

# Krisztián Ádám fullstack developer

adamkrisz@gmail.com +36 30 2327855 hangyas.github.io SKILLS
Elixir/OTP
Tailwind CSS
Scala
Docker
Clojure

## **WORK EXPERIENCE**

## 2022-present Senior Elixir developer @ TSG Global

- Automating the company's billing processes (including stripe integration)
- Fixing bugs and improving the performance of legacy elixir services handling millions of messages per hour

## 2020-2022 Senior Elixir developer @ LIQID Investments GmbH

- Building integrations for various REST API
- Utilizing distributed erlang and OTP features
- Designing a secure public API
- Proposing and delivering an admin dashboard with LiveView and Tailwind CSS

## 2019 Ruby on Rails developer @ Tigra kft

- SQL and HTTP request optimizations
- General full-stack development (ruby, RoR, CSS, Less, ¡Query, PostgreSQL)
- REST API client and server development

## 2016-2020 Freelancing / Student work

- Collaborating with international teams as well as delivering projects on my own
- UpEd: Elixir, Phoenix, Ecto, JSON-API, PostgreSQL
- Telegram bots: NodeJS, MongoDB, docker
- ETL tool: python, xml, yaml, json
- Blackcart.co: Fullstack development, CORS, subdomains, Twillo, Stripe integration
- Serokell: frontend developer (TypeScript, CSS, nix)

## **PUBLICATIONS**

(2021) Depthwise Convolutions using Physicochemical Features of DNA for Transcription Factor Binding Site Classification: Physicochemical Features for DNA-Protein Classification with Depthwise Convolutions (et al) doi:10.1145/3571560.3571563

(2021) Transcription factor binding site detection using convolutional neural networks with a functional group-based data representation (et al) <a href="https://doi.org/10.1088/1742-6596/1824/1/012001">doi:10.1088/1742-6596/1824/1/012001</a>

(2021) Training models employing physico-chemical properties of DNA for protein binding site detection (et al) <a href="https://doi.org/10.1109/ICAPAI49758.2021.9462057">doi:10.1109/ICAPAI49758.2021.9462057</a>

(2018) DNA Readout Viewer (DRV): visualization of specificity determining patterns of protein-binding DNA segments <a href="https://doi.org/10.1093/bioinformatics/btz906">https://doi.org/10.1093/bioinformatics/btz906</a>

- Building a SaaS webapp (https://drv.brc.hu)

#### OTHER ACHIEVEMENTS

## 2018 Teaching Python class for natural science students

- Covering python basics, numpy, matplotlib

# 2016-2022 Telegram Bot API wrapper for crystal

- Developing and maintaining the library in my free time
- It had 68 stars on github
- <a href="https://github.com/hangyas/telegram\_bot">https://github.com/hangyas/telegram\_bot</a>

# 2013-2015 HTML5 apps for Firefox OS

- They were actually successful until Mozilla canceled Firefox OS

#### **EDUCATION**

# 2015-2018 Molecular Bionics Engineering @ University of Szeged

- Including computer science, algorithms, data structures
- physics and biology

#### **CERTIFICATIONS**

2019 Cambridge Assessment English C1

2018 < EPAM> Introduction to Big Data - graduated with honors



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