

Maximilian Lombardo

1.518.707.6805 | maximilian.g.lombardo@gmail.com | github.com/MaximilianLombardo | New York, NY

EXPERIENCE

Biohub (Chan Zuckerberg Initiative)

New York, NY / Remote

Technical Product Lead - AI Research and Cell Science

Nov 2020 – Present

- Defined product vision and technical schemas for **CELLxGENE (10K+ users)**, translating scientific requirements into specs. Partnered with engineering, design, ML/AI, and UX teams to deliver the visualization, data discovery, and API tools for AI models, embeddings, and data access, validating features via prototyping and UX research.
- Orchestrated release of **9 biological foundation models**, bridging AI Research and Engineering to productize inference. For VariantFormer, engineered the technical GTM suite (interactive model playgrounds and Remotion videos) **driving 2.6x increase in engagement rate** (48% vs 18%).
- Led 0-1 build of an automated annotation engine, delivering the model (VAE/Logistic Regression) and UI. Reduced curation time by **99.9%** (6 weeks to <1 min) with high accuracy (F_1 : **0.9**).
- Spearheaded evangelism strategy, developing briefings for the **Frontier Model Forum**. Co-organized severse (with **NVIDIA, TileDB**) and represented CZI at NeurIPS, establishing leadership in the AI x Bio research community.

Kallyope

New York, NY / Remote

Computational Scientist - ML/AI

Sep 2017 – Nov 2020

- Led computational analysis generating a single-cell atlas of a novel tissue, partnering with wet-lab scientists to characterize cell types via graph-based clustering. Delivered insights securing a **\$2M milestone** with Novo Nordisk.
- Engineered multi-omics ML pipelines that supported the company's growth from **Series A (pre-clinical)** to **Series D**, contributing to the identification of lead compounds for the **first human clinical trials**.
- Co-developed probabilistic ML models to map neuronal connections, integrating viral tracing with single-cell sequencing data to decode circuit architecture.
- Presented research progress to the Scientific Advisory Board (including **three Nobel Laureates**), synthesizing complex computational results into strategic updates.

IBM Corporation

Amsterdam, NL

Graduate Machine Learning Research Intern - Center For Advanced Studies

Oct 2016 – Aug 2017

- Implemented and benchmarked machine learning frameworks to identify disease-causing features from multi-omics data.
- Evaluated dimensionality reduction techniques (subspace projections) for visualizing single-cell patient data.

Columbia University Medical Center

New York, NY

Senior Technician - Experimental Therapeutics for Lymphoma

Jul 2013 – Aug 2015

- Uncovered drug resistance via RNA-seq, resulting in **four co-authored publications** (Blood, Clinical Cancer Research).

EDUCATION

University of Amsterdam

Amsterdam, NL

Master of Science in Computational Science

Class of 2018

Columbia University in the City of New York

New York, NY

Master of Arts in Biotechnology

Class of 2014

Siena College

Albany, NY

Bachelor of Science in Biology, Cum Laude

Class of 2012

SKILLS

Languages & AI: Python, PyTorch, SQL, JavaScript, React, RAG, AI Agents, Model Evals, LLMs

Product & Tools: Product Strategy, GTM, Claude Code, AWS, Figma, Marimo, Remotion, Git, User Research

VENTURE PROJECTS

Stealth Biotech Venture Concept (Founder): Defined product vision for an AI-native "Operating System for Biology," leveraging biological foundation models. Recruited early collaborators and pitched the comprehensive SaaS roadmap to venture capital firms.

Bay Bridge Bio Venture Exploration Project: Developed venture concept and MVP for a single-cell analysis SaaS product for the regenerative medicine space, solicited KOL feedback, and pitched to investors from Andreessen Horowitz, Leaps by Bayer, and J&J Innovation.