

Chapter 5

Amorphous State

Structure of polymers

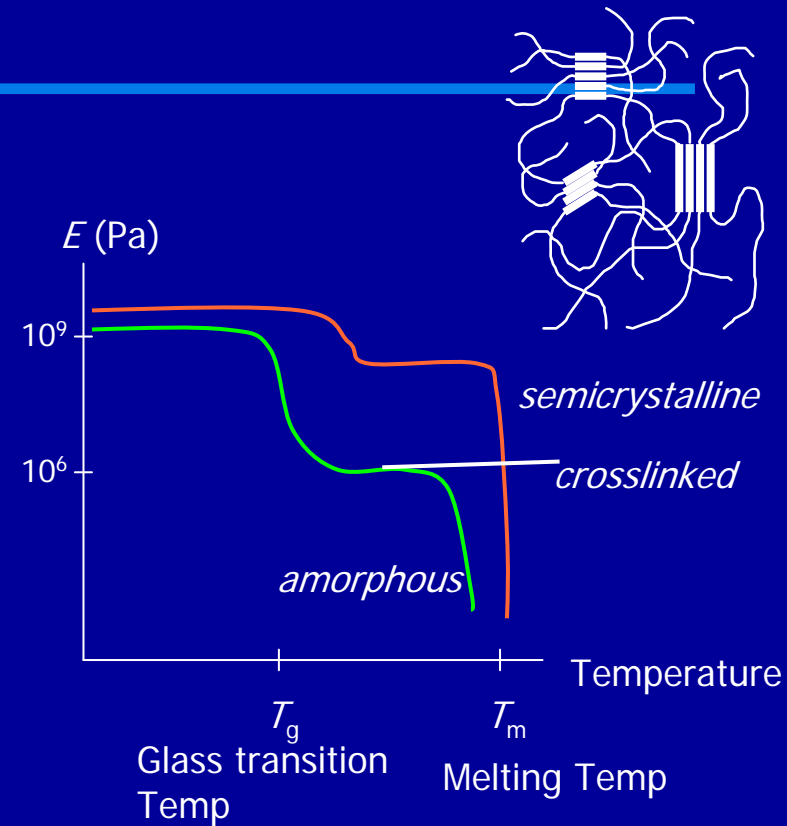
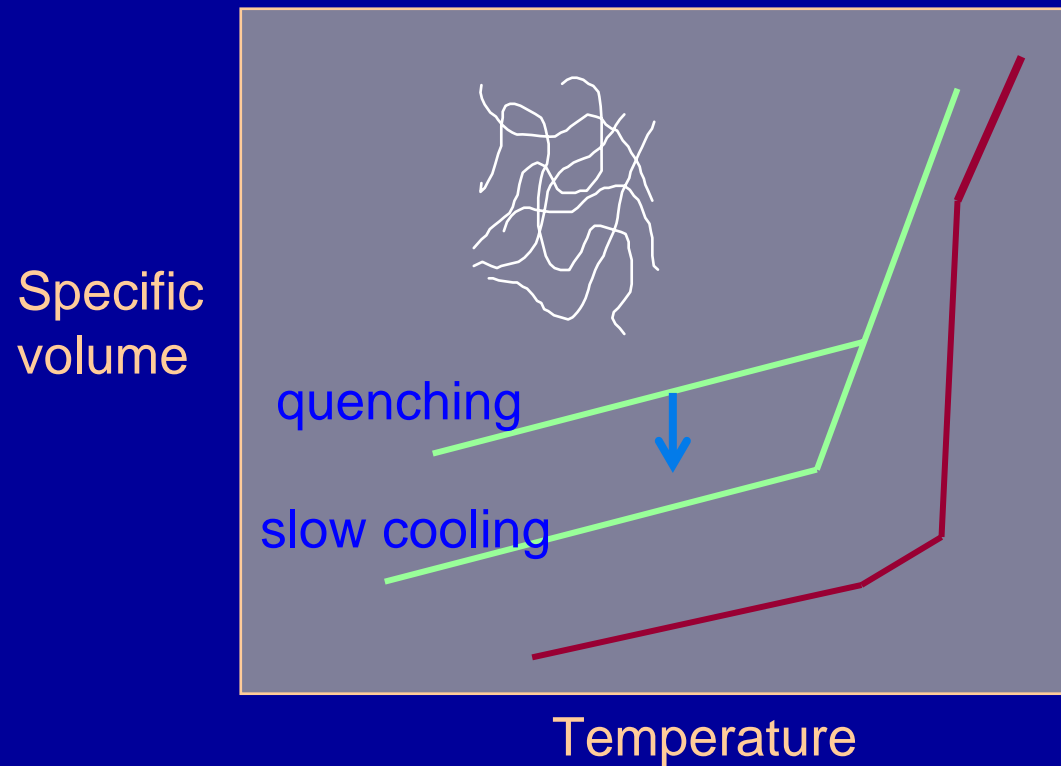
□ 화학적 구조

- 반복단위 구조 atomic structure
- 이성질체 isomers
- 연결 구조 sequence structure

□ 물리적 구조

- 한 사슬의 구조 single chain structure
- 사슬 모임의 구조 aggregation structure
 - » 무정형 구조 amorphous state
 - » 반결정성 구조 semicrystalline state

Polymers upon cooling







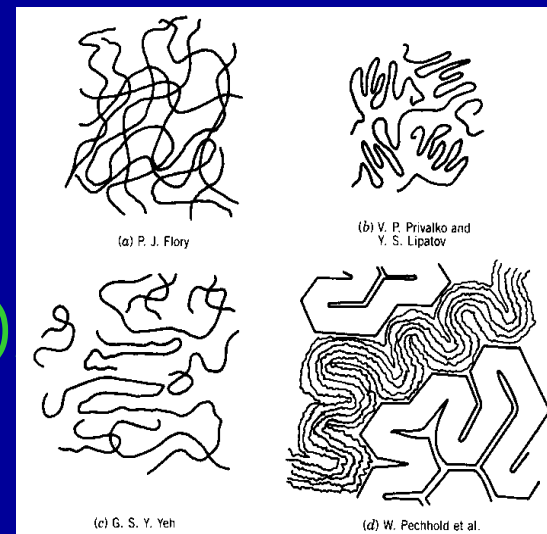
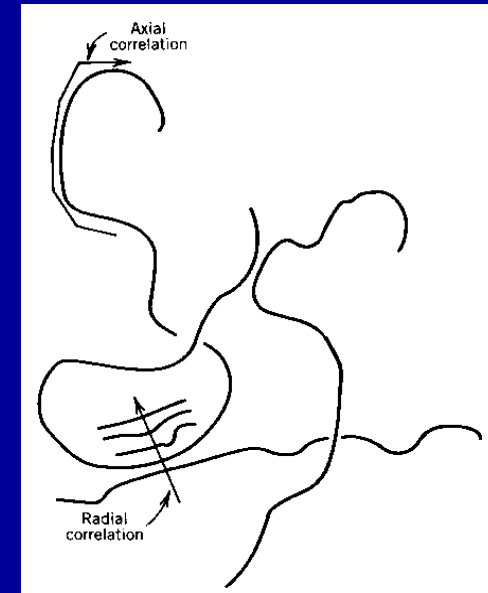
- amorphous vs. semicrystalline
- glass transition, aging, melting, crystallization

Amorphous vs Semicrystalline state

- governing factors
 - chain regularity → crystallizability
 - cooling rate → time for crystallization
- **regular** chain structure
 - » main-chain polymer (PE, nylon, PEster etc)
isotactic or syndiotactic vinyl polymers
 - » semicrystalline when slow-cooled; amorphous upon quenching
- **irregular** chain structure
 - » atactic vinyl polymers
branched polymers
 - » amorphous upon quenching; aged amorphous upon slow-cooling
 - crystallizable when side group is small or crystallizable itself

1&2. Is amorphous completely disordered?

- ❑ Arguing **partial order** ~ parallelization
 - intuition ~ spaghetti ~ space-filing
 - high density (90% of crystal)
 - nodules in electron microscopy
 - lateral (radial) order in ED and WAXS
- ❑ Arguing **complete disorder** ~ random
 - unperturbed dimension by SANS – random coil
 - vanishing radial correlation in WAXS
 - little change in birefringence
 - rubber elasticity works
- **Order should be local, if any.**
- **Short-range order only: No long-range order**
- See discussions  p198-210 (only the results)
 Table 5.6,  p214-216, and  Fig 5.5



3. Single chain structure ~ Conformation

- configuration - breaking single bond

- isomers, copolymers, sequences

- conformation - rotation about single bond

- rotational isomers (conformer, actually)

- Modeling of single chain

- from *artificial* to *real* chain

- freely jointed
- freely rotating
- restricted rotation
- excluded volume effect
- 'Random Coil'

A Polymer Chain

- $n + 1$ atoms
- n bonds (length l)
- $n - 1$ bond angles (angle $180 - \theta$)
- $n - 2$ rotational angles (angle ϕ)

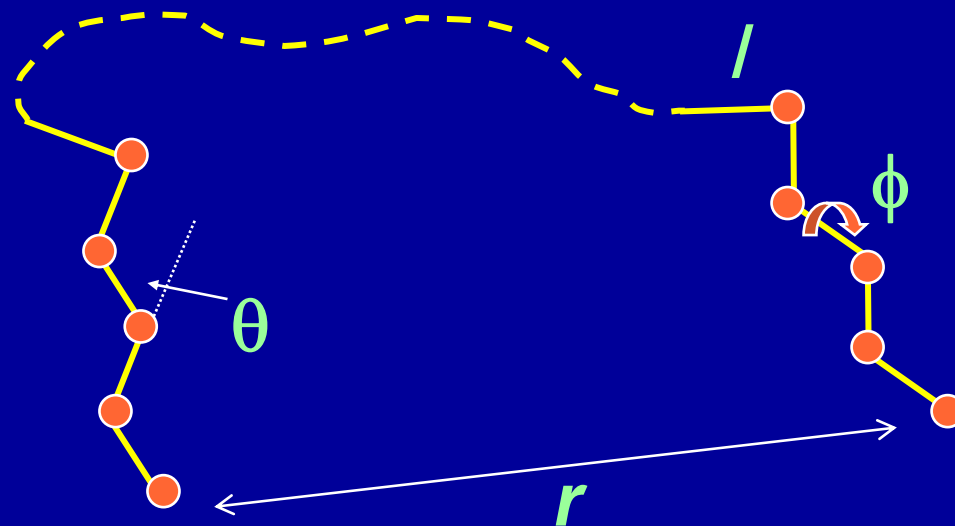


Fig 5.4 p211

- end-to-end distance, r

Freely Jointed Chain

- Bond length only fixed
- Gaussian coil, 'ideal chain'
- mean square end-to-end distance

- a measure of chain dimension

$$\langle r^2 \rangle = n l^2 \quad \text{for FJC}$$

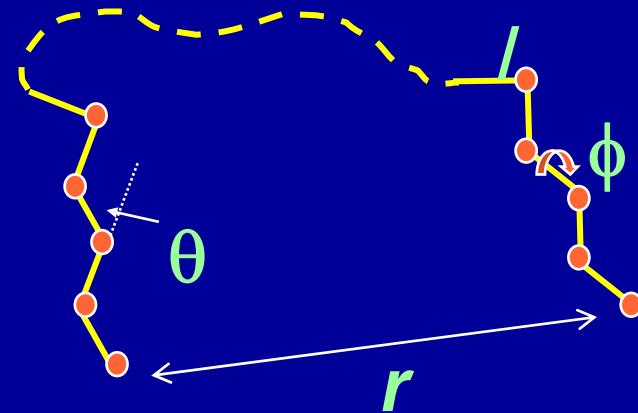
- characteristic ratio

- a measure of chain stiffness

$$C_{\infty} = \langle r^2 \rangle / n l^2 \quad \sim \text{definition} \quad \text{📖 Eqn (5.14) p212}$$

$$C_{\infty} = 1 \quad \text{for FJC}$$

$$C_{\infty} > 1 \quad \text{for } \textit{real} \text{ chains}$$



Freely Rotating Chain

- Bond length and bond angle fixed
 - reduced artificiality

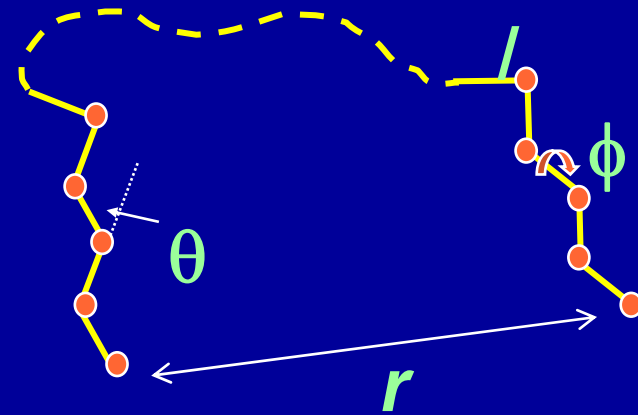
- $\langle r^2 \rangle = n l^2 [(1 + \cos \theta) / (1 - \cos \theta)]$

- For FRC with sp^3 atoms ($\theta = 70^\circ$)

$$\langle r^2 \rangle = 2 n l^2$$

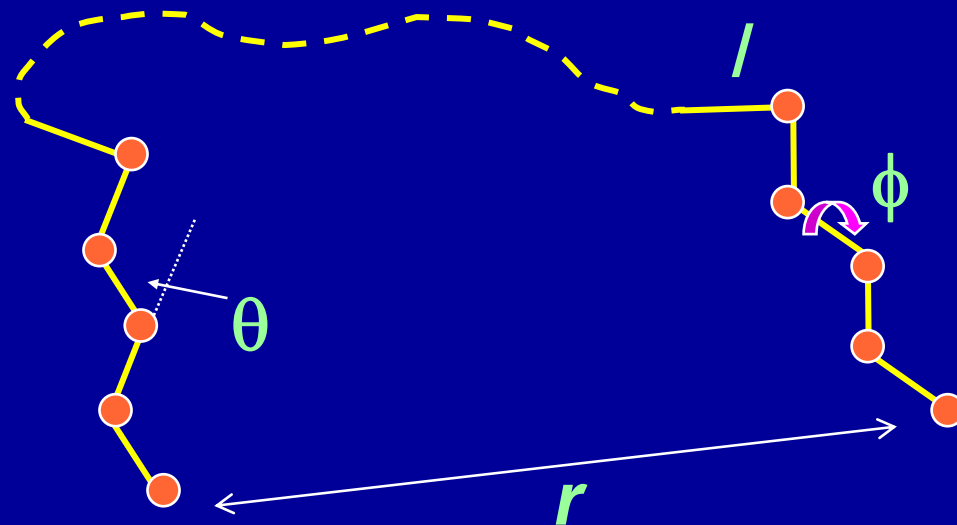
$$C_\infty = \langle r^2 \rangle / n l^2 = 2$$

- FRC is stiffer than FJC, if $\theta < 90^\circ$.



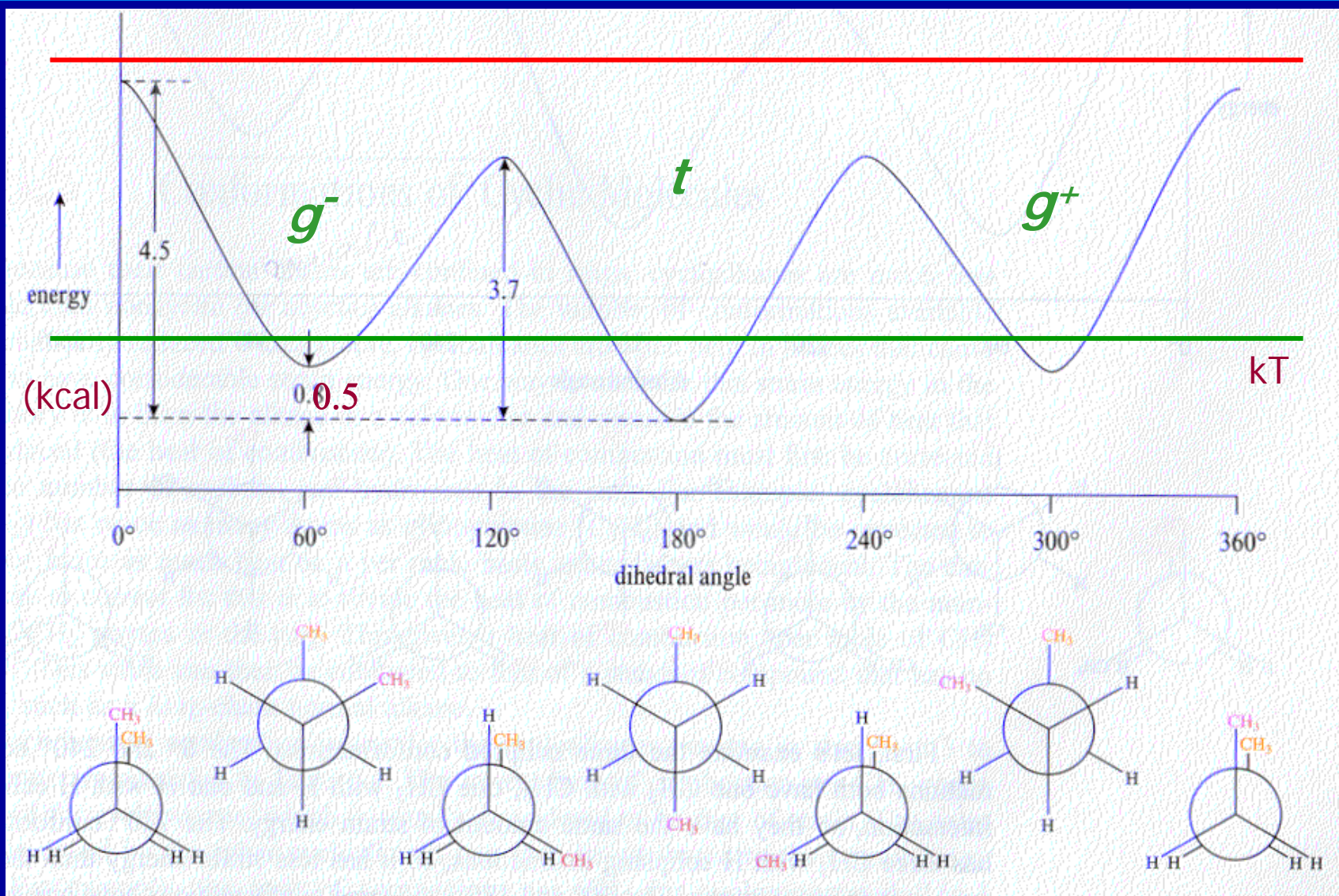
Chain with Restricted Rotations

- Bond length and bond angle fixed
- Not all rotational angles between $0^\circ < \phi < 360^\circ$ allowed.
- Only some discrete states allowed
→ rotational isomeric states (RIS)



Energy map for *n*-butane

Fig 2.11 p49



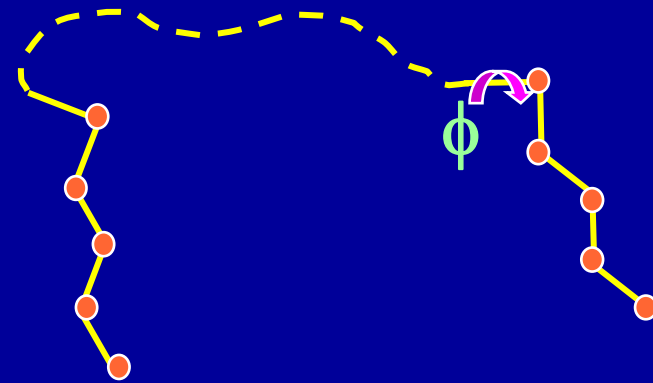
RIS Calculation 1

- 3 RIS (t , g^+ , g) for every bond (except for the first and the last) of the chain
- For a chain with n bonds, $(n - 2) \phi$ gives 3^{n-2} conformations.
- **Statistical weights** for the bonds are determined by Boltzmann distribution (**energy difference** between t , g^+ , and g ; temperature).

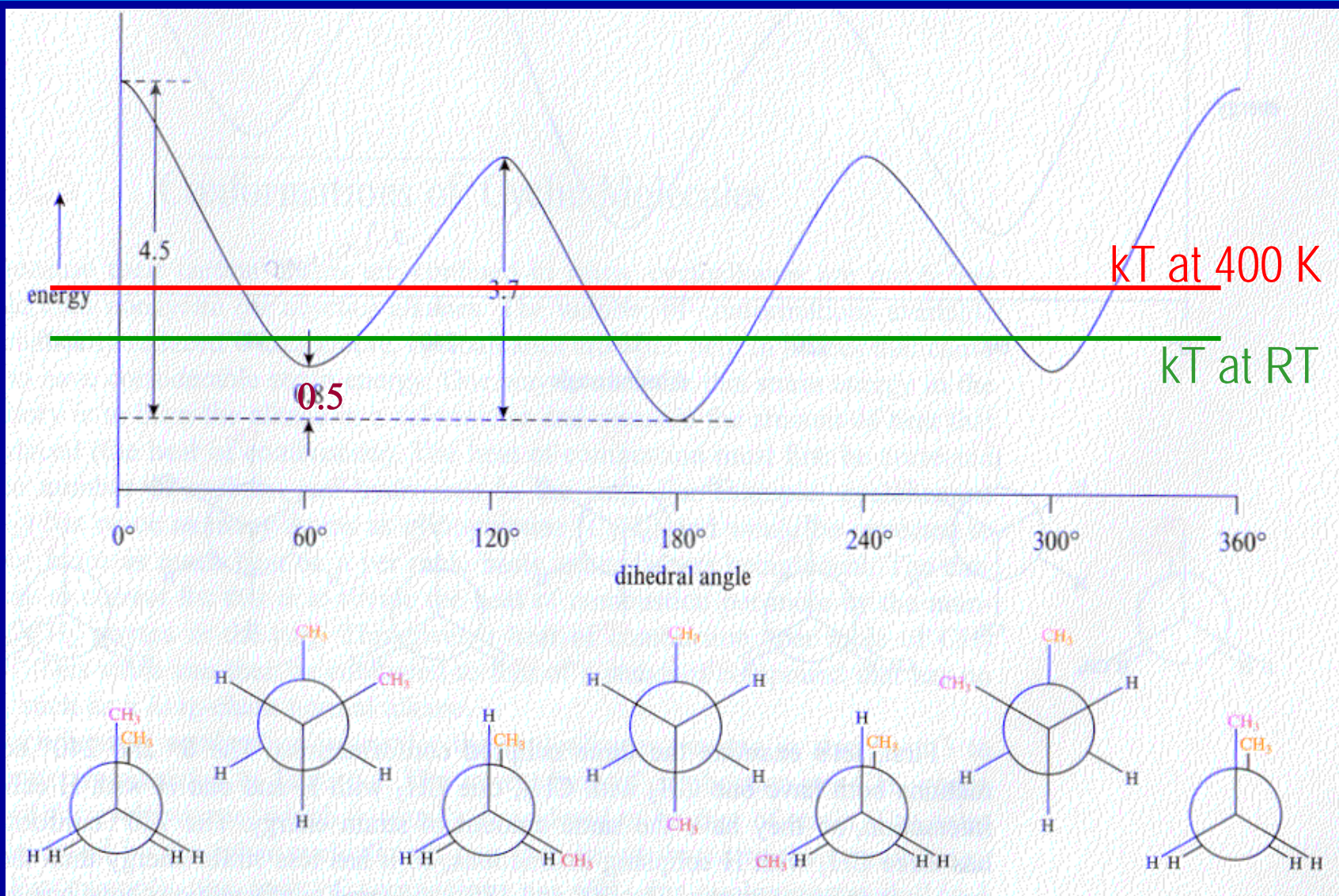
$$u = \exp [-E / kT]$$

$$u(t), u(g^+), u(g)$$

$$\begin{aligned} u(g^+) &= \exp [-E(g^+)/kT] \\ &= \exp [-0.5/0.8] = .53 \text{ at } 400 \text{ K} \end{aligned}$$

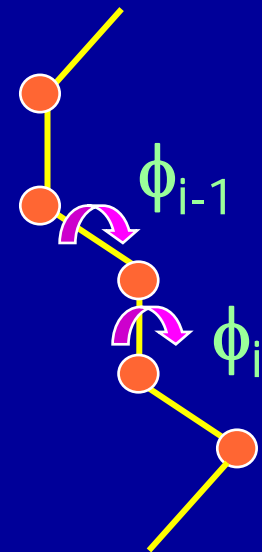


Energy map for *n*-butane



RIS Calculation 2

- Interactions governing **energy**
 - non-bonded interaction (dispersion)
 - dipole interaction
 - intrinsic rotational potential
- Pairwise potential
 - 1st-order interaction
 - » 1-bond rotation ~ 3-bond interaction
 - » t, g^+, g
 - 2nd-order interaction
 - » 2-bond rotation ~ 4-bond interaction
 - » $tt, tg^+, tg, g^+g^+, gg, g^+g$
 - » pentane interaction, **short-range interaction**
 - » Bond rotations should be considered pairwise.



RIS Calculation 3

- By taking 2nd moment (average size) of all the conformations, the chain dimension is calculated.

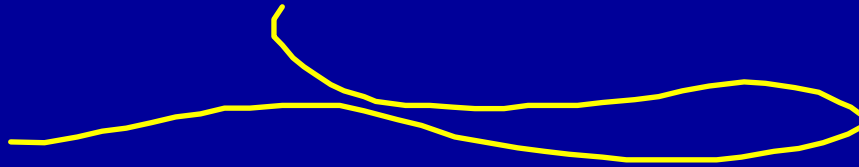
- $C_{\infty} = \langle r^2 \rangle / n l^2 = 5 - 10$  Table 5.7 p212
for chains with restricted rotation

 Eqn (5.13) is totally **wrong**.

1. $\langle \cos \phi \rangle = \sum u \cos \phi / \sum u$
2. 2nd order interaction not considered

- Compare with the experimental results (viscosity, light scattering, SANS, etc.) and adjust parameters

Excluded Volume Effect

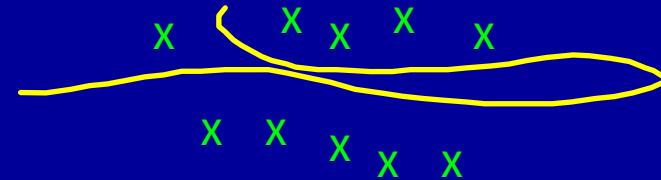


□ Intrachain volume exclusion

- long-range interaction
- gives larger dimension

□ $\langle r^2 \rangle_{EV} = \alpha \langle r^2 \rangle_0$

- in good solvent, $\alpha > 1$
 - » Repulsion(p-p) > Repulsion(p-s)
- in poor solvent, $\alpha < 1$
 - » Repulsion(p-p) < Repulsion(p-s)
- in theta (Θ) condition, $\alpha = 1$ ~ unperturbed (θ)
 - » Repulsion(p-p) = Repulsion(p-s)



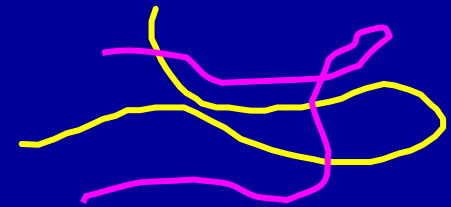
$$\alpha^5 - \alpha^3 = C n^{1/2} \psi (1 - \theta/T)$$

Eqn (3.29) p84

Real Chain in Bulk – 'random coil'

□ In bulk amorphous state

- Repulsion(p-p, **intra**) = Repulsion(p-p, **inter**)
- unperturbed state ($\alpha = 1$)
- 'phantom' or 'ghost' chain
- $\langle r^2 \rangle = \langle r^2 \rangle_0 = r_{\theta}^2 = \langle r^2 \rangle_{\text{RIS}}$
- proposed by Flory; proved by SANS expt
 - also in the **melt state**
 - *probably* also in the **semicrystalline state**





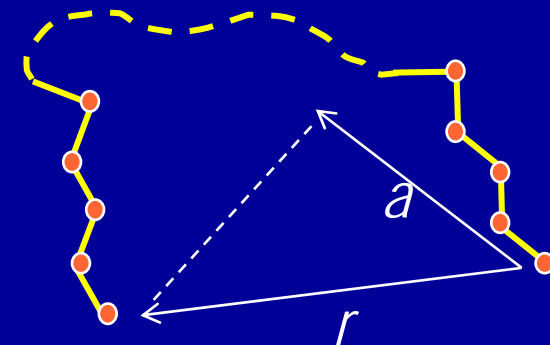
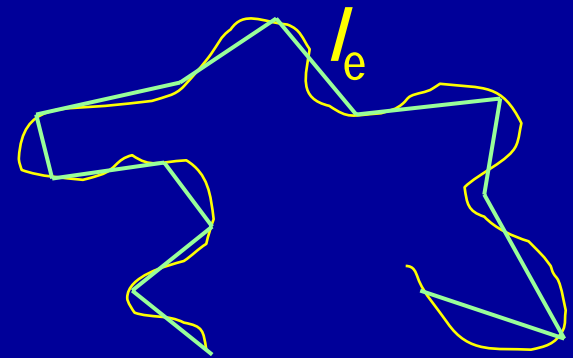
□ RIS model describes the state of single chain in bulk (amorphous, semicrystalline)

- crystal structure ~ conformation with the lowest energy

$$a = l(C + 1)/2 = l_e / 2 \text{ 확인}$$

Kuhn chain and Persistence length

- Kuhn chain = (Statistically) equivalent (freely jointed) chain
- Kuhn (chain) length, l_e
 - $n_e l_e^2 = \langle r^2 \rangle_0 = r_\theta^2 = C_\infty n l^2$
 - $n_e l_e = L$ (contour length)
 - $l_e = r_\theta^2 / L$  Eqn (5.15) p213
 - A measure of axial correlation length
 - For PE, $l_e = 8 l$, $n_e = 0.1 n$  Fig 5.1 p201
- Persistence length, a
 - Projection of r to a bond
 - A measure of axial correlation length also
 - For PE, $a \sim 4 l$



Examples of RIS Application 1

□ Polyethylene

- $\langle r^2 \rangle_0 = 6.8 n l^2$ at 400 K

$$\sigma = u(g^+)/u(t) = 0.5, \omega = u(g^+g)/u(tt) = 0.05$$

$$P(t) = 0.62, P(g^+) = P(g) = 0.19$$

- temperature coefficient

$$d[\ln \langle r^2 \rangle_0] / dT < 0$$

- preferred conformation

all-trans (with the lowest energy)

planar zigzag in crystal structure

Examples of RIS Application 2

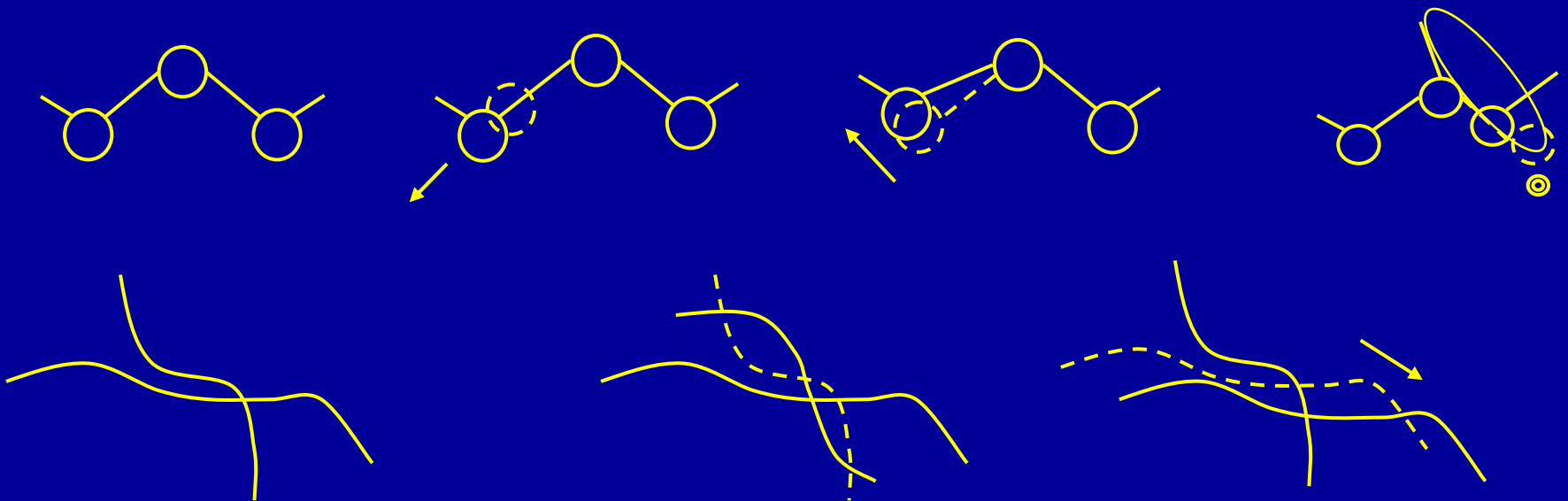
□ Polyoxymethylene

- $\langle r^2 \rangle_0 = 8 n l^2$
 $\sigma = 10, \omega = 0.05$
- $d[\ln \langle r^2 \rangle_0] / dT > 0$
- preferred conformation
 $g^+g^+g^+ \dots$ or $g^-g^-g^- \dots$
 2_1 helix in crystal structure


4. Macromolecular Dynamics




□ Motions in polymers

- Deformation in bond angle and length ~ **elastic**
- Change in conformation ~ segmental motion ~ **viscoelastic**
- Translational motion ~ **viscous**



Models for macromolecular dynamics

- Rouse (– Bueche – Zimm) model  Fig 5.6
 - Bead and spring
 - Single chain, dilute solution

- Reptation (de Gennes (– Doi – Edward)) model  Fig 5.7
 - Effect of entanglements
 - Chain in a tube  Fig 5.9
 - Conc. solution, bulk
 - $\eta = K M^\beta$
 - Effect of branch  Fig 5.10