

# Makhonin Alex

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## Skills

- Golang | Python | Kubernetes | Docker | containerd | Werf | Helm | Buildah | Arduino | Perl | Linux | Bash | Redis | RabbitMQ | Git
- AWS | GCP | Selectel | Yandex Cloud | CI/CD | Gitlab CI | GitHub Actions | Grafana | Prometheus | Zabbix | Nginx | HAProxy | Gunicorn
- MySQL | PostgreSQL | MongoDB | CockroachDB | Terraform | Puppet | Microservices | Distributed Systems | Backend | DevOps
- English - B2 | French - A1 | Russian – Native

## Experience

### Software Engineer

**Flant**

Remote, Russia

02/2023 - Current

- Developed and increased stability of the [Deckhouse Kubernetes platform](#) and the open source projects it uses

### DevOps Engineer

**Flant**

Remote, Russia

12/2021 - 02/2023

- Supported high availability containerized systems.
- Setted up and deployed applications using **Werf, Helm, Buildah, Docker** in **Gitlab CI** or **Github Actions**.
- Implemented CI/CD patterns to simplify product development (review environments, canary deployment, blue-green deployment, multi data center deployment).
- Maintained and developed **Linux** server infrastructure.
- Extended and used the **Kubernetes** API to simplify the routine operations ([annotations-exporter](#) and [database-users-operator](#)).
- Contributed to open-source repositories ([deckhouse](#), [redis-sentinel-proxy](#), [ingress-nginx](#))

### Junior Bioinformatic Scientist

**BostonGene**

Moscow, Russia

09/2021 - 11/2021

- Improved NGS Data Quality Control processes with **Python, FastQC, MultiQC**.
- Created a service for identifying patients by SNPs whose sequencing has already been studied (**PostgreSQL + SQLAlchemy + NumPy**).

### Research Laboratory Technician

**Shemyakin and Ovchinnikov Institute  
of Bioorganic Chemistry**

Moscow, Russia

01/2020 - 09/2021

- Created and maintained web services for internal purposes using a **Linux** server, **Nginx, Gunicorn, Python + Django** ([project link](#)).
- Designed and developed automatic microscope lens rotation using **Arduino** and stepper motor ([project link](#)).
- Constructed fast-drug application system with [Openspritzer](#) for patch-clamp research.
- Attempts have been made to construct a model for predicting whether a protein belongs to the Ly6/uPAR group and test it on the sequencing of two starfish (**Pandas, scikit-learn, NumPy, blast**) ([project link](#)) ([article](#)).
- Wrote Python functions to visualize the alignment of Ly6/uPAR proteins with cysteine-cysteine bonds. (**Matplotlib, biotite, NumPy, Biopython, seaborn**) ([project link](#)) ([article](#)).

## Education

### Bachelor's Degree

**National Research University  
Higher School of Economics**

Moscow, Russia

09/2019 - Current

- Cell and Molecular Biotechnology, Department of Biology and Biotechnology

## Projects

- [regex-dictionary](#): Python type to use regex as keys in dict. (01/2023)
- [cube-rotate](#): golang implementation of [cube.c](#) and [donut-math](#) with user-friendly cli. (11/2022)
- [go-grep](#): A simple library to replace grep functionality in go. (08/2022)
- [Third year coursework](#): "Sequence-Based IsomiR Target Prediction" (repositories that were also used: [mirdbts](#) and [TargetScan](#)) (2022)
- [Second year coursework](#): "Expression of Ly6/uPAR Proteins in Alzheimer Disease". (2021)

## Others

- **Machine learning Summer school "MTS.Theta"**: Result project: [human text recognition](#) model with telegram bot and [simple frontend for browser](#) (08/2021)
- [Article "Just-for-fun experiment: Deploying Kubernetes on two old laptops with Gentoo Linux"](#) (11/2022)
- [Scopus Author profile](#)