

# Eric Alcaide Medicine & Physics, Machine Learning

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## Skills

<b>Multi-Language Coding</b> Python (and scientific stack pytorch, triton, cython, jax, etc), Web, JavaScript, Bash, C++	<b>Cloud &amp; HPC</b> Algorithmic optimization, CPU (1000+ cores) and GPU parallelism (1000+ A/H1/200s), server and cluster computing	<b>Machine Learning</b> LLMs, Geometric deep learning, computer vision, natural language, clustering, graphs, self-supervised learning, etc	<b>Teamwork, Public Speaking, Fast Prototyping, Problem Solving, Strategic Thinking, Operational Excellence, Complex Problems</b>
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## Education

Barcelona, Spain	<b>Medicine Degree, University of Barcelona</b> Medical Degree. Multiple distinctions.
Barcelona, Spain	<b>Physics Degree, University of Barcelona</b> Theoretical Physics Mention

## Professional / Research Experience

2024 – present Switzerland	<b>Founding Engineer, poolside</b> LLMs. (Pre)Training, Inference, ML operations. Distributed workloads on thousands of CPU/GPUs. Algoritgms and Execution. From core research to product.
2021 – 2024 London, United Kingdom	<b>Machine Learning &amp; Translational Scientist, CHARM Therapeutics</b> From Bits to Molecules, and Everything in between: data ingestion pipeline (1000s of CPUs), geometric deep learning research, model training (100s of GPUs), evaluation, drug target research, virtual screening, etc. First employee, architect of DragonFold
2021 – present	<b>Open Source Researcher, EleutherAI, OpenBioML</b> Research at the intersection of Natural Language Processing, Structural Biology and High Performance Computational Methods. Author of RWKV, an RNN Model for the Transformer Era (up to 14B models). RWKV versions 4, 5, 6, and 7. Diverse international research collaboration for the promotion of Open Source AI. Lead efforts of teams of 25+ researchers
2020 – 2021	<b>Machine Learning Researcher, VIR Biotechnology</b> Machine Learning for Structural Biology. Geometric Deep Learning and Natural Language Processing techniques for organic molecules, proteins and monoclonal Antibodies (mAbs).
2019 – 2021	<b>Private Machine Learning Tutor</b> Personalized advice to Masters' students from different backgrounds (from Computational Linguistics to Biomedical Engineering) on Text Classification, Image processing and Information representation. Advised how to carry out Masters' Thesis-level projects.
2019	<b>Non-Profit Health Hackathon Mentor, TV3 - La Marató</b> Advised on healthcare and science X algorithms, including genetic clustering and molecular dynamics for protein conformational changes. Won several prizes.

## Peer Reviewed Papers






2024	<b>Eagle and Finch: RWKV with Matrix-Valued States and Dynamic Recurrence, CoLM, Conference on Language Modelling</b> Next generation RWKV 5 & 6. Best scalable RNNs for their time. 0.1B - 14B models
2023	<b>RWKV: Reinventing RNNs for the Transformer Era, EMNLP</b> Rearrangement of typical RNNs compute graph to allow for GPT training. Trained multilingual Large Language Models from 0.1B to 14B parameters, with chat interface and Open Source release. Future developments sponsored by the Linux Foundation.
2023	<b>UMD-fit: Generating Realistic Ligand Conformations for Distance-Based Deep Docking Models, NeurIPS 2023 Generative AI In Biology Workshop</b> Addressed chemical inaccuracies in Deep Learning Molecular Docking models.

2023	<b>Advancing structural biology through breakthroughs in AI,</b> <i>Current Opinion in Structural Biology</i> Major recent advances driven by technology and applications to novel therapeutics.
2022	<b>InterDocker: End-to-End Cross-Attentive and Geometric Transformers for Efficient Iterative Protein Docking,</b> <i>LMRL-Learning Meaningful Representations of Life, NeurIPS2022</i>
2022	<b>Relevance of myocardial injury biomarkers to the prognosis of COVID-19 patients,</b> <i>Revista Española de Cardiología</i> COVID19 related revision of predictive power of myocardial injury biomarkers (NT-proBNP and hs-TnT) regarding Mechanical Ventilation and Death Events.
2021	<b>MP-NeRF: Massively Parallel Natural Extension of Reference Frame,</b> <i>Journal of Computational Chemistry</i> Parallelized the Natural Extension of Reference Frame for folding polymers (proteins, RNA, etc) from internal angles, 1000x faster. Usage in MD simulations and ML training.

## Courses

2020 – 2020 Barcelona, Spain	<b>HPC-based Computational Biomedicine,</b> <i>Barcelona Supercomputing Centre</i> Impact and Hands-on experience of applied supercomputing to biomedical problems (molecular simulations, genomic analysis, tissue modelling, etc.)
2018 – 2019	<b>Deep Learning, Natural Language Processing and AI for Medicine,</b> <i>Coursera</i> Contents include: foundations of Deep Learning, project management, Computer Vision, sequential data, Natural Language Processing, AI in healthcare, etc
2017 – 2017	<b>Artificial Intelligence Micromasters Program,</b> <i>Columbia University</i> CSMM.101x: Artificial Intelligence (AI) - (through edx.org). Average qualification: 8.1 / 10 Search methods, games, ML introduction, CSPs, NLP, robotics introduction, etc.

## Side Projects

2020 – present	<b>Open Source Contributions</b> Cutting-edge Open Source Software packages (Pytorch Geometric, Fastformers, etc)
2018 – present	<b>Open Source projects</b> Projects and modules for scientific computing which recieved a high degree of community acceptance: <ul style="list-style-type: none"> <li>• <b>2023: Uni-Mol</b>  : Accuracy improvements for ML-based molecular docking.</li> <li>• <b>2021: AlphaFold2 open replication</b>  : Main contributor to the Open Source effort for the replication (and improvement) of the AlphaFold2 architecture (state of the art, deep learning engine for protein structure prediction).</li> <li>• <b>2021: E(n) Equivariant GNN, Geometric Vector Perceptron:</b>  Graph Neural Network architectures on invariant representations in 3D or arbitrary dimensions.</li> <li>• <b>2019: AlphaFold1 imitation: MiniFold:</b>  Predict protein foldings from raw sequences</li> <li>• <b>2018: Keras-WRN:</b>  Wide Residual Networks for image recognition in Keras.</li> </ul>
2017	<b>Deep Learning - Can Computers Learn?</b> Research project focused on the AI and Deep Learning field, subfields and the state of the art techniques. Evolutionary Strategies for architecture optimization in Neural Networks.

## Languages

Spanish	English	German	Mandarin
Native	C2 level	B1 level	HSK 2

## Awards

2019	<b>ESADE - Accenture HealthHackathon Winnner Award,</b> <i>ESADE, Barcelona</i>
2019	<b>AlphaFold v1 Replication Contest Award, Nvidia Titan RTX,</b> <i>Nvidia</i>
2017	<b>Hackathon UPC Winner Award,</b> <i>HackUPC, Barcelona</i>