

Light Scattering

STATIC Light Scattering

Also known as **Rayleigh** or **Classical** Light Scattering

Measures avg. intensity of scattered light for

Absolute Molecular Weight

DYNAMIC

Light Scattering (DLS)

Also known as **Quasi-elastic** Light Scattering (**QUELS**)
or **Photon Correlation Spectroscopy (PCS)**

Measures microsecond fluctuations of single photons

Hydrodynamic Radius (Size)

M.L.Hackert (with figures from Precision Detectors and Wyatt Tech.)

What Do We Mean By ABSOLUTE?

- There are **4 Absolute Methods of Measuring MW**
- 1) Membrane **Osmometry** (Number Average MW)
- 2) **Light Scattering** (Weight Average MW)
- 3) **Sedimentation Equilibrium** (Ultracentrifugation) (z-average MW)
- 4) **Mass spectroscopy**
- **NO** Reference to standards of mass
- **NO** assumptions of molecular model/conformation
- **ALL** parameters measured directly from 1st principles
- **Refractive indices**
- **geometries of cell and detector**
- **wavelength**
- **concentrations**
- **detector response**
- **temperature**
- **dn/dc**

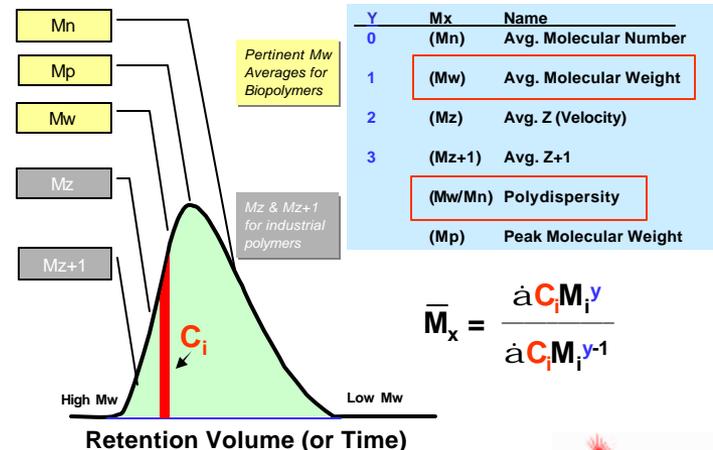
Wyatt Technologies

Light Scattering

Goals for this unit:

- Theory 101 – just the basics – **Know why sky is blue, sunset burnt orange**
- Rayleigh Scattering (Lord Rayleigh ~1871) / **Rayleigh Ratio**
- How does LS yield an “absolute” molecular weight? LS vs. RI
- What is polydispersity? How is it defined?
- Types of “Molecular Weight Averages”
- **LS Instrument / Practical Considerations**
- Static vs. Dynamic vs. X-ray Scattering (info from each)

Calculation of Mw Averages



$$\bar{M}_x = \frac{\sum C_i M_i^y}{\sum C_i M_i^{y-1}}$$

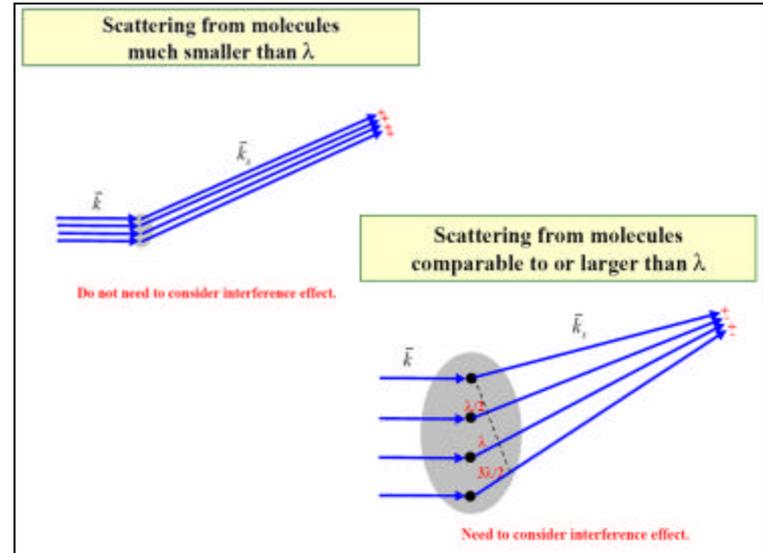
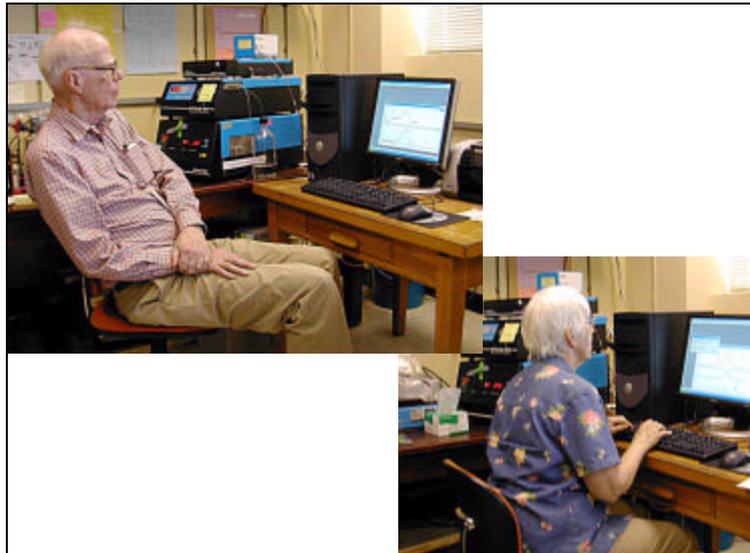
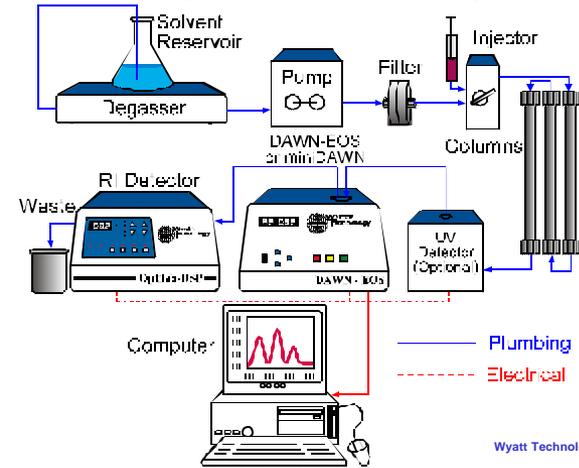
Precision Detectors

Abstract

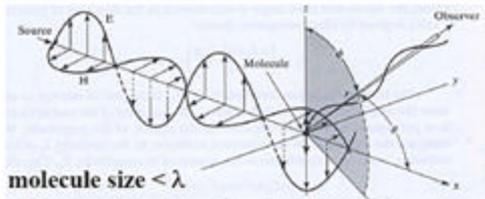
Recent advances in genomics and proteomics have produced a proliferation of new proteins requiring characterization. Mass spectrometry is ideally suited for identification and primary structural purposes but is not well suited for determining conformational structures in solution. As these molecules are expressed in cell culture, purified and then formulated, rigorous production processes must be carefully evaluated to minimize impact on the protein structure and its long-term shelf life. Obtaining a conformational stability profile of protein or antibodies can help weed out bad drug candidates from good ones as environmental factors can change their tertiary and quaternary structure. Environmental factors include pH, ionic strength, temperature, and excipient composition.

HPLC is used in flow injection mode with a detector array composed of laser light scattering (static and dynamic modes) and a concentration source detector (RI or UV). This configuration determines the average molecular weight and average hydrodynamic radius with run times as short as 1 minute. Alternatively, a SEC guard column can be used isolate analyte from excipients (eliminating blank runs) with run times under 3 minutes.

Chromatography with LS Set-up



EM wave scattered by a ^{small} molecule



molecule size $< \lambda$

Dipole induced in the molecule at the origin

$$\vec{p} = \alpha \vec{E} = \alpha \vec{E}_0 e^{i\omega t} \quad E_0: \text{incident field}$$

Electromagnetic wave emitted by the oscillating dipole

$$E = \frac{4\pi^2 \alpha E_0 \sin \phi}{\lambda^2 r} e^{i(\omega t - \vec{k}_s \cdot \vec{r})}$$

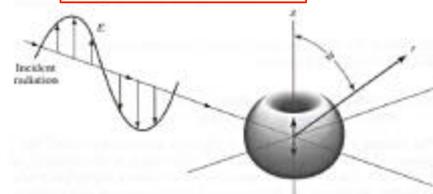
EM wave scattered by a molecule

Electromagnetic wave emitted by the oscillating dipole

$$E = \frac{4\pi^2 \alpha E_0 \sin \phi}{\lambda^2 r} e^{i(\omega t - \vec{k}_s \cdot \vec{r})}$$

Scattering intensity to the incident intensity

$$\frac{I}{I_0} = \frac{16\pi^4 \alpha^2 \sin^2 \phi}{\lambda^4 r^2} \quad \text{for polarized incident light of intensity } I_0$$

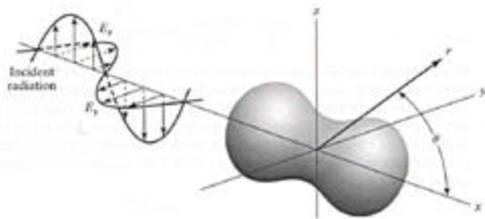


EM wave scattered by a molecule

Scattering intensity to the incident intensity

$$\frac{I}{I_0} = \frac{8\pi^4 \alpha^2 (1 + \cos^2 \theta)}{\lambda^4 r^2} \quad \text{for unpolarized incident light}$$

$I \propto 1/r^2$; $I \propto 1/\lambda^4$; I depends on scattering angle



Scattering from molecules much smaller than λ

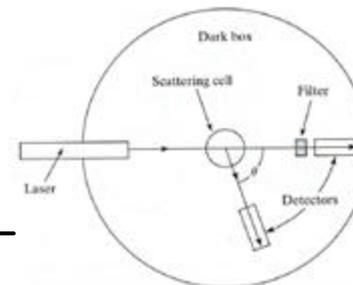
$$\text{molecule size} \ll \lambda \quad \frac{I(\theta)}{I_0} = \frac{2\pi^2 n_0^2}{A\lambda^4 r^2} \left(\frac{dn}{dc} \right)^2 CM (1 + \cos^2 \theta)$$

Light scattering can be used to determine the molecular weight.

Define "Raleigh Ratio" R_q

$$R_q = \frac{I(\theta)}{I_0} \frac{r^2}{(1 + \cos^2 \theta)}$$

$$R_q = KCM \quad \text{or} \quad \frac{K^*c}{R(q)} = \frac{1}{M}$$



Basic Light Scattering Principles

- The **amount of light scattered** is directly proportional to the product of the **molar mass** and the **molecular concentration**

$$I_{LS} = C M_w (dn/dc)^2 P_\theta K_\theta$$

- The **variation of scattered light** with scattering angle is proportional to the **average size** of the scattering molecules.

$$\frac{K^*c}{R(q)} = \frac{1}{M} \left[1 + \frac{16p^2}{3l^2} \langle r_g^2 \rangle \sin^2(q/2) + \dots \right] + 2A_2c$$

Wyatt Technologies

Basic Light Scattering Equation

$$\frac{K^*c}{R(\theta)} = \frac{1}{M_w P(\theta)} + 2A_2c$$

Where:

$$K^* = 4\pi^2 (dn/dc)^2 n_0^2 / (N_A \lambda_0^4)$$

n_0 is the refractive index of the solvent.

c is the concentration of the solute molecules (g/ml).

$R(q)$ is the fraction of light scattered per unit solid angle, in excess of the light scattered by the solvent, divided by the incident intensity.

N_A is Avogadro's number.

λ_0 is the vacuum wavelength of the incident light.

dn/dc is the refractive index increment, which tells how much the refractive index of the solution varies with solute concentration.

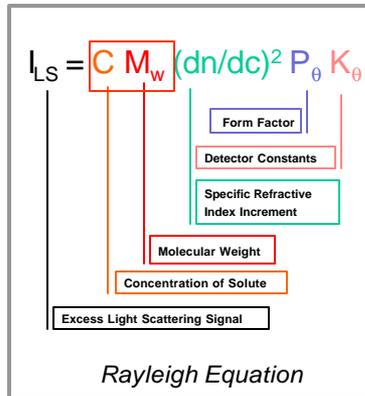
M_w is the weight-average molar mass.

Wyatt Technologies

Static Light Scattering Detection

Determines

- Absolute Molecular Weight**
Independent of Column Calibration
- Radius of Gyration (R_g)**
> 10 nm to 150 nm



Precision Detectors



Accuracy of Molecular Masses of Test Proteins Determined by Light Scattering

Protein	Mass From Structure [Da]	Light Scattering* [Da]	Apparent Error [%]
Carbonic anhydrase	29,023	29,800	+2.7
Alcohol dehydrogenase	145,980	149,000	+1.4
β -Amylase	224,340	228,000	+1.6
Apo ferritin	476,316	484,400	+1.7
Thyroglobulin	669,000	679,000	+1.5
Ornithine decarboxylase	990,684	978,000	-1.3
Octopus Hemocyanin	3,440,000	3,450,000	+0.3

*DAWN detector model-F, 0.19 was used as dn/dc value for all the proteins
Adapted from "Assembly of the Gigantic Hemoglobin of the Earthworm *Lumbricus terrestris* by A. Riggs *et al.* In *J. Bio. Chem.*, Vol. 271, No. 47, pp 30007-30021, 1996.

Wyatt
Technology

RI & Light Scattering Combined for Mw

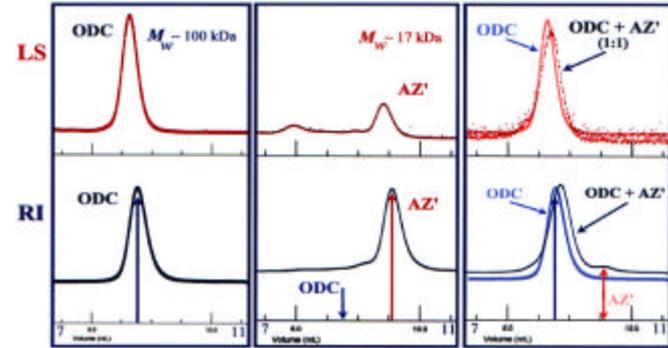
$$R_q = C M (dn/dc)^2 P(q) K(q) \quad \text{Rayleigh Equation}$$

$$RI_{\text{signal}} = K_{RI} (dn/dc) C \quad \text{RI Equation}$$

$$\frac{R_q}{RI_{\text{signal}}} = \frac{K(q) M_w (dn/dc) P(q)}{K_{RI}}$$

$$\frac{R_q}{RI_{\text{signal}}} \sim M_w \quad \left\{ \begin{array}{l} \text{For a truly effective measurement of molecular} \\ \text{weight the static light scattering detector must be} \\ \text{combined with a well matched refractometer} \end{array} \right.$$

- The light scattering signal is directly proportional to the M_w .
- The concentration source signal (e.g. RI) is indirectly proportional to the M_w .



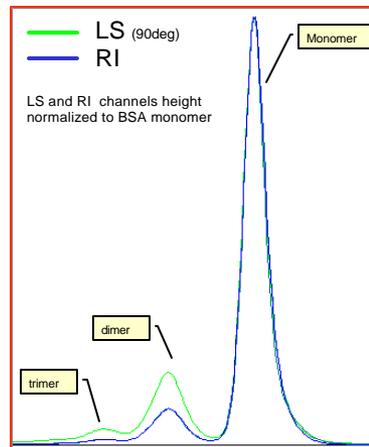
Light Scattering (LS) and Refractive Increment (RI) Results of ODC and AZ'. ODC, AZ' and ODC:AZ' complex were injected onto an HPLC sizing column, separated and analyzed by LS (DAWN EOS) and RI (OptiLab DSP interferometric refractometer). The results shown are for 7 to 11 minutes of elution volume. Frame 1 is for ODC, frame 2 for AZ', and frame 3 for the ODC:AZ' mixture at a 1:1 subunit ratio with the ODC trace from frame 1 superimposed for reference.

Static LS Equation for Proteins

$$M_w = \frac{I_{LS}}{RI} K_{\text{total}}$$

1. M_w is directly proportional to the LS signal
2. M_w is indirectly proportional to the conc. source (RI)
3. The dn/dc is constant for the protein and its aggregates (0.186 mL/g)
4. Degree of aggregation can be approximated from visual inspection

Visualizing Aggregation State



$$M_w = \frac{I_{LS}}{RI} K_{\text{total}}$$

Using visual inspection

- Monomer (LS is equal to RI)
- Dimer (LS is 2X as large as RI)
- Trimer (LS is 3X as large as RI)

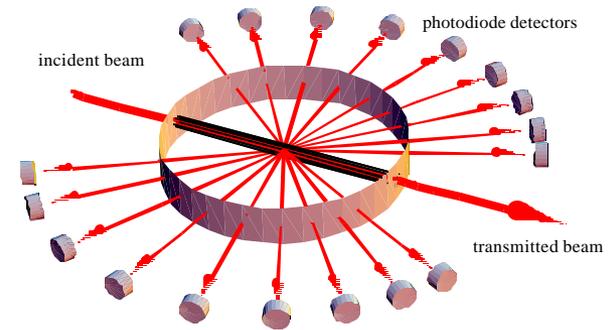
Why Multi-Angle Detection?

Light Scattering Intensity of Particles Shows an Angular Dependence on Size

- Low angles sensitive to large particles
- 90 degree angle more sensitive to smaller particles
- High angles less sensitive to larger particles
- Back angles better suited for opaque matrices

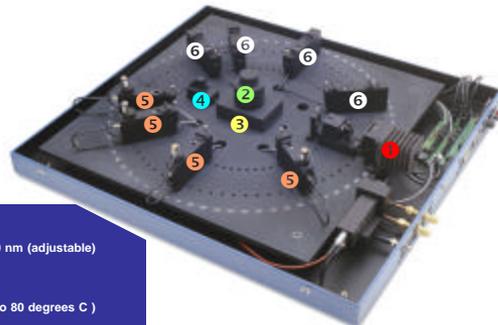


Flow Cell & Detector Geometry

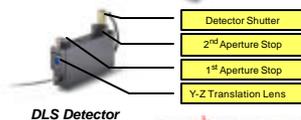


Wyatt Technologies

ALS4000 Optical Platform Features



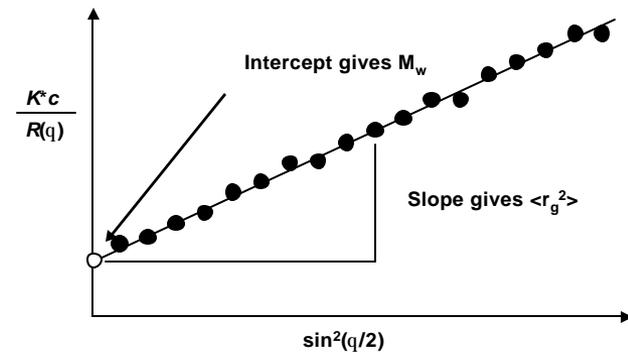
- 1 Diode laser: 230 mW @ 830 nm (adjustable)
- 2 Flow Cell (30 uL)
- 3 Peltier cooling/heating (0 to 80 degrees C)
- 4 Laser Beam Stop
- 5 DLS Detectors (Up to 4)
- 6 Static LS Detectors (Up to 8)



DLS Detector



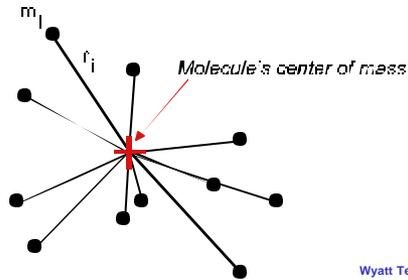
M_w & $\langle r_g^2 \rangle$ determined by MALS



$$\frac{K^*c}{R(q)} = \frac{1}{M} \left[1 + \frac{16p^2}{3l^2} \langle r_g^2 \rangle \sin^2(q/2) + \dots \right] + 2 A_2 c$$

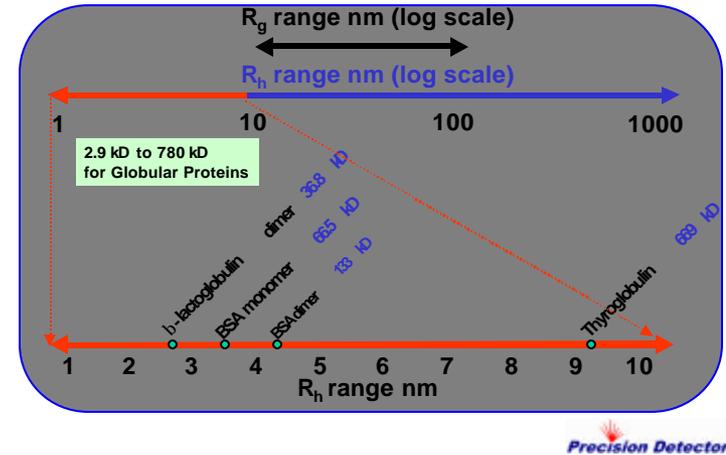
$$\langle r_g^2 \rangle = \frac{\sum r_i^2 m_i}{M}$$

- $\langle r_g^2 \rangle$ is the mean square radius, relating to the distribution of mass within the molecule, given by



Wyatt Technologies

Why use R_h instead of R_g for Biomolecules?



Dynamic Light Scattering Detection

Determines

- Molecular or Particle Size
As Hydrodynamic Radius (R_h)
- Size Range
1 to 1000 nm

$$D_0 = \kappa T (6\pi \eta_0 R_h)^{-1}$$

Stokes-Einstein Equation

Precision Detectors

Hydrodynamic Radius Determination

Precision Detectors

From $g_1(\tau)$ the diffusion coefficient (D) for the scattering particles can be determined. From the diffusion coefficient, the hydrodynamic radius can be calculated.

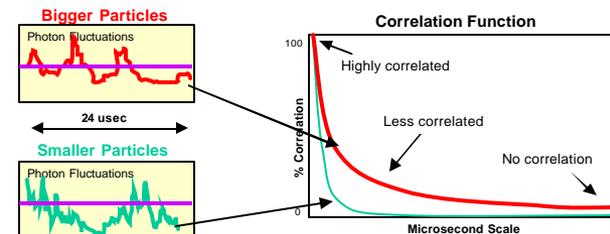
$$D_0 = \kappa T (6\pi \eta_0 R_h)^{-1}$$

Stokes-Einstein Equation

Photon Correlator
Sternian Correlator
Temperature (K) / Solvent Viscosity (Pa·s)
Hydrodynamic Radius (nm)

Applicable DLS Size Range

1.5 to 1000 nm Radius

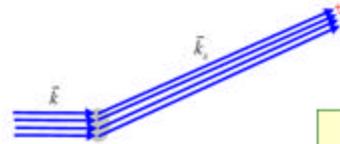


Conclusions on Static and Dynamic LS

- Measures **Hydrodynamic Radius, Radius of Gyration, Molecular Weight, Particle Size Distribution.**
- Instrument Measures
 - Rh size from 1.7 nm to 1000 nm.
 - Rg size from 9 nm to 150 nm.
 - Molecular Weight 200 to 20,000,000 daltons
- Detects branching, aggregates and calculates Mw.
- DLS**, does not require conc or dn/dc measurement for size.
- Batch Mode for non-flow system accessories.

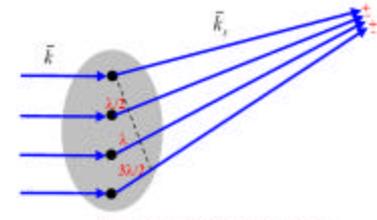
Precision Detectors

Scattering from molecules much smaller than λ



Do not need to consider interference effect.

Scattering from molecules comparable to or larger than λ



Need to consider interference effect.

Scattering from molecules comparable to or larger than λ

$$P(\theta) = \frac{\langle I(\theta) \rangle}{\langle I(0) \rangle} = \exp\left(-\frac{16\pi^2}{3} \cdot \frac{R_G^2}{\lambda^2} \cdot \sin^2 \frac{\theta}{2}\right)$$

Visible light, $\lambda \sim 400 - 700$ nm

For molecules with a few nm, scattering of visible light have a very weak angular dependence.



Difficult to determine the size of the molecules by scattering of visible light.

Biomolecules absorb UV light.



x-ray scattering

X-ray scattering

The major scatterers of x-ray in a molecule are electrons.

$$\frac{I(\theta)}{I_0} = \frac{8\pi^4 \alpha^2 (1 + \cos^2 \theta)}{\lambda^4 r^2} \quad \text{Light scattering by a small molecule}$$



$$\frac{I(\theta)}{I_0} = \left(\frac{e^2}{mc^2}\right)^2 \cdot \frac{(1 + \cos^2 \theta)}{2r^2} \quad \text{X-ray scattering by an electron}$$

X-ray scattering

In a solution of molecules that each has Z electrons, and a molecular weight of M , and concentration of C

$$\frac{I(\theta)}{I_0} = \frac{2\pi^2 n_0^2}{A\lambda^4 r^2} \left(\frac{dn}{dC}\right)^2 CM (1 + \cos^2 \theta) \quad \text{Light scattering}$$



$$\frac{I(\theta)}{I_0} = \frac{1}{2r^2} \left(\frac{e^2}{mc^2}\right)^2 \left(\frac{Z - Z_0}{M}\right)^2 AMC(1 + \cos^2 \theta) \quad \text{X-ray scattering}$$

Z_0 : number of solvent electrons in the volume of a solute molecule

$$\frac{I(\theta=0)}{I_0} = \frac{1}{r^2} \left(\frac{e^2}{mc^2}\right)^2 \left(\frac{Z - Z_0}{M}\right)^2 AMC$$

X-ray scattering can be used to determine the molecular weight.

Small angle X-ray scattering (SAXS)

The size of the molecules are always larger than $\lambda \sim 0.1$ nm.

$$P(\theta) = \frac{\langle I(\vec{S}) \rangle}{\langle I(0) \rangle} = \frac{1}{N^2} \sum_{j=1}^N \sum_{l=1}^N \frac{\sin 2\pi S r_{jl}}{2\pi S r_{jl}} \quad S = \frac{2}{\lambda} \sin \frac{\theta}{2}$$

$Sr_{ij} \ll 1$ only at very low angle

At very low angle $SR_G \ll 1$

$$P(\theta) = \frac{\langle I(\theta) \rangle}{\langle I(0) \rangle} = \exp\left(-\frac{4\pi^2 S^2 R_G^2}{3}\right)$$

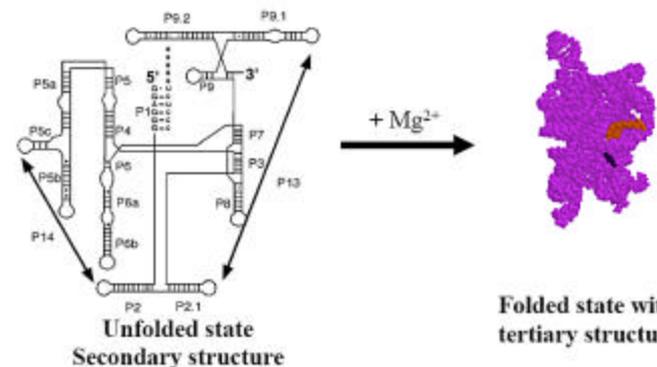
Guinier formula

Molecular weight and size measured by Light scattering or SAXS

Material	M_w	R_G (nm)
Ribonuclease	<i>12,700</i>	<i>1.48</i>
α -Lactalbumin	<i>13,500</i>	<i>1.45</i>
Lysozyme	<i>13,600</i>	<i>1.43</i>
β -Lactoglobulin	36,000	2.17
Serum albumin	70,000	2.98
Myosin	493,000	46.8
Turnip yellow mosaic virus		<i>10.4</i>
Tobacco mosaic virus	39×10^6	92.4

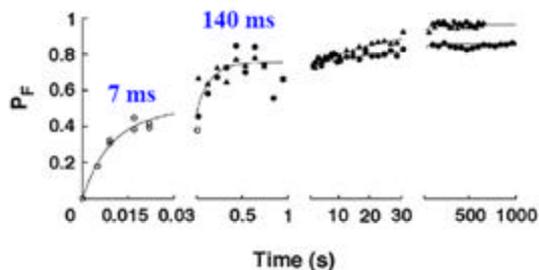
Values in *italic* are from low angle X-ray scattering.

RNA folding



SAXS has a potential to answer how fast RNA molecule compacts.

RNA folding studied by SAXS



- Major compaction happen very fast in two phase: 7 ms and 140 ms

Conclusions Part 1: Static and Dynamic LS

- Measures Hydrodynamic Radius, Radius of Gyration, Molecular Weight, Particle Size Distribution.
- Instrument Measures
 - Rh size from 1.7 nm to 1000 nm.
 - Rg size from 9 nm to 150 nm.
 - Molecular Weight 200 to 20,000,000 daltons
- Detects branching, aggregates and calculates Mw.
- DLS, does not require conc or dn/dc measurement for size.
- Batch Mode for non-flow system accessories.

Conclusions Part 2:

- Static Light Scattering
 - Provides absolute Mw in Solution..
 - Sensitive Aggregation Detection
- Dynamic Light Scattering
 - Provides Rh Sizing down to 1 nm for Biomolecules.
 - “On-the-fly Determinations”
 - Gives more insight on structure of material
 - Can be added to any existing static LS system from any manufacturer
- Flow Injection Analysis is Ideal for Rapid Screening for Avg. Mw and Size

Conclusions Part 3: Why Use Light Scattering Detection?

- *Absolute* Molecular Weight Calculations
- Non Destructive Technique
- *Independent* of Elution Time (no relative standards)
- Radius of Gyration – Rg
- Hydrodynamic Radius - Rh (Dynamic Light Scattering)
- Molecular Geometry