

# Michelle Lynn Gill, PhD

*Applied Research Manager, NVIDIA Virtual Cell Team*

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GitHub | LinkedIn | Personal Website

## Overview

I am an Applied Research Manager on NVIDIA's Virtual Cell team. Our team uses data, deep learning, and HPC to develop models that simulate cellular states. My focus is on the benchmarks and development of supporting tools that help evaluate and improve these models.

Previously I was a Senior Machine Learning Engineer and Data Scientist at BenevolentAI and a member of Arthur Palmer's research group where I developed and applied nuclear magnetic resonance (NMR) spin relaxation experiments to understand how enzyme dynamics are critical to biological function.

## Education

### **Ph.D., Molecular Biophysics & Biochemistry**

2003-2006

Yale University, New Haven, CT

Thesis: Development of  $^{205}\text{Ti}$  NMR methods for the direct study of monovalent metal ions and ligands in nucleic acids

Advisors: J. Patrick Loria and Scott Strobel

### **M.Phil., Molecular Biophysics & Biochemistry**

2001-2003

Yale University, New Haven, CT

### **B.S., Biochemistry**

1997-2001

University of Kansas, Lawrence, KS

Highest Distinction and Honors (Summa Cum Laude)

## Experience

### **Applied Research Manager, NVIDIA, Virtual Cell Team**

2022-Present

Lead benchmarking efforts for virtual cell models, including internal and external collaborations

Developed benchmarking framework and tooling for model assessment and reproducibility

Collaboration with Chan Zuckerberg Initiative on cz-benchmarks for evaluating virtual cell models

### **Scientific Lead, BioNeMo, NVIDIA**

2019-2023

R&D Manager and Scientific Lead for Clara Discovery, NVIDIA's platform for accelerating drug discovery

Developed BioNeMo for pre-training and fine tuning of large language models for cheminformatics and proteomics

|                                                                                                                                                                                              |           |
|----------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|-----------|
| <b>Senior AI and Deep Learning Scientist, NVIDIA</b>                                                                                                                                         | 2019-2023 |
| <p>Focused on proteomics including deep learning models to predict peptide spectral matches with &gt;95% F1</p> <p>Led team using GCNNs to predict molecular properties</p>                  |           |
| <b>Senior Data Scientist and Machine Learning Engineer, BenevolentAI</b>                                                                                                                     | 2018-2019 |
| <p>Matrix factorization and GCNNs for drug mechanism importance in knowledge graphs</p> <p>3D CNNs for ligand pose and affinity prediction</p>                                               |           |
| <b>Senior Deep Learning Consultant, NVIDIA</b>                                                                                                                                               | 2017-2018 |
| <p>Assisted clients in pharmaceutical and materials science in utilizing deep learning</p>                                                                                                   |           |
| <b>Scientist, National Cancer Institute, NIH</b>                                                                                                                                             | 2014-2016 |
| <p>Developed parallelized, compressed sensing methods for NMR data reconstruction</p>                                                                                                        |           |
| <b>Postdoctoral Research Fellow, Columbia University, Department of Biochemistry and Molecular Biophysics</b>                                                                                | 2008-2014 |
| <p>Demonstrated conformational selection in DNA methyltransferase AlkB</p> <p>Developed multiple quantum NMR spin relaxation experiments</p> <p>Advisor: Professor Arthur G. Palmer, III</p> |           |

## Publications

S. Dicks, L. Heumos, S. Jimenez, P. Angerer, I. Gold, I. Virshup, F. Fischer, L. May, C. J. Nolet, M. Gill, M. Boerries, and F. Theis **"Accelerating single-cell analysis with GPU-enabled RAPIDS-singlecell"** *In preparation* (2025).

E. Sevgen, J. Moller, A. Lange, J. Parker, S. Quigley, J. Mayer, P. Srivastava, S. Gayatri, D. Hosfield, M. Korshunova, M. Livne, M. Gill, R. Ranganathan, A.B. Costa, and A.L. Ferguson **"ProT-VAE: Protein transformer variational autoencoder for functional protein design"** *Proceedings of the National Academy of Sciences* (2025).

P. St. John, D. Lin, P. Binder, M. Greaves, V. Shah, J. St. John, A. Lange, P. Hsu, R. Illango, A. Ramanathan, A. Anandkumar, D.H. Brookes, A. Busia, A. Mahajan, S. Malina, N. Prasad, S. Sinai, L. Edwards, T. Gaudet, C. Regep, M. Steinegger, B. Rost, A. Brace, K. Hippe, L. Naef, K. Kamata, G. Armstrong, K. Boyd, Z. Cao, H.Y. Chou, S. Chu, A. dos Santos Costa, S. Darabi, E. Dawson, K. Didi, C. Fu, M. Geiger, M. Gill, D.J. Hsu, G. Kaushik, M. Korshunova, S. Kothén-Hill, Y. Lee, M. Liu, M. Livne, Z. McClure, J. Mitchell, A. Moradzadeh, O. Mosafi, Y. Nashed, S. Paliwal, Y. Peng, S. Rabhi, F. Ramezanghorbani, D. Reidenbach, C. Ricketts, B.C. Roland, K. Shah, T. Shimko, H. Sirelkhatim, S. Srinivasan, A.C. Stern, D. Toczydlowska, S.P. Veccham, N.A.E. Venanzi, A. Vorontsov, J. Wilber, I. Wilkinson, W.J. Wong, E. Xue,

C. Ye, X. Yu, Y. Zhang, G. Zhou, B. Zandstein, A. Chacon, P. Sohani, M. Stadler, C. Hundt, F. Zhu, C. Dallago, B. Trentini, E. Kucukbenli, T. Rvachov, E. Calleja, J. Israeli, H. Clifford, R. Haukioja, N. Haemel, K. Tretina, N. Tadimetri, and A.B. Costa  
**"BioNeMo Framework: a modular, high-performance library for AI model development in drug discovery"** *arXiv* (2024).

M.T. Strauss, I. Bludau, W.F. Zeng, E. Voytik, C. Ammar, J. Schessner, R. Illango, M.L. Gill, F. Meier, S. Willems, and M. Mann  
**"AlphaPept, a modern and open framework for MS-based proteomics"** *Nature Communications* (2024).

D. Reidenbach, M. Livne, R.K. Illango, M.L. Gill, and J.I. Israeli  
**"Improving small molecule generation using mutual information machine"** *arXiv* (2022).

M.L. Gill, A. Hsu, and A.G. Palmer **"Detection of chemical exchange in methyl groups of macromolecules"** *Journal of Biomolecular NMR* (2019).

M.L. Gill, R.A. Byrd, and A.G. Palmer **"Dynamics of GCN4 facilitate DNA interaction: a model-free analysis of an intrinsically disordered region"** *Physical Chemistry and Chemical Physics* (2016).

S. Sun, M.L. Gill, Y. Li, M. Huang, and R.A. Byrd **"Efficient and generalized processing of multidimensional NUS NMR data: the NESTA algorithm and comparison of regularization terms"** *Journal of Biomolecular NMR* (2015).

B. Ergel, M.L. Gill, L. Brown, B. Yu, A.G. Palmer, and J.F. Hunt  
**"Protein dynamics control the progression and efficiency of the catalytic reaction cycle of AlkB"** *Journal of Biological Chemistry* (2014).

M.L. Gill, and A.G. Palmer **"Local isotropic diffusion approximation for coupled internal and overall molecular motions in NMR spin relaxation"** *Journal of Physical Chemistry B* (2014).

M.L. Gill, and A.G. Palmer **"Multiplet-filtered and gradient-selected zero-quantum TROSY experiments for <sup>13</sup>C<sup>1</sup>H<sub>3</sub> methyl groups in proteins"** *Journal of Biomolecular NMR* (2011).

## Patents

### **Automated FEP Path Generation and Optimization by Molecule Foundation Models and Generative AI**

2025

Peng, Y., Kucukbenli, E., Zhou, G., Gill, M.L., Livne, M., Korshunova, M., Rvachov, T., Israeli, Y.

US Patent Application No. 19/208,119; Filing Date: 2025/05/14

### **Extraction of Informative Embeddings from Encoder-Decoder Models**

2024

Livne, M., Gill, M.L.

US Patent Application No. 18/957,301; Filing Date: 2024/11/22

**Contrastive Framework for Unified Generative and Discriminative Representation Learning** 2024

Livne, M., [Gill, M.L.](#)

US Patent Application No. 18/957,294; Filing Date: 2024/11/22

**Efficient Data Loading for Deep Learning Workloads** 2024

Darabi, S., Korshunova, M., Grewal, J., [Gill, M.L.](#), Morkisz, P.

US Patent Application No. 18/885,428; Filing Date: 2024/09/13

**Guardrails for Molecular Generation** 2024

Korshunova, M., Peng, Y., Zhou, G., Rvachov, T., [Gill, M.L.](#), Kucukbenli, E., Israeli, Y.

US Patent Application No. 18/807,805; Filing Date: 2024/08/16

**End to End Deep Learning Workflow for *In Silico* Molecule Design** 2024

Stern, A., [Gill, M.L.](#), Stepniewska-Dziubinska, M., Grzegorzec, T., Nowaczynski, A., Toczydlowska, D., Israeli, Y., Ribalta Lorenzo, P.

US Patent Application No. 18/412,168; Filing Date: 2024/01/12

**Small Molecule Generation Using Machine Learning Models** 2023

Livne, M., Reidenbach, D., [Gill, M.L.](#), Ilango, R., Israeli, Y.

US Patent Application No. 18/450,745; Filing Date: 2023/08/16

**Presentations**

**Scientific Discovery: From the Lab Bench to the GPU, *Andy Byrd Retirement Symposium*** 2024

Invited Keynote, April 19, 2024, Institute for Bioscience & Biotechnology Research, University of Maryland, NIST, Bethesda, MD

[Slides](#) · [Program](#)

**Scientific Discovery: From the Lab Bench to the GPU, *PyData NYC*** 2023

Invited Keynote, November 2, 2023, New York, NY

[Slides](#) · [Video](#) · [Abstract](#)

**NVIDIA BioNeMo: A framework and service for development and use of generative AI in drug discovery, *6th RSC-BMCS / RSC-CICAG Artificial Intelligence in Chemistry Conference*** 2023

Invited Keynote, September 5, 2023, Cambridge, UK

[Slides](#) · [Program](#)

**Exploring Molecular Space and Accelerating Drug Discovery on the GPU with Clara Discovery, *Gates Foundation Grand Challenges: Applications of Artificial Intelligence in Machine Learning*** 2021

Invited Talk, November 10, 2021, Virtual

**Accelerating Drug Discovery with Clara Discovery and MegaMolBART, *4th RSC-BMCS / RSC-CICAG Artificial Intelligence in Chemistry*** 2021

Invited Talk, September 27, 2021, Virtual

[Slides](#)

**Real Time, GPU-Accelerated Analysis and Visualization in the Life Sciences**, *Ken Kennedy Institute Data Science Conference*

2020

[Michelle Gill](#) and Avantika Lal

Invited Keynote, October 26-27, 2020, Virtual

[Slides](#) · [Abstract](#) · [Program](#)

**Artificial intelligence driven drug discovery**, *NYC R Conference*

2019

Invited Presentation, May 10, 2019, New York, NY

[Slides](#)

**Machine learning for target identification and lead optimization in drug discovery**, *New York Area Group for Informatics and Modeling*

2019

Alix Lacoste and [Michelle Gill](#)

Invited Presentation, February 26, 2019, New York, NY

[Abstract](#)

**Accelerating the journey from data to medicine**, *NeurIPS*

2018

Amir Saffari, Dan Neil, Alix Lacoste, and [Michelle Gill](#)

Expo Talk, 2018, Montreal, Canada

[Abstract](#)

**Artificial intelligence as a catalyst for scientific discovery**, *JupyterCon*

2018

Invited Keynote, 2018, New York, NY

[Slides](#) · [Video](#) · [Abstract](#)

**From structural biology to AI: a holistic approach to studying molecular machines**, *Brookhaven National Laboratory*

2018

Invited Presentation, 2018, Upton, NY

[Slides](#)

**Efficient image search and identification: the making of Wine-O.AI**, *SciPy Conference*

2017

Selected Presentation, 2017, Austin, TX

[Slides](#) · [Video](#) · [Code](#)

**Development of  $^{205}\text{TI}$  NMR methods for the direct study of monovalent metal ions and ligands in nucleic acids**, *Ph.D.*

2006

*Thesis Defense, Yale University*

Thesis Defense, 2006, New Haven, CT

[Slides](#) · [Thesis](#)

## Awards

NIH Postdoctoral Research Fellowship (F32 GM089047)

2009-2012

Global BCG Strategy Olympics, Winning Team

2007

NSF Graduate Research Fellowship

2002-2006

|         |                                                                                        |           |
|---------|----------------------------------------------------------------------------------------|-----------|
|         | Barry M. Goldwater Scholar                                                             | 2000-2001 |
|         | Outstanding Undergraduate Honors Research Thesis                                       | 2001      |
|         | Kansas Board of Regents Full Tuition Merit Scholarship                                 | 1997-2001 |
| Service |                                                                                        |           |
|         | <b>Judge, Preliminary and Final Rounds</b> , NYC STEM Fair, New York, NY               | 2022      |
|         | Evaluated submissions in the biochemistry track                                        |           |
|         | <b>Program Chair</b> , PyData NYC, New York, NY                                        | 2018      |
|         | Responsible for conference content, proposal review, speaker selection, and scheduling |           |
|         | <b>Machine Learning Symposium Co-Chair</b> , SciPy Conference, Austin, TX              | 2018      |
|         | Co-chair of machine learning / deep learning symposium                                 |           |