

# KADIR KOCABAŞ

# **BIOINFORMATICIAN BIOMEDICAL EGNINEER**







Address: Istanbul, Pendik, TURKEY Mail: kocabaskadir@gmail.com Phone: +90 534-841-9133

#### **LANGUAGES**

- TURKISH NATIVE
- ENGLISH FLUENT

# **CORE SKILLS**

\*\*\* **PYTHON** MATLAB \*\*\* \*\*\* **DATA** 

SCIENCE



MACHINE LEARNING

**DEEP** 

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LEARNING





#### **CERTIFICATES**

#### **COURSERA**

- NEURAL NETWORKS AND DEEP LEARNING
- IMPROVING DEEP NEURAL NETWORKS: HYPERPARAMETER TUNING, **REGULARIZATION AND OPTIMIZATION**
- R PROGRAMMING
- GETTING AND CLEANING DATA
- THE DATA SCIENTIST'S TOOLBOX

#### **UDEMY**

• THE COMPLETE SQL BOOTCAMP 2020: GO FROM ZERO TO HERO



# **SUMMARY**

I am a PhD student in Bioinformatics and Systems Biology at Gebze Technical University. I am a contributor to The COBRA Toolbox (Linkl, Link2). I have experience in constraint-based mathematical modelling, machine learning, statistics and data analysis. As a highly self-motivated person, I like to improve myself and learn new things. I know how to research and I am good at finding information.



# **EXPERIENCE**

# **Project Assistant**

TUBITAK | 02.03.2021 - Current

120S824, Alzheimer Hastalığı İçin RNA-Seq Verilerinden Faydalanarak Kişiye Özel Moleküler Etkileşim Modelleri Geliştirilmesi ve Hastalık Mekanizmalarının Aydınlatılması

# **Project Assistant**

TUBITAK | 02.12.2019 - 15.12.2020

315S302, Parkinson Hastalığı İçin Transkriptom Verilerinin ve Hücre İçi Ağyapıların Biyoinformatik Analizi: İlaç Hedefi ve İlaç Geliştirilmesi

# **Project Assistant**

TUBITAK | 23.07.2019 - 01.12.2019

316S005, Metabolik Yolakların Kısıt-Tabanlı Modelleme Yaklaşımı ile Încelenmesi ile Klebsiella Pneumoniae ve Salmonella Enterica Kaynaklı Ölümcül Enfeksiyon Hastalıkları İçin Potansiyel İlaç Hedefi Tespiti, Uluslararası



# **EDUCATION**

#### **GEBZE TECHNICAL UNIVERSITY**

### PHD. | FEBRUARY 2021- CURRENT

BIOINFORMATICS AND SYSTEMS BIOLOGY

- MATHEMATICAL MODELLING OF BRAIN CELLS
- NI P BIOLOGICAL DATA ANALYSIS MACHINE LEARNING

#### **GEBZE TECHNICAL UNIVERSITY**

MSC. | JULY 2018- JANUARY 2021

BIOINFORMATICS AND SYSTEMS BIOLOGY

- MATHEMATICAL MODELLING OF BRAIN CELLS
- ANALYSIS OF PARKINSON DISEASE ANALYSIS OF SALMONELLA
- BIOLOGICAL DATA ANALYSIS MACHINE LEARNING -DECONVOLUTION

#### **ERCIYES UNIVERSITY**

#### MSC. | SEPTEMBER 2011 - JUNE 2016

BIOMEDICAL ENGINEERING

GRADUATION PROJECT: DEVELOPING NUTRITION APPLICATION FOR ANDROID PLATFORM



#### **REFERANSLAR**

Tunahan Çakır

Associate Professor at Gebze Technical University

Mail: tcakir@gmail.com

# **P**

# **PUBLICATIONS**

- KOCABAS K., ÇAKIR T. (2020). INTEGRATIVE ANALYSIS OF MULTI-CELLULAR GENOME-SCALE
  METABOLIC NETWORKS WITH CELL TYPE SPECIFIC TRANSCRIPTOME DATA PREDICTED BY
  DECONVOLUTION ALGORITHMS: APPLICATION TO PARKINSON'S DISEASE. POSTER SESSION
  PRESENTED AT THE ANNUAL MEETING OF INTERNATIONAL EURASIAN CONFERENCE ON
  BIOTECHNOLOGY AND BIOCHEMISTRY
- KOCABAŞ K., ÇAKIR T. (2019). IDENTIFICATION OF DRUG TARGET FOR SALMONELLA ENTERICA INDUCED INFECTIONS BY ANALYSIS OF INTEGRATED PATHOGEN HOST METABOLIC NETWORK. POSTER SESSION PRESENTED AT THE ANNUAL MEETING OF THE TÜRK TIP DÜNYASI KURULTAYI.
- KOCABAŞ K., ÇAKIR T. (2019). INTEGRATIVE ANALYSIS OF PATHOGEN HOST METABOLIC NETWORK OF SALMONELLA ENTERICA WITH DUAL RNA SEQ DATA. POSTER SESSION PRESENTED AT THE ANNUAL MEETING OF THE INTERNATIONAL SYMPOSIUM ON HEALTH INFORMATICS AND BIOINFORMATICS.