

A semi-phenomenological approach to the structure and transport properties of macromolecules in solution

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A semi-phenomenological approach to the structure and transport properties of macromolecules in solution

---- *Content*

■ Introduction & Motivation

➤ Macromolecules in Solution

✚ Structure and Transport properties of the solution

■ Macroscopic vs. microscopic theory of molecules in solution

■ Transport properties vs. semi-phenomenological approach

➤ Fokker—Planck equation and friction tensors

■ Results

➤ Translational motion

➤ Rotational motion

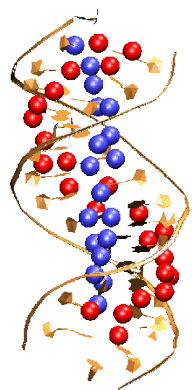
■ Summary and outlook

Macromolecules in solution

---- Introduction & Motivation

IUPAC definition:

A macromolecule is a larger complex molecule consisting of many smaller structural units (molecules, group of atoms, atoms) linked together.



Macromolecules
(polymers, proteins, lipids, PEO, ...)

Biophysics

Chemical physics

Material science

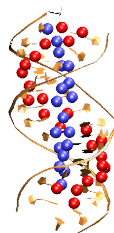
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Macromolecules in solution

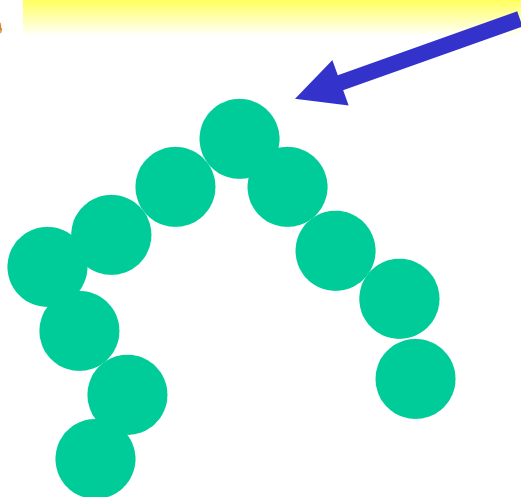
---- Introduction & Motivation

IUPAC definition:

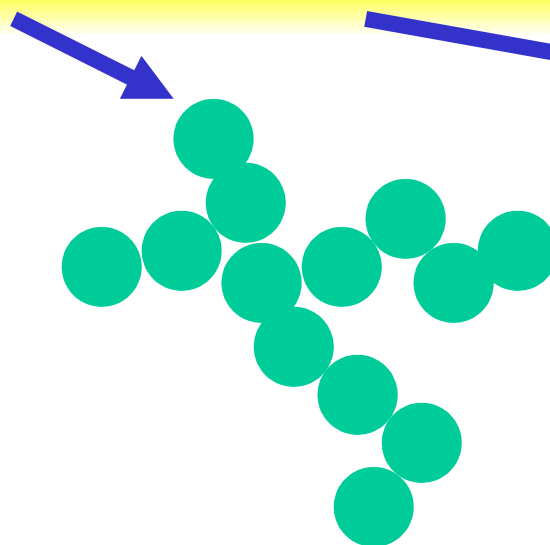
A macromolecule is a larger complex molecule consisting of many smaller structural units (molecules, group of atoms, atoms) linked together.



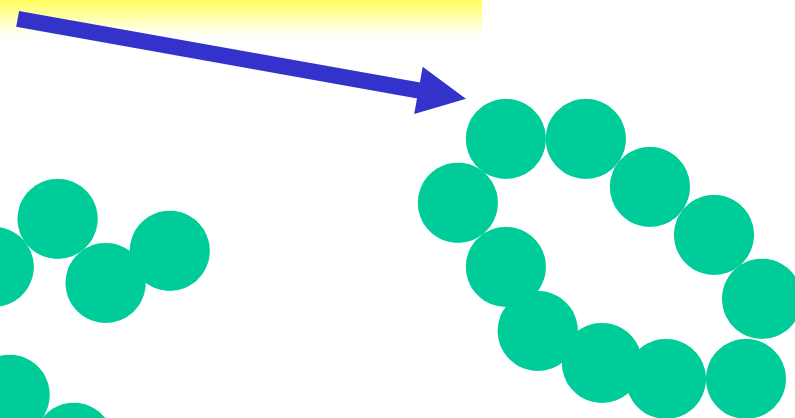
Structural unit (so-called *bead* of the macromolecule)



Chain-shape



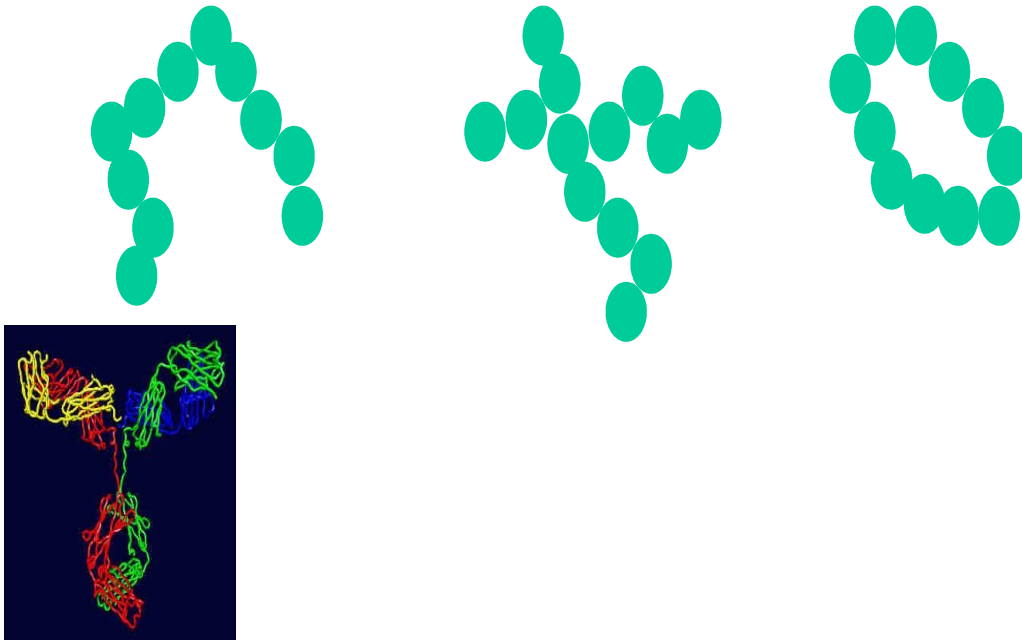
Dendrimer-shape



Circle-shape

Macromolecules in solution

---- Structure & Transport properties



Macromolecules:

- Very massive and complex (10^3 - 10^5 atoms)
- Slow ($v_{\text{Macr}} / v_{\text{H}_2\text{O}} < 10^3$)
- Time scale of relaxation is much larger than the typical relaxation times of simple solvents

Translation diffusion coefficient:

$$D_T = \frac{\langle (R_a(0) - R_a(t))^2 \rangle}{6t}$$

Center-of-mass (CM) diffusion coefficient:

$$D^{(\text{CM})} = \frac{\langle (R_{\text{cm}}(0) - R_{\text{cm}}(t))^2 \rangle}{6t}$$

Rotational diffusion coefficient:

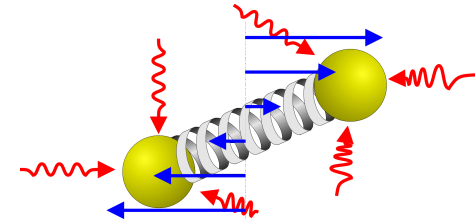
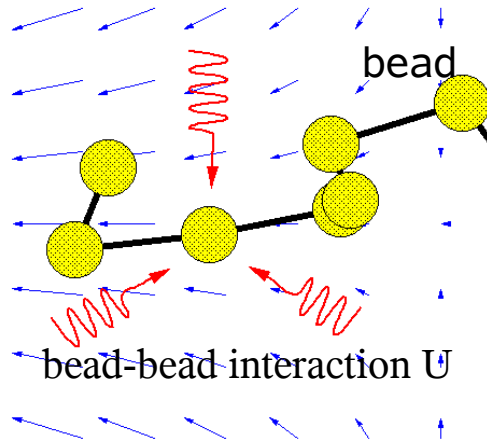
$$D_R = \frac{\langle (\Delta\theta(\Delta t))^2 \rangle}{\Delta t}$$

Time dependent correlation functions:

$$CF(t) = \langle A(0)A(t) \rangle$$

Dynamics of the macromolecules in solution

---- *Macroscopic vs. Microscopic*



Assumptions

- Molecule has internal structure (non-uniform).
- Molecule much more heavy and slower than solvent particles

Macroscopic (phenomenological)

- Solvent is a uniform non-compressible medium.
- Phenomenological boundary conditions
- Bead-solvent interaction: via random forces (Gaussian distributed)

Basic mathematical assumptions:

- The beads are assumed to be “embedded” to the solvent flow

(Stoke’s law: $\mathbf{F} = -\xi \mathbf{v}$)

- The transport and diffusion are described via the Diffusion Eq. (Fick’s laws)

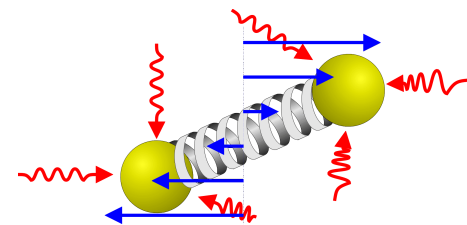
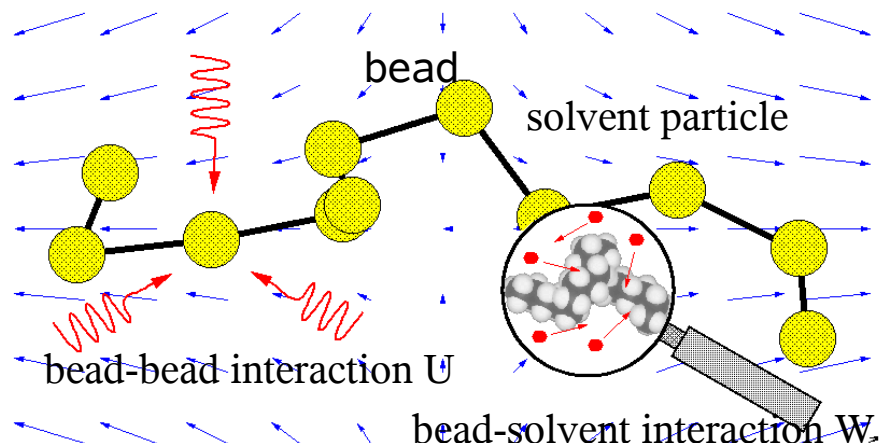
The dynamical behaviour of the macromolecules more complex!

„... these mathematical assumptions must be questioned since macromolecule can not be considered as „embedded“ into the surroundings but dramastically start to „emancipate“ from the solvent“.

B. Dünweg and K. Kremer, *J. Chem. Phys.* 99 (1993) 6984.

Dynamics of the macromolecules in solution

---- *Macroscopic vs. Microscopic*



Assumptions

- Molecule has internal structure (non-uniform).
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Macroscopic (phenomenological)

- Solvent is a uniform non-compressible medium.
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- Bead-solvent interaction: via random forces (Gaussian distributed)

Microscopic

- Solvent as set of particles with given mass m
- Discrete structure of the solvent ('solvent-solvent' interaction V)
- Bead-solvent interaction: via potentials W

Semi-phenomenological approach to the study of the macromolecular dynamics

Total energy of the system (Hamiltonian)

$$H = H_M + H_S + H_{MS}$$

Macromolecular Dynamical Simulations
(Newton's equations)

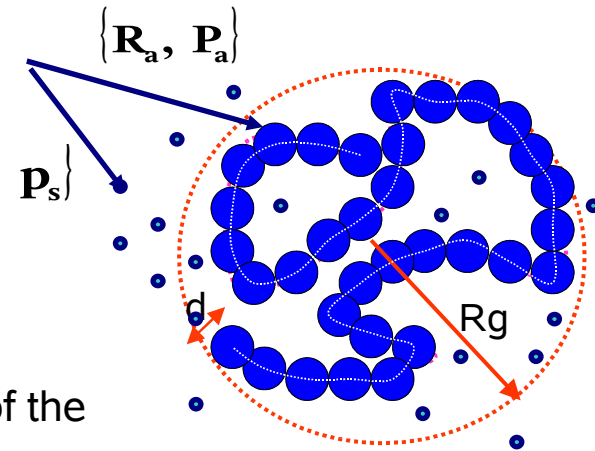
Phase-space formalism
(Liouville equation)

Microscopic theory of molecules in solution

---- Molecular Dynamics Simulation (MDS)

$$\mathbf{H} = \mathbf{H}_M + \mathbf{H}_S + \mathbf{H}_{MS}$$

$$= \underbrace{\sum_{a=1}^N \frac{\mathbf{p}_a^2}{2M}}_M + \underbrace{\sum_{a < b}^N U(|\mathbf{R}_a - \mathbf{R}_b|)}_M + \underbrace{\sum_{s=1}^n \frac{\mathbf{p}_s^2}{2m}}_S + \underbrace{\sum_{s < t}^n V(|\mathbf{r}_s - \mathbf{r}_t|)}_S + \underbrace{\sum_{a=1}^N \sum_{s=1}^n W(|\mathbf{R}_a - \mathbf{r}_s|)}_{MS} \quad \{\mathbf{r}_s, \mathbf{p}_s\}$$



Equations of the motion

Numerical solution of the equations of the motion for given interactions U , V , W

$$\frac{d\mathbf{R}_a}{dt} = \frac{\partial [\mathbf{H}_M + \mathbf{H}_{MS}]}{\partial \mathbf{P}_a}, \quad \frac{d\mathbf{P}_a}{dt} = -\frac{\partial [\mathbf{H}_M + \mathbf{H}_{MS}]}{\partial \mathbf{R}_a}$$

$$\frac{d\mathbf{r}_s}{dt} = \frac{\partial [\mathbf{H}_S + \mathbf{H}_{MS}]}{\partial \mathbf{p}_s}, \quad \frac{d\mathbf{p}_s}{dt} = -\frac{\partial [\mathbf{H}_S + \mathbf{H}_{MS}]}{\partial \mathbf{r}_s}$$

MDS covers only time relaxation processes with duration approx. 100nsec, while relaxation time of the typical macromolecules 10^{-5} - 10^{-3} sec, and longer.

Simulation of the relaxation processes of 1 ms is $\sim 10^9$ more expensive than our abilities in 2006

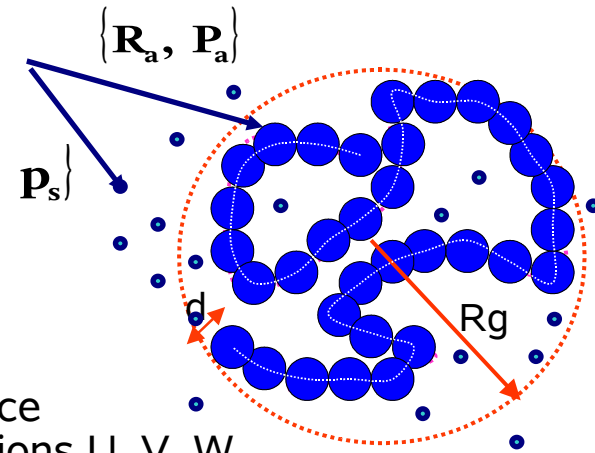
(A recent “larger” MD Simulation:
~10² more expensive than our
(7400 water molecules + macromolecules)-
118 ns duration

Moore's law: this might be possible not before 2050

---- *phase-space dynamics*

$$\mathbf{H} = \mathbf{H}_M + \mathbf{H}_S + \mathbf{H}_{MS}$$

$$= \underbrace{\sum_{a=1}^N \frac{\mathbf{p}_a^2}{2M}}_M + \underbrace{\sum_{a < b}^N U(|\mathbf{R}_a - \mathbf{R}_b|)}_M + \underbrace{\sum_{s=1}^n \frac{\mathbf{p}_s^2}{2m}}_S + \underbrace{\sum_{s < t}^n V(|\mathbf{r}_s - \mathbf{r}_t|)}_S + \underbrace{\sum_{a=1}^N \sum_{s=1}^n W(|\mathbf{R}_a - \mathbf{r}_s|)}_{MS} \quad \{\mathbf{r}_s, \mathbf{p}_s\}$$



Liouville equation (LE)

Liouville equations for the phase-space distribution function for given interactions U, V, W

$$\frac{\partial \rho}{\partial t} = [\hat{L}_M + \hat{L}_S + \hat{L}_{MS}] \rho \left(\{R, P\}, \{r, p\}; t \right)$$

Liouville Eq. describes fully the time evolution of the phase-space df of the overall system 'macromolecule + solvent' and contains all information about dynamics of the system.

Liouville operators: $\hat{L}_i = \sum_i \left[\frac{\partial \mathbf{H}_i}{\partial \mathbf{r}_i} \frac{\partial}{\partial \mathbf{p}_i} + \frac{\partial \mathbf{H}_i}{\partial \mathbf{p}_i} \frac{\partial}{\partial \mathbf{r}_i} \right], \quad i = M, S, MS$

$$\langle \mathbf{B}(\{\mathbf{R}(t), \mathbf{P}(t)\}, \{\mathbf{r}(t), \mathbf{p}(t)\}) \rangle = \int \mathbf{B}(\{\mathbf{R}, \mathbf{P}\}, \{\mathbf{r}, \mathbf{p}\}) \rho(\{\mathbf{R}, \mathbf{P}\}, \{\mathbf{r}, \mathbf{p}\}; t) d\mathbf{R}, \mathbf{P} d\mathbf{r}, \mathbf{p}$$

Integration of the Liouville Eq. has the same difficulties like in the case of MDS

Fortunately, this ansatz can be the starting point for further simplifications.

Microscopic theory of molecules in solution

---- *phase-space dynamics*

Liouville equation

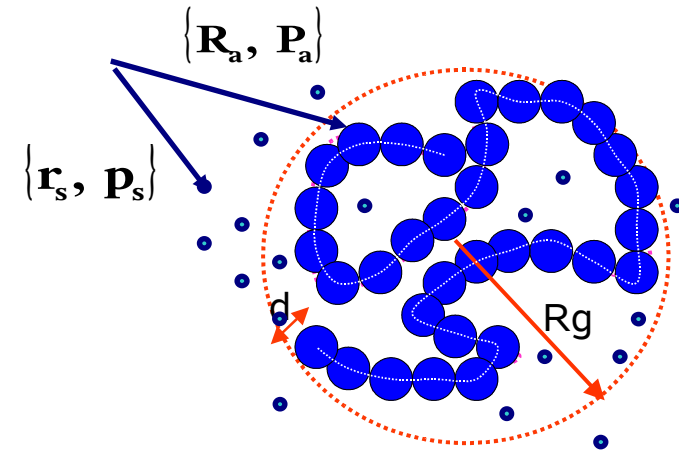
$$\frac{\partial \rho}{\partial t} = [\hat{L}_M + \hat{L}_S + \hat{L}_{MS}] \rho(\{\mathbf{R}, \mathbf{P}\}, \{\mathbf{r}, \mathbf{p}\}; t)$$

Liouville operators

$$\hat{L}_i = \sum_i \left[\frac{\partial H_i}{\partial \mathbf{r}_i} \frac{\partial}{\partial \mathbf{p}_i} + \frac{\partial H_i}{\partial \mathbf{p}_i} \frac{\partial}{\partial \mathbf{r}_i} \right], \quad i = M, S, MS$$

Integration over all coordinates of the solvent

$$\rho_N(\{\mathbf{R}_a, \mathbf{P}_a\}; t) = \int d\{\mathbf{r}_s, \mathbf{p}_s\} \rho(\{\mathbf{R}_a, \mathbf{P}_a\}, \{\mathbf{r}_s, \mathbf{p}_s\}; t)$$



Initial conditions:

- Solvent is in equilibrium state Φ_{eq}
- Macromolecule and solvent do not interact before initial time
- At the initial time, the full distribution can be factorized as $\rho(0) = \rho_N(0) \Phi_{eq}$

$$\frac{\partial \rho_N}{\partial t} = Z(...; U, W)$$

Bead-bead

Bead-solvent

Projection operator formalism:

$$\hat{P}(...) = \int d\{\mathbf{r}_s, \mathbf{p}_s\} (...) \quad \hat{Y} = 1 - \hat{P}$$

Microscopic theory of molecules in solution

---- Fokker-Planck equation & Friction tensor

Liouville equation

$$\frac{\partial \rho}{\partial t} = [\hat{L}_M + \hat{L}_S + \hat{L}_{MS}] \rho(\{\mathbf{R}, \mathbf{P}\}, \{\mathbf{r}, \mathbf{p}\}; t)$$

Liouville operators

$$\hat{L}_i = \sum_i \left[\frac{\partial \mathbf{H}_i}{\partial \mathbf{r}_i} \frac{\partial}{\partial \mathbf{p}_i} + \frac{\partial \mathbf{H}_i}{\partial \mathbf{p}_i} \frac{\partial}{\partial \mathbf{r}_i} \right], \quad i = M, S, MS$$

Integration over all coordinates of the solvent

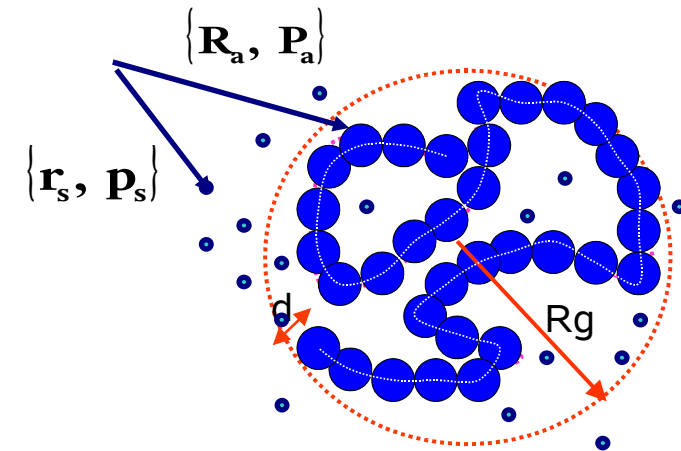
$$\rho_N(\{\mathbf{R}_a, \mathbf{P}_a\}; t) = \int d\{\mathbf{r}_s, \mathbf{p}_s\} \rho(\{\mathbf{R}_a, \mathbf{P}_a\}, \{\mathbf{r}_s, \mathbf{p}_s\}; t)$$

Fokker-Planck equation

$$\frac{\partial \rho_N}{\partial t} + \frac{P_a}{M} \frac{\partial \rho_N}{\partial R_a} - \frac{\partial U(|\mathbf{R}_a - \mathbf{R}_b|)}{\partial P_a} \frac{\partial \rho_N}{\partial P_a} = \frac{\partial}{\partial P_a} \left[\frac{\partial}{\partial P_b} + \frac{P_b}{k_B T M} \right] \rho_N$$

Bead-bead

Friction tensors



dynamical coupling between solvent and molecule

Structure factor of the solution

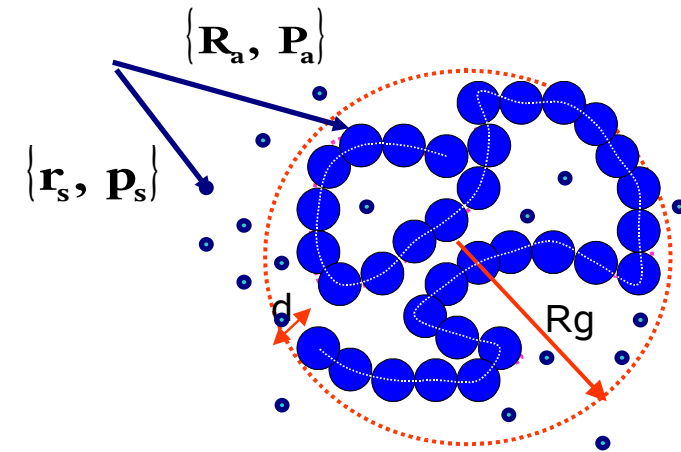
bead-solvent interaction

Fokker-Planck equation & Friction tensors

---- *phase-space dynamics*

$$\frac{\partial \rho_N}{\partial t} + \frac{P_a}{M} \frac{\partial \rho_N}{\partial R_a} - \frac{\partial U(|R_a - R_b|)}{\partial P_a} \frac{\partial \rho_N}{\partial P_a} = \frac{\partial}{\partial P_a} \xi^{ab} \left[\frac{\partial}{\partial P_b} + \frac{P_b}{k_B T M} \right] \rho_N$$

Thermodynamic limit: $n \rightarrow \infty; V \rightarrow \infty; n/V \rightarrow n_0$



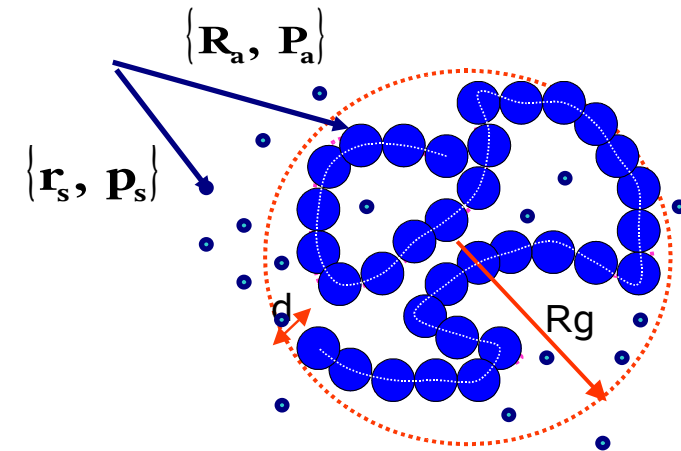
Series expansion of the time evolution operator: $\exp[-t \hat{Y} \hat{L}_{MS}] = \sum_{j=0}^{\infty} \frac{t^j}{j!} [\hat{Y} \hat{L}_{MS}]^j$

$$\xi_{\alpha\beta}^{(ab)} = \sum_j \xi_{\alpha\beta}^{(ab)}{}_{[j]} = \xi_{\alpha\beta}^{(ab)}{}_0 + \xi_{\alpha\beta}^{(ab)}{}_{[1]} + \xi_{\alpha\beta}^{(ab)}{}_{[2]} + \dots$$

Fokker-Planck equation & Friction tensors

---- *phase-space dynamics*

$$\frac{\partial \rho_N}{\partial t} + \frac{P_a}{M} \frac{\partial \rho_N}{\partial R_a} - \frac{\partial U(|R_a - R_b|)}{\partial P_a} \frac{\partial \rho_N}{\partial P_a} = \frac{\partial}{\partial P_a} \xi^{ab} \left[\frac{\partial}{\partial P_b} + \frac{P_b}{k_B T M} \right] \rho_N$$



Translational motion

- ✓ friction tensors parameters
- ✓ single-bead molecules
- ✓ dumbbell molecules
- ✓ N-bead macromolecules
- *various masses, shapes and bead-bead interactions*
- *different topologies*
- ...

Rotational motion

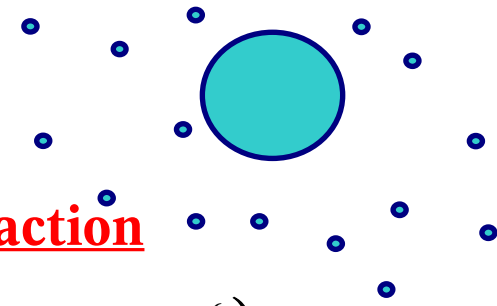
Translational motion of the macromolecule

---- a few, short examples

1. Friction tensor for the various bead-solvent interaction

Charge particles: (Coulomb-like interaction)

Neutral particles: (Born-Mayer and van der Waals interaction)



2. Diffusion coefficient for the „weak“ van der Waals interaction

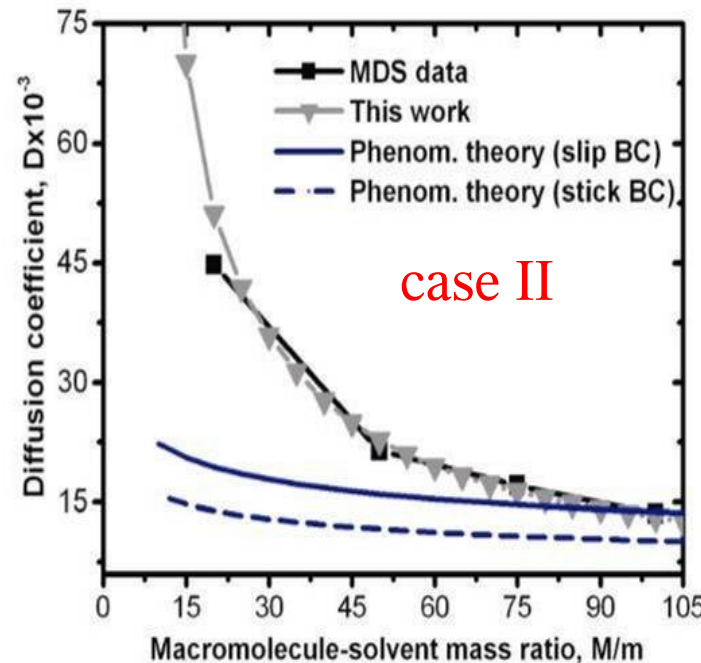
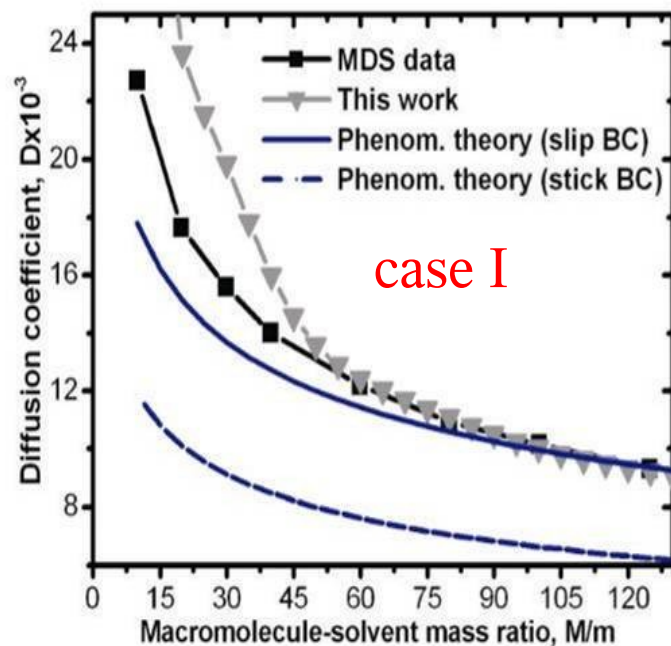
$$\mathbf{D}^{(\text{cm})} = k_B T \xi^{(\text{cm})-1}$$

$$W = \begin{cases} 4\epsilon_{MS} \left(\left[\frac{\sigma}{(r-l_M)} \right]^{12} - \left[\frac{\sigma}{(r-l_M)} \right]^6 \right) & \text{if } r \leq 2.5\sigma \\ 0, & \text{otherwise} \end{cases}$$

A: Single-bead macromolecule:

Diffusion coefficient as function of the mass ratio M/m .

Equal mass density of the bead and solvent particles



case I

$$k_B T = \epsilon_{SS}; \quad n_0 = 0.85 / \sigma^3$$

MD: J. Schmidt and S. Skinner,
J. Chem. Phys. 119, 8062 (2003).

case II

$$k_B T = 2.95\epsilon_{SS}; \quad n_0 = 0.60 / \sigma^3$$

MD: Ould-Kaddour et al.,
Phys. Rev. E 63, 011205 (2001).

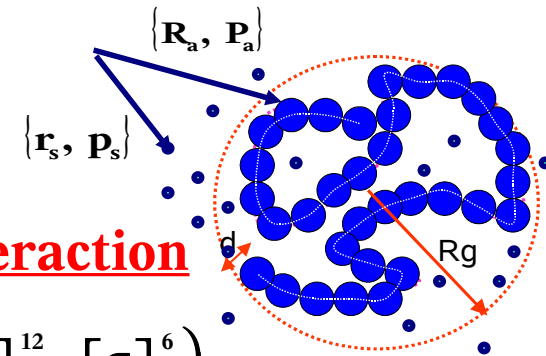
Translational motion of the macromolecule

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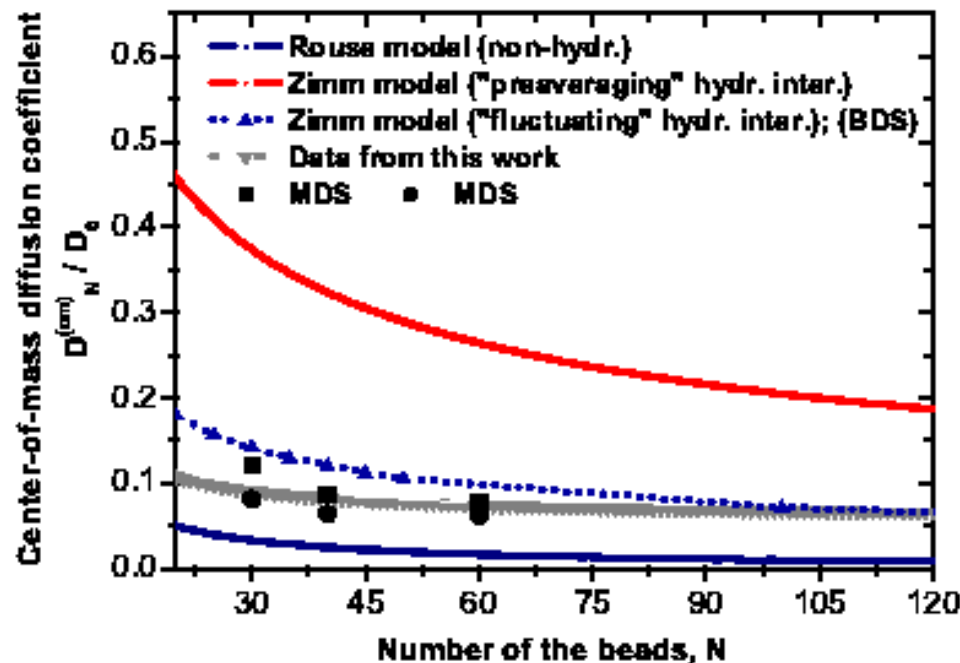
2. Diffusion coefficient for the „weak“ van der Waals interaction

$$\mathbf{D}^{(\text{cm})} = k_B T \xi^{(\text{cm})-1}$$

$$W = \begin{cases} 4\epsilon_{\text{MS}} \left(\left[\frac{\sigma}{r} \right]^{12} - \left[\frac{\sigma}{r} \right]^6 \right) & \text{if } r \leq 2.5\sigma \\ 0, & \text{otherwise} \end{cases}$$

B: *N*-bead chain molecule:

CM Diffusion coefficient as function of the degree of polymerization *N*



$$k_B T = 1.12 \epsilon_{\text{SS}}; \quad n_0 = 0.90 / \sigma^3$$

MD (square): B. Dünweg et al.,
J. Chem. Phys. 117, 930 (2002).

MD (hexagonal): K. Kremer et al.,
J. Chem. Phys. 119, 6983 (1993).

Translational motion of the macromolecule

---- a few, short examples

1. Friction tensor for the various bead-solvent interaction

Charge particles: (*Coulomb-like interaction*)

Neutral particles: (*Born-Mayer and van der Waals interaction*)

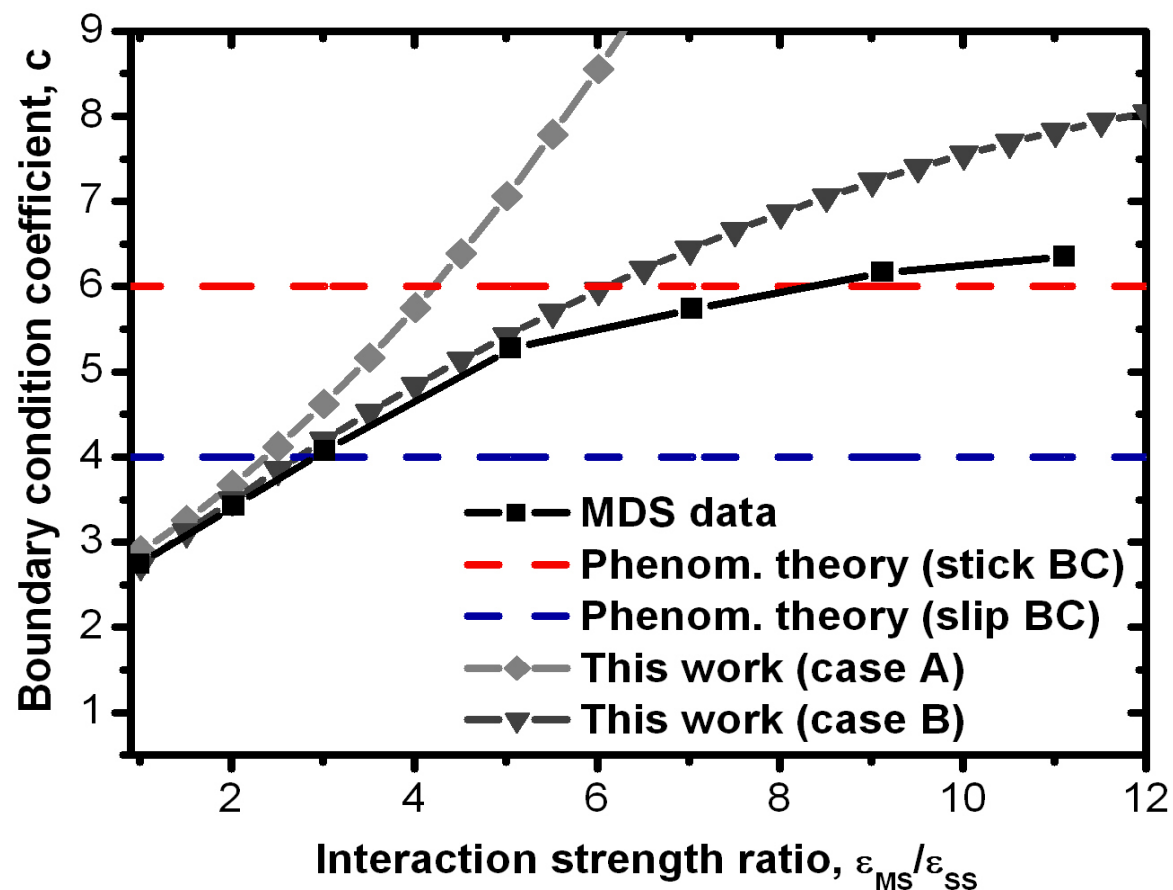
2. Diffusion coefficient for the „weak“ van der Waals interaction

3. Transition from slip to stick boundary condition coefficient

$$c = \frac{k_B T}{D^{(cm)} \pi \eta R_{hydr}}$$

Boundary condition coefficient as
function of the strength ratio $\epsilon_{MS}/\epsilon_{SS}$

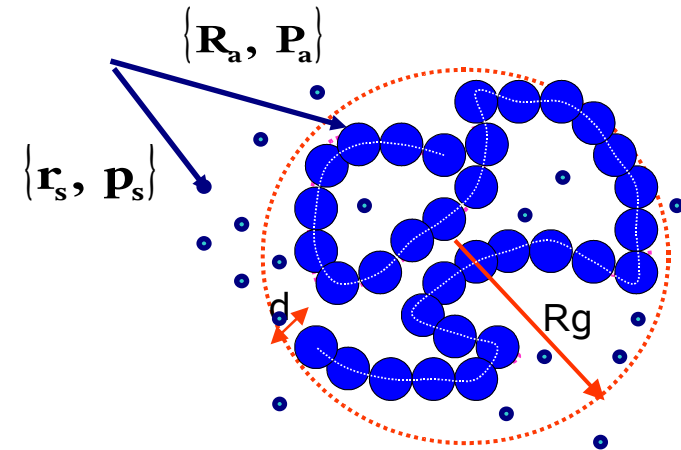
$M/m=200$



Fokker-Planck equation & Friction tensors

---- *phase-space dynamics*

$$\frac{\partial \rho_N}{\partial t} + \frac{P_a}{M} \frac{\partial \rho_N}{\partial R_a} - \frac{\partial U(|R_a - R_b|)}{\partial P_a} \frac{\partial \rho_N}{\partial P_a} = \frac{\partial}{\partial P_a} \xi^{ab} \left[\frac{\partial}{\partial P_b} + \frac{P_b}{k_B T M} \right] \rho_N$$



Translational motion

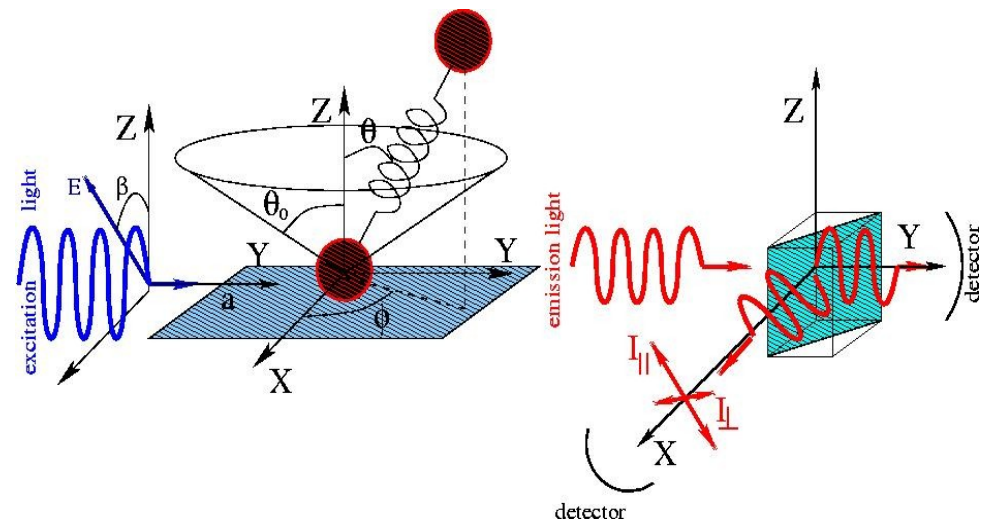
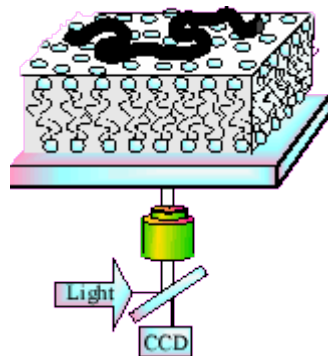
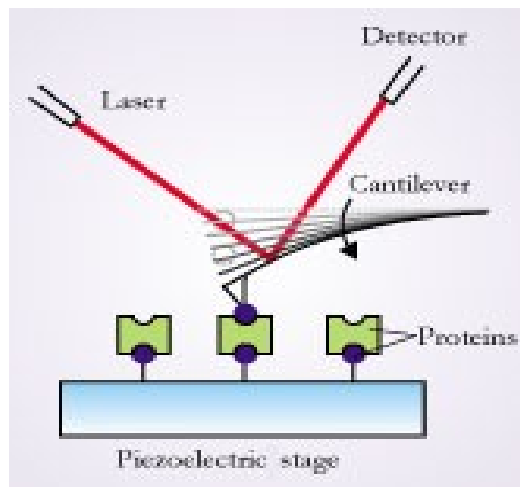
- ✓ friction tensors parameters
- ✓ single-bead molecules
- ✓ dumbbell molecules
- ✓ N-bead macromolecules
- *various masses, shapes and bead-bead interactions*
- *different topologies*
- ...

Rotational motion

- ✓ Restricted rotation of dumbbell molecules
- ✓ Bead-bead and bead-surface interactions
- *3- and 4-bead chains*
- *different topologies*
- ...

Rotational motion of macromolecules

---- *immobilized on the surface*



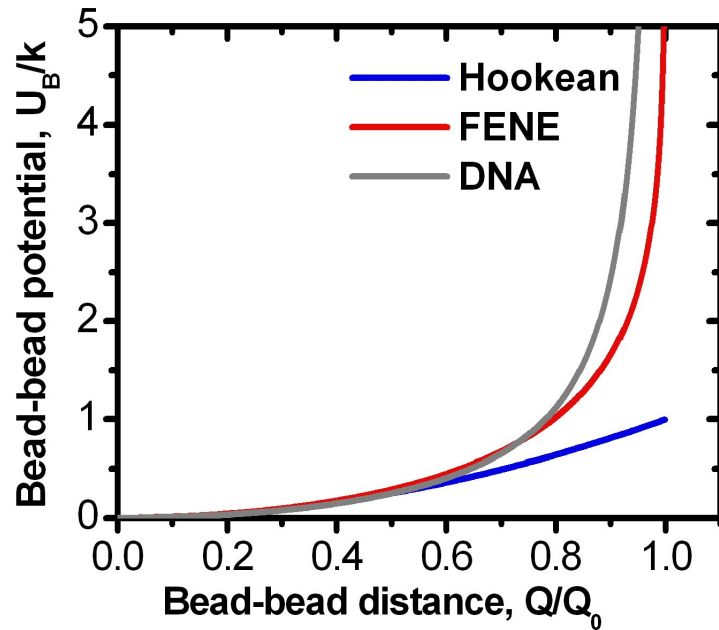
Rotational diffusion coefficient:
$$D_R = \frac{\langle (\Delta\theta(\Delta t))^2 \rangle}{\Delta t}$$

Orientation correlation function:
$$C(t) = \frac{1}{2} \langle 3 \cos^2 \theta(t) - 1 \rangle$$

Phenomenological view point
of the solvent

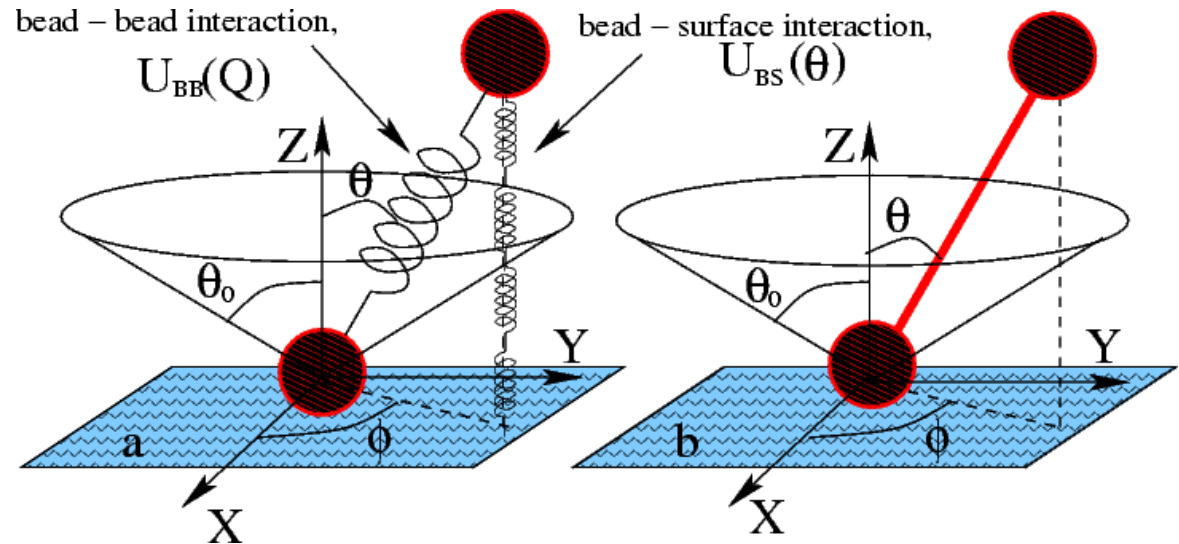
Rotational motion of macromolecule

---- *immobilized on the surface*



Bead-bead interaction:

- Hookean
- FENE
- Fraenkel
- DNA-type



Bead-surface interaction:

- Cone
- Effective double well (Sin)

$$U_B^{\text{DNA}} = k_{\text{DNA}} \left(\frac{1}{4 \left(1 - \frac{Q}{Q_0} \right)} - \frac{1}{4} Q + \frac{Q^2}{2Q_0^2} - \frac{1}{4} \right)$$

Rotational motion of macromolecule

---- Configuration-space distribution

Fokker-Planck equation

$$\frac{\partial \rho_N}{\partial t} + \frac{P_a}{M} \frac{\partial \rho_N}{\partial R_a} - \frac{\partial U(|R_a - R_b|)}{\partial P_a} \frac{\partial \rho_N}{\partial P_a} = \frac{\partial}{\partial P_a} \xi^{ab} \left[\frac{\partial}{\partial P_b} + \frac{P_b}{k_B T M} \right] \rho_N$$

Restricted boundary conditions

$$0 \leq \theta \leq \theta_0 (\leq \pi/2)$$

$$0 \leq \varphi \leq 2\pi$$

$$0 \leq Q \leq Q_0$$

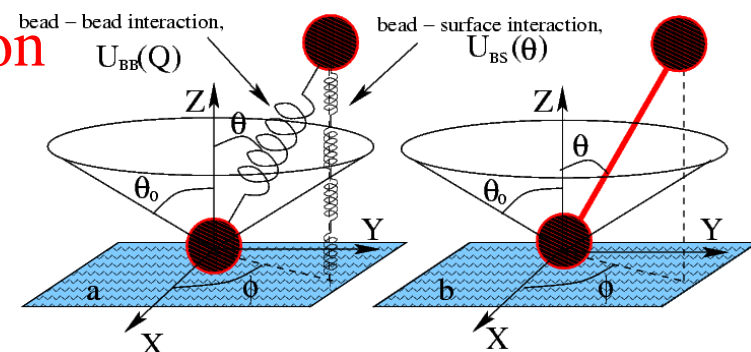
Integration over all moments of the beads

$$\psi(Q = \Delta_{12}, \theta, \varphi; t) = \int dP_1 dP_2 \rho_2(R_1, R_2, P_1, P_2; t)$$

Configuration-space distribution

$$\psi(Q, \theta, \varphi; t) = \sum_{n=1}^{\infty} \sum_{m=1}^{\infty} e^{-v_n^m (v_n^m + 1) D_R t} Y_{v_n^m}^m(\Omega(0)) Y_{v_n^m}^m(\Omega(t)) \Psi_{v_n^m}(Q; U_{BB}, U_{BS})$$

Radial distribution function



Bead-bead

Bead-surface

Rotational motion of the macromolecule

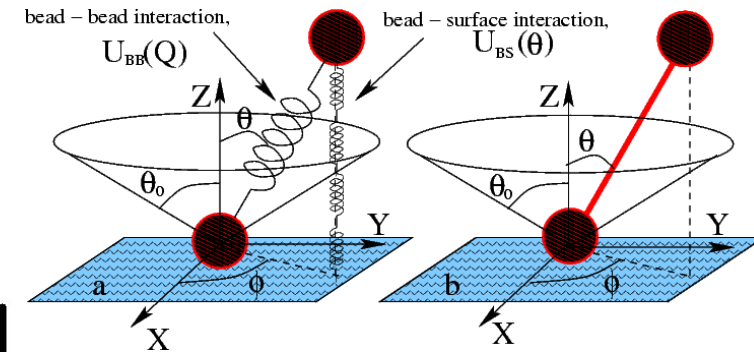
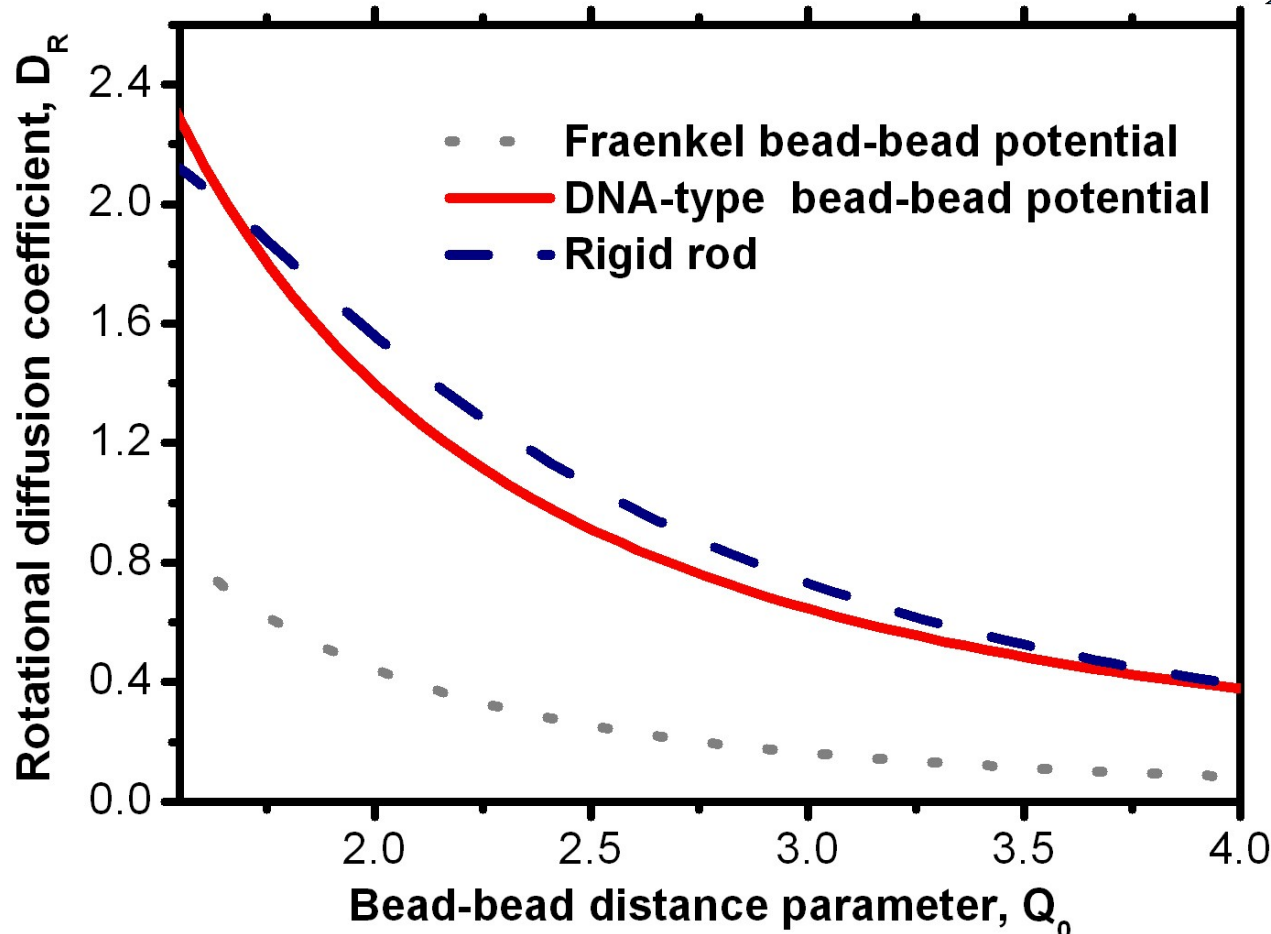
---- a few, short examples

1. Rotational diffusion coefficient

$$\mathbf{D}_R = \mathbf{D}_0 \left\langle \frac{1}{Q^2} \right\rangle$$

Dumbbell-type immobilized macromolecule:

Rotational diffusion coefficient D_R



bead-bead interaction:

● Freankel potential

$$U_{BB}^{Fr} = k_{Fr} (Q - Q_0)^2$$

● DNA-type potentila

$$U_B^{DNA} = k_{DNA} \left(\frac{1}{4 \left(1 - \frac{Q}{Q_0} \right)} - \frac{1}{4} Q + \frac{Q^2}{2Q_0^2} - \frac{1}{4} \right)$$

Rotational motion of the macromolecule

---- a few, short examples

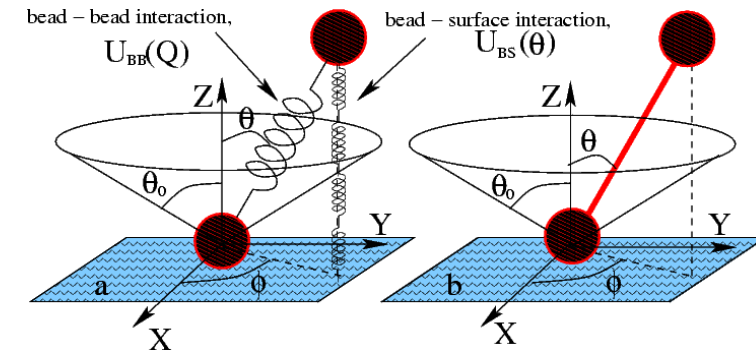
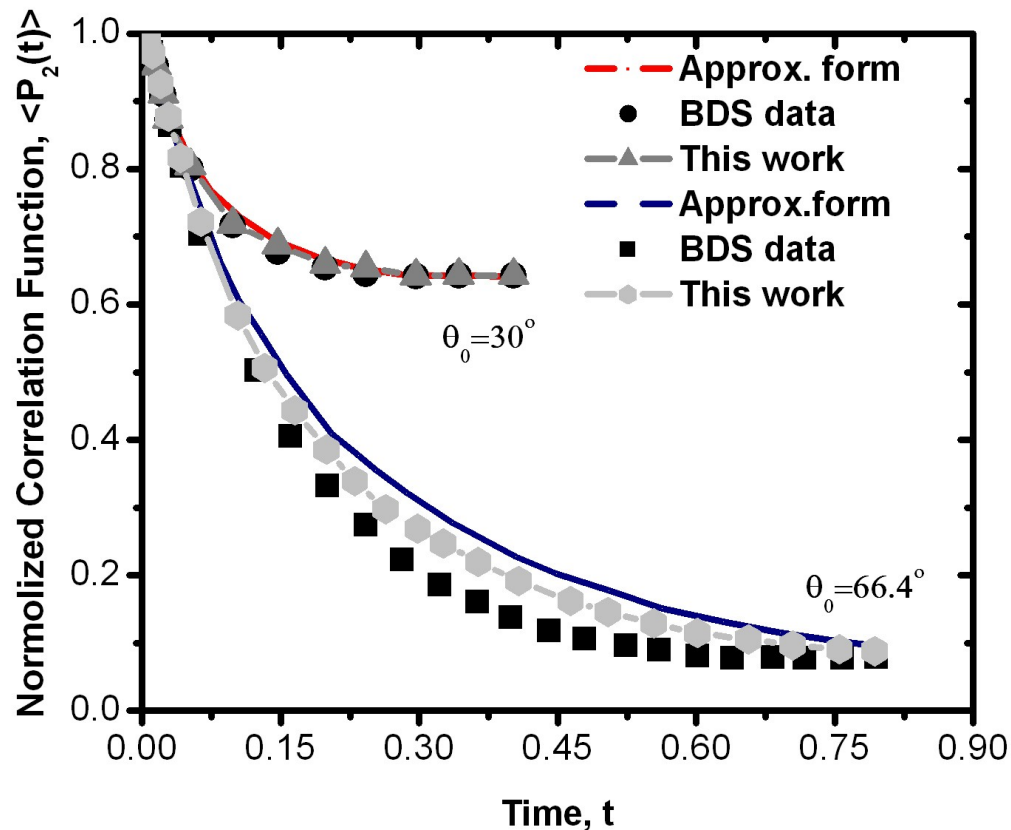
1. Rotational diffusion coefficient

2. Orientational correlation function

$$P_2(t) = \frac{1}{2} (3 \langle \cos^2 \theta(t) \rangle - 1)$$

Dumbbell-type immobilized macromolecule:

Time dependent orientational CF, $P_2(t)$



bead-bead interaction:

● Frenkel potential

$$U_{BB}^{Fr} = k_{Fr} (Q - Q_0)^2$$

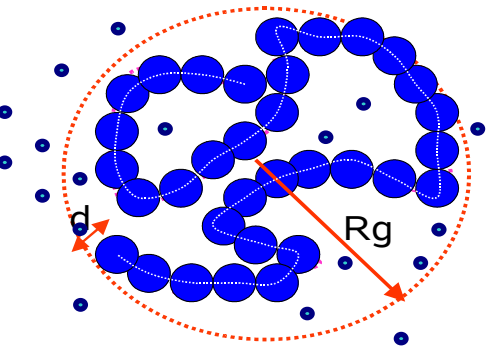
bead-surface interaction:

● Sin+cone potential

$$U_{BS} = \begin{cases} k_{sin} \sin^2 \theta, & \text{if } \theta \leq \theta_0 \\ 0, & \text{otherwise} \end{cases}$$

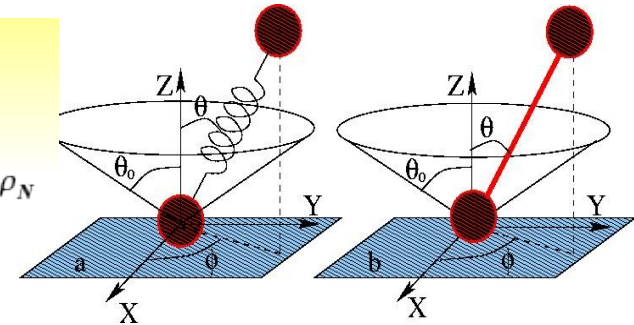
A semi-phenomenological approach to the structure and transport properties of macromolecules in solution

---- Summary



A semi-phenomenological approach

$$\frac{\partial \rho_N}{\partial t} + \frac{P_a}{M} \frac{\partial \rho_N}{\partial R_a} - \frac{\partial U(|R_a - R_b|)}{\partial P_a} \frac{\partial \rho_N}{\partial P_a} = \frac{\partial}{\partial P_a} \xi^{ab} \left[\frac{\partial}{\partial P_b} + \frac{P_b}{k_B T M} \right] \rho_N$$



Translational motion

- ✓ friction tensor parameters
- ✓ single-bead molecules
- ✓ dumbbell molecules
- ✓ N-bead macromolecules
- ✓ various thermo dynamical regimes

Rotational motion

- ✓ Restricted rotation of dumbbell molecules
- ✓ Bead-bead and bead-surface interactions

A. Uvarov, S. Fritzsche, *Macr. Theor. Sim.* **13** (2004), 241;
A. Uvarov, S. Fritzsche, *Chem. Phys. Letter* **401** (2005) 296;
A. Uvarov, S. Fritzsche, *Phys. Rev.* **E73**, (2006) 011111;
A. Uvarov, S. Fritzsche, *Phys. Rev. Letter* (2006), submitted.

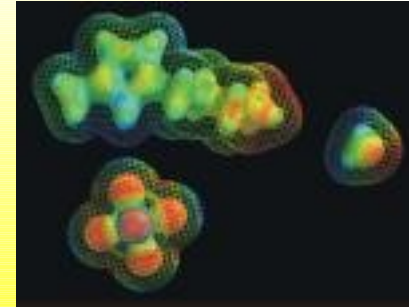
A. Uvarov, S. Fritzsche, *J. Chem. Phys.* **121**, (2004), 6566;
A. Uvarov, S. Fritzsche, *Progr. Colloid Polym Scien* **133**, (2006), **95**;
A. Uvarov, S. Fritzsche, *Chem. Phys. Letter* (2006), in print.

Structure and transport of macromolecules in solution

---- *Goals for future studies*

■ **Salvation dynamics & transport properties of the dense MS**

- *friction, diffusion and conductivity of the solution*
- *different thermodynamics states of the MSs.*
- *behaviour of time-dependent correlation functions.*



■ **Semi-phenomenological approach for the ILs**

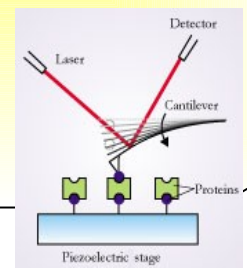
- *interaction among the molecular components of IL (cations and anions)*
- *interplay with various dissolved nanoparticles and clusters*

■ **Choice and parameterization of the interactions**

- *Coulomb*
- *Van der Waals*
- *Hydrogen-bonds*

■ **Orientational properties of free and immobilized macromolecules**

- *Extension of the semi-phenomenological approach on the rotational motion*
- *3- and 4- ... N- bead chains*
- *Complex molecular structures (different topologies)*



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C. Sarpe-Tudoran
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Institute Molecular and Atomic Physics

- **Dr. Alexander Blokhin**
- **Dr. Maxim Gelin**
- **Acad. Prof. Vitaliy Tolkachev**

Thank you for your attention!

Appendix

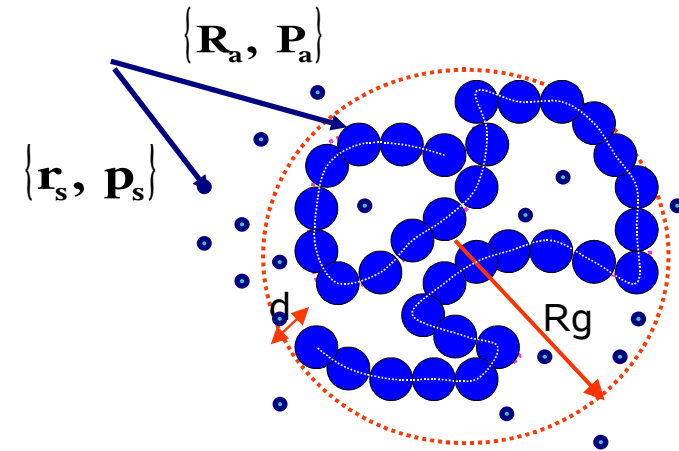
---- a few additional results

Fokker-Planck equation & Friction tensors

---- *phase-space dynamics*

$$\frac{\partial \rho_N}{\partial t} + \frac{P_a}{M} \frac{\partial \rho_N}{\partial R_a} - \frac{\partial U(|R_a - R_b|)}{\partial P_a} \frac{\partial \rho_N}{\partial P_a} = \frac{\partial}{\partial P_a} \xi^{ab} \left[\frac{\partial}{\partial P_b} + \frac{P_b}{k_B T M} \right] \rho_N$$

Thermodynamic limit: $n \rightarrow \infty; V \rightarrow \infty; n/V \rightarrow n_0$



Series expansion of time evolution operator: $\exp[-t \hat{Y} \hat{L}_{MS}] = \sum_{j=0}^{\infty} \frac{t^j}{j!} [\hat{Y} \hat{L}_{MS}]^j$

$$\xi_{\alpha\beta}^{(ab)} = \sum_j \xi_{\alpha\beta[j]}^{(ab)} = \xi_{\alpha\beta_0}^{(ab)} + \xi_{\alpha\beta_{[1]}}^{(ab)} + \xi_{\alpha\beta_{[2]}}^{(ab)} + \dots$$

$$q^{(ab)} = \frac{R_a - R_b}{|R_a - R_b|};$$

$$\xi_{\alpha\beta[j]}^{(ab)} = f_{[j]}(R_a, R_b; W; CF_{[j+2]}^{(solvent)}) = A_{[j]}^{(ab)}(R_a, R_b; W; CF_{[j+2]}^{(solvent)}) \delta_{\alpha\beta} + B_{[j]}^{(ab)}(R_a, R_b; W; CF_{[j+2]}^{(solvent)}) q_{\alpha}^{(ab)} q_{\beta}^{(ab)}$$

Translational motion of the macromolecule

---- a few, short examples

1. Friction tensor for the various „weak“ bead-solvent interaction

$$\xi_{\alpha\beta}^{(ab)} = \sum_j \xi_{\alpha\beta[j]}^{(ab)} \approx \xi_{\alpha\beta[0]}^{(ab)}$$

Charge particles: (Coulomb-like interaction)

Neutral particles: (Born-Mayer and van der Waals interaction)

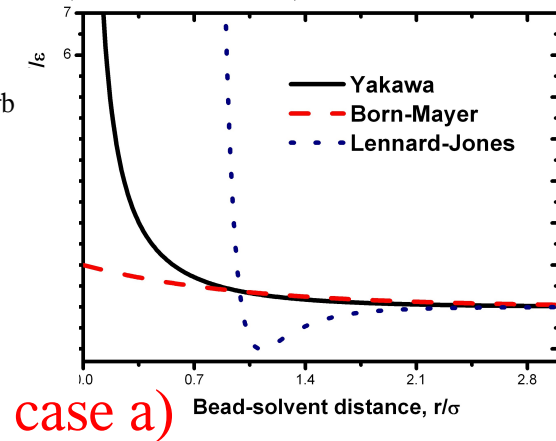
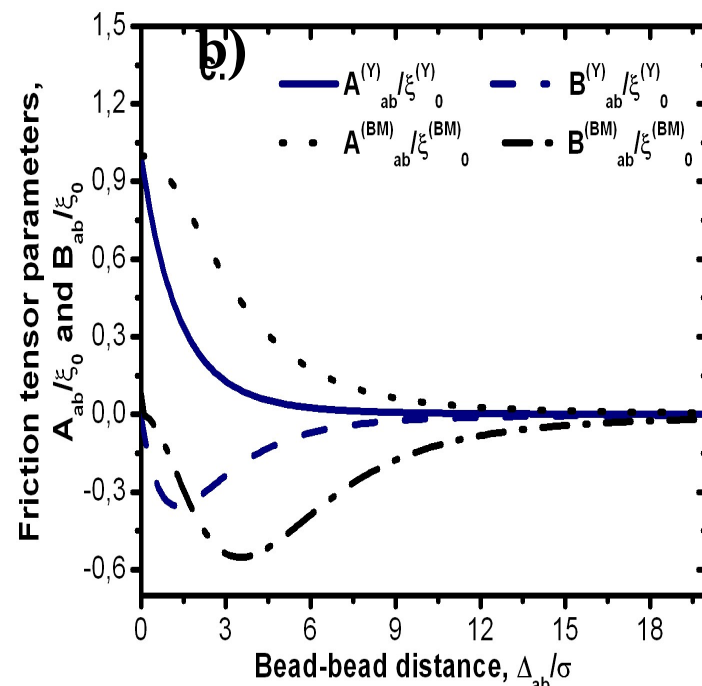
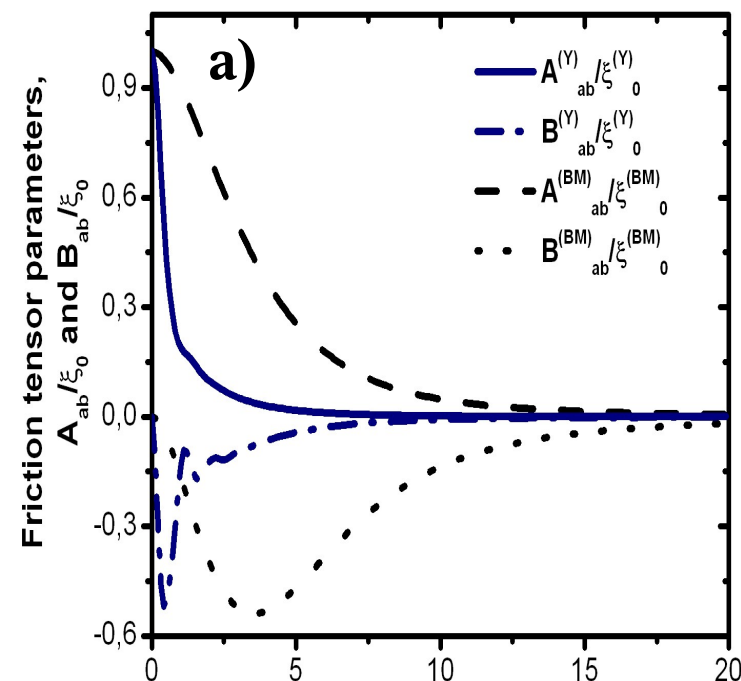
$$W^{(Y)} = \epsilon_{MS}^{(Y)} \frac{\exp[-r/\sigma]}{r/\sigma}$$

$$W^{(BM)} = \epsilon_{MS}^{(BM)} \exp[-r/\sigma]$$

$$W^{(LJ)} = 4\epsilon_{MS}^{(LJ)} \left(\left[\frac{\sigma}{r} \right]^{12} - \left[\frac{\sigma}{r} \right]^6 \right)$$

N-bead macromolecule:

Friction tensor parameters as function of the bead-bead distance $\Delta_{ab} = R_a - R_b$



case a) $k_B T = 1.2\epsilon_{SS}; \quad n_0 = 0.86/\sigma^3$

case b) $k_B T = 1.2\epsilon_{SS}; \quad n_0 = 0.30/\sigma^3$

Rotational motion of macromolecule

---- Configuration-space distribution

$$\Psi_{v_n^m}(Q; U_{BB}, U_{BS})$$

Restricted boundary conditions

$$0 \leq \theta \leq \theta_0 (\leq \pi/2)$$

$$0 \leq \varphi \leq 2\pi$$

$$0 \leq Q \leq Q_0$$

Bead-bead interaction:

- Hookean
- FENE
- Frankel
- DNA-type

$$U_{BB}^{Fr} = k_{Fr} (Q - Q_0)^2$$

$$U_{BB}^{DNA} = k_{DNA} \left(\frac{1}{4 \left(1 - \frac{Q}{Q_0} \right)} - \frac{1}{4} Q + \frac{Q^2}{2Q_0^2} - \frac{1}{4} \right)$$

Bead-surface interaction:

- Cone
- Effective double well (Sin)

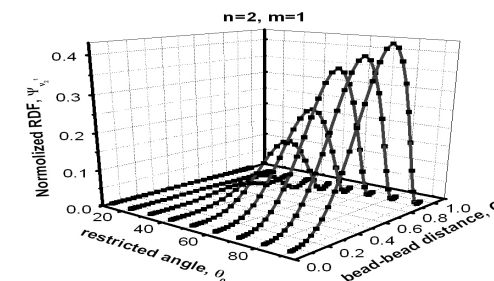
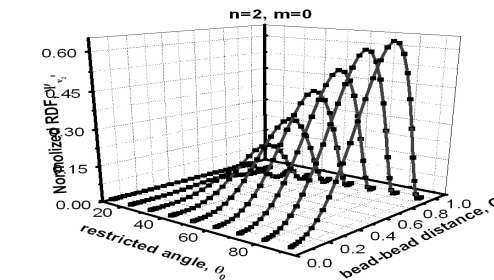
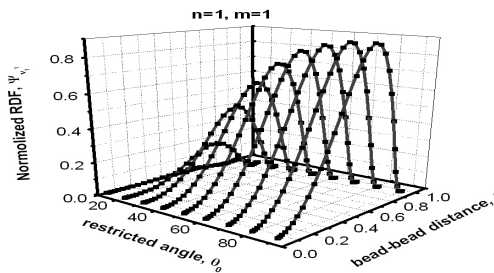
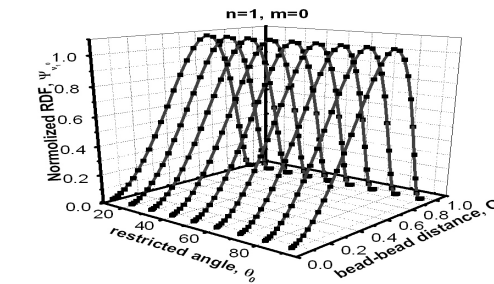
$$U_{BS}^{(Sin)} = k_{BS} \sin^2 \theta$$

Radial distribution

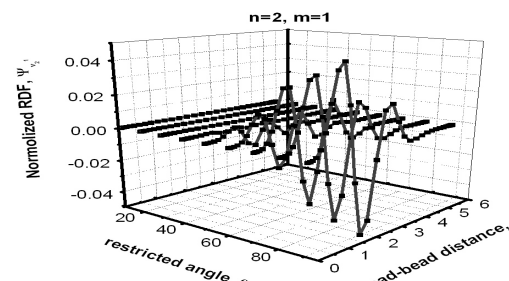
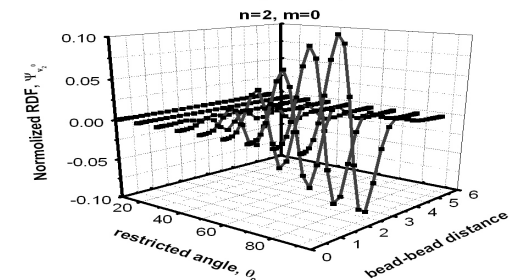
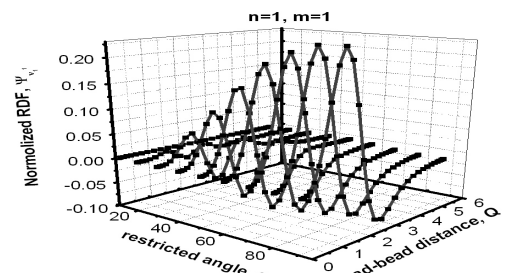
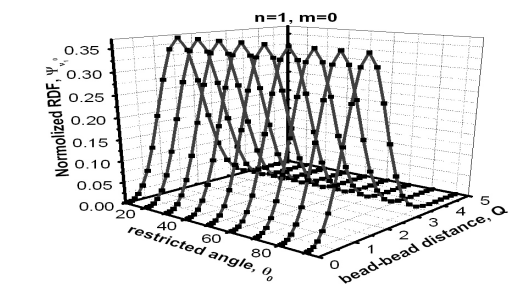
bead – bead interaction,
 $U_{BB}(Q)$

bead – surface interaction,
 $U_{BS}(\theta)$

DNA potential



Frankel potential



Rotational motion of the macromolecule

---- a few, short examples

1. Rotational diffusion coefficient

$$\mathbf{D}_R = D_0 \left\langle \frac{1}{Q^2} \right\rangle$$

Dumbbell-type immobilized macromolecule:

Rotational diffusion coefficient D_R

