

**Svetlana Volkova**  
**CURRICULUM VITAE**

phone +45 42 73 84 11 · svetlanavolkova.v@gmail.com · svevol@biosustain.dtu.dk

## PERSONAL INFORMATION

Born 26 November 1993 in Khabarovsk, Russia

Nationality: Russian

Languages: Russian, English

Now I live in Copenhagen, Denmark

## EDUCATION

|             |  |                              |
|-------------|--|------------------------------|
| 2017-Now    | <b>Novo Nordisk Foundation Center for Biosustainability (DTU Biosustain)<br/>Denmark</b><br><b>NNF Copenhagen Bioscience PhD programme</b><br>PhD student<br>2017-2018, Pre-doc year student (Research Assistant)          |                              |
| 2015 - 2017 | <b>Skolkovo Institute of Science and Technology (Skoltech), Skolkovo</b><br>Biomedical Science and Technology Programme<br>Master of Science in Biotechnology<br>Language of instruction - English<br>Diploma with honours | <b>GPA</b><br><b>5.0/5.0</b> |
| 2011 - 2015 | <b>Lomonosov Moscow State University, Moscow</b><br>Biological Faculty, Division of Biophysics and Bioengineering<br>Bachelor of Science in Biology<br>Diploma with honours  | <b>GPA</b><br><b>4.9/5.0</b> |

## RESEARCH EXPERIENCE

|             |  |
|-------------|--|
| 2017 - Now  | <b>Novo Nordisk Foundation Center for Biosustainability (DTU Biosustain),<br/>Denmark</b><br>Supervisor: Prof. Lars Keld Nielsen<br>PhD project is aimed at metabolomics data analysis with the help of mechanistic model to explain the metabolism of Red Blood Cells and aberrant metabolic phenotype, anaemias. I construct kinetic models, perform data integration and analysis. Big part of my project is dedicated to Bayesian models creation and inference. |
| 2016 - 2017 | <b>Center for Translational Biomedicine, Skoltech, Skolkovo</b><br>Supervisor: Prof. Yuri Kotelevtsev<br>Master thesis: Validation of Egr2 as a target for RNAi based suppression of liver fibrosis.   |
| 2014 - 2015 | <b>Cellular Neurobiology of Learning Lab, Institute of Higher Nervous Activity and<br/>Neurophysiology of RAS, Moscow</b><br>Supervisor: Kolosov PM, PhD<br>Bachelor Thesis: Deep sequencing-based identification of immediate-early genes set during activation of central nervous system of <i>Helix lucorum</i> .   |
| 2012-2014   | <b>Confocal Microscopy Lab, Lomonosov Moscow State University, Moscow</b><br>Supervisor: Moisenovich MM, PhD<br>Project: Silk fibroin application in tissue regeneration and 3D culturing  |

## INDUSTRY EXPERIENCE

|                                |   |
|--------------------------------|---|
| May 2016 -<br>December<br>2016 | <b>BostonGene, Boston MA, USA (Moscow office)</b><br>Bioinformatician, Analyst<br>Responsibilities: SNP/indels calling based on NGS data, variant filter (false positive variants detection) development and usage, variant annotation, tumor neoantigen detection pipeline development |
|--------------------------------|---|

## HONORS, AWARDS, SCHOLARSHIPS

|                |   |
|----------------|---|
| March 2019     | 3rd Advanced Lecture Course on Computational Systems Biology fellowship, selected short talk, Aussois |
| April 2017     | PhD fellowship, Copenhagen Bioscience PhD programme, Novo Nordisk Foundation                          |
| September 2016 | Academic Excellence Award, Skoltech   |
| September 2013 | Heightened state scholarship for the scientific achievements, MSU                                     |

## TECHNICAL SKILLS

**I do metabolic modelling on every day basis. It involves data cleaning, handling, statistical test usage and visualisation, creation of mechanistic models and data integration. For these activities one need a range of programming skills and several programming languages usage. Below, you can find the keywords that can give you a better idea of typical work I do.**

**Computer skills** Unix, R, python, Matlab, statistics, exploratory data analysis, data visualization, Bayesian inference (PyMC3), Approximate Bayesian Computation, MCMC, pandas, scipy

**Metabolic modelling** Metabolic kinetic modelling, Metabolic Control Analysis, CobraPy, Flux Balance Analysis, Metabolic Flux Analysis

**NGS bioinformatics** Transcriptome assembly, transcriptome reads mapping, differential gene expression analyses, pathway/GO enrichment analyses, annotation  
Exome reads mapping, SNP/indels calling and filtering based on NGS data, variant annotation, variants manual revision, neoantigen peptide identification

**Analytical chemistry** Chromatography, Mass Spectrometry, LC-MS, HPLC

**Communication** MS Powerpoint, Adobe Illustrator, Twitter

**Lab biology** **My interests now lie in the area of metabolic modeling and do not require the usage of vlassical molecular and cell biology techniques. However, I have a strong experience in such work that gives me a good feeling about work performed around me and helps in collaboration with both wet-lab and dry-lab colleagues, especially while looking for the “translation”.**

**Molecular biology** Isolation of RNA, DNA, PCR, qPCR, cloning, Gibson assembly, agarose gel electrophoresis, PAGE

**Microbiology** Media preparation and usage, aseptic and sterile techniques, plating methods, bacterial staining

**Animal cell culture** 2D, 3D cell culturing, cells transfection: electroporation, lipofection, cells fixation and staining

**Histology** Basic histological skills: fixation, paraffin embedding, sectioning, immunohistochemistry

**Animals** Mice – injections, handling

## CONFERENCES

**Selected talk, presentation:** Towards personalized mechanistic models of red blood cell metabolism// CompSysBio, Advanced Lecture Course on Computational Systems Biology, Aussois, March 2019

## RESEARCH PUBLICATIONS [Google scholar link](#)

- Chromosomal barcoding as a tool for multiplexed phenotypic characterization of laboratory evolved lineages / Leonie Johanna Jahn, Andreas Porse, Christian Munck, Daniel Simon, **Svetlana Volkova**, Morten Otto Alexander Sommer // Scientific reports - 2018. doi:10.1038/s41598-018-25201-5

- Adaptive changes in the vestibular system of land snail to a 30-day spaceflight and readaptation on return to Earth / Nikolay Aseyev, Aliya Vinarskaya, Matvey Roshchin, Tatiana A. Korshunova, Aleksey Malyshev, Alena Zuzina, Victor N. Ierusalimsky, Maria Lemak, Igor S. Zakharov, Ivan A. Novikov, Peter Kolosov, Ekaterina Chesnokova, **Svetlana Volkova**, Artem Kasianov, Leonid Uroshlev, Yekaterina Popova, Richard D Boyle, Pavel M. Balaban // *Frontiers in Cellular Neuroscience* - 2017. doi: 10.3389/fncel.2017.00348
- Composite scaffolds containing silk fibroin, gelatin, and hydroxyapatite for bone tissue regeneration and 3d cell culturing / M. M. Moisenovich, A. Yu. Arkhipova, A. A. Orlova, M. S Drutskaya, **S. V. Volkova**, S. E. Zacharov, I. I. Agapov, M. P. Kirpichnikov // *Acta naturae* - 2014. PMID: 24772332
- Gelatin concentration impact in composite silk fibroin-based scaffolds on mouse embryonic fibroblasts adhesion and proliferation / A. Orlova, M. Kotlyarova, V. Lavrenov, **S. Volkova**, A. Arkhipova // *Bull Exp Biol Med.* - 2014 (In Russian). doi: 10.1007/s10517-014-2699-2