



## ALINA DENZLER

M.Sc. in Computational Chemistry

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I am a Computational Chemist with a background in molecular modeling and machine learning. I have worked on ligand- and structure-based drug discovery using Python and AI tools. My experience spans academic research across multiple countries. Currently, I explore GenAI.

[LinkedIn](#) • [GitHub](#)

### SKILLS

Python

SQL

LLMs

Drug Discovery

Machine Learning

Data Mining

Statistical Analysis

Cross-Functional Collaboration

### LANGUAGES

English – fluent



Russian – fluent



German – beginner



### EDUCATION

2016 – 2018

**M.Sc. in computational chemistry**

Kazan Federal University, Russia

2012 – 2016

**B.Sc. in pharmaceutical technology**

Saint Petersburg Chemical Pharmaceutical Academy, Russia

### EXTRACURRICULAR ACTIVITIES

➤ Drawing

A space for creative exploration through new techniques, materials, and visual storytelling

➤ Hiking

A way to recharge, stay grounded, and enjoy expansive views

### WORK EXPERIENCE

**GenAI Explorer**

Self-employed

01/2025 — Now

Basel, Switzerland

- Leading AI research pipelines with a focus on data collection, cleaning, analysis, and preparation for modeling.
- GitHub Repo (in progress)

**Sabbatical Break**

Career break

03/2024 — 12/2024

Switzerland & Abroad

- Explored new countries and cultures
- Established a Swiss presence and expanded professional network through biotech and pharma events

**Chemical Data Scientist**

Palacký University

08/2019 — 02/2024

Olomouc, Czech Republic

- Built an end-to-end pipeline to identify selective MARK4 inhibitors for Alzheimer's research, combining ML, docking, and chemical similarity analysis over **1.78M compounds**, resulting in **18% experimental validation** and novel scaffold discovery.

[Article](#) · [GitHub Repo](#)

- Co-developed a virtual screening pipeline based on 4D pharmacophore modeling and molecular dynamics, enabling hit identification across 28,000 compounds and achieving **10–20x better enrichment** than conventional methods.

[Article](#) · [GitHub Repo](#)

- Co-developed a de novo compound generation tool producing **10,000+ synthetically feasible molecules in 12 hours** with **>70% novelty**, validated on **818K structures** and benchmarked across diverse protein families.

[GitHub Repo](#)

- Led data analysis for CACHE Challenge #1, targeting the LRRK2 protein (Parkinson's); applied MD simulations and feature space mapping to prioritize compounds, contributing to a **top 3 finish out of 23 global teams**.

[Article](#)

- Delivered a **Python workshop** for drug discovery scientists (Olomouc, 2023), covering data analysis, QSAR, and machine learning, and translating complex concepts for a multidisciplinary audience.

[GitHub Repo](#)

**Junior Researcher**

Kazan Federal University

02/2017 — 07/2019

Kazan, Russia

- Developed and validated a ligand-based virtual screening tool, outperforming traditional 2D methods across targets like AChE and CYP3A4; tool adopted by **Insilico Medicine** for commercial use.

[Article](#) · [GitHub Repo](#)

- Applied machine learning to pharmacophore-based screening using one-class classifiers, improving compound ranking accuracy and early enrichment; results published and deployed as a **public web tool**.

[Article](#) · [GitHub Repo](#) · [Public Web Tool](#)

**Lab Assistant**

Biocad

06/2015 — 07/2015

Moscow, Russia

- Supported monoclonal antibody production through cell culture; authored documentation for Rituximab manufacturing in accordance with GMP standards.