# Árpád Goretity

Software Architect, Data Scientist

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#### in h2co3 ♂

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- ⊕ goretity.com 
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#### **WORK EXPERIENCE**

# **Lead Bioinformatics Scientist**

Medipredict Ltd.

Dec 2019 - Present

- Designed and implemented internal tools in Python and Rust for microbiome and metabolomics analysis
- Explored data using Pandas and Seaborn
- Built 2 disease prediction models in Scikit-learn
- Automated 2 document generation tasks using Typst
- Advised and mentored 2 junior team members
- Mediated between microbiome researchers, medical doctors, software engineers, and the DevOps team

## **Data Platform Engineer**

PrivátDoktor (S+H Portfolio PLC)

Sept 2020 - Present

- Trained and deployed ML models for a medical decision support system: analyzed ECG, PPG, blood glucose, and blood pressure data using SciPy
- Improved performance of existing ECG signal processing pipeline with Rust and WASM
- Designed normalized database for training data in SQL Alchemy, SQLite, and PostgreSQL

# Site Reliability Engineer

NumberEight Technologies Ltd.

May 2022 - June 2022

- Migrated existing services to Terraform and K8s
- Deployed Google Managed Prometheus
- Created a containerized, reproducible development environment for future SREs

# iOS Application Developer

Freelancer @ Curl (UK), OTP Group, DBIT, Codie (HU) June 2012 - Aug 2018

- Maintained and refactored large, mixed Swift and Objective-C code bases (Curl, iCsekk, Codie)
- Implemented client-side business logic involving local data storage, networking, and cryptography
- Fixed a critical security vulnerability in iCsekk

# **LANGUAGES**

English (C2) French (C1)

Hungarian

Italian

#### **EDUCATION**

#### **MSc in Data Science**

Università degli Studi di Padova, Dip. di Matematica Sept 2018 - Jul 2020

- Thesis: Towards Personalized Disease Risk Prediction from Metagenome Analysis of the Microbiome
- Qualification: 110 / 110, Cum Laude

## **BSc in Bionic Engineering**

Pázmány Péter Catholic University (PPCU FITB) Sept 2013 - Dec 2016

- Thesis: Design and FPGA Implementation of a Protein Structure Comparison Method Based on Alignment of Backbone Conformations
- Qualification: 5 / 5 Excellent

#### **SKILLS**



## **PROJECTS**

#### Parsel 2

Compilers, Parsing, Macros

 Automatically generate a parser from an AST specification, without any mapping boilerplate

# NanoSQL 🗗

Databases, ORMs

 Strongly-typed, very lightweight data mapper for SQLite and Rust

#### SteelSafe 🛂

Cryptography, Security

 TUI password manager using modern cryptography (Argon2id and XChaCha20-Poly1305)