Francoise(Kuan-Hua) A.

IT & Bioinformatics Manager | Al and Full-Stack Systems Architecture

I am an IT and bioinformatics manager skilled in system planning and architecture. My full-stack expertise includes front-end technologies like React, Angular, and TypeScript, and back-end languages such as Java and Python, along with REST/GraphQL API design. I work with databases like PostgreSQL, MySQL, MongoDB, and Neo4j, specializing in AI platforms and data-driven systems, ensuring secure and reliable product delivery throughout development.

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Professional Experience

IT/Bioinformatics Manager | PO/System Architect&Project Designer

Mar 2023 -Present

MedcomAsia Bio-Technology Company, Taiwan

Technologies: Java (Spring Boot/Spring AI)、Python (Flask/FastAPI/Django)、React、Angular、TypeScript、Docker、GitLab CI/CD、PostgreSQL、MySQL、MongoDB、Neo4j、GraphQL、Keycloak、RDF/SPARQL、FHIR

- Product Owner (PO): Defined product vision and managed backlog using Agile.
- WGSpark Architecture: Developed pipelines, APIs, and standardized workflows.
- SciTarget Platform: Created a JCR metric database and submission-strategy engine.
- FHIR ETL & Medical Data Standardization: Led ETL processes and resource modeling.
- Al Decision Support System: Integrated genomic and clinical data for enhanced predictions.
- Genomic Analysis Expertise: Merged NGS, epigenetic, and clinical data for predictive systems.
- Cross-Team Production: Coordinated IT, bioinformatics, and clinical roles; outlined data flows
- Knowledge Graphs Development: Utilized RDF/SPARQL/Neo4j with coding standards.
- Keycloak Integration: Implemented SSO, OIDC, SAML, and RBAC; established security processes.
- Bio-DevOps: Used Snakemake and Nextflow with GitLab Cl and Docker for reproducibility.
- Hybrid SaaS Model: Combined on-premises and edge solutions for data capture and processing.

Intership (Full stack of developer and Bioinformatics analyste

LBBE(Laboratory of Biometry and Evolutionary Biology, France

Technologies: HTML, CSS, JS, jQuery, Python Flask, Docker, Singularity, Unix, Snakmake

- Internship focused on rebuilding the DGINN pipeline with a Snakemake framework and creating an Web API to make it easier for biologists to use.
- Rebuilt the DGINN pipeline with Snakemake
- Built an API using a Python Flask server; front end with HTML, CSS, JS, and jQuery; containerized with Docker
- Edited a research paper in French

Using Machine Learning and Pattern Recognition to Analyze Spotify Music Data (Project)

LBBE(Laboratory of Biometry and Evolutionary Biology, France

Technologies: Python

Machine Learning Course Project — Pair Programming Practice

02/2023 -07/2023

2022/09 -2022/12

Professional Experience

2020/09 -2022/07

Full-Stack Bioinformatics Data Platform Development

GAD (Génétique des Anomalies du Développement) – CHU University Hospital in Dijon Technologies: Python, Flask, JavaScript, HTML, CSS, jQuery, MongoDB, Unix, Apache (internal preprod/server environment), Java, qsub (HPC scheduling)

- Designed and developed a web-based bioinformatics platform for clinical and research use, supporting analysis and visualization of large-scale sequencing data (genome/exome).
- Built an end-to-end data pipeline with Python and MongoDB, processing patient-derived genetic variants.
- Created interactive visuals and dynamic interfaces with JavaScript/jQuery to help geneticists and clinicians interpret results.
- Modified Java tools to strengthen ETL preprocessing and consistency.
- Deployed the system on in-hospital Unix environments and servers to ensure onpremises data security and performance.
- Used the qsub scheduler on HPC clusters for batch computing and optimized scheduling.
- Proficient with Unix commands for automation, monitoring, and maintenance.
- Provided on-call support for the data pipeline, including maintenance, debugging, and tool updates.

06/2016 -06/2019

Administrative Secretary & Project Assistant

Formosa IT Training - Lyon, France

Technologies: HTML, PHP, CSS, JavaScript, SQL

Formosa, an IT training firm, handles course and project management, instructor support, and admin operations.

- Manage admin, HR, and scheduling; help project managers and instructors design training.
- Align clients and tech teams to move projects forward.
- Work with an Agile mindset; set up standardized docs and task-tracking.
- Set the stage for later cross-department work as a Product Owner.

2012-2013

Substitute English Teacher

Jiaojiao Elementary School, Jinmin Branch, Sanxia, New Taipei City, Taiwan

English for Primary Students (7-12). Includes support for school library and information handling

08/2010 -08/2011

Human Resources Handling and Social Work

Church in New Taipei City Sanxia District

Social welfare role focused on documentation and data processing, accompanying young people, and teaching life education on campus.

09/2006 -06/2007

Teaching Assistant at the Biostatistics Center

Public Health Laboratory, China Medical University, Taichung, Taiwan (R.O.C.)

- · Support teaching at the Biostatistics Center:
- Data integration and analysis
- Statistical analysis and development of SAS teaching materials
- · Science research
- Hospital patient surveys to provide data for academic research

09/2006 -06/2007

Clinical Research Work: Practical Data Processing

Biostatistics Center, China Medical University, Taichung, Taiwan (R.O.C.)

Performed data processing and statistical analysis using tools such as SAS, SPSS, and R; also supported course preparation and teaching materials.

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09/2006 -

06/2007

Assistant for Professor, Environmental Sciences

Public Health Laboratory, China Medical University, Taichung, Taiwan (R.O.C.)

Assist microbiology research and participate in applying and analyzing Geographic Information Systems (GIS).

Training and Diplomas

2020 - 2023	Master Degree - Bioinfomatique - University of Claude Bernard Lyon 1		
2019 - 2020	Bachelors of Science, Technology, Health - Bio-Informatique, Statistique et Modelisation - University of Claude Bernard Lyon 1		
2019 - 2020	Professional IT training - Company of Formosa (IT training) program: SQL, Java, Talend, HTML/CSS, JS, Responsive Design		
2011	Teacher Training Program in Chinese as a Foreign Language		
2006 - 2007	Teaching Assistant Service - Practical Data Processing - China Medical University, Taichung, Taiwan (R.O.C)		
2007	Bachelors of Science - Department of Risk Management - College of Public Health - China Medical University, Taichung, Taiwan (R.O.C)		
2002	High School Degree - Taipei Municipal Ching-Mei Girls' Senior High School, Taiwan (R.O.C)		
С	onferences & Projects		
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09/2022- 12/2022	Master's Project – EcoFood API Technologies: Python, Flask, Web API, PostgreSQL Public health research: developed an API to calculate the carbon footprint of Crous meals.		
	Technologies: Python, Flask, Web API, PostgreSQL		
12/2022 09/2022-	Technologies: Python, Flask, Web API, PostgreSQL Public health research: developed an API to calculate the carbon footprint of Crous meals. Data Visualization of Mega-Data Technologies: Python, Web API, R Shiny, SQL		
12/2022 09/2022- 11/2022 01/02-	Technologies: Python, Flask, Web API, PostgreSQL Public health research: developed an API to calculate the carbon footprint of Crous meals. Data Visualization of Mega-Data Technologies: Python, Web API, R Shiny, SQL Interactive visualization of air pollution data in France Conference: French Annual Meeting of Genetics and Medical Genetics, Rennes "Innovative Human Genetics Database: SCNVBase for storing human genomic variant data." Authors: Kuan-Hua Artignan, Yannis Duffourd, et al. (Poster N°28146)) SCNVBase Project Technologies: MongoDB, Python Flask, HTML, CSS, JavaScript, jQuery, Apache; quality-control server Developed a Python API using a three-tier architecture to visualize genomic variant data, implemented JWT authentication and security mechanisms, and presented project results to		
12/2022 09/2022- 11/2022 01/02- 04/02/2022 2020 -	Technologies: Python, Flask, Web API, PostgreSQL Public health research: developed an API to calculate the carbon footprint of Crous meals. Data Visualization of Mega-Data Technologies: Python, Web API, R Shiny, SQL Interactive visualization of air pollution data in France Conference: French Annual Meeting of Genetics and Medical Genetics, Rennes "Innovative Human Genetics Database: SCNVBase for storing human genomic variant data." Authors: Kuan-Hua Artignan, Yannis Duffourd, et al. (Poster N°28146)) SCNVBase Project Technologies: MongoDB, Python Flask, HTML, CSS, JavaScript, jQuery, Apache; quality-control server Developed a Python API using a three-tier architecture to visualize genomic variant data,		

Technologies: HTML、CSS、Bootstrap、JavaScript、MySQL、PHP

foundation.

Developed the website front-end interface and content management system for a

Technical Skills Overview

Al-Driven Healthcare Platform / System Architecture	****
NGS Analysis (Next Generation Sequencing)	****
Al / Large Language Models (RAG/GraphRAG, Prompt Engineering, Model Evaluation)	***
Knowledge Graphs and Data Interoperability (RDF/SPARQL, GraphQL, FHIR)	***
Data Analysis & Machine Learning	****
Front-End (React/Angular/TypeScript)	****
Back-End (Java Spring Boot/Spring AI; Flask/FastAPI/Django)	****
Data Visualization (matplotlib / Dash / R Shiny / JS)	****

Technical Domain	Skills & Expertise
AI & Large Language Model Engineering	RAG/GraphRAG architecture design and implementation; prompt engineering; model fine-tuning and evaluation pipeline; AI model governance and version tracking; integration with full-stack web platform for decision-support systems; design of multilingual medical dialogue systems
Product Ownership & System Architecture	Agile/Scrum/Kanban project management; product backlog, user story, and sprint planning; layered/modular architecture design; cross-department collaboration and stakeholder communication (development, bioinformatics, AI); business model and operational strategy design; UX/UI wireframing and requirements analysis
Data Analysis & Machine Learning	Scikit-learn, Pandas, NumPy; feature engineering and pattern mining; statistical analysis (R/SPSS/SAS); big data analytics (PySpark/Node.js); model visualization and interpretability
Database & Knowledge Graph	PostgreSQL, MySQL, MongoDB, Neo4j (graph database); RDF, SPARQL, GraphQL; graph-based reasoning and semantic integration
FHIR & Interoperability	FHIR Resource/Terminology(ontology mapping (SNOMED/LOINC/ICD/RxNorm); FHIR API tools; FHIR-based ETL; RDF semantic transformation and standard mapping; FHIR applications
Biomedical/Bioinformatic s Data Analysis (NGS & Omics Integration)	RNA-seq, ChIP-seq, WGS, WES, Metagenomics; Snakemake/Nextflow pipelines; integrative analysis of genomic and clinical data (Omics + Clinical)
Full-Stack Development	Frontend: HTML, CSS, JavaScript, Angular, React, TypeScript, Jinja2, jQuery; Backend: Python (Flask/FastAPI/Django), Java (Spring Boot/Spring AI)
DevOps / MLOps / System Deployment	Docker, GitLab CI/CD, Conda, Linux/Unix, Windows, Kubernetes (K8s), unit testing, automation monitoring; system log analysis and environment governance
Identity & Access Management (IAM)	Keycloak (SSO/OIDC/SAML/RBAC); IAM/OAuth2/Zero Trust; token-based API security and access control; multi-layer data security and compliance design
Business Intelligence & Data Integration	Talend (ETL, field mapping, scheduling); Python ETL (pandas, FHIR API, REST import and transformation); report visualization (Dash/R Shiny); ETL automation and data workflow monitoring