

# Jeremy Li

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Summary	I am an accomplished researcher and engineer with extensive experience in fast-paced startup environments, leading and rapidly scaling data/research teams, and interfacing with executive, engineering, and product teams. I have a strong technical background in statistical modeling, machine learning, computational genomics, and software engineering.	
Skills	<b>Languages:</b> Python, R, bash, some C++, Rust <b>Tools:</b> Docker, AWS (EC2, S3, ECR, Batch), git, GNU/Linux, cluster computing	
Experience	<b>Consultant, Research &amp; Engineering</b> Stealth biotech startup, USA	12/2024 – present
	<ul style="list-style-type: none"><li>• Consulting work for a seed-stage biotech startup, engaged to optimize a class of highly specialized machine learning models for genomic data</li><li>• Unifying disparate multi-TB genomic datasets comprising &gt;500k whole genome sequences</li></ul>	
	<b>Director of Research</b> <b>Head of Research</b> <a href="#">Gencove</a> , New York, NY	02/2022 – 12/2024 10/2021 – 02/2022
	<ul style="list-style-type: none"><li>• Oversaw company-wide ML and R&amp;D strategy, scaling a team from 1 to 7 PhD-level researchers in under 12 months to develop novel computational genomics solutions at scale</li><li>• Spearheaded deep learning initiatives (PyTorch) to replace expert-driven models, reducing manual overhead by 30%+ while accelerating analysis pipelines</li><li>• Collaborated with product and engineering teams to build proof-of-concept solutions, resulting in multiple \$100k-\$1M partnerships</li><li>• Alongside engineering, designed and maintained genomics pipelines processing petabytes of genomic data in a high-throughput AWS environment, decreasing compute costs by &gt;25%</li></ul>	
	<b>Senior Research Engineer</b> <b>Research Scientist</b> (employee # 5) <a href="#">Gencove</a> , New York, NY	03/2021 – 10/2021 02/2019 – 02/2021
	<ul style="list-style-type: none"><li>• Developed and implemented novel statistical and computational methods for sparse NGS data, enabling POC demonstrations that helped secure 15+ enterprise deals</li><li>• Collaborated with engineering teams to build out the initial production workflows for genomic data, achieving stable parallel processing of multi-TB datasets</li><li>• Led massive-scale biobank analyses, designing novel pipelines and methods to handle 100s of thousands of whole-genome sequences to extract biological insights</li><li>• Authored three first-author publications and presented at major genomics conferences</li></ul>	
	<b>Lead Research Scientist</b> (employee # 4) <a href="#">Genomic Prediction</a> , North Brunswick, NJ	07/2017 – 02/2019
	<ul style="list-style-type: none"><li>• Led 3-person team to develop and deploy the cloud-based NGS analysis pipeline underpinning the first revenue-generating product of the company</li></ul>	
Publications	Published <a href="#">14 research articles</a> ( <i>h</i> -index: 5), presented findings at major genomics conferences	
Education	<b>University of Washington</b> for a B.S. in Physics and B.A. in Chemistry	2013 – 2017