

# Erwin ERDEM

 [linkedin.com/in/erwin-erdem](https://www.linkedin.com/in/erwin-erdem)    [erwin.erdem@outlook.com](mailto:erwin.erdem@outlook.com)  
 [github.com/erwinerdem](https://github.com/erwinerdem)    [erwinerdem.github.io](https://erwinerdem.github.io)  
 Schiedam, The Netherlands    Chamber of Commerce : 77877373

## >\_ PROGRAMMING SKILLS

**Languages** Python (**pandas**, **scipy**, **numpy**, **scikit-learn**, **scanpy**, **tensorflow**, **pytorch**, **plotly**, **matplotlib**, **seaborn**, **dash**) • R (**tidyverse**, **ggplot2**, **DESeq2**) • JavaScript (**D3**, **plotly**) • Bash • MATLAB • Julia • Java • Perl

**Development Tools** Visual Studio Code • Project Jupyter • Git • Conda • Snakemake • Docker

## WORK EXPERIENCES

<b>Current</b> <b>Feb 2020</b>	<b>Bioinformatics Consultant, PROQR THERAPEUTICS, Leiden, The Netherlands</b> Consultancy in bioinformatics for the development of new RNA editing technologies at the Innovation Unit. <ul style="list-style-type: none"><li>&gt; QC, map and analyze (single cell) RNA-seq data on a server using <b>MultiQC</b>, <b>STAR(solo)</b>, <b>samtools</b>, <b>snakemake</b>, <b>scanpy</b>, <b>DESeq2</b> and <b>plotly</b> among others</li><li>&gt; Employing and training deep learning models with <b>keras</b> and <b>tensorflow</b></li><li>&gt; RNA folding and protein interaction prediction (<b>ViennaRNA</b>, <b>UFold</b>, <b>AlphaFold</b> etc.)</li></ul>
<b>Apr 2020</b> <b>Mar 2017</b>	<b>Student Assistant, DELFT UNIVERSITY OF TECHNOLOGY, Delft, The Netherlands</b> Assisted students in MATLAB exercises and homework assignments for the Biophysics course (NB1132) given by prof.dr. Chirlmin Joo of the Bionanoscience department at TU Delft for five weeks every year. <ul style="list-style-type: none"><li>&gt; Modeling various biophysical processes</li><li>&gt; Communicating complex biophysics topics and teaching basic programming to students</li></ul>
<b>Jan 2019</b> <b>May 2018</b>	<b>Tutor, STUDENTSPLUS, The Netherlands</b> Aided high school students with chemistry, physics, mathematics and biology
<b>Sep 2018</b> <b>Jul 2018</b>	<b>API Tester, DATA ARCHIVING AND NETWORKED SERVICES, DANS, Den Hague, The Netherlands</b> Created test cases for archiving research data onto a server using <b>curl</b> and Git for version control. <ul style="list-style-type: none"><li>&gt; API improvement through regular communication with the developers</li></ul>

## INTERNSHIPS & THESES

<b>Jun 2021</b>	<b>Master Thesis, ERASMUS MC, COMPUTATIONAL POPULATION BIOLOGY GROUP, Rotterdam, The Netherlands</b>
<b>May 2020</b>	Developed a computational framework which elucidates more on the complex genomics of brain (imaging) phenotypes by combining GWAS with scRNA-seq data. Supervised by dr. Gennady Roshchupkin and prof.dr. Steven Kushner. <ul style="list-style-type: none"><li>&gt; Incorporate data from various (public) resources (NHGRI-EBI GWAS Catalog, GWAS atlas, descartes) to assess the framework</li></ul>
<b>Jan 2020</b> <b>Sep 2019</b>	<b>Intern, PROQR THERAPEUTICS, Leiden, The Netherlands</b> Established a computational platform in Python to accelerate the development of Trident™, an RNA pseudouridylation editing technology. Supervised by dr. Pedro Morais. <ul style="list-style-type: none"><li>&gt; Development of novel analysis tool specific for the Trident™ technology</li></ul>
<b>Jul 2017</b> <b>Feb 2017</b>	<b>Bachelor Thesis, TU DELFT BIONANOSCIENCE DEPARTMENT, CHIRLMIN JOO LAB, Delft, The Netherlands</b> Observed the dynamics of FnCas9 from the CRISPR-Cas9 system with a custom guide RNA and DNA complex using single molecule fluorescence resonance energy transfer with a total internal reflection fluorescence (TIRF) microscopy. Supervised by dr. Viktorija Globyte and prof.dr. Chirlmin Joo. <ul style="list-style-type: none"><li>&gt; Expression, isolation and purification of the FnCas9 DNA endonuclease protein and gRNA</li></ul>

## EDUCATION

2018-2021	<b>MSc Nanobiology</b> (TU Delft and Erasmus MC)
2014-2018	<b>BSc Nanobiology</b> (TU Delft and Erasmus MC)
	<b>Minor</b> Computational Approaches to Disease, Signaling and Drug Targets (Leiden University)