jun.ling90@gmail.com | 0452 266 922

EDUCATION

UNIVERSITY OF QUEENSLAND

BE IN CHEMICAL & BIOLOGICAL Engineering

Discontinued | St Lucia, QLD

BE IN SOFTWARE ENGINEERING Expected July 2015 | St Lucia, QLD

BSC IN GENETICS

Expected July 2015 | St Lucia, QLD

Grad. Dec 2007 | Brisbane, Australia

LINKS

Github://jling90 LinkedIn:// Jun Ling

COURSEWORK

UNDERGRADUATE

Algorithms & Data structures Computer Graphics Data Analysis Information Systems Operating System Architecture Machine Learning Software Design Process Design Computing **Embedded Systems** Biochemistry & Molecular Biology Physical Chemistry Genomics & Bioinformatics **Advanced Genetics**

SKILLS

PROGRAMMING

Python • R • C • C++ Java • BASH • HTML JavaScript • CSS • SQL MATLAB

TOOLS & IDES

Unix • Microsoft Windows Git • SVN • LaTEX • Eclipse IPvthon Notebook • Vim Microsoft Visual Studio 2010

EXPERIENCE

CSIRO | PROGRAMMER

Sep 2011 - Dec 2013 | St Lucia, QLD

- Produced scripts for parsing and processing .csv files containing weather data and temperature sensor readings.
- Automated filtering and flagging of erroneous or suspicious data and generation of timeseries plots from the cleaned data.
- Field work: aided in assembly and placement of temperature sensors.
- R, HPC Job Manager

Reference: Scott Chapman · +61 (7) 3214 2254 · scott.chapman@csiro.com

ST. PETERS LUTHERAN COLLEGE ACPFG APPLIED BIOINFORMATICS | VOLUNTEER INTERN

Jun 2013 - Aug 2013 | St Lucia, QLD

- Worked independently on a project to perform de novo assembly of Illumina sequence data from seagrass. This involved developing a pipeline in Bash and Python for read trimming, mapping and assembly.
- Gained experience in running jobs on UQ's Barrine HPC cluster using PBS.
- Python, Bash, PBS, High-performance Computing, Linux

Reference: Kenneth CHAN · c.chan10@uq.edu.au

RIO TINTO ALCAN | Summer Intern + Casual contractor

Nov 2013 - June 2014, Nov 2014 - Dec 2014 | Pullenvale, QLD

- Developed .dll plugins for Microsoft Excel and SysCAD to provide functions for simulating chemical processes. The plugins were distributed among R&D facilities within Rio Tinto Alcan, and to other affiliated research groups.
- Participated in meetings with clients to evaluate and direct project to meet user requirements.
- Produced documentation for end-users and future developers.
- C++, Visual Studio 2010, Git

Reference: Alistair GILLESPIE +61 (7) 3107 4610 · alistair.gillespie@riotinto.au

OTHER PROJECTS

METNET | PROJECT CO-FOUNDER

September 2013

- Pipeline for analysis and identification of metastatic cancer sub-networks from Gene Ontology and microarray data.
- Python, R, Bash

SEQFINDR-WEB | AUTHOR

July 2014

- Web tool for dynamic and interactive visualisation of genomic data using Flask and d3.is.
- Public deployment expected Jan 2015.
- Python, HTML, CSS, JavaScript