Curriculum Vitae

Skills

Core skills

Design and development of desktop and web applications, focused on lifesciences and devoted to clean, sustainable architecture, future-proof technologies and usability, covering every lifecycle phase from application development to testing, deployment, system administration and every layer between database design and user interface.

Technologies, programming languages and software

devops, Domain Driven Design, Object Oriented Design, JEE, REST, (R)DBMS, functional programming, reactive programming, CI/CD, Scrum, Grid Engines, datascience

Python, Java, JavaScript, HTML5, CSS3, Bash, SQL, PHP

Docker, Linux, git, pandas, jupyter notebooks, AWS, VSCode, Eclipse, Oracle DB, VSCode, PyCharm, Glassfish, Maven, HTCondor, Firebird DB, Vim, Jenkins, Gitlab, Jira, Confluence, celery, Adobe Creative Suite

Spoken languages

German native proficiency **English** professional working proficiency **Russian, Swedish** elementary proficiency

Systems Engineer

since January 2021

AnyDesk Software GmbH Stuttgart, Deutschland

Technical Product Management for the development of the user- and device management software my.anydesk. Analysis of previous work, development of a new architecture and establishment of an proof-of-concept.

Development of a new Software Team consisting of Frontend-, Backend- and Full-Stack-Engineers. On- and offboarding and mentoring. Technical and managerial Leadership.

Establishment of **Software Development Life Cycles and Documentation processes**, customized for challenges and requirements particular to this team.

Continuous **contribution to the product development, database and interface designs**, requirement engineering and stakeholder communication.

Consultancy of adjecent teams on topics of automation and interface design for established and novel products and tools.

Quarkus, Vert.X, React, Next.js, Postgresql, Gitlab, Docker, Jira, Confluence

Software Engineer and IT Allrounder

March 2015 - December 2020Insilico Biotechnology AG Stuttgart, Germany

Developing a machine learning pipeline to create customized predictive hybrid models for the analysis of bioprocess performance and various optimization scenarios.

Developing enterprise grade software for the design and simulation of metabolic network models, quantitative comparison of bioprocess performance and feed media optimization.

Contribution of software engineering expertise throughout the software development lifecycle, from requirements, through design and implementation to deployment and in all layers between database, middleware, grid engine and user interface.

Maintenance and administration of IT infrastructure including multiple testing and deployment machines running CentOS and SUSE Linux, Oracle 11g database, the CI pipeline, an HTCondor compute cluster and containerized applications on premise and using AWS.

Python, jupyter notebooks, gitlab, AWS, HTCondor, Oracle DB, Postgresql, Jira, Confluence

Internship

May 2013 - Jul 2013

Insilico Biotechnology AG

(3 months)

Stuttgart, Germany

Developing a genome-scale metabolic flux model of an antibiotic producing bacteria including extensive literature and database research and modelling with proprietary software.

Insilico Discovery, LaTeX, BENDA, EMBL, ExPASy, GenBank, NCBI

Research Assistant

Nov 2012 - Apr 2013

Institute of Technical Biochemistry University of Stuttgart, Germany

Development of a User Interface for the novel enzyme sequence-structure-function database BioCatNet.

PHP, Firebird DB, Perl, HTML, JavaScript, CSS, Git

Research Assistant

Jan 2011 - Jun 2011

Department of Physics, Chemistry and Biology Linköping University, Sweden

Gene Expression Analysis of the late flowering gene in the common pea *Pisum sativum*. Working with PCR/rtPCR machines, DNA/RNA-extraction kits and Gelelektrophoresis.

Certified Java EE Workshop

March 2017

PC College, Stuttgart

Enterprise Java Beans 3.1, Session- and MD Beans, CDI, JPA, Lifecycle

Dipl. Biologe, t.o.

technically oriented Biology diploma

Oct 2007 - Nov 2014

University of Stuttgart, Germany Focus: Bioinformatics, Biophysics

Jan 2011 - Jun 2011

Linköping University, Sweden Focus: Neuroscience, Gene Expression

Diploma Thesis

Development of a novel Protein Database System for Sequence-Structure-Function Relationships: BioCatNet

Oct 2013 - Dec 2014

Institute of technical Biochemistry University of Stuttgart, Germany

Development of a bioinformatic data warehouse application including middleware, API and UI, capable of holding and managing protein sequence, structure and function information, using PHP, Perl, HTML, CSS and JavaScript. The system is publicly available for scientific use on https://biocatnet.de

Maintenance and administration of IT infrastructure including Apache HTTP and Firebird database servers on Debian Linux machines.

Student Research Paper

Analyse thermokinetischer Potential-Strom-Kennlinien an einem Modell von *Escherichia coli*

Oct 2011 - Jun 2012

Institute for System Dynamics University of Stuttgart, Germany

Assessing the predictive quality of the novel modelling technique *Thermokinetic Modelling* through automated simulations, using Mathematica, R and bash scripts on a Linux system.