

## EDUCATION

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**PhD Biology**, New York University, *New York, NY* Sept. 2018 – May 2023 (expected)  
Thesis: Intraspecies sequence diversity and evolution of ribosomal RNA  
Dissertation advisor: Andreas Hochwagen, PhD

**MSc Molecular Biology**, Moscow State University, *Moscow, Russia* June 2018  
Thesis: PARP-1 interaction with DNA breaks in chromatin  
Thesis advisor: Vasily Studitsky, PhD

**BSc Molecular Biology & Biochemistry**, Moscow State University, *Moscow, Russia* June 2016  
Thesis: Factor PARP-1 and recognition of DNA breaks in the nucleosome  
Thesis advisor: Vasily Studitsky, PhD

## RESEARCH EXPERIENCE

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**Genomics Researcher** (PhD student) New York University, *New York, NY* Sept. 2018 – Present

- Lead a multidisciplinary project involving genomics, bioinformatics, structural and molecular biology
- Discovered and characterized ubiquitous sequence variation of ribosomal RNA in budding yeast, as a potential source of phenotypic diversity and novel translational regulation mechanisms
- Described previously understudied aspects of the evolution of multi-copy ribosomal RNA genes
- Develop large-scale computational and experimental approaches to streamline the exploration of new ribosomal variants with further validation as possible therapeutic targets and for future use in biotechnology and synthetic biology

**Biochemistry Researcher** Fox Chase Cancer Center, *Philadelphia, PA* Aug. 2016 – Aug. 2018

- Discovered a new epigenetic mode of protein PARP1 using FRET microscopy
- Initiated collaborative work on a previously undescribed mechanism of action of anti-cancer drugs
- Co-generated 2 successful grant proposals

## TEACHING & MENTORSHIP EXPERIENCE

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**Teaching Assistant in Protein Biochemistry**, New York University, *New York, NY* Sept. 2020 – Dec. 2022  
*Graduate-level course*

- Taught 2 groups of students both online and in-person
- Prepared study material to help students better understand theory and techniques in protein biochemistry
- Organized midterm and final exams

**Teaching Assistant in Principles of Biology**, New York University, *New York, NY* Sept. 2019 – May 2020  
*Undergraduate-level course*

- Facilitated discussions by using an interactive approach to teach groups of 15-25 students twice a week
- Adapted and taught online recitations to 2 groups of 24 students to broaden course accessibility

**Research Supervisor** Sept. 2021 – Nov. 2022

- Co-developed and oversaw large-scale whole genome sequencing research conducted by 2 graduate students

## AWARDS & FELLOWSHIPS

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- Fleur Strand Graduate Fellowship (NYU, USA, 2022)  
⇒ **Awarded to a student who shows promise in becoming a leader in biomedical research**
- Gladys Mateyko Research Award (NYU, USA, 2021)  
⇒ **Awarded to a student who demonstrated excellence in research**
- Best Graduate Poster Award (ASBMB Protein Data Bank Symposium, USA, 2021)
- Henry M. MacCracken Fellowship (NYU, USA, 2018-2023)
- Best Undergraduate Thesis Award (MSU, Russia, 2016)

## PUBLICATIONS

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- **Sultanov, D.** and Hochwagen, A. Varying strength of selection contributes to the intragenomic diversity of rRNA genes. *Nature Communications* 13, 7245 (2022).

## PREPRINTS

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- Molinar, T., **Sultanov, D.**, Klein, H., Hochwagen, A. Topoisomerase I promotes formation of a novel DNA intermediate in the ribosomal DNA. <https://www.biorxiv.org/content/10.1101/2022.05.24.493248v1>
- Rendleman, J., Mohammad, M., Pressler, M., Maity, S., Hronová, V., Gao, Z., Herrmannová, A., Lei, A., Allgoewer, K., **Sultanov, D.**, Hinckley, W., Szkop, K., Topisirovic, I., Larsson, O., Hatzoglou, M., Valášek, L., Vogel, C. Regulatory start-stop elements in 5' untranslated regions pervasively modulate translation. <https://www.biorxiv.org/content/10.1101/2021.07.26.453809v2>

## INVITED TALKS

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- Rockefeller University - Genome integrity meeting, USA 2022

## CONFERENCE TALKS

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- Biochemical society - Translation, UK 2021
- Temple University - Fels Institute Trainee Day, USA 2018

## POSTER PRESENTATIONS

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- Deep selection shapes the intragenomic diversity of ribosomal RNA. RNA 2022 (Annual meeting of the RNA Society), USA 2022
- A workflow to investigate sequence diversity of highly repetitive rRNA genes in eukaryotes. International Plant and Animal Genome Conference, USA 2022
- Mining for functional ribosomal variants in *Saccharomyces cerevisiae*. ASBMB Protein Data Bank Symposium – 50<sup>th</sup> year celebration, USA 2021  
⇒ **Received an award for “Best graduate poster”**

## PEER REVIEWS

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- **BMC Cancer** (1 manuscript)
- **Cancer Biology and Therapy** (2 manuscripts)
- **Gene** (2 manuscripts)
- **STAR Protocols** (6 manuscripts)

## PROFESSIONAL AFFILIATIONS

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- RNA Society
- Society for Molecular Biology and Evolution