CECAM and IUPAP workshop on High density DNA arrays: models, theories and multiscale simulations



Book of abstracts

Ljubljana, Slovenia July 24 2019 – July 26 2019

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Contents

General Information	7
Description	13
Schedule	15
Abstracts – invited lectures	19
Modeling high density DNA arrays: Mission accomplished! (Aleksei Aksimentiev)	20
Lyotropic cholesteric phases of DNA and its assemblies ($Jonathan$ $Doye$)	$\frac{n}{21}$
Super resolution microscopy of DNA packaging in mammalian cell nuclei (<i>Christoph Cremer</i>)	22
Open boundary molecular dynamics of star-polymer melt and DNA in salt solution $(Jurij Sablić)$	24
Multiscale simulations of DNA aggregation by multivalent ions (Alexander Lyubartsev, Tiedong Sun, Alexander Mir- zoev, Vishal Minhas, Nikolay Korolev and Lars Norden- skiöld)	26
Nucleosome conformational variability in interphase nuclei and in solution explored by cryo-electron microscopy and tomography of vitreous sections (<i>Mikhail Eltsov</i> ,	20
$Diana\ Grew,\ Françoise\ Livolant,\ Amélie\ Leforestier)$.	27
Polymer globules in bulk and in confinement (<i>Stanard Mbebwe</i>	20
Pachong, Jan Smrek, Kurt Kremer)	29
Toward controlled self-assembly of peptides (<i>Achille Giacometti</i>)	30
DNA without DNA: Watson-Crick selectivity controls the self-assembly of mononucleotides (<i>Tommaso Fraccia, Marce</i>)
Todisco, Giuliano Zanchetta, Greg Smith, Noel Clark,	-
Tommaso Bellini)	31

Spontaneous domain formation in spherically-confined elastic	2
filaments (Tine Curk, James D. Farrell, Jure Dobnikar,	,
Rudi Podgornik)	. 32
How to model DNA translocation through nanopores – A	
multiscale simulational exploration (<i>Christian Holm</i>)	. 33
Effect of knots during viral DNA packaging and ejection (<i>Cris</i>	-
$tian Micheletti) \ldots \ldots \ldots \ldots \ldots \ldots \ldots \ldots$. 34
Density-nematic coupling in isotropic linear polymers: acous-	
tic and osmotic birefringence (Aleksandar Popadić, Dana	iel
Svenšek, Rudolf Podgornik, and Matej $Praprotnik$)	. 35
Homology recognition without proteins [DNA sequence-struct	Jure
relationship, helical coherence, and a universal homol-	-
ogy recognition mechanism-theory vs experiments] $(A.A)$	•
Kornyshev, $D.J.(O')$ Lee, S.Leikin, A.A. Wynveen).	. 36
From chromosome territories to ring polymers: Physical prop-	-
erties of untangled polymer melts (Angelo Rosa)	. 38
"Genetics" in two dimensions: DNA base pairing in thin sur-	-
face films (Matjaž Ličen, Lucija Čoga, Stefano Maseiro,	,
Lea Spindler, Irena Drevenšek-Olenik)	. 39
Atomistic MD simulations as a tool to elucidate mechanical	l
properties of DNA/RNA systems (<i>Franci Merzel</i>)	. 42
DNA-based dendrimers: From a single molecule to the dense	Ś
solution description (N. Adžić, C. Jochum, E. Stiakakis,	
G. Kahl, C. N. Likos)	. 43
Free energy landscapes for dendrimer-like DNAs via neural	I
networks (Florian Buchner, Clemens Jochum, Gerhard	l.
Kahl, Andreas Singraber, Christoph Dellago)	. 45
	. 10
Abstracts – contributed lectures	47
Analysis structures and forces defining chromatin condensa-	-
tion (Nikolay Korolev, Wahyu Surya, Sook Yi Wong,	,
Qinming Chen Alexander P. Lyubartsev, John van Noort	t,
Lars Nordenskiöld)	. 48
Genetic outbreak from the endosome:	:
A molecular perspective on gene therapy (Bart Bruinink	ks) 49
Multiscale simulations of double-stranded DNA with sequence	- -
dependent mechanical and conformational properties ($Second Second Seco$	al-
vatore Assenza)	. 50
Dynamics and organization of cyclic polymers melt in a con-	-
finement (S.M. Pachong, J. Smrek, K. Kremer)	. 51

DNA-based dendrimers: From a single molecule to the dense solution description

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We present a joint theoretical-experimental study of a novel class of macromolecules, the so-called DNA-based dendrimers. They have recently been synthesized from the enzymatic ligation of Y-shaped DNA unit, a three-armed structure consisting of double-stranded DNA (ds-DNA), formed via hybridization of three single-stranded DNA chains (ss-DNA), each of which has partially complementary sequences to the other two [1]. In order to describe such dendrimers of various generations we have employed two independent models: a bead-spring model and the oxDNA model. In the bead-spring model, base-pairs of a single DL-DNA molecule are modeled by charged monomers, whose interactions are chosen to mimic the equilibrium properties of DNA correctly. On the other hand, the oxDNA model allows us to take a closer look into the DNA structure, treating DNA as a string of rigid nucleotides which interact through potentials that depend on the position and orientation of the nucleotides. We have performed Molecular Dynamics Simulations and we have also employed dynamic/static light scattering in order to determine equillibrium properties and conformational characteristics of all-DNA dendrimers as well as the behavior of their solutions. We have investigated their behavior in ionic solution, paying particular attention on their salt-responsiveness. Our computational and experimental results reveal that the DL-DNA are rigid objects with low internal monomer concentration, regular voids in their interior, with high persentage of absorbed counterions, and that show high resistance to stimuli-responsiveness [2]. These properties shape the behaviour of their solutions. Namely, both experimental as well as computational results show anomalous structure factor of dense DL-DNA solutions, as it had been predicted theoretically in Ref [3]. In this way we have found the object which was a missing puzzle in understanding the full phase diagram of star polymer solutions.

^[1] Y. Li, Y. Tseng, and D. Luo, Nat. Mater. 3, 38, (2004).

^[2] C. Jochum, N. Adžić, E. Stiakakis, T. L. Derrien, D. Luo, G. Kahl, and C. N. Likos, Nanoscale, 11, 1604 (2019).