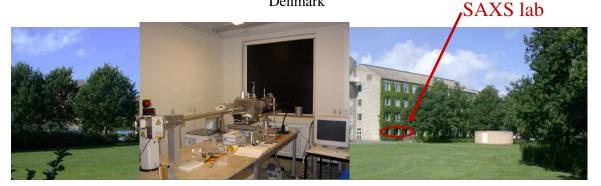




## Form and Structure Factors: Modeling and Interactions

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#### **Outline**

- Model fitting and least-squares methods
- Available form factors ex: sphere, ellipsoid, cylinder, spherical subunits... ex: polymer chain
- Monte Carlo integration for form factors of complex structures
- Monte Carlo simulations for form factors of polymer models
- Concentration effects and structure factors
   Zimm approach
   Spherical particles
   Elongated particles (approximations)
   Polymers

#### Motivation

- <u>not</u> to replace shape reconstruction and crystal-structure based modeling we use the methods extensively
- alternative approaches to reduce the number of degrees of freedom in SAS data structural analysis (might make you aware of the limited information content of your data !!!)
- provide polymer-theory based modeling of flexible chains
- describe and correct for concentration effects

3

#### Literature

Jan Skov Pedersen, *Analysis of Small-Angle Scattering Data from Colloids and Polymer Solutions: Modeling and Least-squares Fitting* (1997). *Adv. Colloid Interface Sci.*, **70**, 171-210.

Jan Skov Pedersen

Monte Carlo Simulation Techniques Applied in the Analysis of Small-Angle Scattering Data from Colloids and Polymer Systems

in Neutrons, X-Rays and Light

P. Lindner and Th. Zemb (Editors) 2002 Elsevier Science B.V.

p. 381

Jan Skov Pedersen

Modelling of Small-Angle Scattering Data from Colloids and Polymer Systems

in Neutrons, X-Rays and Light

P. Lindner and Th. Zemb (Editors) 2002 Elsevier Science B.V. p. 391

Rudolf Klein

#### Interacting Colloidal Suspensions

in Neutrons, X-Rays and Light

P. Lindner and Th. Zemb (Editors) 2002 Elsevier Science B.V.

p. 351

#### Form factors and structure factors

#### Warning 1:

Scattering theory – lots of equations!

= mathematics, Fourier transformations

#### Warning 2:

Structure factors:

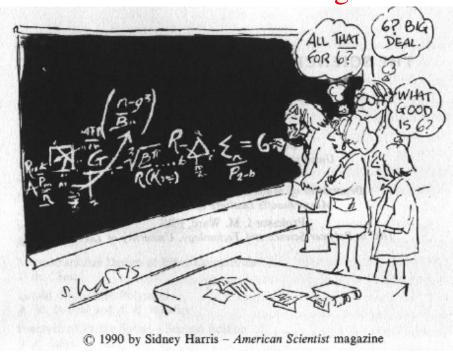
Particle interactions = statistical mechanics

Not all details given

- but hope to give you an impression!

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# I will outline some calculations to show that it is not black magic!



#### Input data: Azimuthally averaged data

$$q_i, I(q_i), \sigma[I(q_i)]$$
  $i = 1,2,3,...N$ 

 $q_i$  calibrated

- $I(q_i)$  calibrated, i.e. on absolute scale noisy, (smeared), truncated
- $\sigma[I(q_i)]$  Statistical standard errors: Calculated from counting statistics by error propagation do not contain information on systematic error !!!!

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#### Least-squared methods

Measured data:  $I^{\exp}(q_i)$ , i = 1,...,N

Model:  $I^{\text{mod}}(q_i)$   $a_i$ , i = 1,...,M.

Chi-square:  $\chi^2 = \sum_{i=1}^{N} \left( \frac{I^{\exp}(q_i) - I^{\operatorname{mod}}(q_i)}{\sigma_i} \right)^2$ 

Reduced Chi-squared:  $\chi_r^2 = \frac{\chi^2}{N-M}$  = goodness of fit (GoF)

Note that  $\chi_r^2 = 1$  for N >> M corresponds to  $|I^{\exp}(q_i) - I^{\operatorname{mod}}(q_i)| = \sigma_i$  i.e. statistical agreement between model and data

#### Cross section

 $d\sigma(q)/d\Omega$ : number of scattered neutrons or photons per unit time, relative to the incident flux of neutron or photons, per unit solid angle at q per unit volume of the sample.

For system of monodisperse particles

$$\frac{d\sigma(q)}{d\Omega} = I(q) = n \, \Delta \rho^2 V^2 P(q) S(q) = c \, M \, \Delta \rho_m^2 P(q) S(q)$$

*n* is the number density of particles,

 $\Delta \rho$  is the excess scattering length density, given by electron density differences

V is the volume of the particles,

P(q) is the particle form factor, P(q=0)=1

S(q) is the particle structure factor,  $S(q=\infty)=1$ 

- $V \propto M$
- n = c/M
- $\Delta \rho$  can be calculated from partial specific density, composition

Form factors of geometrical objects

#### Form factors I Homogenous rigid particles

- 1. Homogeneous sphere
  - 2. Spherical shell:
  - 3. Spherical concentric shells:
- 4. Particles consisting of spherical subunits:
  - 5. Ellipsoid of revolution:
  - 6. Tri-axial ellipsoid:
  - 7. Cube and rectangular parallelepipedons:
  - 8. Truncated octahedra:
  - 9. Faceted Sphere:
  - 9x Lens
  - 10. Cube with terraces:
- 11. Cylinder:
  - 12. Cylinder with elliptical cross section:
  - 13. Cylinder with hemi-spherical end-caps:
  - 13x Cylinder with 'half lens' end caps
  - 14. Toroid:
  - 15. Infinitely thin rod:
  - 16. Infinitely thin circular disk:
  - 17. Fractal aggregates:

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#### Form factors II

#### 'Polymer models'



- 18. Flexible polymers with Gaussian statistics:
- 19. Polydisperse flexible polymers with Gaussian statistics:
- 20. Flexible ring polymers with Gaussian statistics:
- 21. Flexible self-avoiding polymers:
- 22. Polydisperse flexible self-avoiding polymers:
- 23. Semi-flexible polymers without self-avoidance:
- 24. Semi-flexible polymers with self-avoidance:
  - 24x Polyelectrolyte Semi-flexible polymers with self-avoidance:
  - 25. Star polymer with Gaussian statistics:
  - 26. Polydisperse star polymer with Gaussian statistics:
  - 27. Regular star-burst polymer (dendrimer) with Gaussian statistics:
  - 28. Polycondensates of A<sub>f</sub> monomers:
  - 29. Polycondensates of AB<sub>f</sub> monomers:
  - 30. Polycondensates of ABC monomers:
  - 31. Regular comb polymer with Gaussian statistics:
  - 32. Arbitrarily branched polymers with Gaussian statistics:
  - 33. Arbitrarily branched semi-flexible polymers:
  - 34. Arbitrarily branched self-avoiding polymers: (Block copolymer micelle)
  - 35. Sphere with Gaussian chains attached:
  - 36. Ellipsoid with Gaussian chains attached:
  - 37. Cylinder with Gaussian chains attached:
  - 38. Polydisperse thin cylinder with polydisperse Gaussian chains attached to the ends:
  - 39. Sphere with corona of semi-flexible interacting self-avoiding chains of a corona chain. 12

#### Form factors III

$$P(q) = P_{cross-section}(q) P_{large}(q)$$

40. Very anisotropic particles with local planar geometry:

Cross section:

- (a) Homogeneous cross section
- (b) Two infinitely thin planes
- (c) A layered centro symmetric cross-section
- (d) Gaussian chains attached to the surface

Overall shape:

- (a) Infinitely thin spherical shell
- (b) Elliptical shell
- (c) Cylindrical shell
- (d) Infinitely thin disk

#### 41. Very anisotropic particles with local cylindrical geometry:

Cross section:

- (a) Homogeneous circular cross-section
- (b) Concentric circular shells
- (c) Elliptical Homogeneous cross section.
- (d) Elliptical concentric shells
- (e) Gaussian chains attached to the surface

Overall shape:

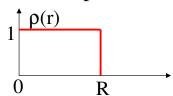
- (a) Infinitely thin rod
- (b) Semi-flexible polymer chain with or without excluded volume

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#### From factor of a solid sphere





$$A(q) = 4\pi \int_{0}^{\infty} \rho(r) \frac{\sin(qr)}{qr} r^{2} dr = 4\pi \int_{0}^{R} \frac{\sin(qr)}{qr} r^{2} dr$$

$$= \frac{4\pi}{q} \int_{0}^{R} \sin(qr) \ r dr =$$

$$\left( \int f'g \, dx = [fg] - \int fg' dx \right)$$

(partial integration)...

r

$$= \frac{4\pi}{q} \left( -\frac{R\cos qR}{q} + \left[ \frac{\sin qr}{q} \right]_{0}^{R} \right)$$

$$=\frac{4\pi}{q}\left(-\frac{R\cos qR}{q}+\frac{\sin qr}{q^2}\right)$$

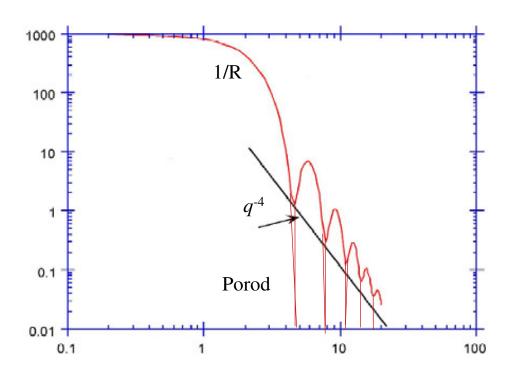
$$=\frac{4\pi}{q^3}\big(\sin qR - qR\cos qR\big)$$

$$= \frac{4}{3}\pi R^{3} \frac{3[\sin(qR) - qR\cos(qR)]}{(qR)^{3}}$$

spherical Bessel function

#### Form factor of sphere

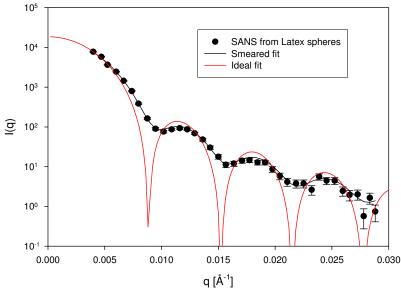
$$P(q) = A(q)^2/V^2$$



C. Glinka

#### Measured data from solid sphere (SANS)

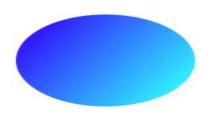
$$I(\langle q \rangle) = \int R(\langle q \rangle, q) \frac{\mathrm{d}\sigma(q)}{\mathrm{d}\Omega} \, \mathrm{d}q \qquad \frac{d\sigma}{d\Omega}(q) = \Delta \rho^2 V^2 \left[ \frac{3[\sin(qR) - qR\cos(qR)]}{(qR)^3} \right]^2$$



Instrumental smearing is routinely included in SANS data analysis

Data from Wiggnal et al.

#### Ellipsoid



5. Ellipsoid of revolution: This expression was determined by Guinier (1939). The averaging over orientations has to be done numerically. For the semi-axes  $R, R, \epsilon R$ :

$$P_5(q, R, \epsilon) = \int_0^{\pi/2} F_1[q, r(R, \epsilon, \alpha)]^2 \sin \alpha \, d\alpha, \tag{27}$$

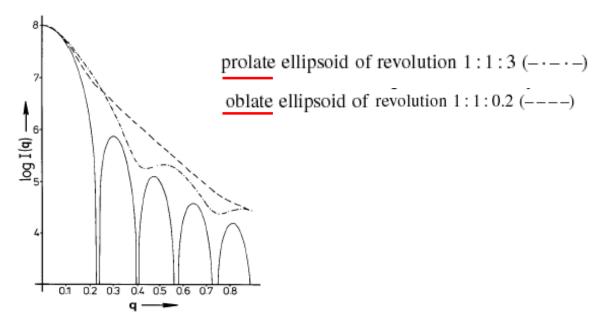
where  $r(R, \epsilon, \alpha) = R(\sin^2 \alpha + \epsilon^2 \cos^2 \alpha)^{1/2}$ .

And:

$$F_1(q, R) = \frac{3[\sin(qR) - qR\cos(qR)]}{(qR)^3}.$$

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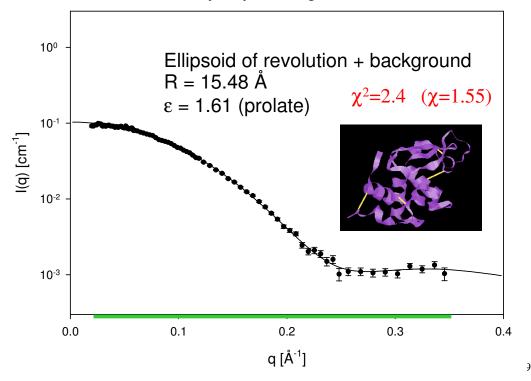
#### P(q): Ellipsoid of revolution



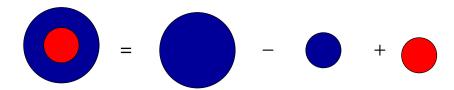
Glatter

Lysosyme

#### Lysozyme 7 mg/mL



#### **Core-shell particles:**



$$A_{core-shell}(q) = \Delta \rho_{core} \left[ \Delta \rho_{shell} V_{out} \Phi(qR_{out}) - (\Delta \rho_{shell} - \Delta \rho_{core}) V_{in} \Phi(qR_{in}) \right]$$

where

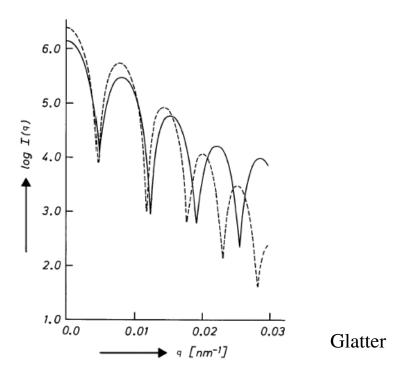
 $V_{out} = 4\pi R_{out}^{3}/3$  and  $V_{in} = 4\pi R_{in}^{3}/3$ .

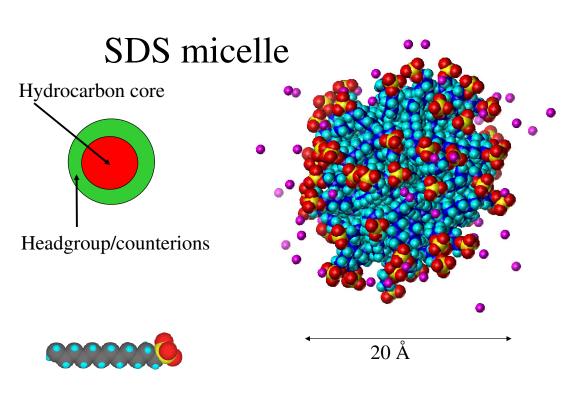
 $\Delta 
ho_{core}$  is the excess scattering length density of the core,

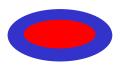
 $\Delta \rho_{shell}$  is the excess scattering length density of the shell and:

$$\Phi(x) = \frac{3[\sin x - x\cos x]}{x^3}$$

#### P(q): Core-shell

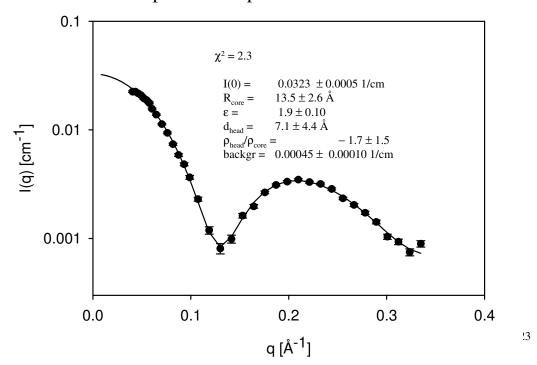






#### SDS micelles:

#### prolate ellispoid with shell of constant thickness



#### **Molecular constraints:**

$$\Delta \rho_{tail}^e = \frac{Z_{tail}^e}{V_{tail}} - \frac{Z_{H2O}^e}{V_{H2O}}$$

$$V_{core} = N_{agg}V_{tail}$$

$$\Delta \rho_{head}^{e} = \frac{Z_{head}^{e} + nZ_{H2O}}{V_{head} + nV_{H2O}} - \frac{Z_{H2O}^{e}}{V_{H2O}}$$

 $V_{shell} = N_{agg} \left( V_{head} + n V_{H2O} \right)$ 

$$n_{micelles} = \frac{c}{N_{agg} M_{surfactant}}$$

#### Cylinder



11. Cylinder: The expression for a cylinder with radius R and length L was given by Fournet (1949):

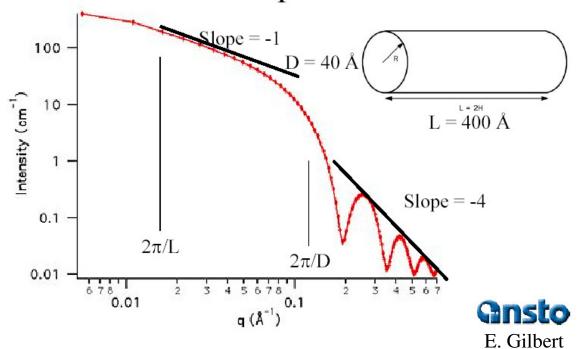
$$P_{11}(q) = \int_{0}^{\pi/2} \left[ \frac{2J_1(qR\sin\alpha)}{qR\sin\alpha} \frac{\sin(qL\cos\alpha/2)}{qL\cos\alpha/2} \right]^2 \sin\alpha \,d\alpha, \tag{30}$$

where  $J_1(x)$  is the first order Bessel function.

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P(q) cylinder

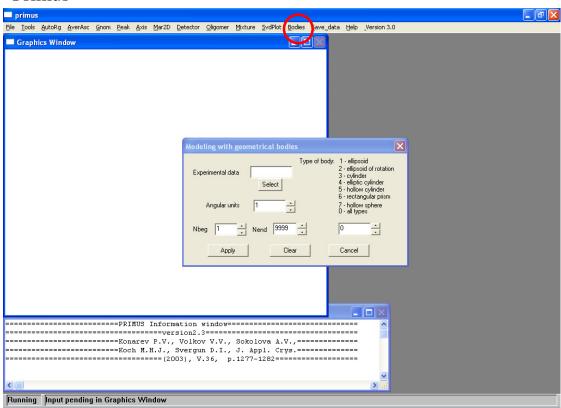
# Dilute randomly oriented monodisperse rods



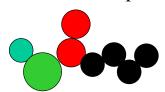
# Glucagon Fibrils R=29 Å R=16 Å

Cristiano Luis Pinto Oliveira, Manja A. Behrens, Jesper Søndergaard Pedersen, Kurt Erlacher, 27 Daniel Otzen and Jan Skov Pedersen J. Mol. Biol. (2009) 387, 147–161

#### **Primus**



Collection of particles with spherical symmetry



center of sphere 
$$\vec{r} = r' + \vec{s}$$

$$A(\vec{q}) = \sum b_j \exp(-i\vec{q} \cdot \vec{r}_j) = \sum b_{ij} \exp(-i\vec{q} \cdot (\vec{r}_i' + \vec{s}_j))$$

$$\begin{split} I(q) = & \sum b_{ij} b_{kl} \exp(-i\vec{q} \cdot (\vec{r}_i' + \vec{s}_j - \vec{r}_k' - \vec{s}_l)) \\ = & \sum b_{ij} \exp(-i\vec{q} \cdot \vec{r}_i') b_{kl} \exp(i\vec{q} \cdot \vec{r}_k') \exp(-i\vec{q} \cdot (\vec{s}_j' - \vec{s}_k')) \\ = & \sum V_j \Delta \rho_j \Phi(qR_j) V_k \Delta \rho_k \Phi(qR_k) \frac{\sin qd_{jk}}{qd_{jk}} \\ = & \sum V_j^2 \Delta \rho_j^2 P(qR_j) + \sum_{j \neq k} V_j \Delta \rho_j \Phi(qR_j) V_k \Delta \rho_k \Phi(qR_k) \frac{\sin qd_{jk}}{qd_{jk}} \end{split}$$

self-term

interference term

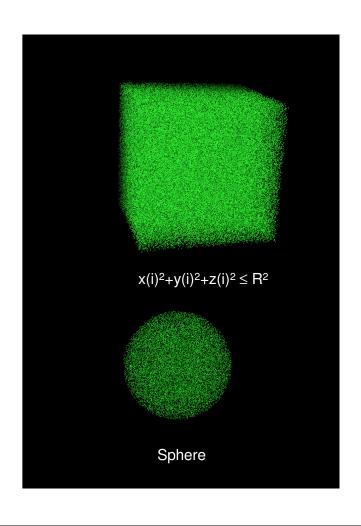
phase factor

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Debye, 1915

# Monte Carlo integration in calculation of form factors for complex structures

- •Generate points in space by Monte Carlo simulations
- •Select subsets by geometric constraints
- •Caclulate histograms p(r) functions
- •(Include polydispersity)
- •Fourier transform

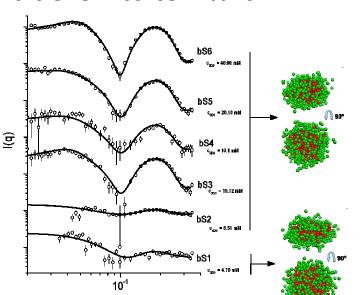


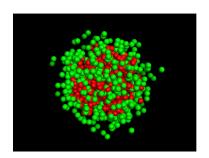
#### The Role of Decorated SDS Micelles in Sub-CMC Protein Denaturation and Association

Kell K. Andersen<sup>1,2</sup>, Cristiano L. Oliveira<sup>3</sup>, Kim L. Larsen<sup>2</sup>, Flemming M. Poulsen<sup>4</sup>, Thomas H. Callisen<sup>5</sup>, Peter Westh<sup>6</sup>, Jan S. Pedersen<sup>6</sup> and Daniel Otzen<sup>1\*</sup>

J. Mol. Biol. (2009) 391, 207-226

#### Pure SDS micelles in buffer:





 $C > CMC (\sim 5 \text{ mM})$ 

 $N_{agg}$ = 66±1 oblate ellipsoids R=20.3±0.3 Å  $\epsilon$ =0.663±0.005

 $C \leq CMC$ 

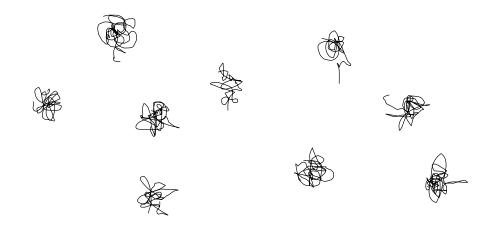
31

#### Polymer chains in solution

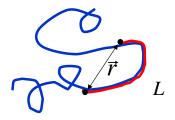
#### Gigantic ensemble of 3D random flights

- all with different configurations

 $q \, [\text{\AA}^{\text{-1}}]$ 



#### Gaussian polymer chain



Look at two points: Contour separation: *L* Spatial separation: *r* 

Contribution to scattering:  $I_2(q) = \frac{\sin(qr)}{qr}$ 

 $\exp\left[-q^2Lb/6\right]$ 

For an ensemble of polymers, points with

L has 
$$\langle r^2 \rangle = Lb$$

and *r* has a Gaussian distribution:

$$D(r) \propto \exp\left[-\frac{3r^2}{2 < r^2 > }\right]$$

#### Add scattering from all pair of points

'Density of points':  $(L_o - L)$ 

 $L_o$  L33

#### Gaussian chains: The calculation

$$P(q) = \frac{1}{L_o^2} \int_0^\infty dr \int_0^{L_o} dL_1 \int_0^{L_o} dL_2 D(r, | L_2 - L_1 |) \frac{\sin(qr)}{qr} r$$

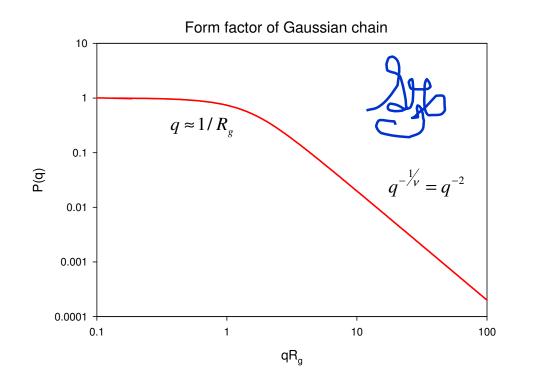
$$= \frac{1}{L_o} \int_0^\infty dr \int_0^{L_o} dL (L_o - L) D(r, L) \frac{\sin(qr)}{qr} r^2$$

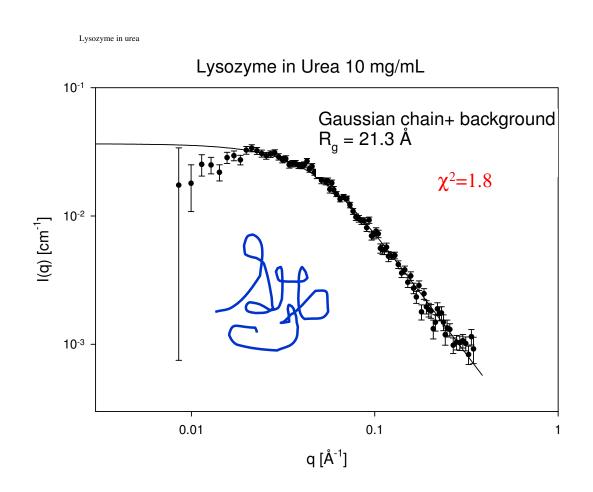
$$= \frac{1}{L_o} \int_0^{L_o} dL (L_o - L) \exp[-q^2 Lb / 6]$$

$$= \frac{2[\exp(-x) - 1 + x]}{r^2} \qquad x = R_g^2 q^2 \qquad R_g^2 = Lb / 6$$

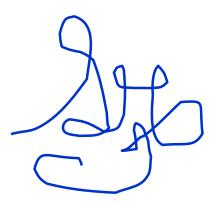
How does this function look?

#### Polymer scattering



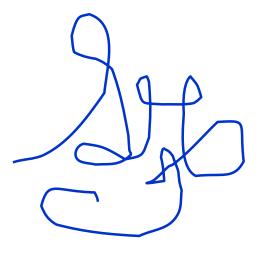


#### Self avoidence



No excluded volume

$$P(x = R_g^2 q^2) = \frac{2[\exp(-x) - 1 + x]}{x^2}$$



No excluded volume => expansion

no analytical solution!

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#### Monte Carlo simulation approach

Pedersen and Schurtenberger 1996

- (1) Choose model
- (2) Vary parameters in a broad range Generate configs., sample P(q)
- (3) Analyze P(q) using physical insight
- (4) Parameterize P(q) using physical insight
- (5) Fit experimental data using numerical expressions for P(q)

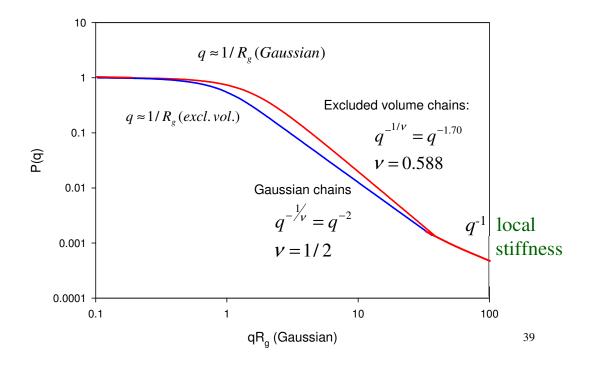


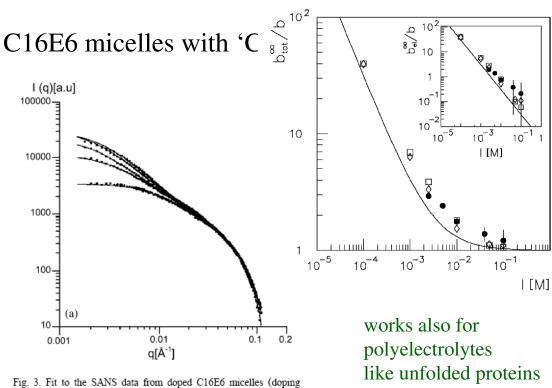
L = contour length

b = Kuhn ('step') length

#### Expansion = 1.5

#### Form factor polymer chains





level of 6% of ionic surfactant) with different salt concentrations at low surfactant concentrations (c = 0.4 mg/ml) [32 $^{\bullet}$ ]. From top to bottom: 0.1, 0.01, 0.005, 0.0025 and 0.001 M salt.

#### Models II

Polydispersity: Spherical particles as e.g. vesicles

No interaction effects: Size distribution D(R)

$$\frac{\mathrm{d}\sigma(q)}{\mathrm{d}\Omega} = \Delta \rho^2 \int_0^\infty D(R) V(R)^2 P(q, R) \, \mathrm{d}R.$$

Oligomeric mixture: Discrete particles

$$\frac{d\sigma(q)}{d\Omega} = c \,\Delta \rho^2 \sum_i M_i f_i P_i(q)$$

Application to insulin:

Pedersen, Hansen, Bauer (1994). European Biophysics Journal 23, 379-389. (Erratum). *ibid* 23, 227-229.

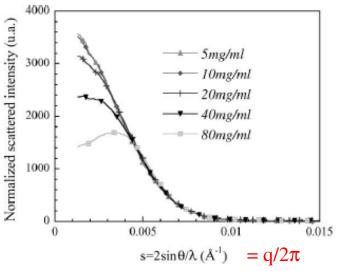
$$f_i = \text{mass fraction} \quad \sum_i f_i = 1$$

Used in PRIMUS 'Oligomers'

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#### Concentration effects

#### Concentration effects in protein solutions



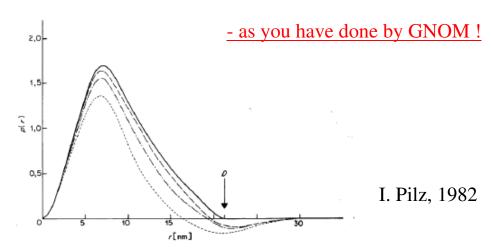
### Fig. 1. Variation of normalized scattered intensity of $\alpha$ -crystal-lin solutions as a function of protein concentration, at 20°C, in 150 mM phosphate buffer at pH=6.8.

## $\alpha$ -crystallin eye lens protein

S. Finet\*,1, A. Tardieu

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#### p(r) by Indirect Fourier Transformation (IFT)



At high concentration, the neighborhood is different from the average further away!

- (1) Simple approach: Exclude low q data.
- (2) Glatter: Use Generalized Indirect Fourier Transformation (GLFT)

#### Low concentraions Zimm 1948 – originally for light scattering

$$P'(q) = \bigcirc$$

Subtract overlapping configuration

$$P'(q) = P(q) - vP(q)^{2}$$
  
=  $P(q)[1 - vP(q)]$ 

 $v \sim$  concentration

Higher order terms:

$$P'(q) = P(q)[1 - vP(q)\{1 - vP(q)\}]$$
  
=  $P(q)[1 - vP(q) + v^2P(q)^2]$ 

$$= P(q)[1 - vP(q) + v^{2}P(q)^{2} - v^{3}P(q)^{3}....]$$

$$= \frac{P(q)}{1 + vP(q)}$$

= Random-Phase Approximation (RPA)

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#### Zimm approach

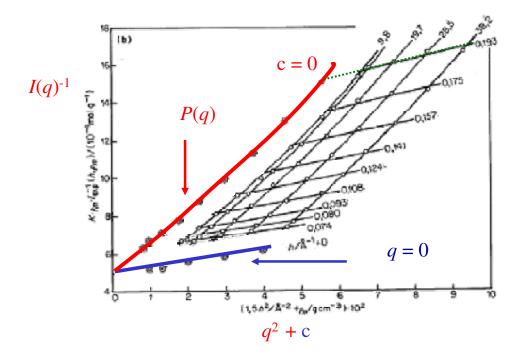
$$I(q) = K \frac{P(q)}{1 + vP(q)}$$

$$I(q)^{-1} = K^{-1} \frac{1 + vP(q)}{P(q)} = K^{-1} \left[ \frac{1}{P(q)} + v \right]$$

With 
$$P(q) \approx \frac{1}{1 + q^2 R_g^2 / 3}$$

With 
$$P(q) \approx \frac{1}{1 + q^2 R_g^2 / 3}$$
  $I(q)^{-1} = K^{-1} [1 + q^2 R_g^2 / 3 + v]$ 

Plot  $I(q)^{-1}$  versus  $q^2 + c$  and extrapolate to q=0 and c=0!



My suggestion:

( - which includes also information from what follows)

- Minimum 3 concentrations for same system.
- Fit data simultaneously all data sets

$$\frac{I(q_i)}{c} = \frac{P_i}{1 + ca_1 \exp(-q_i^2 a_2^2 / 3)}$$

With  $a_1$ ,  $a_2$ , and  $P_i$  fit parameters

# But now we look at the information content related to these effects...

Understand the fundamental processes and principles governing aggregation and crystallization

Why is the eye lens transparent despite a protein concentration of 30-40%?

49

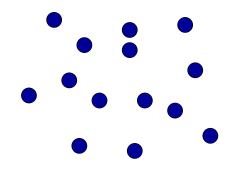
#### Structure factor

Spherical monodisperse particles

$$I(q) = V^{2} \Delta \rho^{2} P(q) \sum_{j} \frac{\sin q r_{jk}}{q r_{jk}}$$

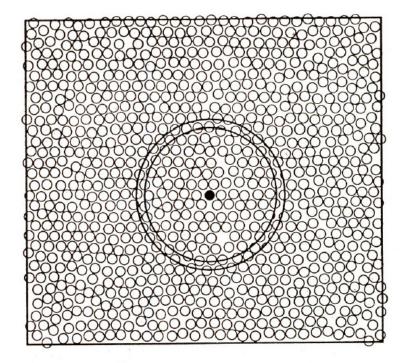
$$= V^{2} \Delta \rho^{2} P(q) S(q)$$

$$= V^{2} \Delta \rho^{2} P(q) \sum_{j} p(r_{j}) \frac{\sin q r_{j}}{q r_{j}}$$



S(q) is related to the probability distribution function of inter-particles distances, i.e. the pair correlation function g(r)

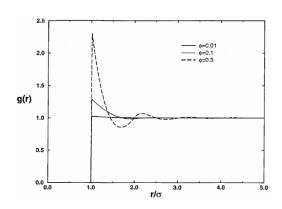
#### Correlation function g(r)

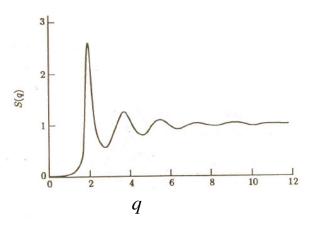


 $g(r) = \frac{Average}{}$  of the normalized density of atoms in a shell [r; r+dr] from the center of a particle

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#### g(r) and S(q)





$$S(q) = 1 + n 4\pi \int (g(r) - 1) \frac{\sin(qr)}{qr} r^2 dr$$

#### **GIFT**

Glatter: Generalized Indirect Fourier Transformation (GIFT)

$$I(q) = 4\pi \int p(r) \frac{\sin(qr)}{qr} dr$$

With concentration effects

$$I(q) = S_{eff}(q, \eta, R, \sigma(R), Z, c_{salt}) 4\pi \int p(r) \frac{\sin(qr)}{qr} dr$$

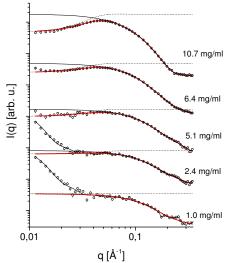
Optimized by constrained non-linear least-squares method

- works well for globular models and provides p(r)

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## A SAXS study of the small hormone glucagon: equilibrium aggregation and fibrillation

#### Home-written software



A SAXS Study of Glucagon Fibrillation

Hexamers

trimers

monomers

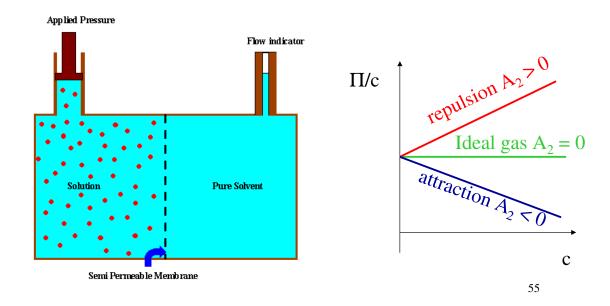
r [Å]

Cristiano Luis Pinto Oliveira 1\*, Manja Annette Behrens 1, Jesper Søndergaard Pedersen 2, Kurt Erlacher 1, Daniel Otzen 2, 3 and Jan Skov Pedersen 1

J. Mol. Biol. (2009) 387, 147-161

#### Osmometry (Second virial coeff A<sub>2</sub>)

$$\Pi = c (RT/M) (1 + A_2c + A_3c^2 + ...)$$



#### S(q), virial expansion and Zimm

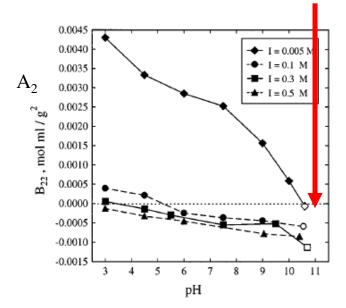
From statistical mechanics...:

$$S(q=0) = RT \left(\frac{\partial \Pi}{\partial c}\right)^{-1} = \frac{1}{1 + 2cMA_2 + 3c^2MA_3 + ...}$$

→ In Zimm approach  $v = 2cMA_2$ 

#### A<sub>2</sub> in lysozyme solutions

#### Isoelectric point

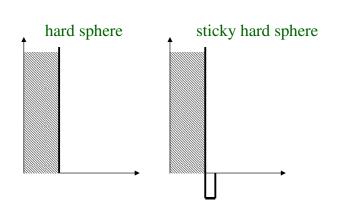


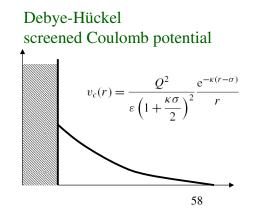
and A. M. Lenhoff

FIGURE 2 A summary of virial coefficients of lysozyme obtained from the SLS data at varying pH and four electrolyte concentrations (each point O. D. Velev, E. W. Kaler, corresponds to one data set similar to those plotted in Fig. 1). The open symbols denote experimental conditions where the onset of aggregation was detected.

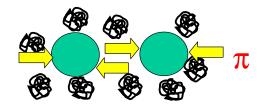
#### Colloidal interactions

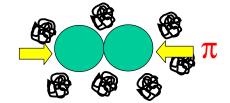
- Excluded volume 'repulsive' interactions ('hard-sphere')
- Short range <u>attractive</u> van der Waals interaction ('stickiness')
- Short range attractive hydrophobic interactions (solvent mediated 'stickiness')
- •Electrostatic repulsive interaction (or attractive for patchy charge distribution!) (effective Debye-Hückel potential)
- <u>Attractive</u> depletion interactions (co-solute (polymer) mediated )





#### Depletion interactions





$$v_{\mathrm{eff}}(r;\rho_1,\rho_2,T) = \begin{cases} \infty; & r < \sigma_1 \\ v_{A0}(r); & \sigma_1 < r < \sigma_1 + \sigma_2 \\ 0; & r > \sigma_1 + \sigma_2 \end{cases}$$

where

$$v_{A0}(r) = -k_B T \rho_2 \frac{\pi \sigma_1^3 (1+q)^3}{6} \left[ 1 - \frac{3r}{2(1+q)\sigma_1} + \frac{r^3}{2(1+q)^3 \sigma_1^3} \right]$$

,

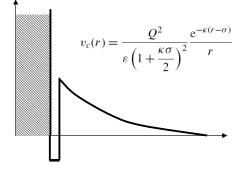
with  $q = \sigma_2/\sigma_1$  the ratio of diameters.

Asakura & Oosawa, 58<sub>59</sub>

#### Theory for colloidal stability

DLVO theory: (Derjaguin-Landau-Vewey-Overbeek)

Debye-Hückel screened Coulomb potential + attractive interaction



#### Integral equation theory

Relate g(r) [or S(q)] to V(r)

At low concentration  $g(r) = \exp(-V(r) / k_B T) \approx 1 - V(r) / k_B T$ Boltzmann approximation (weak interactions)

Make expansion around uniform state [Ornstein-Zernike eq.]

$$g(r) = 1 - n c(r) - [3 \text{ particle}] - [4 \text{ particle}] - \dots$$

$$= 1 - n c(r) - n^2 c(r) * c(r) - n^3 c(r) * c(r) * c(r) - \dots$$
[\* = convolution]  $c(r)$  = direct correlation function

$$S(q) = 1 - n c(q) - n^2 c(q)^2 - n^3 c(q)^3 - \dots = \frac{1}{1 - nc(q)}$$

 $\Rightarrow$  but we still need to relate c(r) to V(r) !!!

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#### Closure relations

Systematic density expansion....

Mean-spherical approximation MSA:  $c(r) = -V(r) / k_B T$  (analytical solution for screened Coulomb potential

- but not accurate for low densities)

Percus-Yevick approximation PY:  $c(r) = g(r) \left[ \exp(V(r) / k_B T) - 1 \right]$  (analytical solution for hard-sphere potential + sticky HS)

Hypernetted chain approximation HNC:

$$c(r) = -V(r) / k_B T + g(r) - 1 - \ln(g(r))$$

(Only numerical solution

- but quite accurate for Coulomb potential)

Rogers and Young closure RY:

Combines PY and HNC in a self-consistent way

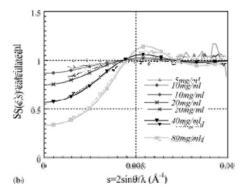
(Only numerical solution

- but very accurate for Coulomb potential)

#### $\alpha$ -crystallin

#### 

Fig. 1. Variation of normalized scattered intensity of  $\alpha$ -crystallin solutions as a function of protein concentration, at 20°C, in 150 mM phosphate buffer at pH = 6.8.

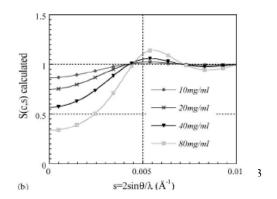


#### S. Finet, A. Tardieu

#### **DLVO** potential

$$v(r) = \begin{cases} Z_{\rm p}^2 e^2 / \{4\pi \varepsilon_0 \varepsilon k \operatorname{Tr} (1 + 0.5\sigma/\lambda_{\rm D})^2\} \\ \times \exp\left[-(r - \sigma)/\lambda_{\rm d}\right] \\ -J\sigma/r \exp\left[-(r - \sigma)/d\right] & \text{for } r > \sigma, \\ +\infty & \text{for } r \leqslant \sigma. \end{cases}$$

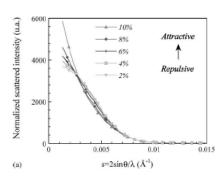
#### HNC and numerical solution



#### α-crystallin + PEG 8000

#### S. Finet, A. Tardieu

#### Depletion interactions:

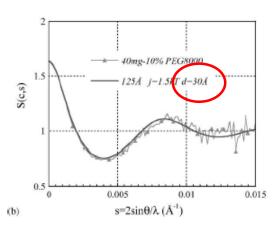


40 mg/ml  $\alpha$ -crystallin solution pH 6.8, 150 mM ionic strength.

#### **DLVO** potential

$$v(r) = \begin{cases} Z_{\rm p}^2 e^2 / \{4\pi \varepsilon_0 \varepsilon k \operatorname{Tr} (1 + 0.5\sigma/\lambda_{\rm D})^2\} \\ \times \exp\left[-(r - \sigma)/\lambda_{\rm d}\right] \\ -J\sigma/r \exp\left[-(r - \sigma)/d\right] & \text{for } r > \sigma, \\ +\infty & \text{for } r \leqslant \sigma. \end{cases}$$

#### HNC and numerical solution



#### **Anisotropy**

$$\frac{d\sigma(q)}{d\Omega} = \Delta \rho^2 V^2 \left[ \sum_{i} F_i(q, e_i)^2 + \frac{1}{N} \sum_{i,j} F_i(q, e_i) F_j(q, e_j) [S_{i,j}(q, e_i, e_j) - 1] \right]$$

Small: decoupling approximation (Kotlarchyk and Chen,1984):

$$\frac{\mathrm{d}\sigma(q)}{\mathrm{d}\Omega} = \Delta \rho^2 V^2 P(q) [1 + \beta(q)(S(q) - 1)],$$

$$\beta(q) = \langle F(q) \rangle_o^2 / \langle F^2(q) \rangle_o,$$

Measured structure factor:

$$S_{meas}(q) = \frac{\frac{\partial \sigma(q)}{\partial \Omega}}{n\Delta \rho^2 P(q)} = 1 + \beta(q) [S(q) - 1] \neq S(q) \quad !!!!!!$$

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#### Large anisotropy

Polymers, cylinders...

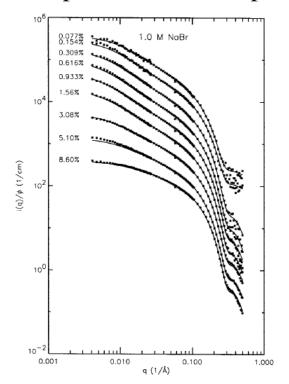
Anisotropy, large: Random-Phase Approximation (RPA):

$$\frac{d\sigma}{d\Omega}(q) = n\Delta\rho^{2}V^{2} \frac{P(q)}{1 + vP(q)}$$
  $v \sim concentration$ 

Anisotropy, large: Polymer Reference Interaction Site Model (PRISM) Integral equation theory – equivalent site approximation

$$\frac{d\sigma}{d\Omega}(q) = n\Delta \rho^2 V^2 \frac{P(q)}{1 - nc(q)P(q)}$$
 c(q) direct correlation function related to FT of  $V(r)$ 

#### **Empirical PRISM expression**



SDS micelle in 1 M NaBr

$$\frac{d\sigma}{d\Omega}(q) = n\Delta\rho^2 V^2 \frac{P(q)}{1 - nc(q)P(q)}$$

c(q) = rod formfactor - empirical from MC simulation

Arleth, Bergström and Pedersen

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#### Overview: Available Structure factors

- 1) Hard-sphere potential: Percus-Yevick approximation
- 2) Sticky hard-sphere potential: Percus-Yevick approximation
- 3) Screened Coulomb potential:

Mean-Spherical Approximation (MSA).

Rescaled MSA (RMSA).

Thermodynamically self-consistent approaches (Rogers and Young closure)

- 4) Hard-sphere potential, polydisperse system: Percus-Yevick approximation
- 5) Sticky hard-sphere potential, polydisperse system: Percus-Yevick approximation
- 6) Screened Coulomb potential, polydisperse system: MSA, RMSA,
- 7) Cylinders, RPA
- 8) Cylinders, `PRISM':
- 9) Solutions of flexible polymers, RPA:
- 10) Solutions of semi-flexible polymers, 'PRISM':
- 11) Solutions of polyelectrolyte chains 'PRISM':

#### **Summary**

- Model fitting and least-squares methods
- Available form factors
   ex: sphere, ellipsoid, cylinder, spherical subunits...
   ex: polymer chain
- Monte Carlo integration for form factors of complex structures
- Monte Carlo simulations for
- •form factors of polymer models
- Concentration effects and structure factors
   Zimm approach
   Spherical particles
   Elongated particles (approximations)
   Polymers

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#### Literature

Jan Skov Pedersen, *Analysis of Small-Angle Scattering Data from Colloids and Polymer Solutions: Modeling and Least-squares Fitting* (1997). *Adv. Colloid Interface Sci.*, **70**, 171-210.

Jan Skov Pedersen

Monte Carlo Simulation Techniques Applied in the Analysis of Small-Angle Scattering Data from Colloids and Polymer Systems

in Neutrons, X-Rays and Light

P. Lindner and Th. Zemb (Editors) 2002 Elsevier Science B.V.

p. 381

Jan Skov Pedersen

Modelling of Small-Angle Scattering Data from Colloids and Polymer Systems

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P. Lindner and Th. Zemb (Editors) 2002 Elsevier Science B.V.

p. 391

Rudolf Klein

#### Interacting Colloidal Suspensions

in Neutrons, X-Rays and Light

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p. 351