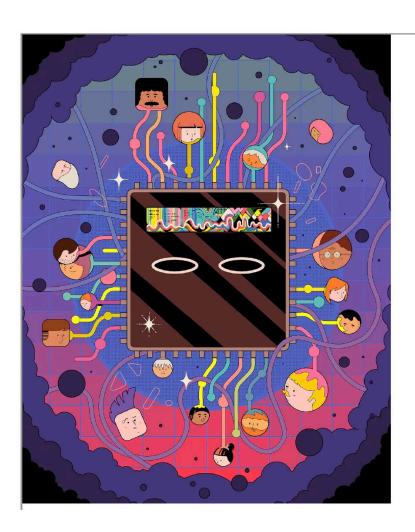
Teaching Machines to Doubt: Building Humble and Curious AI for Healthcare

Leo Anthony Celi (we/us)
MIT Critical Data







Opinion

AI isn't smart enough to destroy us

The real threats Al poses will come from the humans who wield it.

Today at 6:30 a.m. EDT











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Article Published: 26 February 2024

scGPT: toward building a foundation model for single-cell multi-omics using generative AI

Haotian Cui, Chloe Wang, Hassaan Maan, Kuan Pang, Fengning Luo, Nan Duan & Bo Wang ⊠

Nature Methods 21, 1470–1480 (2024) | Cite this article







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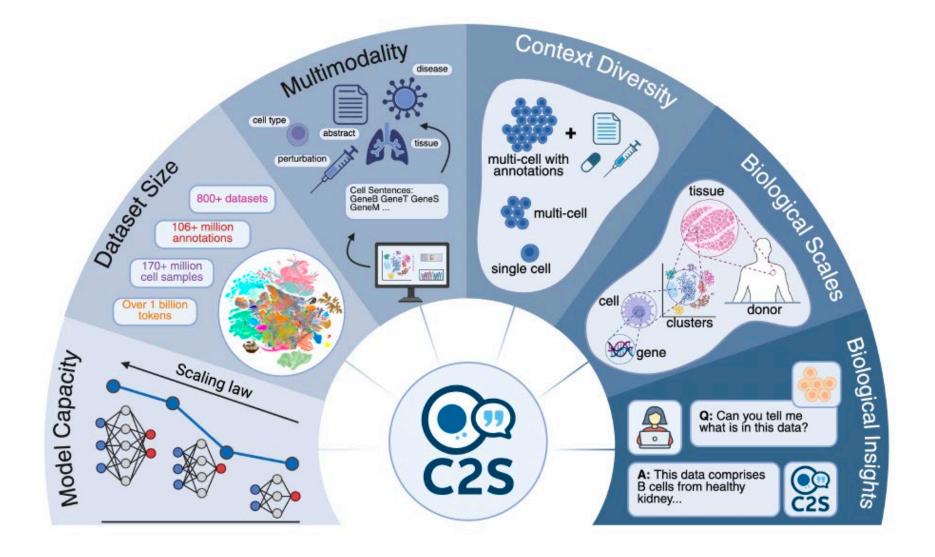
Scaling Large Language Models for Next-Generation Single-Cell Analysis

© Syed Asad Rizvi, Daniel Levine, Aakash Patel, Shiyang Zhang, Eric Wang, Curtis Jamison Perry, Nicole Mayerli Constante, Sizhuang He, David Zhang, Cerise Tang, Zhuoyang Lyu, Rayyan Darji, Chang Li, Emily Sun, David Jeong, Lawrence Zhao, Jennifer Kwan, David Braun, Brian Hafler, © Hattie Chung, Rahul M. Dhodapkar, Bryan Perozzi, Jeffrey Ishizuka, Shekoofeh Azizi, © David van Dijk doi: https://doi.org/10.1101/2025.04.14.648850















ARTICLE · Online now, September 26, 2025

Single-cell nascent transcription reveals sparse genome usage and plasticity

Shaoqian Ma <a>^ Ma · Yantao Hong · Junhan Chen · Jingzhao Xu · Xiaohua Shen <a>^ 2 Ma

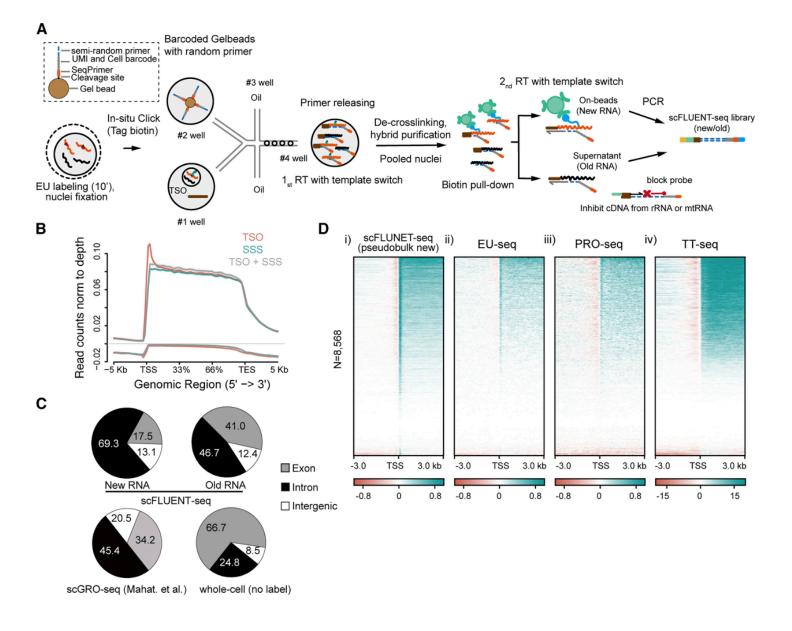




Sparse, Stochastic Single-cell Nascent Transcription Newly synthesized RNA Old nuclear RNA 10-min EU labeling scFLUENT-seq Single-nucleus barcoding Consistent activity Fluctuated activity in euchromatin in heterochromatin Cell 1 **Developmental Plasticity Transcription Diversity** Metastable mRNA/ncRNA state CytoTRACE Pseudo-time **Cell-state Progression**

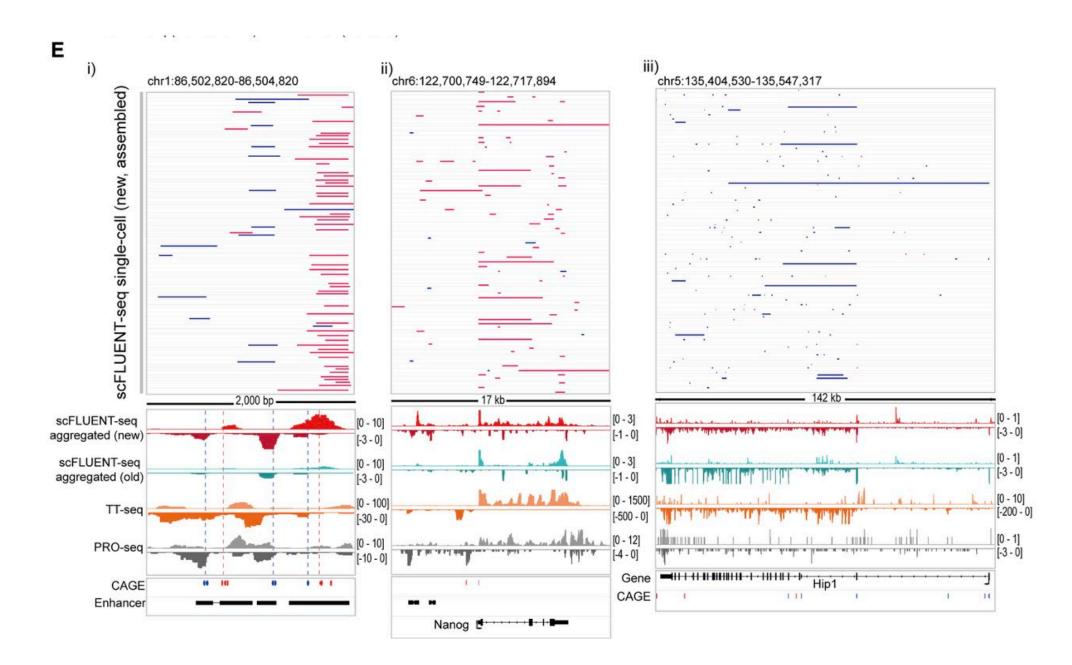








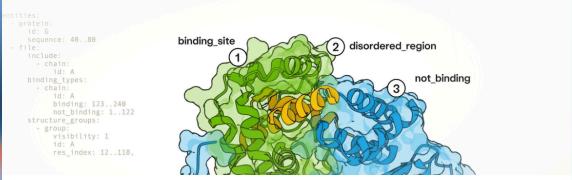












Hi,

As noted on Wednesday during MoML 2025 @ MIT, it's our pleasure to announce the official release of BoltzGen.

BoltzGen is a new, generative model for designing proteins and peptides of any modality that can bind to a wide range of biomolecular targets.

As with Boltz-1 and Boltz-2, BoltzGen is fully open source under the MIT license for unrestricted commercial and academic use, meaning it can be freely used for real-world problems and applications.





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Short Report Open access Published: 18 April 2025

Zero-shot evaluation reveals limitations of single-cell foundation models

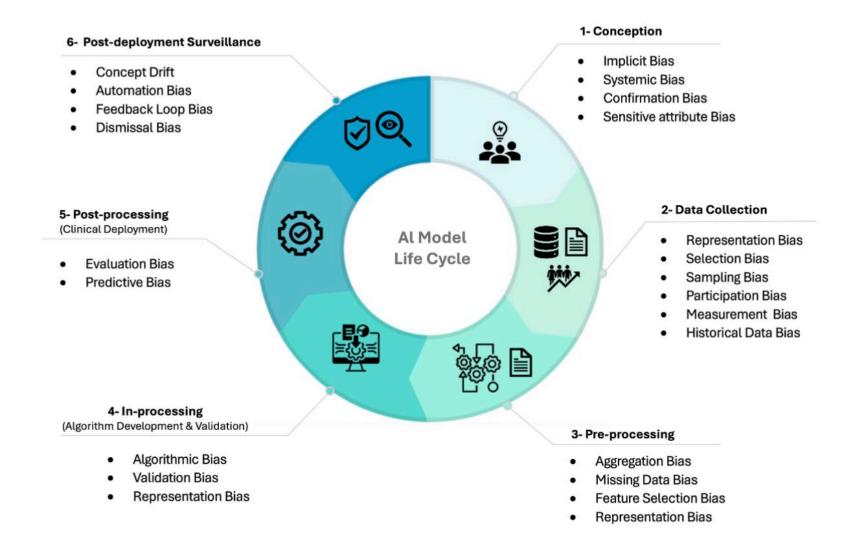
Kasia Z. Kedzierska, Lorin Crawford, Ava P. Amini & Alex X. Lu ⊠

Genome Biology 26, Article number: 101 (2025) Cite this article





Bias Pervades Every Level of Omics Research







The Solution - Integrated Mitigation Strategies

- Harmonize and standardize
 - instrumentation protocols
 - data collection protocols
 - analytical workflows across platforms
- Transparency is key
 - Open proprietary systems
 - Document all preprocessing decisions
 - Ensure reproducible computational pipelines





The Solution - Integrated Mitigation Strategies

- Equitable infrastructure
- Rigorous validation
- Robust governance frameworks





Al's Three Horizons: Retrieval (Current Focus)

- Automates access to existing digitized knowledge
- Democratizes information for larger groups—but not universally
- Success metric: Equity of access, benchmark scores





Al's Three Horizons: Recombination (Emerging)

- Fuses siloed, non-digitized, non-English knowledge into new hypotheses
- Connects religious texts, indigenous wisdom, multilingual sources previously inaccessible
- Success metric: Recombination yield—number of testable cross-domain hypotheses generated





Al's Three Horizons: Discovery (The Promise)

- Co-creates new knowledge from weak signals and rare data about underserved populations
- Solves problems that have evaded human brains by interfacing previously disconnected knowledge
- Success metric: Discovery lift—impact on underserved, multilingual, multimodal communities





