# **Nicolas Arning**

n.arning@gmx.de

github.com/narning1992

linkedin.com/in/nicolas-arning

bit.ly/nicolas\_arning

# PhD Computational Genomics

## Education

M.Sc Biosciences | 2016 | Grade 1.4 WWU Münster

Secondment | 2015 | Grade 1.1 University of British Columbia

**B.Sc Biosciences** | 2014 | Grade 1.8 WWU Münster

Secondment | 2014 | Grade 1.6 University of Sheffield

#### Skills

Python of the original of the Bash R Matlab • Slurm

### Extra-curricular

#### Participant in

Young Entrepreneur Science

- Conceived hypothetical startup • Developed financing
- and marketing strategies

#### Lecturing at **Doctoral Training Centre**

- Developed and held introduction into LaTeX
- Lectured for Python, C and Phylogenomics

## Referees

Prof. Daniel J. Wilson

Big Data Institute University of Oxford daniel.wilson@bdi.ox.ac.uk

Prof. David C. Clifton

Institute for Biomedical Engineering University of Oxford davidc@robots.ox.ac.uk

# Relevant Experience

Since Oct 2022

Health Data Scientist

Big Data Institute, University of Oxford

Developing disease independent risk factor prediction using Bayesian model averaging on UK Biobank data

Sep 2017 Jul 2022

**Doctor of Philosophy** 

Big Data Institute, University of Oxford

Developed Campylobacter source attribution algorithm using Machine learning on whole genomes

Uncovered genetic underpinnings of *Campylobacter* infectivity through genome wide association studies

Jun 2018 Jul 2022

Founder and leader DTC Coding Dojo

Doctoral Training Centre, University of Oxford

Developed peer-support network for Bioinformatics

Nov 2019

Course co-organiser

Doctoral Training Centre, University of Oxford

Conceived and held deep learning course with two peers

Jan 2019 Mar 2019

Oxford Nanopore Technologies

Intern

Developed taxonomic classification deep learning algorithm

# Selected publications

Oct 2021

Machine learning to predict the source of campylobacteriosis using whole genome data

PLOS Genetics

N Arning, S Sheppard, S Bayliss, DA Clifton, DJ Wilson

Jul 2021

Mapping the human genetic architecture of COVID-19 Nature

COVID-19 Host Genetics Initiative

Oct 2020

Genomewide association study of severe Covid-19 with respiratory failure

New England Journal of Medicine

Severe Covid-19 GWAS Group

Jul 2020

The past, present and future of ancient bacterial DNA

Microbial Genomics

N Arning, DJ Wilson

Mar 2018

Hemimetabolous genomes reveal molecular basis of termite eusociality

Nature Ecology and Evolution

MC Harrison, E Jongepier, HM Robertson, N Arning [...]