

Statistical study of linear heteropolymers

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1996 Fiscal Year Final Research Report Summary

Statistical study of linear heteropolymers

Research Project

Project/Area Number

07640508

Research Category

Grant-in-Aid for Scientific Research (C)

Allocation Type

Single-year Grants

Section

一般

Research Field

物性一般(含基礎論)

Research Institution

Kanazawa University

Principal Investigator

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Project Period (FY)

1995 – 1996

Keywords

diblock polyampholyte / protein folding / Monte-Carlo simulation / pairing transition

Research Abstract

The understanding of the conformational transitions of biopolymers is a fundamental problem of statistical physics. Protein as well as nucleic acid folding are the main puzzling problems of biological importance. In order to tackle such problems, we undertook a systematic study of polyampholytes so as to understand the relation between the sequence of charges and the observed conformations. The term polyampholyte (designed as PA hereafter) is a generic name describing a polymer chain bearing positively and negatively charged monomers in a given sequence. A coil-globule transition was observed in an alternate polyampholyte. It was recognized that alternate sequences were highly specific and that random sequences should not exhibit any coil-globule transition. For globally neutral sequences, which is the relevant situation to proteins, our general aim is to understand what, in a given sequence, is relevant to conformational transitions. Note that nucleic acids are not PA but polyelectrolytes (any nucleotide bears the same negative charge), so that the electrostatic interactions do not depend on the sequence of bases but merely on the ionic force.

Diblocks and alternate sequences are the two extreme cases among random sequences. For an alternate PA, it is just 0 (except of course for the two monomers in the middle of the chain, but their contribution becomes negligible as the chain length grows). Although long diblocks are far from being the rule among biopolymers (and hereafter it will clearly appear why they don't exist in nature at ordinary temperature) they are nonetheless relevant to the physics of biopolymers for the following reason: any sequence of charges generally contains small diblocks spread throughout the chain. The longer the chain, the longer the diblocks. These small diblocks should undergo the earliest transitions (i. e. at the highest temperatures). It seems therefore important to understand the nature of the transitions occurring in diblocks as a function of their size. This is the main issue of the present work. [▲ Less](#)

Research Products (50 results)

All Other

All Publications (50 results)

- [Publications] K.Uehara: "First-principle molecular dynamics calculation of selenium clusters" J. Mol. Simulation. 18. 385-394 (1997) ▼
- [Publications] K.Uehara: "First-principles molecular dynamics simulations for Se₈ and Se₈⁺ clusters" J. Mol. Simulation, in press. ▼
- [Publications] S.Makino: "Classical Molecular Dynamics for the Formation Process of a Fullerene Molecule" J. Phys. Chem. Solids, in press. ▼
- [Publications] T. Hashimoto: "Structural Transformations of Ice at High Pressures via Molecular Dynamics Simulations" J. Mol. Simulation. 18. 115-132 (1996) ▼
- [Publications] T.Hashimoto: "Structural transformations of ice at normal and high pressures via molecular dynamicssimulations" Proc. PCI'96, in press. ▼
- [Publications] T.Hashimoto: "Structural transformation of ice at high pressures via molecular dynamics simulation II" J. Mol. Simulation. 18. 395-406 (1997) ▼
- [Publications] J.Habasaki: "Fracton Excitation and Levy Flight Dynamics in Alkali Silicate Glasses" Phys. Rev. B, in press. ▼
- [Publications] J. Habasaki: "Relaxation processes and the mixed alkali effect in alkali metasilicate glasses" Proc. MRS Fall Meetings, in press. ▼
- [Publications] J. Habasaki: "MD study of the mixed alkali effect in terms of the potential surface in the lithium-potassium metasilicate glass" J. Non-Cryst. Solids, in press. ▼
- [Publications] J. Habasaki: "Mixed Alkali Effect in Alkali Metasilicate Glasses" Proceedings of YKIS96, in press. ▼
- [Publications] 中崎潤子: "アルカリケイ酸塩ガラスの構造とダイナミクス-ガラス転移,緩和現象と混合アルカリ効果-" 高分子論文集. 53. 774-787 (1996) ▼
- [Publications] T. Muranaka: "Study on the β peak of χ " for a two-dimensional supercooled fluid state via molecular dynamics simulation" J. Mol. Simulation. 16. 387-397 (1996) ▼
- [Publications] T.Muranaka: "Coupling between Jump Motions and Correlated Motions" Proceedings of YKIS96, (in press). ▼
- [Publications] 橋本保: "氷に関する最近の分子動力学シミュレーション-氷の構造相転移-" 日本雪氷学会誌 雪氷. 58. 422-424 (1996) ▼
- [Publications] Y. Hiwatari: "Molecular-dynamics simulation for the formation process of a fullerene molecule" RIKEN Rev. 14. 7-8 (1996) ▼

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- [Publications] Y.Hiwatari: "Molecular-Dynamics Study of Supercooled Liquids and the Theory of the Glass Transition" J. Phys. Soc. Jpn., Comp. Physics as a New Frontier in Condensed Matter Res.329-339 (1995) ▼
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- [Publications] J. Habasaki: "Origins of the two-step relaxation and the boson peak in an alkali silicate glass studied by molecular-dynamics simulation" Phys Rev.E52. 2681-2687 (1995) ▼
- [Publications] J. B. Imbert: "Conformational Transitions of a Diblock Polyampholyte in 2 and 3 Dimensions" submitted to Europhysucs Letters. ▼
- [Publications] N. Urakami: "Multicanonical Monte Carlo Simulation of a Polymer with Stickers" J. Phys. Sci. of Japan. 65. 2694-2699 (1996) ▼
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- [Publications] N.Urakami: "Monte-Carlo simulation of a polymer with stickers · · · Initial relaxation and free energy calculations · · ·" Proceedings of YKIS96, in press. ▼
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