Annaëlle BRUNET 1, chemin Reboul

PhD in Biophysics

31100 TOULOUSE (FRANCE) PhD student from 01/09/2012in the LPT & IPBS labs, CNRS (Toulouse-France), defense expected at the end of October

Date of Birth: 13/01/1987

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Keywords: Single molecule approaches, Kinetic Monte-Carlo simulation, statistical physics, DNA dynamics, soft matter



Nationality: French Driver's license, B

SKILLS					
	Modeling	Microscopy			
Approach:	Dynamic Kinetic Monte-Carlo simulation	Techniques:	STM, MEB, fluorescence and dark field		
Data analyses procedure			microscopy		
Developing an computing procedure to check the physical		Analysis:	ImageJ		
coherence (symmetry factor, correlation function) of gigaoctet			Characterization technique		
of raws data and to applied first step of analyses		DLS, Zetasizer, UV spectroscopy, quantification PCR			
Informatics		Surface treatment process			
Tools:	Mathematica, Matlab, Labview, Scribus (PAO),	Chemical treatment: epoxydation, thiolisation, piranha			
	Pymol, gnuplot, office tools	Physical treatment : Plasma cleaner, UV ozone			
Languages:	Fortran, C, R, LaTeX, Beamer, bash	Method: spin coating, deposit convective self-assembly			
Office tools:	GIMP, Inkscape, Microsoft office and		Languages		
	OpenOffice (type Word, Exel, Powerpoint)	English:	TOEIC Score 775		
Systems:	Linux (Ubuntu), Windows	German:	Conversational basics		

TRAINING AND DIPLOMA				
2012-Now	PhD in Physic (defense planned the October 16th 2015)	University of Toulouse III (France)		
	Doctoral School : Matter of Science,	Laboratories LPT et IPBS, CNRS,		
	Specialty: Physique	Toulouse (France)		
	Single molecule study of DNA molecules conformations with local defects or under a large set of			
	physicochemical conditions - Advisor : Destainville N. (LPT) and Tardin C. (IPBS)			
	 Measure the impact of intrinsic bending, local denaturation or variation of ion concentration in 			
	solution on the DNA conformations with high-throughput Tethered Particle Motion (HT-TPM)			
	 Perform a kinetic Monte-Carlo Simulation based on a mesoscopic statistical model of DNA 			
	 Develop a computational procedure for the analysis of the 	large data set from HT-TPM		
2011-2012	Master 2R, Specialty: Nanosciences, Nanomesures	University of Toulouse III (France)		
	Master's training period in nanotechnologies (6 months)	LAAS-CNRS, Toulouse (France)		
	Formation of nano-energetic material made of Al/CuO alloy driven by DNA auto-assembly and chip			
	integration – Advisor : Bancaud A. and Rossi C.			
	 Construct heterogeneous advances material structured on 1D, 2D or 3D-dimensions by using the 			
	complementarity of the double strand DNA, and optimize its stability and its energetic response			
2010-2011	Master 1, Specialty: Fundamental Physic	University of Toulouse III (France)		
	Master's training period in microscopy (2 months)	CEMES-CNRS, Toulouse (France)		
	Studies of the 2-(3-perylene) ethanoic acid molecular by Scanning Tunneling Microscopy (STM) at			
	Low Temperature and Ultra High Vacuum – Advisor : Coratger R.			
	 Measure the value of the single negative charge appearing during the process 			

2009-2010 Licence 3, Specialty: Physics et Applications **University of Toulouse III (France)** Training period at the Braley company **Braley Company, Bozouls (France)**

> Realization of a solar furnace coupled with a Stirling engine Realize the solar furnace and animate a stand on the energies at the open day of the company

SCIENTIFIC PUBLICATIONS

2015 Brunet, A., Tardin, C., Salomé, L., Rousseau, P., Destainville, N., Manghi, M., Dependence of DNA persistence length on ionic strength of solutions with monovalent and divalent salts: a joint theoryexperiment study, Macromolecule, DOI: 10.1021/acs.macromol.5b00735 2015 Brunet, A., Chevalier, S., Destainville, N., Manghi, M., Rousseau, P., Salhi, M., Salomé, L., Tardin, C.,

Probing a label-free local bend in DNA by single molecule tethered particle motion, Nucleic acids research, DOI: 10.1093/nar/gkv201

In preparation Brunet, A., S., Destainville, N., Manghi, M., Rousseau, P., Salomé, L., Tardin, C., Dependence of DNA

apparent length and persistence length on temperature of solutions, a joint theory-experiment study

CONGRESSES AND THEMATICS SCHOOLS

Oral Communication

In coming, 8-9 Aout 2015 Gordon Research Seminars (GRS): Soft Condensed Matter Physics - New London (USA)

November 2014 3rd meeting of FRBT – **Toulouse (France)**, January 2014 Seminar IRSAMC – **Toulouse (France)**,

May 2013 3rd edition of the "Les Houches School" in computational physics: DNA, from molecules to

evolution - Les Houches (France)

Thematic School

August 2014 Summer school: SOFT-FIRE-2014 – Cargèse (France), (2 weeks)

May 2013 3rd edition of the "Les Houches School" in computational physics: DNA, from molecules to

evolution – Les Houches (France), (2 weeks)

Poster Communication

In coming, 9-14 Aout 2015 Gordon Research Conference (GRC): Soft Condensed Matter Physics - New London (USA)

August 2014 Summer school : SOFT-FIRE-2014 – Cargèse (France)

May 2013 GDR Cell Tiss 2013 - Lyon (France)

May 2013 3rd edition of the "Les Houches School" in computational physics: DNA, from molecules to

evolution - Les Houches (France)

STUDENT SUPERVISION

Master's training period : Juliette Wilhem

(2 months)

Summer 2014 Probing the experimental effect of the ionic strength on the DNA conformation release by TPM, at the single molecule level

REFERENCES

Pr. Destainville Nicolas

Laboratoire de Physique Théorique,

Team : Physique Statistique des Systèmes Complexes

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Dr. Bancaud Aurélien

Laboratoire d'analyse et d'architecture des systèmes, Team : Nano Ingénierie et Intégration des Systèmes

Toulouse, France

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Dr. Tardin Catherine

Institut de Pharmacologie et de Biologie Structurale,

Team: Membrane and DNA dynamics

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Dr. Salomé Laurence

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Team: Membrane and DNA dynamics

Toulouse, France

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OTHER ACTIVITIES AND INTERESTS

Other professional experience:

2012 Technique staff at the Symposium J de l'E-MRS 2012 Spring Meeting - Strasbourg (France)

Associative activities:

Since 2009 Co-founder and editor of the inter-university newspaper: Le Lapin Blanc – University of Toulouse I, II and

III (France)

2007-2008 University of Toulouse III site organizer for the AMIDONS association – Association Inter-university for

donation of blood – University of Toulouse III (France)

Since 2006 AMIDONS association volunteer - Toulouse (France)

2003 First aid qualification - Attestation de Formation aux Premiers Secours - Œuvre hospitalière française de

l'ordre de Malte (France)

Entertainment: Reading (Polar, Heroic fantasy, Anticipation), climbing, volleyball, badminton, cinema, theater, rugby

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Toulouse, 26th may 2015

<u>Subject</u>: Applications of a research engineer position in computational imaging and super-resolution microscopy

Dear Dr. Christophe Zimmer,

I am delighted to apply for a research engineer position in computational imaging and superresolution microscopy

I am very interested by your project on the genome architecture and the dynamic of chromosome folding. If DNA is well-known for containing the genetic code essential to the cell physiology, its mechanical characteristics are now also under close study as they may actively participate to the different biological processes. Studying the spatial architecture and dynamics of the genomic material compose an important field of research to fully understand of genome conformation and function. To probe the role played by the statistical and mechanical properties of the chromosome in the nucleus, acquisition of experimental data with high-accuracy and resolution at high-throughput is needed in one hand. In the other hand, computational models become a crucial and decisive complement in order to make sense of the experimental data. I truly believe that combining computational model and experimental data will allow a better understanding of how chromosomes fold, move and interact.

Indeed during my PhD project (defense planned at the end of October 2015), I have studied at the single molecule level the conformational changes of DNA molecules caused by the presence of local defects on the molecule or changes in global physicochemical conditions. My work aims at both observing and modeling the physical mechanisms which regulate the rigidity of doublestranded DNA to improve the understanding and the theoretical description of soft matter. For that, I have performed in one hand experimental measurements at the single molecule level by using the techniques called Tethered Particle Motion (TPM) (IPBS, Toulouse) and on the other hand a Kinetic Monte Carlo simulations using an ad'hoc mesoscopic statistical model (LPT, Toulouse). I'm particularly interested by the biophysic field and sensitive to the complementarity of the combining of experimental, theoretical and computational approaches. During this interdisciplinary project, I have experienced the work with molecular biologists (LMGM, Toulouse), biophysicists and physicists teams and learned how to deal with such multidisciplinary approaches. Accustomed to use imaging methods and optical microscopy, I am aware and mindful of the experimental constrains and as well as the experimental limitations, and particular with single-molecule requirements. Furthermore, I stay turned to revise and develop possible improvements.

Moreover, the IPBS group, where I carried out my TPM measurements, has not only a recognized expertise in TPM techniques but they also developed a biochip that enables the high parallelization of TPM measurements resulting in the analysis of several hundred molecules simultaneously. On part of my job has consisted in coding and developing a robust and reliable procedure to analyses this large acquisition data sets. For this, I have established a computing procedure in Mathematica language to check the physical of gigaoctet of raws data and perform first step of analyses. Thanks to the ensuing high-throughput data acquisition and the analyses procedure, I have accumulated a large number of individual statistics that enabled me to reveal and characterize a local bend present in DNA and discriminate between different models of polymer physics that describe the salt effect on the DNA persistence length.

Participating in a project dealing with the biology, biophysics, physics and the computational biology of DNA, would perfectly fit with my will to continue in the condensed matter field at the frontier with biology. Besides, I am extremely eager to explore the conformations of the DNA in its native state such as it is organized in chromatin. As a result, I am strongly motivated to participating in a project which prospected the spatial architecture and dynamics of the genomic material in the nucleus and the implication in various aspects of gene expression, DNA repair, recombination, and replication. I believe that I have the computational and experimental skills, that match your requirements of your project and that this project offer me the opportunity to develop new ones.

czimmer@pasteur.fr

Dear Dr. Christophe Zimmer,

I am writing to you to apply for the position of a research engineer in computational imaging and super-resolution microscopy.

My current PhD (defense planned in the end of October 2015) let me to use both experimental and numerical approaches at the single molecule level. I acquired computational skills as I developed a program (Mathematica) for the fast and efficient analysis of large data sets and preformed simulations (.c) of DNA conformations based on statistical models. In addition, during this project I have successfully collaborate with multidisciplinary teams such as molecular biologists, biophysicists and physicists.

I am very interested by your project. I believe that my qualifications would match your requirements, and would offer me he opportunity to develop new ones.

Please find the details of my CV and cover letter as an attachment,

I am available for any further information that you would need. Yours sincerely

Annaël Brunet