シンクロトロン放射光を用いた単分散ミセルの構造解析 -プラト ニックミセルの提案-

Characterizing Self-Assembled Nanoparticles Employed in DDS

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奈良先端大院 12月2日

Outline

Self-assembly and Drug Delivery System (DDS) Strength of SAS technique in exploring DDS particles Examples from Our Recent Studies

- Polymeric Micelle for Delivering Hydrophobic Drugs

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Example: Therapeutic Efficacy Strongly Depends on its Size



Tumour accumulation with different diameters.

In vivo real-time microdistribution of DACHPt/m with different diameters in tumours

The sizes less than 100 nm are very good for SAS.

Accumulation of sub-100 nm polymeric micelles in poorly permeable tumours depends on size K. Kataoka et al., *Nature Nanotechnology* (2011)





Diameter / nm

Goal: Precise Structural Analysis and Visualization of DDS nano-particles by use of SAXS with combination of FFF/MALS

Molecular Design Product Control FDA approval Fundamental Physics and Chemistry

People are really care about what is going to be injected into your blood vessel.

FFF: filed flow fluctuation, MALS: multi-angle light scattering

what I am afraíd ís what's ínsíde !!.



aFFF coupled with MALS



Asymmetric-Flow Field-Flow-Fractionation (aFFF or FFF) is a one-phase chromatography.

Channel flow + cross flow. i.e., No matrix material (do not care adsorption)

Optical purification of SLS in aqueous solutions.

aFFF (Asymmetric flow field flow fractionation)



Dendron-Stabilized Gold Nanoparticles



AFFF fractograms of M5 (10, 20, 30, 40, and 60 nm) AuNP mixture in 0.02 % NaN₃ mobile phase. 90° MALS (LS) / UV traces (-)/(--)

De-Hao TSai (in Hackley's group) JACS 2011, 8884

DLSとaEFF/MALSの比較

Figure 6. Examples of size distribution for PS-latex nanoparticle suspensions determined by DLS using cumulant analytical method. (a) STADEX SC-0110-D and (b) T0625.



Figure 7. Examples of size distribution for PS-latex nanoparticle dispersions determined by AFFFF-MALS. (a) STADEX SC-0110-D and (b) T0625.



Nanomaterials 2012, 2, 15-30

SAXS at SPring-8 and our set-up



Scattering angle < 2 degree => 1 – 100 nm Small angle X-ray scattering (SAXS)

$$q = \frac{2\pi}{\lambda} \sin \theta \quad d \propto \frac{1}{q}$$

WAXS vs. SAXS

Scattering angle 5 – 25 degree => less than 0.5 nm Wide angle X-ray scattering (WAXS, XRD)

Structural Factor (構造因子): distances between atoms.

Scattering angle < 2 degree => 1 – 100 nm Small angle X-ray scattering (SAXS)

Structural Factor (構造因子): distance between particles

Form Factor (形状因子): shape



Spherical waves from many points = > Structural factor

Intuitive understanding of diffraction



Diffraction from a slit with finite width => form factor

q: magnitude of the scattering vector => distance from the scattering center







sample



Scattering angle < 2 degree => 1 – 100 nm Small angle X-ray scattering (SAXS)

$$q = \frac{2\pi}{\lambda} \sin \theta \quad d \propto \frac{1}{q}$$

Determining the structure of matter



SAXS 15nm と10nmを協用できる



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Case 1: Polymeric Micelles

PEG; Hydrophilic block



Hydrophobic drug (or equivalent Br-modified)

SAXS+FFF/MALS

- ➔ Aggregation number
- ➔ Core and corona sizes
- → Nature of the interface: Overcrowding of PEG

ASAXS (anomalous SAXS)

- \rightarrow How the drugs are encapsulated in the core
- ➔ Difficult for normal SAXS
- → Br attached molecule as a probe

FFF: filed flow fluctuation, MALS: multi-angle light scattering



poly(ethylene glycol)-*block*-poly(α , β -benzyl-*L*-aspartate) (PEG-PBLA)



n of PEG >> n of Asp \rightarrow Stable Spherical micelle

SLS vs. SAXS from dilute solutions

$$I(\theta) = \frac{M_W}{N_A} C_M \overline{\nu}^2 (\overline{\rho} - \rho_0)^2 F(\theta)$$

- M_W : weight average molecular weight
 - : partial specfic volume

 $\overline{\rho} - \rho_0$: electron density difference between solvent and solute

 $F(\theta) \rightarrow 1$ at low θ and low concentration

SLS (static light scattering) or MALS (multi-angles light scattering)

 ν

- $\overline{v}^2(\overline{\rho}-\rho_0)^2$ can be determined as $\partial n / \partial c$, thus M_W can be evaluated.
- From $F(\theta)$, the radius of gyration ($\langle S^2 \rangle$ or R_g) is determined.
- •Can be coupled with GPC or FFF.

SAXS (small angle X-ray scattering)

- \overline{v}^2 and $(\overline{\rho} \rho_0)^2$ should be determined separately and thus M_W cannot be determined.
- $\mathrm{From}F(\theta)\,$, inner structures can be evaluated.



n = 118

M_w/M_n <1.05

Molecular Characters

code	Asp number	Benzylation / %				
HH9-31-1	26.4	76.9				
HH9-31-2	26.4	81.4				
HH9-24-2	26.4	88.6 (highest)				
HH9-22-2	20.2 (shortest)	83.7				
HH9-22-4	23.8	84.0				
HH9-22-3	29.7	82.2				
HH9-22-5	32.0 (longest)	77.2				
HH9-24-1	26.9	65.8 (lowest)				
HH9-18-1	26.9	83.3				
HH9-32*	26.9	83.3				

Benzylation



Film was sonicated in PBS

* Dialysis

From NMR

FFF coupled with MALS



What determines the aggregation number ?



SAXS from a PEG-Asp/Bzl micelle



•

Optical impurities in LS and SAXS



LS \rightarrow impossible to obtain accurate Mw in batch measurement

Fitting Model for SAXS



SAXS from a PEG-Asp/Bzl micelle



$$G = G_{core}(R_e, \varphi) + G_{in}(\gamma) + G_{shell}(\chi)$$

 G_{core} the entropic elasticity of the core chains determined by $R_{_{\!\!P}}\,/\,R_{_{\!\!C}}$





ell the osmotic energy: balance of the elastic stretching and the excluded volume effect.

The chain length dependence of the core size.

-Sailing theory tells the aggregation number is determined by the entropic packing of the core chains. -No dependence of the shell chain.



 DP_{Asp} dependence of R_C for the same R_{Bzl} .

 DP_{Asp}

20

30

40

 $R_{\rm C}$

/ nm

Scaling theory for polymeric micelles: Zhulina *Macromolecules* **2005**, Halperin, A., *Macromolecules* **1987**, de Gennes, P. G., 1979.

Crowding Nature of the PEG chains in the Corona







When the loading ratio exceeds 9-10 %, LE starts leaking. => What is the inclusion mechanism ?

Neuroblastoma: Pre-incubation of SH-SY5Y human neuroblastoma cells with either RAR-pan-antagonist LE540 or MAP kinase kinase 1 (MEK-1) inhibitor PD98059.

Breast Neoplasms (Breast Cancer) In ZR-75-1 human breast cancer cells, cotreatment of LE135 and LE540 with all-trans-RA inhibited all-trans-RA-induced apoptosis.

SAXS from a PEG-Asp/Bzl micelle



Density change upon loading



What's ASAXS?

Anomalous Small-Angle X-ray Scattering X線小角異常散乱

Resonance Small-Angle X-ray Scattering X線小角共鳴散乱



Reflective index

absorbance

Electron Adsorption Edge

	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18
1						1 H	4											2 He
2	3 Li	4 Be							F 1 .				5 B	6 C	7 N	8 0	9 F	10 Ne
3	11 Na	12 Mg					K-6	edge	5 KE	ev- 20	J Kev		13 Al	14 Si	15 P	16 S	17 CI	18 Ar
4	19 K	20 Ca	21 Sc	22 Ti	23 V	24 Cr	25 Mn	26 Fe	27 Co	28 Ni	29 Cu	30 Zn	31 Ga	32 Ge	33 As	34 Se	35 Br	36 Kr
5	37 Rb	38 Sr	39 Y	40 Zr	41 Nb	42 Mo	43 Tc	44 Ru	45 Rh	46 Pd	47 Ag	48 Cd	49 In	50 Sn	51 Sb	52 Te	53 I	54 Xe
6	55 Cs	56 Ba		72 Hf	73 Ta	74 W	75 Re	76 Os	77 Ir	78 Pt	79 Au	80 Hg	81 TI	82 Pb	83 Bi	84 Po	85 At	86 Rn
7	87 Fr	88 Ra		104 Du	105 Jo	106 Rf	107 Bh	108 Ha	109 Mt									







Photon Energy [eV]

ASAXS Measurement

SAXS measurements were carried out with several X-ray energy near absorption edge.

From the difference of spectra of some X-ray energy, scattering property of specific atom can obtain.

$$I(q)_{Br} \propto C \times \left[\frac{I(q)_{E_1} - I(q)_{E_2}}{f'_{E_1} - f'_{E_2}} - \frac{I(q)_{E_1} - I(q)_{E_3}}{f'_{E_1} - f'_{E_3}} \right]$$
$$C = f'_{E_2} - f'_{E_3} + \frac{f''_{E_1}^2 - f''_{E_2}^2}{f'_{E_1} - f'_{E_2}^2} - \frac{f''_{E_1}^2 - f''_{E_3}^2}{f'_{E_1} - f'_{E_3}} \right]$$



















Increase the S/N ratio by $10^2 - 10^3$.

Masunaga et al, submitted.



-S/N 10-100 times. -Low BG at high *q*. -No peaks from the kapton window -Low noise around beam stopper



10⁶

10⁵

10⁴

10³

10²

air

vacuum

ŗ١

1.5

Intensity / a.u.





13.75

-4 -6

-8 -10 -12 13.25 f'

13.35

Scattering between Overall Micelle and Br

- (1) v^2 term can be fitted with a simple hard-sphere model
- (2) The core is smaller than the Br- hard sphere



SAXS profiles (black circles), resonant terms of the polymeric micelles (blue circles), and theoretical curves calculated from hard sphere models





TCB is Infiltrating into the PEG Domain





TBC distribution in the micelle. TBC is smearing into the PEG densely-packed interface. Probably, because PEG is partially dehydrated due to the overcrowding.



Overall:

- The core-chain's hydrophobicity is the major factor to determine the aggregation number and thus core size.
- *N*_{agg} = 30- 40 at BI = 80% → 100 at 90%
- At the same hydrophobicity, the scaling theory describes the chain length dependence of *R*c and *R*s.

Long period exists. Drug-loading erases the ordering.

The drugs are uniformly distributed in the core.



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Topoic 2: Shape Persistence Micelles for DNA Delivery



Cholic Acid Micelle (or Aggregate): no rule without an exception



Cholic acid, along with chenodeoxycholic acid, is one of two major bile acids produced by the liver.





→ cationic lipids for DNA delivery

Synchrotron Small Angle X-ray Scattering









Na Specificity of the sharp minimum of the intensity

artificial effect in the solvent subtraction.

10¹

• Micelle peak is only one \rightarrow No aggregation

- The red and blue data are almost completely overlapping with each other.
- Molar mass = 5.7×10^3 \rightarrow Aggregation number is 6

FFF for Cal[4]C6

$$I(0) = NV_{M}^{2} (\bar{\rho} - \rho_{0})^{2} = \frac{M_{M}}{N_{A}} C_{M} \bar{v}^{2} (\bar{\rho} - \rho_{0})^{2}$$

 $\overline{
ho}ho_{\scriptscriptstyle 0}$ can be calculated from the atomic scattering factor .

 \overline{v} by measuring the density increment

Need an assumption ! No information about the distribution

The molar masses of C3 and C6 are 600 and 14800.

The molar masses determined with different methods and the aggregation numbers.

sample	SAXS	FFF+N	IALS	AUC	Aggregation	
	$M_{ m w}$ / 10 ³	$M_{_W}$ / 10 ³	M_w/M_n	<i>M_w</i> / 10 ³	M_z/M_w	number
CaL[4]C3	6.00 ± 0.20	5.69 ± 0.93	1.007	6.10 ±0.20	1.0 ₇	6
CaL[4]C6	14.8 ± 0.95	13.0 - 20	1.5	14.7 ± 0.90	1.5	10 - 16

SAXS: synchrotron small-angle X-ray scattering. LS: static light scattering.

AUC: analytical ultracentrifugation

Shape Determination with Dummy Atom Model

(Cross section)

Ab initio Shape Determination by Simulated Annealing using Bead Model Restriction: cubic symmetry (Dmitri Svergun)

At low pH and presence of Na⁺ or K⁺

Fujii et al., Langmuir, 2012, 28 (6), 3092–3101