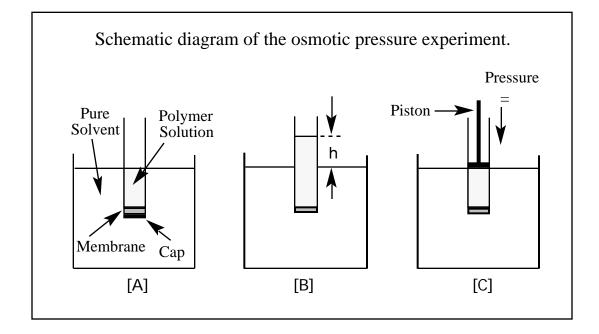
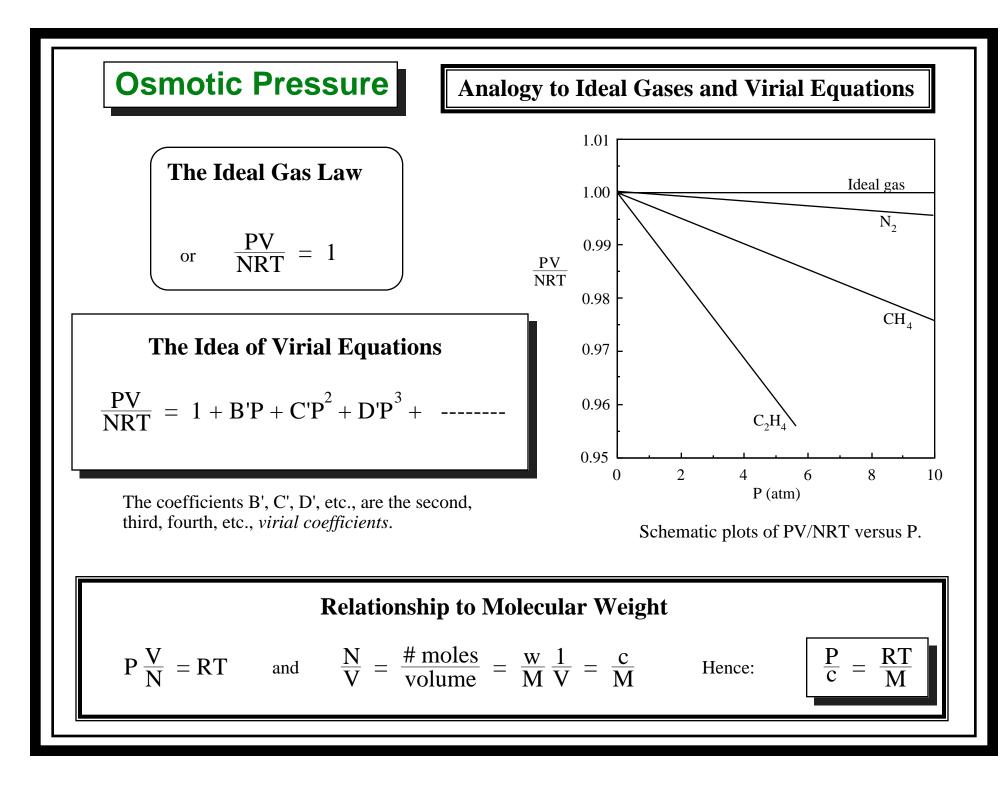
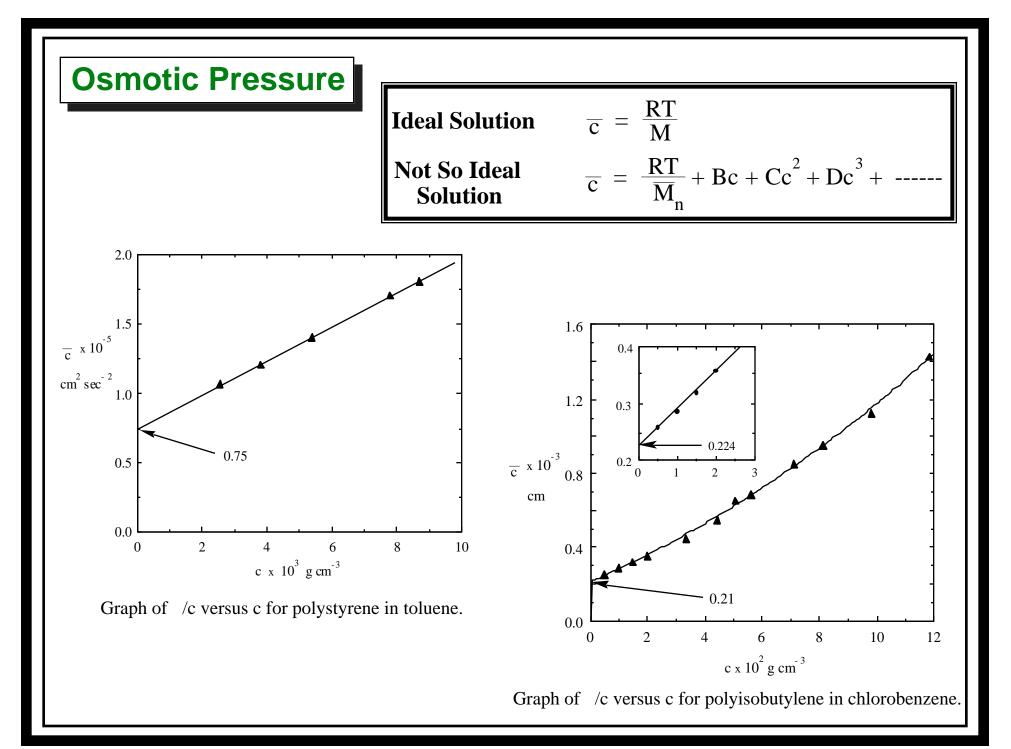


## **Osmotic Pressure**

Osmotic pressure—belongs to a family of techniques that come under the heading of *colligative property measurements*.







## **Osmotic Pressure**

#### **Derivation of a Virial Equation from the Flory–Huggins Equation**

Osmotic Pressure can be related to the chemical potential via the Flory—Huggins equation:

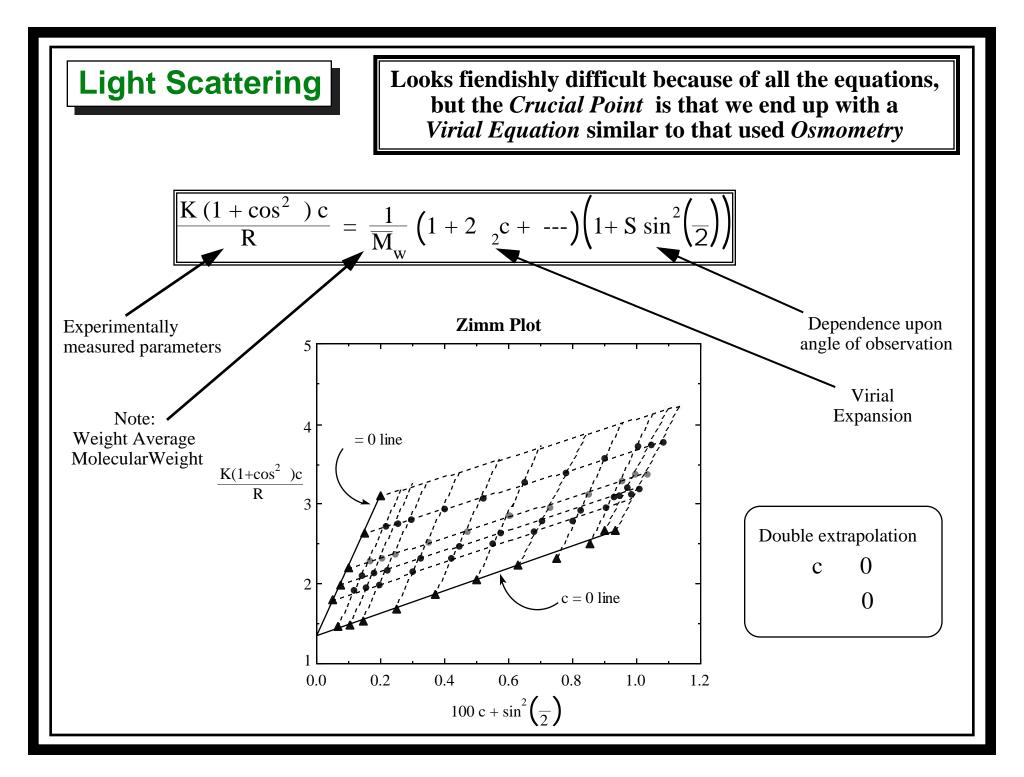
$$\frac{\mu_{s} - \mu_{s}^{0}}{RT} = \ln_{s} + \left(1 - \frac{1}{\overline{M}_{n}}\right)_{p} + \frac{2}{p}$$

a

and:  
= 
$$-\frac{RT}{V_s} \left[ \ln_s + p \left( 1 - \frac{1}{\overline{M}_n} \right) + p \right]$$

Expanding the Ln term:  $\ln_{s} = \ln (1 - p) = -p - \frac{p}{2} - \frac{p}{3} - \cdots$ Leads to:  $= \frac{\mathrm{RT}}{\mathrm{V}_{\mathrm{s}}} \left| \frac{\mathrm{p}}{\mathrm{\overline{M}}_{\mathrm{n}}} + \frac{\mathrm{p}}{\mathrm{p}} \left( \frac{1}{2} - \right) + \frac{\mathrm{p}}{3} + \dots \right|$ 

This has the same form as the virial equation, but uses the concentration variable <sub>p</sub> instead of c. However, we must be careful because the Flory-Huggins theory does not strictly apply to dilute solutions.



#### Measuring the Viscosity of Polymer Solutions

Most common method used to determine the viscosity of a polymer solution is to measure the time taken to flow between fixed marks in a capillary tube under the draining effect of gravity. The (volume) rate of flow, , is then related to the viscosity by Poiseuille's equation:

$$= \frac{\mathbf{P} \mathbf{r}^4}{\mathbf{8} \mathbf{l}}$$

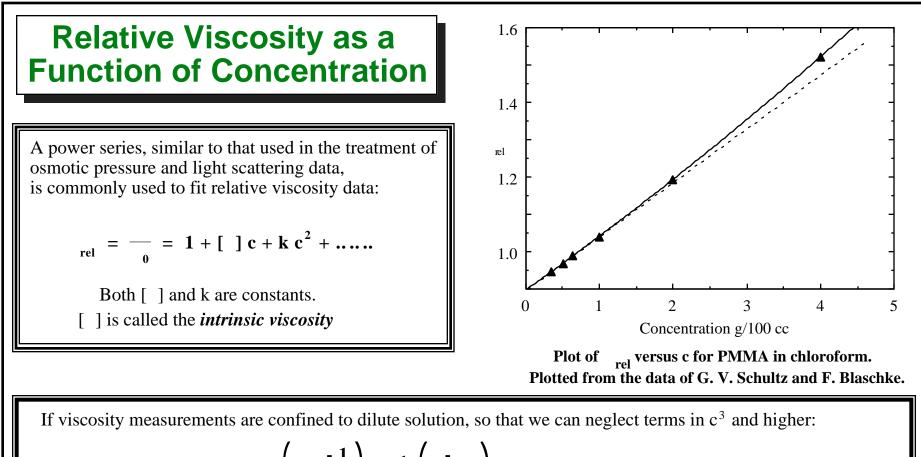
where P is the pressure difference maintaining the flow, r and l are the radius and length of the capillary and is the viscosity of the liquid.

#### **Relative Viscosity**

Defined as the viscosity of a polymer solution divided by that of the pure solvent and for dilute solutions:

$$_{\rm rel} = \frac{1}{0} \frac{t}{t_0}$$

where t is the time taken for a volume V of solution (no subscript) or solvent (subscript 0) to flow between the marks.



$$\left(\frac{-rel}{c} - \frac{1}{c}\right) = \frac{1}{c} \left(\frac{-0}{0}\right) = \begin{bmatrix} 1 + k \end{bmatrix} + k$$

The *Specific Viscosity* is defined as:  $_{sp} = _{rel} - 1$ 

Note also that as c goes to zero (infinite dilution), then the intercept on the y-axis of a plot of  $\binom{sp}{c}$  against c is the intrinsic viscosity, []:

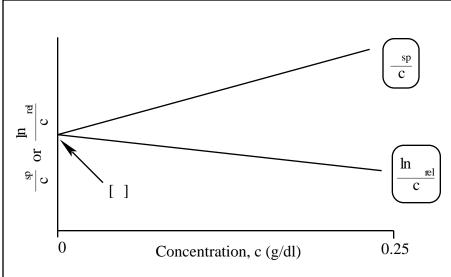
$$\begin{bmatrix} \end{bmatrix} = \left(\frac{sp}{c}\right)_{c} = 0$$

#### Measuring the Intrinsic Viscosity

In practice, we use two semi-empirical equations suggested by Huggins and Kraemer

$$\frac{sp}{c} = [] + k' []^2 c$$

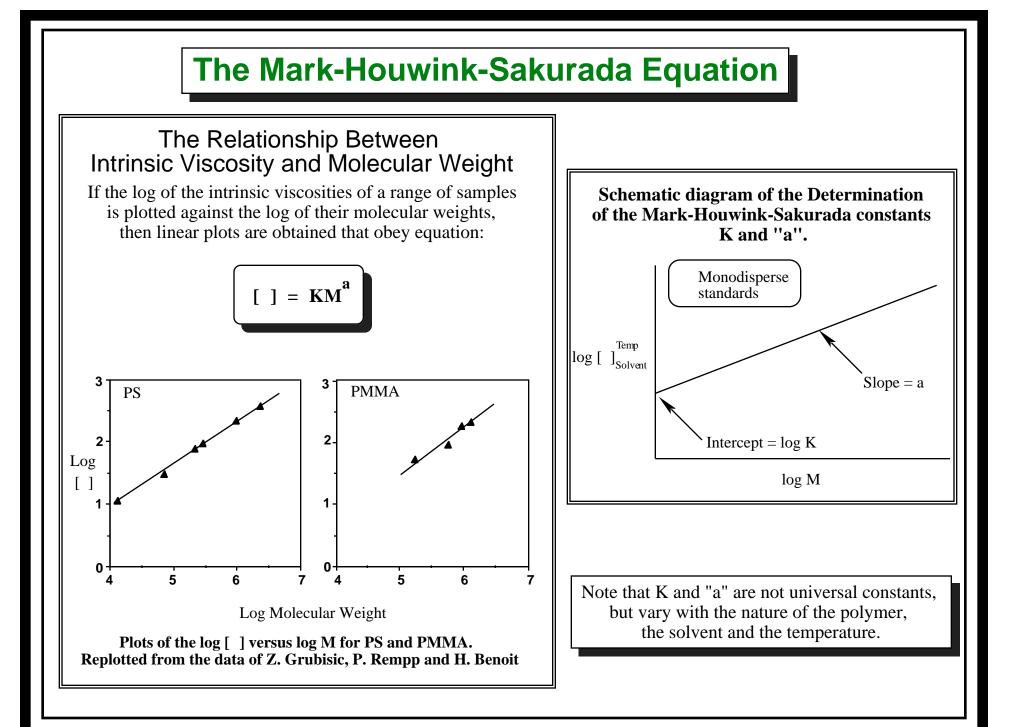
$$\frac{\ln}{c} = [] + k''[]^2 c$$



Schematic diagram illustrating the graphical determination of the intrinsic viscosity.

Schematic diagram illustrating the effect of strong intermolecular interactions.

Most "extrapolation to zero concentration" procedures have a serious limitation. Where one would like to perform measurements is at the lowest concentrations possible, but this is generally where the greatest error in measurement occurs.



#### The Viscosity Average Molecular Weight

For Osmotic Pressure and Light Scattering we saw that there is a clear relationship between experimental measurement and the number and weight molecular weight average, respectively. Viscosity measurements are related to molecular weight by a semi-empirical relationship and a new average, the Viscosity Average for polydisperse polymer samples is defined.

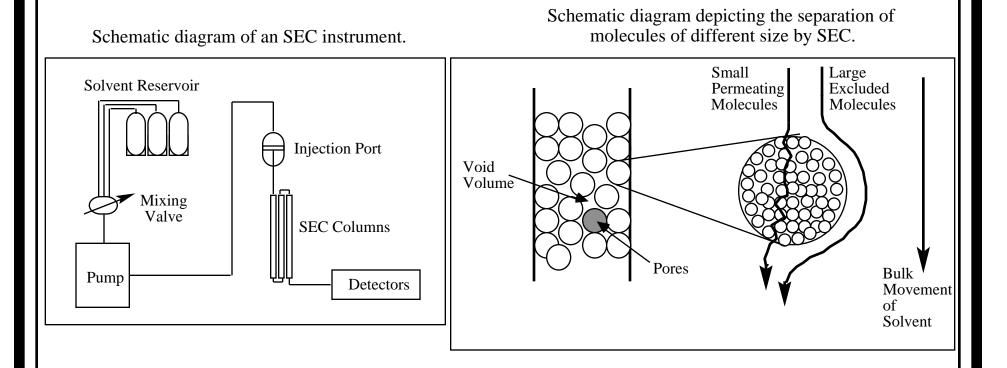
In very dilute solutions 
$$_{sp} = \left( \begin{array}{c} _{sp} \right)_{i}$$
  
Now:  $\frac{\left( \begin{array}{c} _{sp} \right)_{i}}{c_{i}} = K M_{i}^{a}$  Hence:  $_{sp} = K M_{i}^{a} c_{i}$  And:  $[] = \frac{_{sp} - sp}{c} = \frac{K M_{i}^{a} c_{i}}{c}$ 

By substitution and rearranging we obtain:

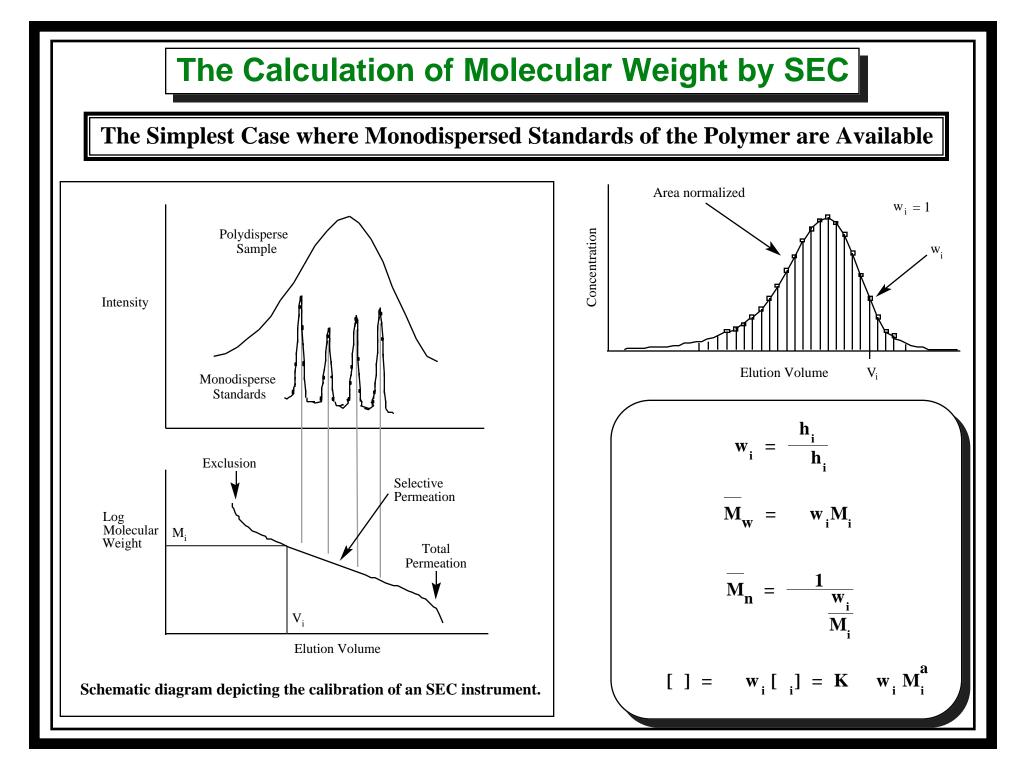
$$\overline{\mathbf{M}}_{\mathbf{v}} = \begin{bmatrix} \mathbf{N}_{i} \mathbf{M}_{i}^{(\mathbf{a}+1)} \\ \frac{\mathbf{i} & \mathbf{N}_{i} \mathbf{M}_{i}}{\mathbf{N}_{i} \mathbf{M}_{i}} \end{bmatrix}^{\frac{1}{\mathbf{a}}}$$

Note that the Viscosity Molecular Weight is <u>Not</u> an Absolute Measure as it is a function of the solvent through the Mark-Houwink parameter "a".

## Size Exclusion (or Gel Permeation) Chromatography



For a given volume of solvent flow, molecules of different size travel different pathlengths within the column. The smaller ones travel greater distances than the larger molecules due to permeation into the molecular maze. Hence, the large molecules are eluted first from the column, followed by smaller and smaller molecules.



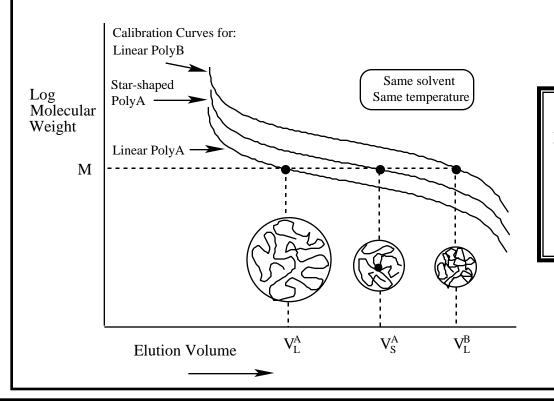
# **How Does SEC Separate Molecules ?**

If the molecular weight of monodisperse polystyrenes of different molecular architecture(e.g., linear, star-shaped, comb-like, etc.) are plotted against elution volume they do not fall on a single calibration curve.

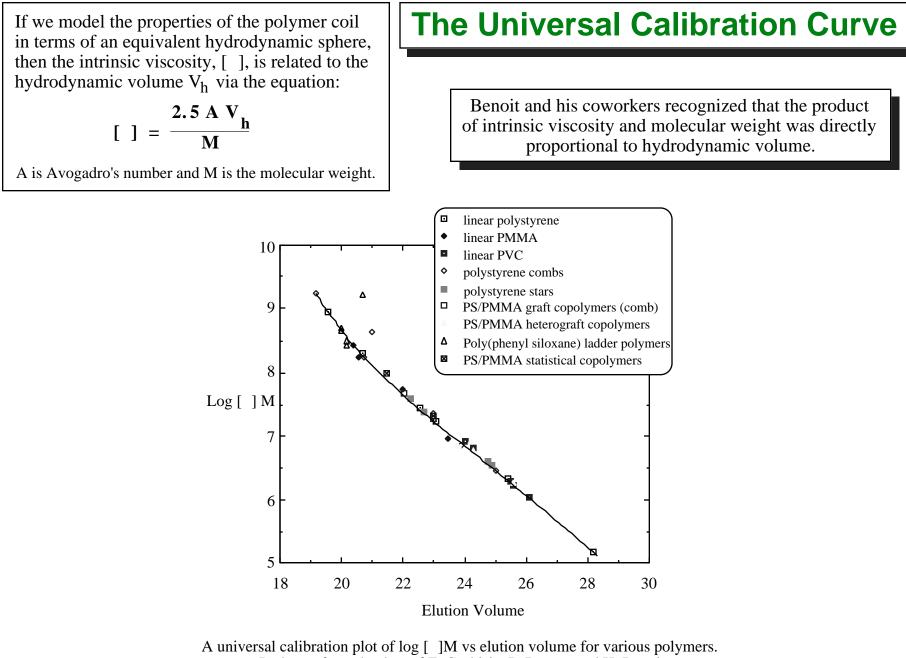
In other words, if we had three *monodisperse* polystyrenes, one linear, one star -shaped and one comb-like, *all with the same molecular weight*, they would not come off the column at the same time (elution volume).

Similarly, *different* monodisperse polymers <u>of the same molecular weight</u> generally elute at different times. Thus, for example, monodisperse samples of polystyrene and PMMA <u>having the same molecular weight</u> might come off the column at different times.

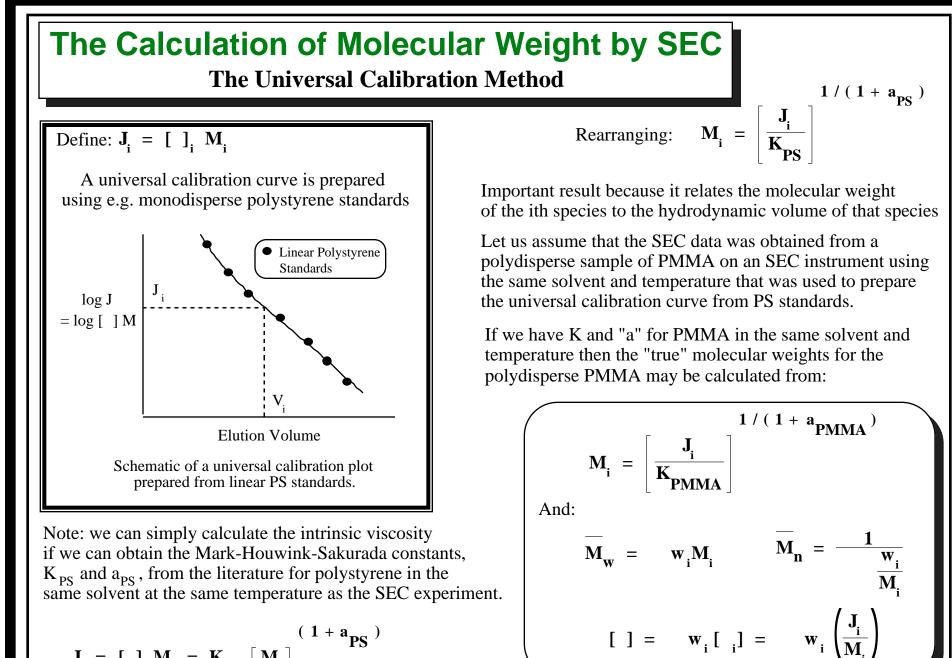
In effect, this means we would require different calibration curves for different polymers and even the same type of polymer if the architecture is different.



Benoit and his coworkers recognized that SEC separates not on the basis of molecular weight but rather on the basis of hydrodynamic volume of the polymer molecule in solution.

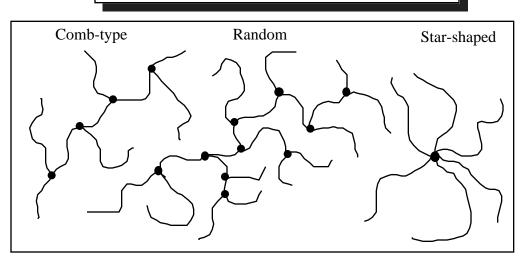


Redrawn from the data of Z. Grubisic, P. Rempp and H. Benoit.



$$\mathbf{J}_{i} = []_{i} \mathbf{M}_{i} = \mathbf{K}_{PS} [\mathbf{M}_{i}]$$

## Long Chain Branching



Schematic representation of different long chain branches.

Long chain branching can have a major effect upon the rheological and solution properties of polymers. Difficult to quantitatively determine the amount of long chain branching using conventional analytical techniques, such as NMR or vibrational spectroscopy.

Very low concentration of any species that can be attributed to the presence of a long chain branch.

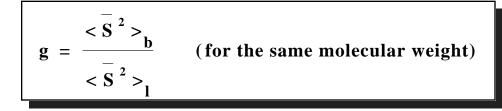
АААААААААААААААААА
B
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B
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#### Long Chain Branching and Mean Square Dimensions

The introduction of only one or two long chain branch points leads to a significant decrease in the mean-square dimensions of macromolecules compared to linear molecules of the <u>same molecular weight</u>.

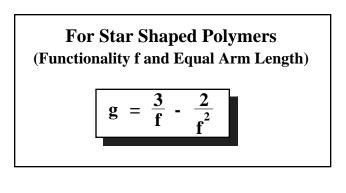
This statement may be expressed in terms of the ratio of the respective radii of gyration, g.

B. H. Zimm and W. H. Stockmayer, J. Chem. Phys., 17, 1301 (1949).
B. H. Zimm and R. W. Kilb, J. Polvm. Sci., 37, 19 (1959).



Subscripts b and l denote branched and linear molecules

g is a function of the *number and type* of long chain branch points in the molecule.





$$g_4 = \left[ \left( 1 + \frac{\overline{m}_n}{6} \right)^{1/2} + \frac{4 \overline{m}_n}{3} \right]^{-1/2}$$

 $m_n$  is the number average number of branch points per molecule

# Long Chain Branching—Relation to Intrinsic Viscosity

g, the ratio of the radii of gyration of the branched to linear polymer chain of the *same molecular weight*, is related to the intrinsic viscosity by a branching function, g' :

g' = 
$$\frac{\begin{bmatrix} 1 \\ b \end{bmatrix}}{\begin{bmatrix} 1 \\ 1 \end{bmatrix}}$$
 (for the same molecular weight)

For Star Shaped Polymers (Theoretical Relationship) g' = g<sup>0.5</sup>

For Randomly Branched Monodisperse Polymers (Empirical Relationship-Kurata et al.)

$$\mathbf{g'} = \mathbf{g}^{0.6}$$

The experimentally determined intrinsic viscosity of a branched polymer will be less than that calculated from the SEC data using the universal calibration curve (which assumes that the polymer chains are perfectly linear).

An appropriate branching function, g'(, M) that contains a branching parameter, , is defined such that:

$$[]_{b} = g'(, M) []_{l} = g'(, M) KM^{a}$$

