## ABHISHEK DEY, Ph. D 112 NC 54, Apt J6

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#### **Education**

Jawaharlal Nehru University/ CSIR-CDRI New Delhi/Lucknow, India

Ph. D, Biochemistry and Structural Biology December 2014

Kurukshetra University Kurukshetra, Haryana, India

Master of Science, Biochemistry

July 2006

Kurukshetra University Kurukshetra, Haryana, India

Bachelor of Science, Biochemistry, Zoology, Chemistry April 2004

**Professional Experience** 

# University of North Carolina-Chapel Hill Postdoctoral Research Associate

PI- Dr. Alain Laederach

Chapel Hill, NC, USA May 2020- present

- Identified conformational landscape of RNA Frame Shifting Element (FSE) responsible for regulating protein translation in SARS-COV 2 virus.
- Investigating the effect of RNA secondary structure at alternatively and constitutively spliced exon-intron junctions of MAPT precursor mRNA in Alzheimer's Disease.
- Mentored a potential graduate student in RNA Structural biology.

# **University of North Carolina-Charlotte Postdoctoral Research Associate**

PI- Dr. Kausik Chakrabarti

Charlotte, NC, USA August 2017-April 2020

- Analyzed and characterized folding dynamics and functional aspects of *T. brucei* telomerase RNA *in vivo*.
- Developed functional characterization assay for Telomerase ribonucleoprotein.
- Examined secondary structure of various long non-coding RNA in parasites of eukaryotic origin through RNA-SHAPE assay.
- Developed and characterized water-soluble small molecules as a novel RNA-SHAPE reagent.
- Responsible for recruitment and mentoring junior scientists and graduate students.

## Carnegie Mellon University Postdoctoral Research Associate

Pittsburgh, PA, USA March 2017-August 2017

PI- Dr. Kausik Chakrabarti

• Determined different secondary structure domains in *in vitro* transcribed *T. brucei* telomerase RNA.

#### **University of Connecticut health**

Farmington, CT, USA

#### **Postdoctoral Research Associate**

October 2015-March 2017

PI- Dr. Justin Radolf

- Explored and characterized biophysical and immunologic properties of rare outer membrane proteins in spirochetes.
- Analyzed three-dimensional model of Oligopeptide Permease Protein A (OPPA) from *Borrelia Burgdorferi* which is essential for the peptide uptake and virulence of pathogen.

## PREMAS Biotech Analytical Scientist- QC

Gurugram, Haryana, India April 2014 – July 2015

- Established analytics and assay development division within the organization and trained junior personnel for same.
- Developed assays characterizing Trypsin/Chymotrypsin inhibitors for the development of oral anti-diabetic drug.
- Analyzed and examined final products before delivery.
- Quality check for the final project deliverables.
- Managed and edited Standard Operating Procedures (SOPs), Manuals, Master formats and reports while working along with Quality Assurance (QA) personnel.

## CSIR-Central Drug Research Institute Doctoral Student

Lucknow, UP, India May 2009-April 2014

**Thesis Title:** Structural studies on transcriptional regulatory protein(s) from *Mycobacteria*.

PI- Dr. Ravishankar Ramachandran

- Characterized the overall three-dimensional structure and function for Rv2779c from *M. tuberculosis*.
- Proposed a mechanistic model of transcriptional regulation by Rv2779c in M. tuberculosis.
- Identified small molecule inhibitors against Feast famine regulatory proteins (FFRPs) which can act as novel drug targets in *M. tuberculosis*.

## CSIR-Central Drug Research Institute Project Assistant-II

Lucknow, UP, India June 2007- May 2009

PI- Dr. Ravishankar Ramachandran

- Cloning, Overexpression and Purification of Feast famine regulatory proteins (FFRPs) *form M. tuberculosis*.
- Biochemical and Biophysical characterization of Rv2779c protein from *M. tuberculosis*.

# **CSIR-National Botanical Research Institute Project Trainee**

Lucknow, UP, India July 2005- August 2005

PI- Dr. Sunita Kochhar

• Developed functional assays of superoxide dismutase enzyme present in mung beans.

### **Awards and Fellowships**

Postdoctoral Fellowship (NIH), University of North Carolina-Chapel Hill	2020-present
Postdoctoral Fellowship (NSF), University of North Carolina-Charlotte	2017-2020
Postdoctoral Fellowship (NSF), Carnegie Mellon University	2017

Postdoctoral Fellowship (NIH), University of Connecticut Health ICMR-Senior Research Fellowship, CSIR-CDRI ICMR- Junior Research Fellowship, CSIR-CDRI David Blow Scholarship, CCP4 study weekend, UK  Teaching Experience	2015- 2017 2011- 2014 2009- 2011 2010
BIOL 4000/5000 (Gene Silencing) Assistant Instructor, UNC-Charlotte Topics: Chromatin structure, Transcription and DNA methylation, Techniques	2020
BIOL 4000/5000 (Gene Silencing) Assistant Instructor, UNC-Charlotte Topics: Structural and Functional importance of Ribozymes	2019
BIOL 6000/8000 (Introduction to RNA Molecular Biology) Assistant Instructor, UNC-Charlotte Topics: RNA structure, Telomerase RNP complex	2018
Proteins: Structure and Function Assistant Instructor, CSIR-CDRI Topics: Hierarchy in Protein folding and stabilizing forces	2012-2014

#### **Peer-Reviewed Publications**

Sequence and tissue targeting specificity of ZFP36L2 reveals *Elavl2* as a novel target with coregulation potential. Redmond IC, Ardizzone M, Hekimoglu H, Hatfield BM, Waldern JM, **Dey** A, Montgomery SA, Laederach A, Ramos SBV. (2022) *Nucleic Acid Research*, **50** (7), 4068-4082.

In vivo Architecture of the Telomerase RNA Catalytic Core in *Trypanosoma brucei*. **Dey A**, Eklund AM, Klotz K, Saha, A., Davis J, Li, B. Laederach A, Chakrabarti K. (2021) *Nucleic Acid Research*, **49** (21), 12445-12466.

To knot or not to knot: Multiple conformations of the SARS-CoV-2 frameshifting RNA element. Schlick T, Zhu Q, **Dey A**, Jain S, Yan S, and Laederach A. (2021) *Journal of American Chemical Society*, **143** (30), 11404-11422.

The RNA Structurome in the Asexual Blood Stages of Malaria Pathogen *Plasmodium falciparum*. Alvarez DR, Ospina A, Barwell T, Zheng B, **Dey A**, Li C, Basu S, Shi X, Kadri S, Chakrabarti K. (2021) *RNA Biology*, **18**(12), 2480-2497.

Innately water soluble isatoic anhydrides with modulated reactivities for RNA SHAPE analysis Fessler A\*, **Dey A\***, Finis DS, Flower AJ, Chakrabarti K, Ogle C (2020). *Bioconjugate Chemistry*, **31**, **(3)**, 884-888. \* **Equal Contribution** 

The *Non-Coding RNA* Journal Club: Highlights on Recent Papers—7. **Dey A,** Chakrabarti K, et. al. (2019), *Non-coding RNA*, **2019**, **5**, 40.

Water-soluble isatoic anhydrides: a platform for rna-shape analysis and protein bioconjugation. Fessler A, **Dey A**, Garmon CB, Finis DS, Saleh N, Fowler AJ, Jones DS, Chakrabarti K and Ogle C (2018), *Bioconjugate Chemistry*, **29**, 3196-3202.

Sequence variation of rare outer membrane protein  $\beta$ -barrel domains in clinical strains provides insights into the evolution of *Treponema pallidum* subsp. pallidum, the Syphilis Spirochete. Kumar S, Caimano M.J, Anand A, **Dey A**, Hawley K.L, et.al (2018), *mBio*, **9**, (3), *e01006-18*.

Current perspectives of telomerase structure and function in eukaryotes with emerging views on telomerase in human parasites. **Dey A,** Chakrabarti K (2018), *International Journal of Molecular Sciences*, **19**, 333.

Peptide uptake is essential for *Borrelia burgdorferi* viability and involves structural and regulatory complexity of its oligopeptide transporter. Groshong A.M, **Dey A**, Bezsonova I, Caimano M, Radolf J.D. (2017), *mBio*, **8**, **(6)**, *e02047-17*.

The major outer sheath protein forms distinct conformers and multimeric complexes in the outer membrane and periplasm of *Treponema denticola*. Puthenveetil R, Kumar S, Caimano M.J\*, **Dey A**\*, Anand A\*, Vinogradova O, Radolf, J.D. (2017), *Nature Scientific Reports*, **7**, (**13260**), *13550-13556*. \***Equal contribution** 

Crystal Structure of *Mycobacterium tuberculosis H37Rv* AldR (rv2779c), a regulator of the ald gene: DNA-binding and identification of small-molecule inhibitors. **Dey A,** Shree, S, Pandey, S.K., Tripathi, R. P., Ramachandran, R. (2016), *Journal of Biological Chemistry*, **291**, **(23)**, *11967-11980*.

Cloning, Overexpression, Purification, and preliminary X-ray analysis of a feast/famine regulatory protein (Rv2779c) from *Mycobacterium* tuberculosis H37Rv. **Dey A,** Ramachandran, R. (2014) *Acta Crystallographica Sect F, F70*, 97–100. (Crystal image on cover of ActaF 2015)

Ligand-induced structural transitions, mutational analysis and 'open' quaternary structure of the *M. tuberculosis* Feast/Famine Regulatory protein (Rv3291c). Shrivastava, T, **Dey A**, Ramachandran, R (2009) *Journal of Molecular Biology*, *392*, *1007–1019*.

#### **Future Publications**

Quantitative integration of RNA structure and splicing elements to explain alternative splicing of Microtubule-Associated Protein Tau gene. Kumar J, Lackey L, Waldern JM, **Dey A**, Mathews DH, Laederach A. *eLife* (*In Review*)

#### Conference/Meetings and Invited talks

Klotz K, **Dey A**, Eklund AM, Saha, A, Davis J, Li B, Laederach A, Chakrabarti K, High-resolution profiling of Telomerase RNA structure dynamics in the eukaryotic pathogen *Trypanosoma brucei*, **Rustbelt RNA meeting-2021**, Virtual meeting, (abstract selected for podium presentation)

**Dey A**, Eklund AM, Saha, A, Li B, Laederach A, Chakrabarti K, Novel features of telomerase RNA folding and interactions in *Trypanosoma brucei* telomerase ribonucleoprotein complex, **RNA Society-2020**, Virtual meeting.

**Dey A**, Saha, A, Eklund AM, Laederach A, Li B, Chakrabarti K, Structural and Mechanistic insight into *T. brucei* Telomerase RNA, **Kinetoplastid Molecular and Cell Biology** meeting-2019, MA, USA.

Chakrabarti K, **Dey A**, Novel aspects of Telomerase RNA regulation in Kinetoplastid pathogen, *Trypanosoma brucei*, **Telomeres and Telomerase-2019**, Cold Spring Harbor Laboratory, NY, USA.

**Dey A**, Fessler A, Ogle C, Chakrabarti K, Structural imprints of in vivo RNA folding in human parasites, **Rustbelt RNA meeting-2018**, OH, USA.

**Dey A,** Ravishankar R, Crystal Structure of the Feast-famine regulatory protein (Rv2779c) form *M. tuberculosis*, **42nd National Seminar on Crystallography- 2013**, New Delhi, India (abstract selected for podium presentation)

**Dey A,** Ravishankar R, Structural and Functional studies of transcriptional regulatory protein (Rv2779c) from form *M. tuberculosis* H37Rv, **EMBO global exchange lecture course-2012**, Hyderabad, India.

**Dey A,** Shrivastava T, Ravishankar R, Feast-famine regulatory protein of *M. tuberculosis* H37Rv, **Mini symposium on macromolecular cyrtsallography-2011**, Hyderabad, India (**abstract selected for podium presentation**)

#### **Courses and Training**

CCP4 study weekend- From Crystals to Structure with CCP4-2010, University of Nottingham, Nottingham, United Kingdom.

#### **Mentoring Experience**

Elizabeth Abrash, PhD student, UNC-Chapel Hill	2020
Tiffany Barwell, (NSF-REU), UNC-Charlotte	2019
Sarah Catherine Paschall (NSF-REU), UNC-Charlotte	2019
Diana Renteria Alvarez (Honors Student), UNC-Charlotte	2018-2019
Justin Davis, (NSF-REU), UNC-Charlotte	2018
Isabella Sabato, UNC-Charlotte	2017-2018
Debarati Bhanja, (NSF-REU), CMU	2017
Jyoti, (Project Assistant-II), CSIR-CDRI	2012-2013

## **Professional affiliation**

RNA Society-Postdoc member American Heart Association member Indian Crystallographic Association member (SM 191)  Scientific Outreach	2020-present 2018-2021 2013-present
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Guest Editor, Journal of Visualized Experiments (JoVE) (RNA Biology: Current Methods and Protocols)	2022
Reviewer, RNA Biology	2022
Reviewer, RNA	2022
Reviewer, BioChem	2022
Reviewer, Healthcare	2022
Reviewer, Non-Coding RNA	2021-present
Reviewer, International Journal of Molecular Sciences (IJMS)	2021-present
Reviewer, Current Issues in Molecular Biology (CIMB)	2021-present
Reviewer, Antibiotics	2021-present
Reviewer, Pharmaceutics	2021-present
Judge, North Carolina Student Academy of Sciences (NCSAS)	2021
Judge, North Carolina Science and Engineering Fair (NCSEF)	2021
Judge, ENVISION Research Competition by Women in Stem (WiSTEM)	2021-2022

### 1) Dr. Alain Laederach

## **Professor**

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## 2) Dr. Kausik Chakrabarti

#### **Assistant Professor**

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## 3) Dr. Ravishankar Ramachandran

### **Chief Scientist (Ph. D Advisor)**

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