

►Higher frequency (elastic) gives higher T_g . (See [stress-strain curve at different extension rate.](#))

►Empirically

$$T_g = 0.6 T_m \quad (2.4)$$

►Remember the T_g of copolymer (Fox eq) – Inverse additivity

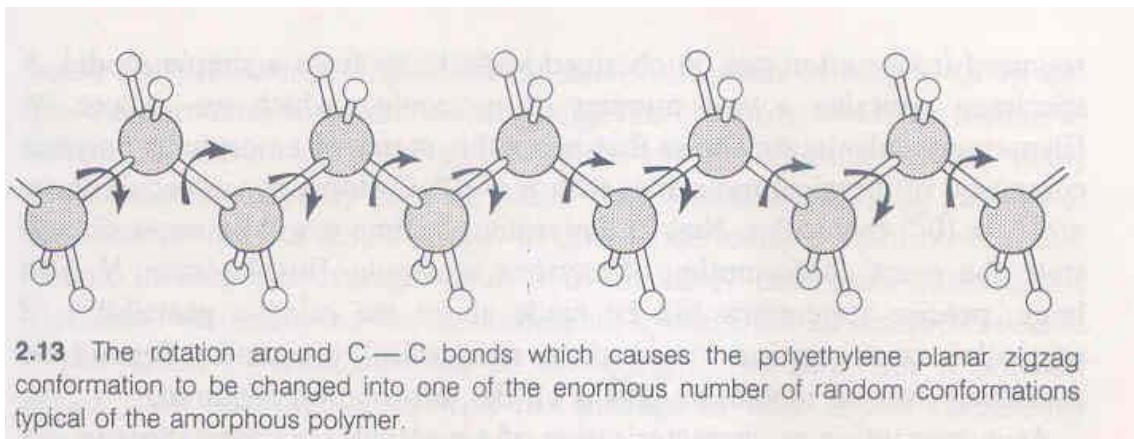
2.7 Molecular Conformation in the Amorphous Polymer

►Molecular conformation in crystal:

Well defined by the crystalline structure.

Ex. Fully extended planar zig-zag conformation, being caged in by its neighbors (See [Figure 2.1b](#)) for PE.

►In amorphous state (glassy, rubbery state) or in dilute solution, molecule assumes a random conformation (See the rotation C—C b in [Figure 2.13](#)).



Number of random conformation (Figure 2.14)

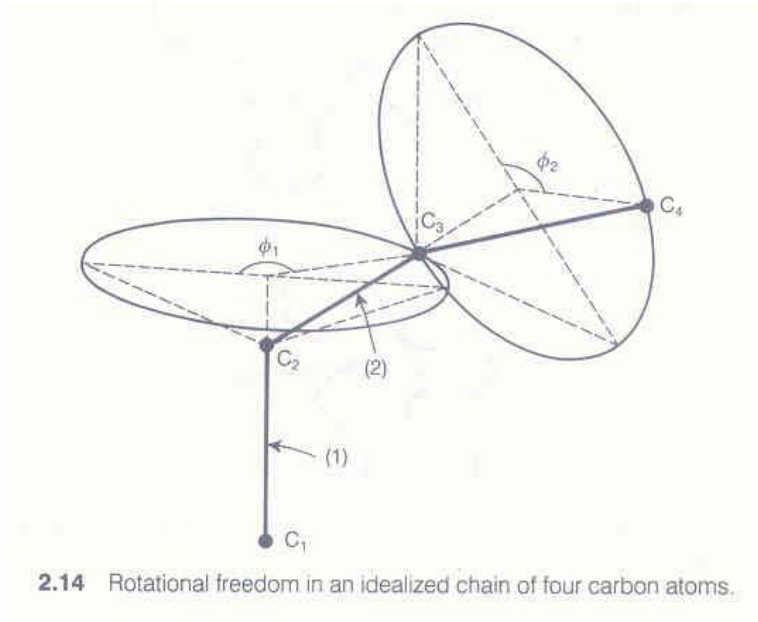


Figure 2.14

Assume bond (1) is fixed.

Position of C₃ = $f(\phi_1)$, $0 \leq \phi_1 \leq 2\pi$

Position of C₄ = $f(\phi_1, \phi_2)$, $0 \leq \phi_2 \leq 2\pi$

.....

Position of C_n = $f(\phi_1, \phi_2, \dots, \phi_{n-2})$, $0 \leq \phi_{n-2} \leq 2\pi$

The position of last C atom (C_n) depends on $\phi_1, \phi_2, \dots, \phi_{n-2}$, each varies from 0 to 2π .

Typical value of n = $O(10^3) \sim O(10^4)$

► Number of random conformation $\rightarrow \infty$

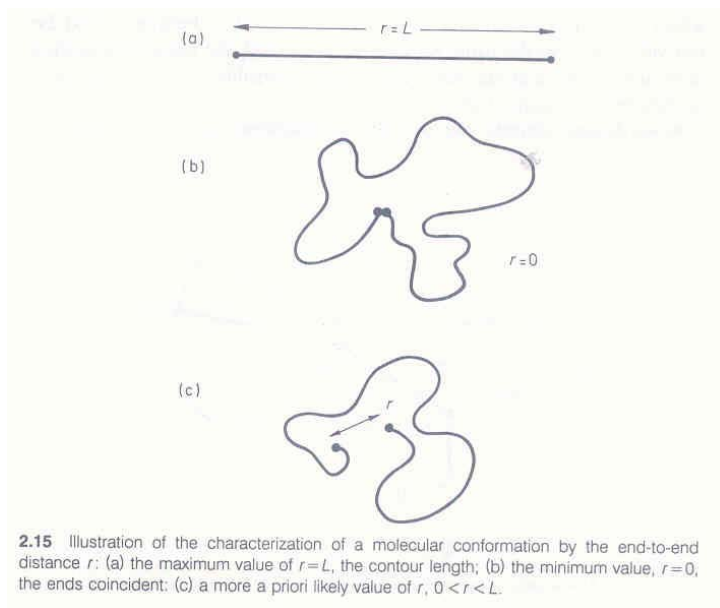
So, statistical analysis applies.

Ideas for polymer molecular size

► Radius of gyration

$$\langle s^2 \rangle = \langle \frac{r^2}{6} \rangle$$

► End to end distance (r) (See [Figure 2.15](#))



$$0 \leq r \leq L \text{ (=Contour length, fully extended length)}$$

We are interested in the average value of r .

Two types of average r :

- i) Time average of one polymer molecule (rubbery, melt state)
- ii) Average over N molecules at a specific instance (glassy polymer)

For statistically large number of samples, i) = ii)

2.8 Freely Jointed Chain

Contour length

$$L = n l \quad \circ \circ (2.5)$$

$n = \#$ of segment

$l =$ length of one segment (Same for all segments.)

The n rigid segments are joined flexibly one to another ($0 \leq \theta_{ij} \leq 180^\circ$)

(θ_{ij} = bond angle between segments i and j).

Then the end-to-end vector

$$\mathbf{r} = \sum \mathbf{l}_i \quad \circ \quad (2.6)$$

Magnitude of a vector is obtained by taking its scalar product.

So,

$$\begin{aligned} r^2 &= \mathbf{r} \cdot \mathbf{r} = (\mathbf{l}_1 + \mathbf{l}_2 + \dots + \mathbf{l}_n) \cdot (\mathbf{l}_1 + \mathbf{l}_2 + \dots + \mathbf{l}_n) \\ &= (\mathbf{l}_1 \cdot \mathbf{l}_1 + \mathbf{l}_2 \cdot \mathbf{l}_2 + \dots + \mathbf{l}_n \cdot \mathbf{l}_n) \\ &\quad + 2(\mathbf{l}_1 \cdot \mathbf{l}_2 + \mathbf{l}_1 \cdot \mathbf{l}_3 + \dots + \mathbf{l}_n \cdot \mathbf{l}_{n-1}) \\ &= nl^2 + 2l^2(\cos\theta_{12} + \cos\theta_{13} + \dots + \cos\theta_{n, n-1}) \quad (2.7) \end{aligned}$$

(Note $l_1 = l_2 = \dots = l_n$)

For N molecules, the average r is obtained as:

$$\begin{aligned} \langle r^2 \rangle &= \frac{1}{N} \sum_1^N [nl^2 + 2l^2(\cos\theta_{12} + \cos\theta_{13} + \dots + \cos\theta_{n, n-1})] \\ &= nl^2 + \frac{2l^2}{N} \sum_1^N (\cos\theta_{12} + \cos\theta_{13} + \dots + \cos\theta_{n, n-1}) \quad (2.8) \end{aligned}$$

Note that

$$\sum_1^N (\cos\theta_{12} + \cos\theta_{13} + \dots + \cos\theta_{n, n-1}) = 0 \quad (2.9)$$

\therefore Angle was taken randomly in the range of $0 \leq \theta \leq 180^\circ$ (See the cosine curve for the range.).

Then,

$$\langle r^2 \rangle = nl^2 \quad (2.10)$$

The root mean square end-to-end distance (R)

$$R = \langle n l^2 \rangle^{1/2} = n^{1/2} l \quad (2.11)$$

The ratio of R to the contour length

$$\frac{R}{L} = \frac{n^{\frac{1}{2}} l}{n l} = \frac{1}{n^{\frac{1}{2}}} \quad (2.12)$$

As n increases ratio of end-to-end distance to contour length becomes small - The molecules exist in highly coiled state!!!

Note that Eq (2.10) in general form is:

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

C=1 for Freely jointed model

C≥2 for Freely rotating model

C≥3 for Hindered rotating model

Freely rotating model (See details in Lecture # 203)

Bond angle is fixed. ($\tau \sim 110^\circ$ for C-C bond)

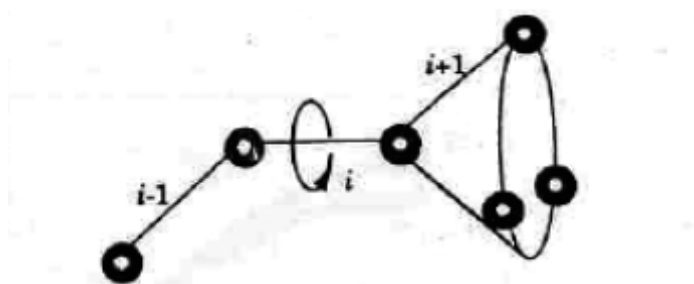


Figure 2.11 Three different rotational isomers are generated by torsion about bond i . If the three states are equally populated, the average vector of bond $i + 1$ has no component perpendicular to a vector parallel to the i th bond.

$$\begin{aligned}\langle r^2 \rangle &= nl^2 \left(1 + \frac{2\alpha}{1-\alpha} \right) = nl^2 \left(\frac{1+\alpha}{1-\alpha} \right) \\ &= nl^2 \frac{1 + \cos(180 - \tau)}{1 - \cos(180 - \tau)} \quad (2.19)\end{aligned}$$

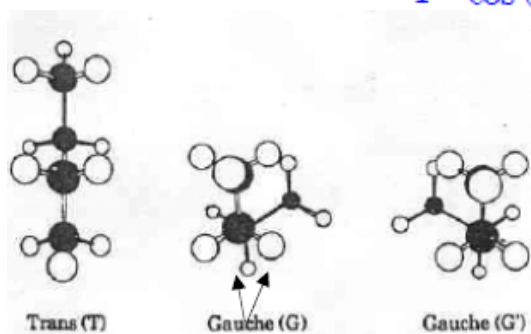


Figure 2.3 Conformational states of *n*-butane. Note that the views of the gauche conformers are along the middle carbon-carbon bond. Carbon – shaded; hydrogen – white.

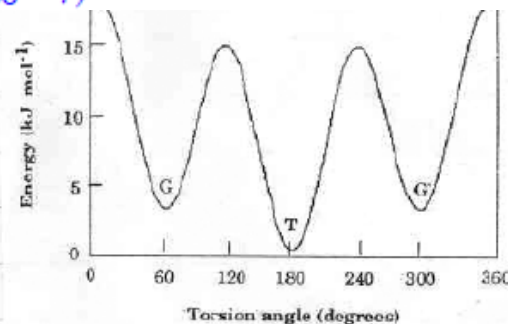


Figure 2.4 Conformational energy of *n*-butane as a function of torsion angle of the central carbon-carbon bond. The outer carbon-carbon bonds are assumed to be in their minimum energy states (staggered positions).

Subbing $\tau \sim 110^\circ$ gives the value of the second term of RHS becomes approx 2.

Hindered rotating model

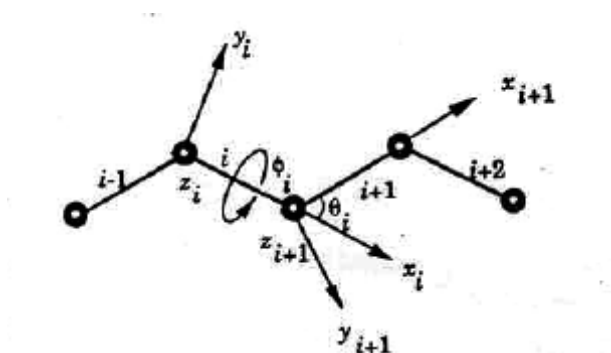


Figure 2.12 Definition of CS_i and CS_{i+1} ; the torsion angle ϕ is set to zero for the planar trans conformation. Axis y_i is in the plane of bonds i and $i-1$, and z_i completes the right-handed Cartesian coordinate system.

$$\langle r^2 \rangle = nl^2 \left(\frac{1 + \cos(180 - \tau)}{1 - \cos(180 - \tau)} \right) \left(\frac{1 + \langle \cos \phi \rangle}{1 - \langle \cos \phi \rangle} \right) \quad (2.29)$$

$\langle \cos \phi \rangle =$ origin of temperature dependence

This model assumes three rotational isomers, viz. T, G, and G'.