

#### **CONTACT**

**Q** Ludwigsfelde, Germany

+4915216878666

in www.linkedin.com/in/fmachani

https://machanig.github.io/

### **TECHNICAL SKILLS**

**Programming Languages** 

- Python
- SQL (PostresSQL, MySQL)
- R

# Data Analysis and Manipulation

 Pandas, Numpy, Statsmodel, Scipy, Tidyverse

### **Data Visualization Tools**

• Matplotlib, Seaborn, ggplot2, Power BI

### Machine Learning

- ML Libraries and Frameworks
  - TensorFlow, Scikit-Learn, XGBoost, MLextend, LightGBM
- Supervised Learning
  - Regression & Classification
  - Anomaly Detection
- Unsupervised Learning
  - Clustering (K-means, Hierarchical, K-medoids)
  - Dimensionality Reduction (PCA, t-SNE, PCoA)

# FRIDAH GECHEMBA MACHANI

# **DATA SCIENTIST**

#### **SUMMARY**

A data-driven professional with experience in using advanced analytical methods to uncover hidden patterns, and transform complex, high-dimensional datasets, into actionable strategic insights. I am skilled in predictive modeling, machine learning, and advanced analytics, and eager to create impactful solutions using Python, R, SQL, Power BI, and modern ML libraries and frameworks.

#### WORK EXPERIENCE

# Additional Training / Full-time Parenting Mar 2024 – April 2025

• I deepened my data science and machine learning expertise, learned querying databases with SQL, and Power BI for data visualization.

# Research Assistant Aug 2020 – Feb 2024 Max Planck Institute MPIMP – Golm, Germany

- Developed and calibrated regression models (R<sup>2</sup> = 0.93) in R to estimate nitrogen concentrations and enzymatic activity from known dilutions; deployed models and made predictions for over 500 samples, demonstrating end-to-end ML model development.
- Implemented multifactor ANOVA models with interactions in R to quantify the effect of mutations and nutrient availability on multiple plant features, showcasing experimental design and statistical modeling.
- Conducted unsupervised learning using K-means, PCA, and PERMANOVA to cluster multivariate metabolomics data (66 features); uncovered tissue-specific and nutrient-specific profiles.
- Utilized threshold-based outlier detection to identify anomalous gene expression profiles in 91K+ high-dimensional Affymetrix data and validated anomalies through wetlab quantification.
- Automated pre-processing pipelines in R to transform raw outputs from qPCR, GCMS, and spectrophotometer systems into structured datasets; by applying custom normalization, log transformation and feature extraction, streamlining data analysis workflows.

# Research Assistant Oct 2019 – May 2020 Universität Münster IEB – Münster, Germany

• Applied time-series analysis to determine optimal exudate collection time for maximizing seed germination rates.

# Lecturer Aug 2018 – May 2019 Machakos University – Machakos, Kenya

• Designed and delivered interdisciplinary courses to students from 3 faculties, adapting content to the audience needs and feedback.

- Ensemble Learning
  - Bagging, Voting
  - Boosting, Stacking

### Statistical Analysis

- Descriptive statistics
- Inferential statistics
- Hypothesis testing
- Diagnostic and Prescriptive analysis
- Predictive analytics
- Causal analytics

#### Version Control

• Git & GitHub

# **Professional Strengths**

- Analytical thinking & datadriven decision making
- Strong attention to detail
- Effective communicator & collaborative team player
- Creative problem solver with a growth mindset
- Project management

#### **LANGUAGES**

- English Native / bilingual
- German A2

#### **PROJECTS**

- Stacking ML models and feature engineering stacked regressors for house price prediction and combined it with advanced feature engineering including polynomial features.
- Comparing ML models for TB detection compared various classifiers in their ability to predict TB from chest X-ray data.
- Scaling up sentiment prediction with TensorFlow built and scaled up a sentiment classifier on 4 million product reviews using TensorFlow and batch processing.

# **Genomics Data Analyst**

Jan 2017 – May 2018

# World Agroforestry Center - Nairobi, Kenya

- Constructed UPGMA and neighbor-joining dendrograms to analyze genetic divergence (max distance = 0.30) and assessed clustering robustness via 1,000 bootstrap replicates.
- Quantified intra- and inter-cluster variances using ANOVA, revealing 86% molecular variation within populations (p < 0.001).

#### **EDUCATION**

# **Trainee in Data Analytics**

Mar 2024 - Jun 2024

ReDi School of Digital Integration, Berlin, Germany Coursework:

- Data analytics with Python fundamentals of the Pandas toolkit, principles of data filtering and the groupby method, data exploration and storytelling.
- Data analytics with SQL fundamentals of SQL, data analysis and visualization in SQL, creation and presentation of data dashboard.

#### **Dr.rer.nat. Molecular Genetics**

**Sept 2020 – Nov 2023** 

Universität Potsdam, Potsdam, Germany Gained expertise in:

- R programming, data cleaning, data wrangling, data analysis (EDA, PCA, clustering, regression, ANOVA, t-tests, post-hoc tests, anomaly detection), data visualization with ggplot2, and statistical inference using R.
- Creating data reports that effectively meet stakeholder needs.

# **Master of Science Biotechnology**

May 2015 - Aug 2018

Kenyatta University, Nairobi, Kenya Coursework:

- Introduction to Statistics: covered statistical methods such as, analysis of variance, Chi-square, t-test, correlation, regression, probability distributions, post-hoc tests, and nonparametric tests.
- Scientific Data Analysis: covered descriptive, diagnostic, and causal analyses, and making inferences to a larger population.
- Research Methodology for Pure and Applied Science: focused on qualitative / quantitative methods, hypothesis testing and p values.

### ADDITIONAL TRAINING

- IBM **Data Science Specialization** Coursera. Aug, 2024
- Machine Learning Specialization by Stanford University & DeepLearning.AI Coursera. Mar, 2025
- Machine Learning Fundamentals with Python Skill Track DataCamp. Mar, 2025
- **Supervised Machine Learning in Python** Skill Track DataCamp. Mar, 2025
- University of California Irvine **Predictive Modeling, Cluster Analysis & Association Mining** Coursera. Apr, 2025
- Power BI Fundamentals Skill Track (5 courses) DataCamp