Owais Siddiqi

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Profile

I am a data scientist specialising in bioinformatics, data analytics, and multi-omics, with proven expertise in statistical modelling, clinical data analytics, and large-scale multi-omic dataset analysis. My experience includes developing and optimising automated bioinformatics workflows, integrating AIdriven analytics and computational biology to enhance accuracy, efficiency, and reproducibility in biomarker discovery and precision oncology. I excel in collaborative, cross-disciplinary teams and continuously advance my skills to remain at the cutting edge of genomics research and AI-driven healthcare innovation.

Education

Imperial College London

Oct. 2022 - Oct. 2023

MSc in Biomedical Engineering

- Developed expertise in Reinforcement Learning (DQN, PPO, SAC), Bayesian Probability, and Markov Decision Processes.
- Applied deep learning models for image classification, object detection, and segmentation.
- Conducted advanced time series analysis, neural decoding, and machine learning for brainmachine interfaces.

Queen Mary University of London

Sept. 2019 - Sept. 2022

BEng (Hons) in **Biomedical Engineering** – 1st Class

- Mastered calculus, differential equations, eigenvalues, vectors, and Gaussian elimination for engineering applications.
- Strengthened skills in multivariable calculus, linear algebra, and vector analysis for biomedical signal processing.

Work Experience

ANGLE plc.

Jan. 2025 - Present

R&D Data Analyst

- Clinical and Bioinformatics Data Management: Analyse large-scale imaging and multi-omic datasets for biomarker discovery in liquid biopsy and circulating tumour cell (CTC) applications. Implement automated, reproducible bioinformatics pipelines for sequencing data analysis, enhancing scalability and consistency.
- Workflow Automation and Optimisation: Develop Python and Bash scripts automating data extraction, transformation, analysis, and reporting, significantly reducing manual workload.
- Statistical and Computational Analysis: Apply statistical methods and machine learning approaches to validate experimental data, optimise research design, and enhance biomarker selection for precision oncology.
- Cross-Functional Collaboration: Work closely with scientists and IT teams to integrate data solutions and ensure regulatory compliance with ISO 13485:2016+A11:2021 standards.
- **Precision Oncology Advancement:** Leverage multi-omic technologies to advance biomarker identification, accelerate drug development, and improve clinical decision-making.

Junior Genomic Data Scientist

- Cancer Biomarker Prediction: Developed AI models using PyTorch for neoantigen biomarker prediction, facilitating personalised vaccine production for cancer immunotherapy.
- Bespoke VCF Production: Designed a specialised pipeline converting raw FASTQ data into customised VCF formats, enhancing downstream genetic analyses of tumour-specific biomarkers.
- Bioinformatics and Computational Tools: Processed RNA-seq data and performed variant calling using Docker containers and managed Linux-based virtual machines on Google Cloud Platform.
- Optimised Data Processing: Accelerated genetic downstream analysis of neoantigen biomarkers from one week to a few days.
- Continuous Learning: Regularly participated in workshops and online courses to stay current in genomics, AI, and bioinformatics advancements.

PDUK Ltd. Aug. 2023 – Jan. 2024

Data Scientist Intern

- Data Engineering: Utilised SQL queries in MySQL for data cleaning and management, reducing data processing time by 25% across 3000+ customer records.
- Data Visualisation: Created visualisations using Pandas, Matplotlib, and Seaborn, improving insight and decision-making through clear, data-driven reports.
- **Predictive Modelling:** Leveraged recurrent neural network (RNN) models in PyTorch for predictive time-series forecasting, improving accuracy by 20% and reducing inventory costs by 10%.
- Customer Segmentation: Implemented K-means clustering for customer segmentation, resulting in a 15% increase in targeted marketing conversion rates.

Imperial College London (MSc Project)

Mar. 2023 - Sept. 2023

AI Researcher (Deep Learning)

- Bespoke Model Development: Created a novel regression AI model using microscopy data for cell differentiation, achieving over 80% accuracy.
- Data Management and Augmentation: Implemented advanced data preprocessing and augmentation methods, improving model generalisation by 30% and improved predictive accuracy.
- Tool Integration: Integrated Pandas, Matplotlib, OpenCV, TensorFlow, Keras, and NumPy, optimising data preprocessing, parameter extraction, and model implementation, ensuring an efficient and effective workflow.

Technical Skills

- Programming: Python, MySQL, Bash, MATLAB
- Machine Learning: TensorFlow, PyTorch, Scikit-learn, Keras
- Bioinformatics: Genomics, Transcriptomics, FASTQ-VCF Pipelines, Nextflow
- Cloud/DevOps: Google Cloud, Docker, Git, CI/CD, VMs, Linux
- Data Analysis: Statistics, Statistical Modelling, Visualisations

Publications

Siddiqi, O., Winfield, J., Sormpas-Petridis, K., Harris, E., Ramkumar, A., Candito, A., Görner, S., Messiou, C., Blackledge, M., & Thrussell, I. (2022). *Investigating the Repeatability of Multifrequency Magnetic Resonance Elastography applied to a Soft Gelatine Phantom*. Presented at the ISMRM 31st Conference & Exhibition, London, 2022.