# KIRILL GRIGOREV / КИРИЛЛ ГРИГОРЬЕВ

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COMPUTATIONAL GENETICS, EPIGENOMICS, TRANSCRIPTOMICS, GENOME ASSEMBLY, GENOMICS ALGORITHMS, DATA SCIENCE



## PHD CANDIDATE

Weill Cornell Medicine, New York, NY Mason Lab, Institute for Computational Biomedicine

M.S. in Biology, University of Puerto Rico B.S. in Biotechnology, Saint Petersburg Chemical and Pharmaceutical Academy

## PRINCIPAL AREAS OF ACADEMIC INTEREST

Genomics algorithms Translational and personalized genomics Epigenomics and epitranscriptomics

#### PRINCIPAL SKILLSET

Genomic data analysis Algorithm development Advanced Python, essential R, Perl, and C++

#### **RESEARCH SUMMARY**

2017 –	Weill Cornell Medicine, Institute for Computational Biomedicine, Mason Lab Assembly algorithms, nanopore sequencing data algorithms, epigenomics, comparative transcriptomics, phylogenetics algorithms
2018 –	<b>GeneLab</b> . Analysis of the effect of microgravity on the animal transcriptome
2015 – 2017	University of Puerto Rico, Caribbean Genome Center
	Methods of genome assembly, conservation genetics, Genome 10K
2014 - 2017	Dobzhansky Center for Genome Bioinformatics
	Methods of genome assembly, GWAS visualization tools,
	epigenomics of early childhood development
2013 - 2014	iBinom inc. Medical genome analysis, cloud SaaS

#### PUBLIC SPEAKING & OTHER ACADEMIC EXPERIENCE

2019 <i>talk</i>	Comparative circadian transcriptomics: novel and conserved features of the mammalian pineal gland OU Genomics Symposium, Oakland University, MI
2017 workshop	<b>Development of robust bioinformatics pipelines</b> Fifth annual Bioinformatics Summer School, Moscow, Russia
2017 <i>talk</i>	Genomics and conservation of the Hispaniolan Solenodon  IX Caribbean Biodiversity Congress, Santo Domingo, Dominican Republic
2016 <i>TA</i>	Bioinformatics pipelines Recent Advances in Conservation Genetics, Tihany, Hungary
2015 workshop	Linux toolset for bioinformatics Third annual Bioinformatics Summer School, Moscow, Russia
2015 lectures	Introduction to genetics Biotechnology Stepik.org online course

### **PUBLICATIONS**

- F Gaiti, R Chaligne, H Gu, RM Brand, S Kothen-Hill, RC Schulman, K Grigorev, D Risso, K Kim, A Pastore, KY Huang, A Alonso, C Sheridan, ND Omans, E Biederstedt, K Clement, L Wang, JA Felsenfeld, EB Bhavsar, MJ Aryee, JN Allan, R Furman, A Gnirke, CJ Wu, A Meissner, DA Landau. Epigenetic evolution and lineage histories of chronic lymphocytic leukaemia. Nature 569 (7757), 576. doi:10.1038/s41586-019-1198-z
- 2. ABR McIntyre, N Alexander, K Grigorev, D Bezdan, H Sichtig, CY Chiu, CE Mason. **Single-molecule sequencing detection of N6-methyladenine in microbial reference materials**. Nature Communications 10 (1), 579. doi:10.1038/s41467-019-08289-9
- S Kolchanova, S Kliver, A Komissarov, P Dobrinin, G Tamazian, K Grigorev, WW Wolfsberger, AJ Majeske, J Velez-Valentin, R Valentin de la Rosa, JR Paul-Murphy, DSM Guzman, MH Court, JL Rodriguez-Flores, JC Martinez-Cruzado, TK Oleksyk. Genomes of three closely related Caribbean amazons provide insight for species history and conservation. Genes 10 (1), 54. doi:10.3390/genes10010054
- 4. K Grigorev, S Kliver, P Dobrynin, A Komissarov, W Wolfsberger, K Krasheninnikova, YM Afanador-Hernandez, AL Brandt, LA Paulino, R Carreras, LE Rodriguez, A Nunez, JR Brandt, F Silva, JD Hernandez-Martich, AJ Majeske, A Antunes, AL Roca, SJ O'Brien, JC Martinez-Cruzado, TK Oleksyk. Innovative assembly strategy contributes to understanding the evolution and conservation genetics of the endangered Solendon paradoxus from the island of Hispaniola. GigaScience 7 (6), giy025. doi:10.1093/gigascience/giy025
- 5. OY Naumova, M Dozier, PV Dobrynin, K Grigorev, A Wallin, I Jeltova, M Lee, A Raefski, EL Grigorenko. **Developmental dynamics of the epigenome: a longitudinal study of three toddlers**. Neurotoxicology and teratology 66, 125-131. doi:10.1016/j.ntt.2017.12.006
- AL Brandt, K Grigorev, YM Afanador-Hernandez, LA Paulino, WJ Murphy, A Nunez, A Komissarov, JR Brandt, P Dobrynin, JD Hernandez-Martich, R Maria, SJ O'Brien, LE Rodriguez, JC Martinez-Cruzado, TK Oleksyk, AL Roca. Mitogenomic sequences support a north-south subspecies subdivision within Solendon paradoxus. Mitochondrial DNA Part A 28 (5), 662-670. doi:10.3109/24701394.2016.1167891