

KARIM MEZHOUD

I am Data Scientist and Associate Professor in Nuclear Science Center of Tunisia - Tunis. I am teaching Bioinformatics and Biostatistics to Master degrees in Biomedical. The courses are based on GNU/Linux environment and R programming language. I started wrangling Cancer genomics data and developing tools for Analysis, visualization and prediction. Today, I extended my skills to multiple data types for business solutions, nuclear physics or Medical image processing. I am familiar with Python and the most known machine learning algorithms. Mainly I use R for EDA and Python for Prediction.



EXPERIENCE

2022
|
Present

Charles Nicolle Hospital of Tunis

System Administration and Data Scientist

📍 Tunisia, Tunis

- HPE Data server Storage
- Illumina Dragen Server for NGS data analysis
- Illumina NextSeq 550

2018
|
Present

National Center for Nuclear Sciences and Technologies

Data Scientist

📍 Tunisia, Tunis

- Dockerise shiny App in cloud
- Web service, security, cloudflare, auth0, freemom
- Dashboard, shiny, shiny-mobile
- R package
- Data cleaning, Processing, exploring, and visualization
- Modeling and Prediction with R/Python
- Data auto-Reporting (Rmarkdown)
- SQLite, MySQL, MSSQL
- TABLEAU
- github, gitlab, dockerhub
- DigitalOcean
- Agility: Scrum

2016
|
Present

National Center for Nuclear Sciences and Technologies

Associate Professor

📍 Tunisia, Tunis

- R programming Courses for Master degrees (Tunis, e-learning)
- Soft Structural Bioinformatics courses for Master Degree (Tunis, e-learning)
- Nuclear education network (e-learning, moodle)
- Medical images classification using Deep Learning
- Dosimetry data management and analysis

2015
|
Present

National Center for Nuclear Sciences and Technologies

Data Analyst

📍 Tunisia, Tunis

- Data analysis (Biology, health, Medicine, Pharma)
- R Programming, Shell, Regex, Shiny App
- Statistics: Pre-processing, Processing, & Data mining
- GNU/Linux & OSX environment
- Deep user of Bioinformatics tools (Proteomics, Chemoinformatics, Pharmacophore, Toxicology)

2009
|
2016

National Center for Nuclear Sciences and Technologies

Assistant Professor

📍 Tunisia, Tunis

- Teaching Bioinformatics Master Degree
- Research in Proteomics, Toxicology, & Mass Spectrometry

CONTACT INFO

✉ kmezhoud@gmail.com

🎓 [Portfolio](#)

🌐 [github](#)

[in linkedin](#)

[k kaggle](#)

📞 +216 25-909-901

SKILLS

System Administration
Data Analysis
Data Processing
Data Viz
GNU/Linux, OSX
R Shiny, Python, Bash,
Regex
Docker
github, gitlab
Agility: Scrum, Kanban

LANGUAGE

English
French
Arabic

CERTIFICATES

Coursera (Bioinformatics & Biostatistics)
DataCamp (R)
Dataquest (R)
LinkedIn learning (Cloudera, Hadoop)
Analytics Vidhya & HackerRank (Python)

2004
|
2006

Freelance Self-employed

Consultant in safety foods

📍 Paris, France

- Consultant in hygiene and food safety in collective catering and restaurant in Paris
- Hygiene good practice analysis, Hazard Analytic Critical Control point (HACCP)
- Trainer in Food Safety



SELECTED PACKAGES

2020

Orders and Stores web/mobile management

Mobile App: R/Docker

📍 [github](#)

2020

COVID Tracker

RShiny Dashboard

📍 [gitlab](#)

2019

Business Analysis of Laptops

Rshiny Dashboard

📍 [gitlab](#)

2017

Radiant.dose A shiny app for Dosimetry Data management

Randiant.dose

📍 [github](#)

2018

PapSmear: Uterine Cervical test image classification using Deep learning

PapSmear

📍 [github](#)

2016

BioCancer is a Shiny App to explore interactively Multi-Assay of Cancer Genomic Data

bioCancer

📍 [Bioconductor](#)

2015

cancerR - A Graphical User Interface for accessing and modeling the Cancer Genomics Data of MSKCC

cancerR

📍 [Bioconductor](#)



EDUCATION

2009
|
2013

Habilitation à Diriger des Recherches (HDR)

Faculté des Sciences de Tunis

📍 Tunisia, Tunis

- Graphical identification of cancer-associated gene sub-networks based on small proteomics data sets.
- Teaching
- Structural Bioinformatics (3D protein structure modeling, Docking, interactomics)
- GNU/Linux for Biology (Shell, regex)
- Introduction to R in Biology (How to choice of statistic tests, graphics)

2004
|
2007

PhD in Proteomics and Bioinformatics

Museum National d'Histoire Naturelle

📍 France, Paris

- Proteomic and phosphoproteomic analysis of cellular responses in medaka fish (*Oryzias latipes*) following oral gavage with microcystin-LR
- Animal assay: Mouse, Fish, Rat
- Cellular assay: CHOK1
- Electrophoresis, mass spectrometry, imaging, Western blot

2003
|
2004

Master's degree in Biotechnology and Toxins

Museum National d'Histoire Naturelle, Paris

📍 France, Paris

- Environmental Toxicology
- Fish Model
- Apoptosis
- Stress Oxydative
- Acid Okadaic

2002
|
2003

Master's degree in Total Quality and Management

Ecole Nationale Vétérinaire d'Alfort, Paris

📍 France, Paris

- Veterinary
- Safety food
- Agro-food
- Industry
- Catering

1995
|
2000

Engineer's degree in Agronomy: Food technology

Institut National Agronomique de Tunisie, Tunis

📍 Tunisia, Tunis

SELECTED PUBLICATIONS

2014

Radiobiology Worker Risk Assessment Using Stress Indicators and Proteomics.

Karim Mezhoud, Amina Sakly, Hassen Ben Cheikh, Mouldi Saïdi, and Marc Edery.
📍 *Int. J. Low Radiation* Vol.9, No.3, 2014.

2013

Graphical identification of cancer-associated gene sub-networks based on small proteomics data sets

Karim Mezhoud 📍 *OMICS: A Journal for Integrative Biology* 2013 17(7): 393-397.

2011

Toxins and stress in fish: proteomic analysis and response network.

Karim Mezhoud, Puiseux-Dao S., Edery M. 📍 *Toxicon*, 57(7-8), 959-969.

2009

Deciphering peculiar protein-protein interacting modules in *Deinococcus radiodurans*

Mezhoud K., Sghaier H. and Barkallah I. 📍 *Biology Direct*

2009

Toxicoproteomics: a way to unravel response pathways induced by a toxin

Malécot M., Mezhoud K., Marie A., Praseuth D., Puiseux-Dao S. et Edery M.
📍 *Aquatic Toxicology*, (2009) 94(2):153-61.

2008

Global quantitative analysis of protein expression and phosphorylation status in the liver of the medaka fish (*Oryzias latipes*) exposed to microcystin-LR I. Balneation study

Mezhoud K., Praseuth D., François J.C., Bernard C., and Edery M.
📍 *Aquatic Toxicology*, (2008) 86 : 166 - 175

2008

Proteomic and phosphoproteomic analysis of cellular responses to microcystin-LR in medaka fish

Mezhoud K., Bauchet A.L., Praseuth D., Marie A., François J.C., Fontaine J.J., Jaeg J.P., Cravedi J.P., Puiseux-Dao S., Bernard C. and Edery M.
📍 *Toxicon* 51 (2008) 1431 - 1439

Last updated on 2023-02-17.