

# Saksham Malhotra

+91 9899338414 | [saksham2196@gmail.com](mailto:saksham2196@gmail.com) | [linkedin.com/in/saksham219/](https://linkedin.com/in/saksham219/) | [github.com/saksham219](https://github.com/saksham219) | [saksham219.github.io](https://saksham219.github.io)

Data Scientist with 5+ years of experience building large-scale forecasting, optimization, and machine learning systems in production.

## EXPERIENCE

<b>PriceLabs</b> <i>Senior Data Scientist</i>	Jan 2025 – Present
	<i>Remote</i>
<ul style="list-style-type: none"><li>Developing a market-demand forecasting model for 500k+ short-term rental properties, integrating pricing, booking, and event signals to drive demand-elasticity-based revenue optimization.</li><li>Designed and deployed a dynamic pricing pipeline for 25k mid-term rental properties.</li><li>Built a compliance-aware pricing framework for properties affected by Los Angeles fire regulations.</li><li>Mentoring a Data Scientist on the forecasting framework, including ongoing maintenance and improvements.</li></ul>	
<b>PriceLabs</b> <i>Data Scientist</i>	March 2024 – Dec 2024
	<i>Remote</i>
<ul style="list-style-type: none"><li>Optimized forecasting pipelines through vectorization, reducing runtime by 60%.</li><li>Built an experimentation framework to evaluate pricing algorithm changes across 500k+ properties.</li><li>Led LLM-based real-time event detection, including prompt design and evaluation to identify novel market events.</li></ul>	
<b>Elucidata</b> <i>Data Scientist</i>	June 2018 – June 2021
	<i>New Delhi, India</i>
<ul style="list-style-type: none"><li>Applied statistical analysis (differential gene expression, dimensionality reduction, clustering) on large-scale RNA-seq datasets to support research labs and therapeutic companies in disease mechanism and drug target discovery.</li><li>Built an NLP-based ML pipeline classifying control vs. perturbation samples using metadata from 90k+ GEO samples, enabling cross-dataset differential gene discovery.</li><li>Developed Python/R data pipelines transforming public RNA-seq repositories into 100k+ FAIR-compliant ML-ready datasets for the Polly platform; packaged pipelines with Docker for AWS EC2 deployment.</li><li>Built a parallelized Python pipeline for single-cell BAM processing and automated RNA count report generation.</li></ul>	

## EDUCATION

<b>Master of Science, Applied Mathematics</b> <i>Technical University of Munich</i>	Oct 2021 – Oct 2023
	<i>Munich, Germany</i>
<ul style="list-style-type: none"><li>Relevant coursework: Probability theory, Non-linear optimization, Machine learning, Introduction to deep learning, Deep generative models, Computer vision, Case studies in scientific computing.</li><li>Thesis– Fast Eigensolvers for Koopman Operator Approximation: Developing an eigensolver algorithm that utilises the algebraic structure of Koopman eigenfunctions.</li></ul>	
<b>Bachelor of Engineering, Electronics and Communication</b> <i>Maharaja Surajmal Institute of Technology, GGSIPU</i>	April 2014 – April 2018
	<i>New Delhi, India</i>

## PROJECTS

- Optimal race line for autonomous race car:** Solved an optimal control problem for autonomous racing path planning using IPOPT.
- Training ML models:** Trained CNNs, VAEs, and diffusion models in PyTorch for generative and vision tasks.
- cmapPy:** Open-source contributor to a Python package for parsing GCT biological data files.
- Blog:** Mathematics blog with articles on probability, optimization, and applied ML.

## TECHNICAL SKILLS

- Programming:** Python, R, Javascript, C++, Matlab, Git, Docker, SQL
- ML/DS frameworks:** Pytorch, Tensorflow, Scikit-learn
- Data/Scientific Computing:** Numpy, Pandas, Polars