) - Wr(C_1)$$

Simplified Model (Swigon, 2009)

Elastic Energy

$$\Psi = \frac{1}{2} \int_0^L \overbrace{A |\mathbf{t}'|^2}^{\text{bending}} + C \underbrace{\left([\mathbf{t}_1 \times \mathbf{d}] \cdot \mathbf{d}'(s) - \bar{\Omega} \right)^2}_{\text{squared difference of twist density}} ds \implies A (\mathbf{t} \times \mathbf{t}'') + C \Delta \Omega \mathbf{t}' = \mathbf{F} \times \mathbf{t}$$

Under standard condition: $A \approx 50kT \cdot \text{nm}$, $C \in [25, 100]kT \cdot \text{nm}$

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Electrostatic Energy

For M charged sites,

$$\Phi = \frac{4\delta^2}{4\pi\epsilon} \sum_{m=1}^{M-1} \sum_{n=m+1}^M \frac{\exp(-\kappa |\mathbf{x}_m - \mathbf{x}_n|)}{|\mathbf{x}_m - \mathbf{x}_n|}$$

Generalized Kirchhoff's Rod Dynamics

$$\partial_t \mathbf{P} = \mathbf{F}' + \mathbf{f} \quad \text{(linear momentum)}$$

$$\partial_t \mathbf{R} = \mathbf{M}' + \mathbf{x}' \times \mathbf{F} + \mathbf{m} \quad \text{(angular momentum)}$$

Initial Ideas

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- **Problem:** How can we know K ? Local VS Global?

Naive Ideas (con't)

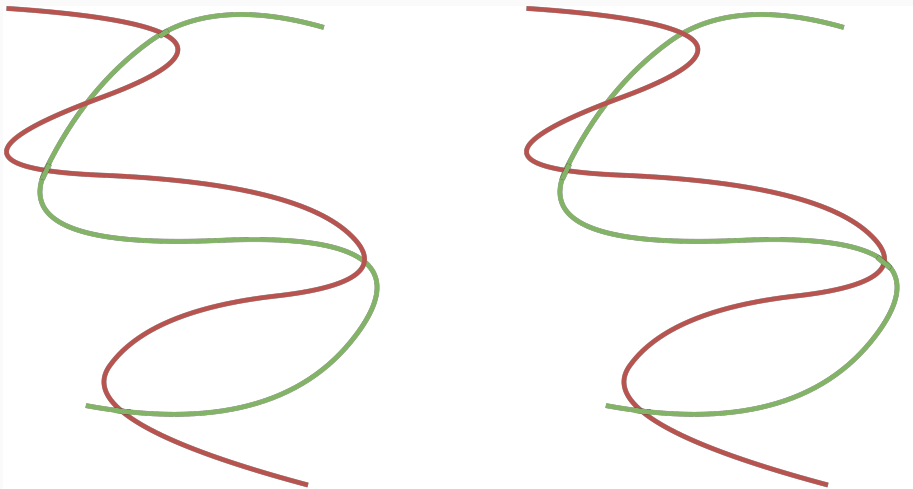


Figure 1: Visualize the process. Left side is original and right side is randomly transformed

- How can *TOP2* know the cleavage site?