

# KIRILL GRIGOREV / КИРИЛЛ ГРИГОРЬЕВ

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COMPUTATIONAL GENETICS, GENOMICS ALGORITHMS, GENOMICS DATABASES, SPACE GENOMICS, TELOMERES, GENOME ASSEMBLY, DATA SCIENCE



## PHD CANDIDATE

**Weill Cornell Medicine, New York, NY** | [weill.cornell.edu](http://weill.cornell.edu)

**Mason Lab**, Institute for Computational Biomedicine | [masonlab.net](http://masonlab.net)

**GeneLab** Multi-Omics Analysis Working Group and Visualization Working Group

[genelab.nasa.gov/awg/charter](http://genelab.nasa.gov/awg/charter) | [visualization.genelab.nasa.gov](http://visualization.genelab.nasa.gov)

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  
Space genetics

## PRINCIPAL SKILLSET

Genomic data analysis  
Graph and numerical algorithms  
Advanced Python (SciPy stack, Numba, Dask)  
Lua/LuaJIT, R, essential C and Perl

## RESEARCH SUMMARY

- 2017 – ... **Weill Cornell Medicine, Institute for Computational Biomedicine, Mason Lab**  
Assembly algorithms, telomere bioinformatics, sequencing data analysis  
Developed frameworks for identification of novel telomeric motifs and haplotype inference [Gri21], studied the effects of prolonged spaceflight on dynamics of telomeres [Lux20] and cell-free DNA [Bez20], carried out sequence analyses in translational [Wes20] and theoretical [Mci19] applications.
- 2018 – ... **GeneLab**. Cross-dataset analysis platform for space biology data  
Developed an API for transparent programmatic, as well as browsable, access to the space flight biological data stored in the GeneLab repository, powering data visualization and enabling a range of downstream applications  
[Ber21, Sco20, Ber20, [visualization.genelab.nasa.gov/GLOpenAPI](http://visualization.genelab.nasa.gov/GLOpenAPI)].
- 2018 **New York Genome Center**. Epigenetic evolution of cancers, phylogenetics algorithms  
Developed a computationally faster, scalable algorithm for the calculation of the four-gamete test [Gai19].
- 2015 – 2017 **University of Puerto Rico, Caribbean Genome Center**  
Methods of genome assembly, conservation genetics, Genome 10K  
Carried out *de novo* whole-genome and mitogenome assemblies for a number of endangered Caribbean species, contributing to the understanding of conservation status and strategies [Kol19, Gri18, Bra17].
- 2014 – 2017 **Dobzhansky Center for Genome Bioinformatics**  
Methods of genome assembly, GWAS visualization tools, human epigenetics  
Studied epigenomics of early childhood development [Nau18], processed data for visualization in the Genome-Wide Association Tracks Chromosome Highway ([gen-watch.org](http://gen-watch.org)).
- 2013 – 2014 **iBinom inc.** Medical genome analysis, cloud SaaS

## PUBLICATIONS

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- Gri21 K Grigorev, J Foux *et al.* **Haplotype diversity and sequence heterogeneity of human telomeres.** *Genome Research* 31 (7), 1269. doi:[10.1101/gr.274639.120](https://doi.org/10.1101/gr.274639.120)
- Ber21 D Berrios *et al.* **NASA GeneLab: interfaces for the exploration of space omics data.** *Nucleic Acids Research* 49 (D1), D1515. doi:[10.1093/nar/gkaa887](https://doi.org/10.1093/nar/gkaa887)
- Lux20 J Luxton *et al.* **Temporal Telomere and DNA Damage Responses in the Space Radiation Environment.** *Cell Reports* 33 (10), 108435. doi:[10.1016/j.celrep.2020.108435](https://doi.org/10.1016/j.celrep.2020.108435)
- Bez20 D Bezdan *et al.* **Cell-free DNA (cfDNA) and exosome profiling from a year-long human spaceflight reveals circulating biomarkers.** *IScience* 23 (12), 101844. doi:[10.1016/j.isci.2020.101844](https://doi.org/10.1016/j.isci.2020.101844)
- Sco20 R Scott *et al.* **Advancing the Integration of Biosciences Data Sharing to Further Enable Space Exploration.** *Cell Reports* 33 (10), 108441. doi:[10.1016/j.celrep.2020.108441](https://doi.org/10.1016/j.celrep.2020.108441)
- Ber20 D Berrios *et al.* **Visualizing Omics Data from Spaceflight Samples using the NASA GeneLab Platform.** In *Proceedings of the 12th International Conference on Bioinformatics and Computational Biology* (Vol. 70, pp. 89-98). doi:[10.29007/rh7n](https://doi.org/10.29007/rh7n)
- Wes20 C Westover *et al.* **Engineering Radioprotective Human Cells Using the Tardigrade Damage Suppressor Protein, DSUP.** *bioRxiv* (2020). doi:[10.1101/2020.11.10.373571](https://doi.org/10.1101/2020.11.10.373571)
- Gai19 F Gaiti, R Chaligne, H Gu *et al.* **Epigenetic evolution and lineage histories of chronic lymphocytic leukaemia.** *Nature* 569 (7757), 576. doi:[10.1038/s41586-019-1198-z](https://doi.org/10.1038/s41586-019-1198-z)
- Mci19 ABR McIntyre *et al.* **Single-molecule sequencing detection of N6-methyladenine in microbial reference materials.** *Nature Communications* 10 (1), 579. doi:[10.1038/s41467-019-08289-9](https://doi.org/10.1038/s41467-019-08289-9)
- Kol19 S Kolchanova, S Kliver *et al.* **Genomes of three closely related Caribbean amazons provide insight for species history and conservation.** *Genes* 10 (1), 54. doi:[10.3390/genes10010054](https://doi.org/10.3390/genes10010054)
- Gri18 K Grigorev, S Kliver *et al.* **Innovative assembly strategy contributes to understanding the evolution and conservation genetics of the endangered *Solenodon paradoxus* from the island of Hispaniola.** *GigaScience* 7 (6), giy025. doi:[10.1093/gigascience/giy025](https://doi.org/10.1093/gigascience/giy025)
- Nau18 OY Naumova *et al.* **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](https://doi.org/10.1016/j.ntt.2017.12.006)
- Bra17 AL Brandt, K Grigorev *et al.* **Mitogenomic sequences support a north-south subspecies subdivision within *Solenodon paradoxus*.** *Mitochondrial DNA Part A* 28 (5), 662-670. doi:[10.3109/24701394.2016.1167891](https://doi.org/10.3109/24701394.2016.1167891)

## PUBLIC SPEAKING, ROUNDTABLES

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- talk, 2020 **GeneLab sequencing data analysis and visualization.** Virtual (USRA / NASA Ames Research Center)
- talk, 2019 **Comparative circadian transcriptomics: novel and conserved features of the mammalian pineal gland.** OU Genomics Symposium, Oakland University, MI
- workshop, 2019 **GeneLab Visualization Working Group meeting.** 35th Annual Meeting of ASGSR, Denver, CO
- workshop, 2019 **GeneLab visualization workshop.** Broad Institute, Cambridge, MA
- workshop, 2017 **Development of robust bioinformatics pipelines** Fifth annual Bioinformatics Summer School, Moscow, Russia
- talk, 2017 **Genomics and conservation of the Hispaniolan *Solenodon*** IX Caribbean Biodiversity Congress, Santo Domingo, Dominican Republic
- TA, 2016 **Bioinformatics pipelines.** Recent Advances in Conservation Genetics, Tihany, Hungary
- workshop, 2015 **Linux toolset for bioinformatics.** 3rd annual Bioinformatics Summer School, Moscow, Russia
- instructor, 2015 **Introduction to genetics.** Biotechnology Stepik.org online course ([stepik.org/course/94](https://stepik.org/course/94))