

# MARK C ZIELINSKI, Ph.D.

Boston, MA | (708) 539-4138 | [mzielinski@gmail.com](mailto:mzielinski@gmail.com)  
[github.com/mczielinski](https://github.com/mczielinski) | **mcz.fyi** | [linkedin.com/in/mczie](https://linkedin.com/in/mczie)

## SKILLS

---

<b>Languages and tools</b>	Python, MATLAB, R, HTML, CSS, Javascript, Bash/Zsh, SQL, Unix/Linux/HPCC, Git/GitHub/Bitbucket, Docker, AWS, GCP, LLMs, MLflow, DVC
<b>Packages</b>	Jupyter, Pandas, NumPy, SciPy, Scikit-learn, Matplotlib, Seaborn, Bokeh, BeautifulSoup, Selenium, PyTorch, TensorFlow, Tensorboard, DESeq2, Hugging Face, PySpark, Dask, XGBoost
<b>Skills &amp; Techniques</b>	regression, classification, clustering, resampling, dimensionality reduction, modeling, machine learning, experimental design, parametric/nonparametric/circular/Bayesian statistics, causality inference, class imbalance, time series analysis, digital/analog signal processing, time delay embedding, dynamical systems, manifold learning, graph/information theory, explainable AI (xAI)  RNASeq and scRNASeq, eQTL, differential expression,  transformers, attention mechanisms, prompt engineering, few-shot learning, transfer learning, MLOps, CI/CD for ML, A/B testing, feature engineering, model monitoring, data versioning, reinforcement learning, GANs, recommendation systems
<b>Leadership and Business</b>	mentorship, project management, JIRA, Confluence, internal/external stakeholder communication, AI powered automation, ETL/data pipeline development, product benchmarking/development, cross-functional team leadership, technical documentation, cost-benefit analysis, ROI assessment, AI strategy development, responsible AI practices

## EXPERIENCE

---

<b>Scipher Medicine</b> <i>Data Scientist, Senior Data Scientist</i>	03/2021-Present <i>Boston, MA</i>
<ul style="list-style-type: none"><li>Used Bayesian inference, deep learning, and graph theoretic techniques in R and Python to infer causality/directionality for drug re-purposing, using single cell and bulk RNASeq machine learning techniques, pipelines, and data sources.</li><li>Ingested, organized, and streamlined clinical and genomic data/metadata from internal clinical trials and external EMR/EHR data sources (40% of all US RA patients), sharing these clean and GitHub version controlled data pipelines/ETLs to an AWS-native data lake, and piloting CI/CD and modern DevOps department wide for reproducible code and data collaboration.</li><li>Developed analyses, presentations, and materials for stakeholders to pursue data monetization and drug re-purposing partnerships using aforementioned EHR and genomic data sources and pipelines.</li><li>Developed a clinical endpoint simulation algorithm, improving ML outcomes with probabilistic labels, delivering conference abstracts and communicating with internal and external stakeholders on commercialization and medical/academic rheumatology impact.</li><li>Implemented and currently productionizing a SOTA DL model in Pytorch to predict drug response and non-response in RA, using fuzzy/probabilistic labels and bulk RNAseq + clinical data, MLOps, and model/data versioning.</li><li>Mentored three full-time summer interns and one MPH student practicum, designing and leading adoption of a formal mentorship framework in the DS department consisting of Python, git, SQL, AWS, statistics, and ML/AI.</li></ul>	
<b>Brandeis University</b> <i>Graduate Researcher, Teaching Assistant, and Postdoctoral Scholar</i>	09/2013 - 03/2021 <i>Boston, MA</i>
<ul style="list-style-type: none"><li>Collected and analyzed 1GB/min time series data to study neural interactions between the hippocampus and prefrontal cortex, two interconnected brain regions important for learning and decision making.</li><li>Designed multiple, multi-year long studies and projects, including interpreting literature for gaps and fit, designing hardware, software, and novel analytical and mathematical techniques in MATLAB and Python.</li><li>Used PCA, generalized linear models, unsupervised learning techniques, and Bayesian methods to decode brain cell responses and brain area communication, providing published new insights into representations of memory.</li><li>Mentored graduate and undergraduate students in analytical techniques; wrote and directed a yearly internal course on computer science, continuous and discrete data analysis, and common statistical methods.</li></ul>	

**Freelance Data Science Consulting**

10/2020 - 02/2021

*Neuroscience/Data Science Consultant for Wave Neurosciences**Boston, MA*

- Analyzed double-blind clinical trial data of veterans with PTSD, consisting of 84 21-channel EEGs x 3 longitudinal time points.
- Used supervised and unsupervised machine learning techniques, information theory, and graph theory for comparisons and longitudinal trends in functional connectivity between sham and neuromodulation groups in wide and narrow-band power and coherence.
- Contracted for 80hrs, with deliverables including study and statistical design, python code, hosted data, notebooks, visualizations, presentations, and a study report outlining analyses.

**Insight Data Science**

08/2019 - 01/2020

*Data Science Fellow**Boston, MA*

- Consulted with PyrAmes Inc. to identify, cluster, and clean movement artifacts from a wireless, non-invasive wearable device collecting continuous blood pressure diagnostics.
- Parsed over 100 hours of labeled and 1000 hours of unlabeled time series data, used spectral methods to engineer features and perform unsupervised clustering / blind signal source separation.
- Delivered well-documented code to PyrAmes Inc, a report on possible further optimization techniques, and a pipeline to implement the detection, cleaning, and clustering algorithm.

**University of Chicago Medical Center**

12/2011 - 09/2013

*Research Technologist- Pancreatic Islet Research Lab**Chicago, IL*

- Investigated human pancreatic cells and their changes due to Type 1/2 Diabetes, fetal and neonatal development, and other endocrine disorders, resulting in 8 published papers with physicians and scientists
- Collected and analyzed 50TB+ of image data, overhauled legacy image analysis library into MATLAB, created specimen analysis routines using automated image processing and computer vision techniques
- Sectioned human and animal tissues, prepared specimens, analyzed results, and presented findings to collaborating labs, physicians, scientists, guest lecturers, and visiting companies

**EDUCATION**

---

**Brandeis University**

2013 - 2020

Ph.D. in Neuroscience, Certificate in Quantitative Biology

**University of Chicago**

2007 - 2011

B.A. in Biology, Specialization in Neuroscience, Minor in Computational Neuroscience