

# 絡み合いとプラトー弾性率

佐々木 裕<sup>1</sup>

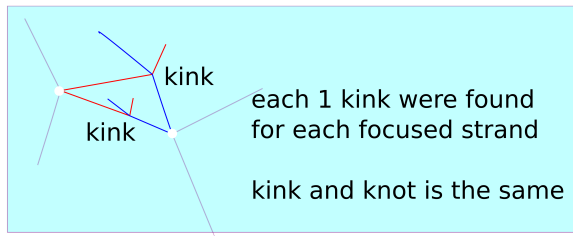
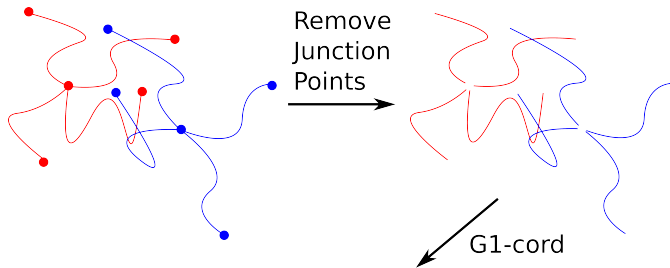
東亜合成株式会社

April 13, 2023

---

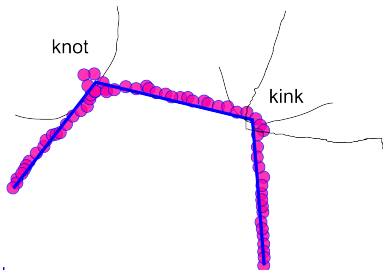
<sup>1</sup>hiroshi\_sasaki@mail.toagosei.co.jp

# Z1 code



# Comparison of Z1 vs. PPA

- 右図のような比較から、二倍程度は妥当かと。
- 次ページにあるネットワークでの比較でも明らかに PPA 由来の経路長は長い
- これと、プラトー弾性率との関係はよくわかりません。



Z1code:

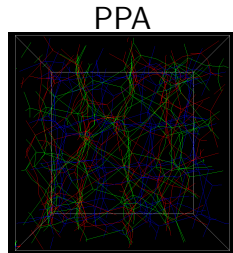
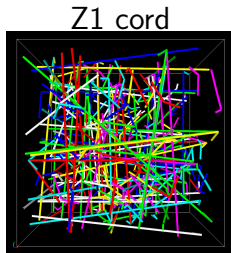
1. Using G1 cord algorithm, strand is shrunk as a straight line preserving kinks.
2. Calc.  $z$  by counting kinks on each strand
3. Calc.  $Me$  by dividing Number of segments by  $z$ .

PPA:

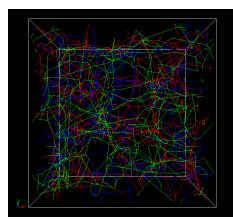
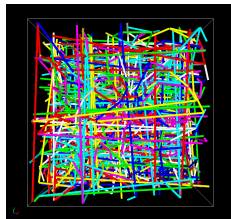
1. Shrink every bond preserving kinks and knots.
2. Calc. contour length by summing each bond length.
3. Calc.  $l_k$  from contour length

# Example for NW

NPT  
絡み合い小



NVT  
自然な絡み合い



# Classical Theory of Rubber Elasticity

## Free Energy Density of Rubbers against Strain invariant

$$\frac{F}{V} = W = C_0 + \underbrace{C_1(I_1 - 3) + C_2(I_2 - 3)}_{\text{Mooney-Rivlin Model}} + \sum_{i,j=1}^{\infty} C_{ij}(I_1 - 3)^i(I_2 - 3)^j$$

### Neo-Hookean Model

$$W = C_1(I_1 - 3)$$

against Uniaxial elongation

$$\sigma_{nom} = 2C_1 \left( \lambda - \frac{1}{\lambda^2} \right) = G \left( \lambda - \frac{1}{\lambda^2} \right)$$

### Mooney-Rivlin Model

$$W = C_1(I_1 - 3) + C_2(I_2 - 3)$$

against Uniaxial elongation

$$\sigma_{nom} = 2 \left( C_1 + C_2 \frac{1}{\lambda} \right) \left( \lambda - \frac{1}{\lambda^2} \right)$$

## With or without Junction Points fluctuation

### Affine Network Model <sup>a</sup>

$$G_{affine} = \nu k_B T$$

$\nu$ : Number density of strands in the system

<sup>a</sup>P.J. Flory, Principles of Polymer Chemistry, (1953)

### Phantom Network Model <sup>a</sup>

$$G_{phantom} = \nu k_B T \left( 1 - \frac{2}{f} \right)$$

$f$ : Functionality of Junction Points

<sup>a</sup>H.M. James, E.J. Guth, Chem. Phys., 21, 6, 1039 (1953)

# Constraint Factors for Junction Points and Strands

## Vicinity of Junction Point

- Junction points are surrounded by many of **adjacent strands**(x in fig.).
- Fluctuation of junctions are **suppressed**.



Storage modulus  $G$  is **combination** of  $G_c$  and  $G_e$

- Constrained Junction Model
  - Constraints are reduced and  $G$  approaches to  $G_c$ .<sup>a</sup>
- Topological relationships
  - Contribution of entanglement.<sup>b</sup>

$$G_e = T_e G_N^0$$

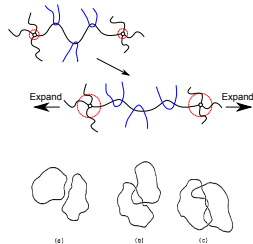


Figure 4. Three topological relationships between two closed loops: (a) not entwined, (b) once entwined, (c) twice entwined.

<sup>a</sup>P.J.Flory, J.Chem.Phys., 66, 12, 5720 (1977)

<sup>b</sup>D.S.Pearson and W.Graessley, Macromol., 11, 3, 528 (1978)

# Recent approach for Constraints (Entanglements)

- Diffused-Constraint Model
  - Confining potential affect all points along the chain.<sup>1</sup>
- Nonaffine Tube Model
  - Improved model of "Edwards' Tube Model".<sup>2</sup>
- Slip-tube Model
  - A pairwise interaction of chains is introduced.<sup>3</sup>

$$f^*(\lambda^{-1}) = G_c + \frac{G_e}{0.74\lambda + 0.61\lambda^{-1/2} - 0.35}$$

$$G_c = \nu k_B T \left(1 - \frac{2}{\phi}\right), \quad G_e = \frac{4}{7} \nu k_B T L$$

$L$  is the number of slip-links per network chain

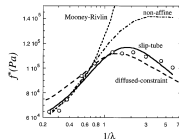


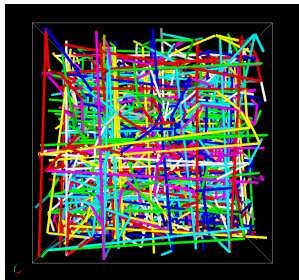
Figure 5. Fit of the data by Pak and Flory<sup>10</sup> on cross-linked poly(dimethylsiloxane) (open circles) by the diffused-constrained model (solid line), Mooney-Rivlin expression (dotted line), nonaffine tube model (dash-dotted line), and the slip-tube model (solid line).

<sup>1</sup> A. Kloczkowski, J.E. Mark, B. Erman, Macromol., 28, 5089 (1995)

<sup>2</sup> M. Rubinstein, S. Panyukov, Macromol., 30, 25, 8036 (1997)

<sup>3</sup> M. Rubinstein, S. Panyukov, Macromol., 35, 6670 (2002)

# ランダムネットワークの絡み合い解析: Z1-code



Z1-code での絡み合い

## ホモポリマーとの比較

- $Z$  は一本鎖あたりの絡み合い
- 今回のネットワークは、  
ホモポリマーと同等

	Homo	4 Chain NW
Segments	50	48
Chains	200	768
Entanglements	204	800
Entangled Chains	134	557
$\langle Z \rangle_{Z1}$	1.02	1.04

## Z1-code とは

- 絡み合いを可視化するアルゴリズム<sup>a</sup>

<sup>a</sup>M. Kröger, Comput. Phys. Commun. 168, 209 (2005)



# 絡み合いの効果について

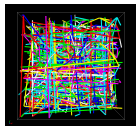
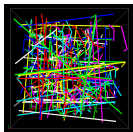
## Entanglement in Slip-tube Model

Rubinstein らの先行研究<sup>a</sup>

$$G_c = \nu k_B T \left(1 - \frac{2}{\phi}\right), \quad G_e = \frac{4}{7} \nu k_B T L$$

and L is the number of slip-links per network chain

<sup>a</sup>M. Rubinstein, S. Panyukov, Macromolecules, 35, 6670 (2002)



	NPT	NVT
Chains, $\nu$	768, 0.018	
$G_c = \nu \times (1 - 2/4)$	0.009	
Entanglements	278	800
Entangled Chains	249	557
$L$	278/768=0.36	800/768=1.04
$G_e = 4/7 \times \nu \times L$	0.004	0.011
$G_{calcd.} = G_c + G_e$	0.013	0.020
$G_{measd.}$	0.013	0.022