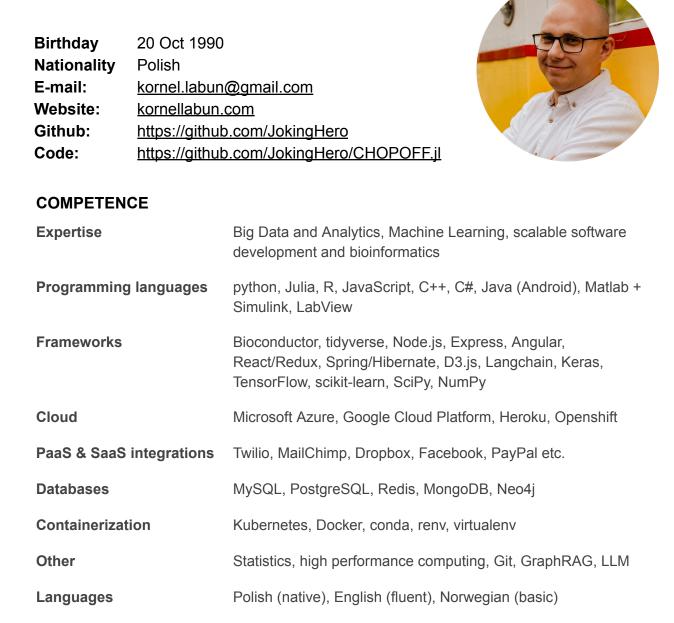
Kornel Labun



INTERESTS

PERSONALITY

Jujitsu, judo, chess and books	Lighthearted
Blockchain and design patterns	Persistent
Abstraction and Reasoning Corpus	Curious
Machine learning and generative models	Helpful

EDUCATION

Sep 2015 - Oct 2019 PhD: University of Bergen, Department of Informatics, Computational Biology Unit

Title: "In silico design and analysis of gene editing experiments." Defense date: 27th January 2020

Oct 2009 - Sep 2014 MSc Eng: Silesian University of Technology, Faculty of Automatic Control, Electronics and Computer Science; Biotechnology, spec. Bioinformatics

> Title: "Spatial evolutionary games as a tool for modeling inter-population interactions." Defense date: 16th September 2014

EXPERIENCE

Dec 2022 -

Researcher at University of Bergen

- <u>CHOPOFF</u> tool for off-target detection (10x speed and 100% precision, 96% code coverage)
- Analysis of datasets and tool creation (GUIDE-seq, CRISPR etc.) for precision pediatrics and gene editing
- RCP-seq (RNA/80S/SSU sequencing) for mouse neuronal cells
- <u>https://ribocrypt.org/</u> interactive visualization in genomics

Jan 2021 - Nov 2022

- Postdoctoral fellow at University of Bergen
 - Development of tools for off-target detection for genome editing with CRISPR, enabling precision medicine
 - Applying Oxford Nanopore technology for modification detection
 - Analyzing Ribo-seq and TCP-seq datasets for metastasis signature
- Nov 2019 Dec 2020 Senior Consultant at Sonat Consulting AS
 - Youwell/Helse I Hardanger Software engineering Android development and Bluetooth integration: RxJava, Dagger2, Retrofit, MVVM, backend: Azure/C#, Redux/React
 - Cutters Software engineering and data analysis Back-end website development, building marketing platform in Google Cloud Platform, Python, SQL, Facebook API integration and data analysis.
 - Absa Software engineering and data analysis Build analysis service in SQL Server Analysis Services (SSAS multidimensional cube). Big data ETL, data warehouse with schema design and dimensional data modeling.

Sep 2015 - Oct 2019	 PhD student: "In silico design and analysis of gene editing experiments." supervisor: Eivind Valen Analysis of RNA-Seq, Ribo-Seq, Cage-Seq, ATACK-Seq, SMS-seq and other *-Seq kind of datasets, as well as Oxford Nanopore sequencing data (noisy signal over time) Development of pipelines and tools (ORFik, ampliCan, RareVariantVis, CHOPCHOP, tailfindR) Machine learning related projects: prediction of polyA tail from nanopore data, prediction of repair profile after CRISPR edits Teaching for courses: INF207 (Social Network Theory), INF109 (Computer Programming for Science), and <u>R Crash Course</u> for Molecular and Computational Biology Research School and NORBIS
Jul 2014 - Jul 2015	Software Developer/data analyst at rspective (renamed to Voucherify)
	Front-end and back-end website development, machine learning solutions and data analysis, quality assessment, architectural design and deployment.
2013 - 2014	MSc project: "Spatial evolutionary games as a tool for modeling inter-population interactions". supervisor: Prof. Andrzej Świerniak
	Evolutionary game theory in the cellular automaton to solve differential equations
2012 - 2013	Eng project: "Seeking for the signature of radiosensitivity with the use of data on single nucleotide polymorphisms and the results of γ -H2AX test with the expressions of the chosen DNA repair genes." supervisor: Prof. Joanna Polańska
	Developing a novel statistical model for the signature of radiosensitivity.
Aug 2011	Internship at Maria Sklodowska-Curie Memorial Cancer Center and Institute of Oncology Gliwice Branch, Department of Nuclear Medicine and Endocrine Oncology
	Designing microRNA discovery pipeline.

SELECTED PUBLICATIONS

Total number of publications: **15** h-index: **10**

CHOPCHOP v3: expanding the CRISPR web toolbox beyond genome editing.

Labun K, Montague TG, Krause M, Torres Cleuren YN, Tjeldnes H, Valen E Nucleic acids research, 47(W1):W171–W174. (2019) Citations: 1411

CHOPCHOP v2: a web tool for the next generation of CRISPR genome engineering.

Labun K, Montague TG, Gagnon JA, Thyme SB, Valen E Nucleic acids research, 44(W1):W272–6 (2016) Citations: 990

tailfindr: alignment-free poly(A) length measurement for Oxford Nanopore RNA and DNA sequencing.

Krause M, Niazi AM, <u>Labun K</u>, Torres Cleuren YN, Müller FS, Valen E RNA , 25(10):1229–1241. (2019) Citations: 80

Accurate analysis of genuine CRISPR editing events with ampliCan.

Labun K, Guo X, Chavez A, Church G, Gagnon JA, Valen E Genome Research , 29 (5) (2019) Citations: 65

ORFik: a comprehensive R toolkit for the analysis of translation

Tjeldnes H, <u>Labun K</u>, Torres Cleuren YN, Chyżyńska K, Świrski M, Valen E BMC bioinformatics 22 (1), 336 Citations: 26

TEACHING EXPERIENCE

- May 2021Organizer and teacher of R Crash Course for NORBIS (national research
school in bioinformatics, biostatistics and systems biology) and MCB
(Molecular and Computational Biology) research school. Course website:
r-crash-course.github.io
- Spring 2018INF109 (Computer Programming for Science) for Institute ofSpring 2017Informatics/University of Bergen. Designing homework assignments and
grading them.

Autumn 2018INF207 (Social Network Theory) for Institute of Informatics/University ofAutumn 2016Bergen. Teaching exercises on a blackboard.