



International Centre for
**Cancer Vaccine
Science**

Bioinformatics Analyst - Mass spectrometry based -omics

Job Description:

The International Center for Cancer Vaccine Science is hiring a data-science team to solve the many computationally complex problems we encounter in the search for the cure for cancer. We work towards the development of new immunotherapies that leverage the patient's own immune-system to eradicate the disease. Our work and findings help to improve the treatment and care for cancer patients around the world.

The data we deal with is truly huge: individual datasets can be hundreds of terabytes, and can include numerical, textual and imaging data. The team we are building is incredibly diverse, with biologists, statisticians, software engineers, bioinformaticians, mathematicians and data scientists. Together, we are all working as a team toward the common goal of developing new therapies for cancer patients. Use your technical skills to tackle one of the hardest problems of our time as you continue to grow into data-science.

We are searching for a Bioinformatics Analyst passionate about using mass-spectrometry to gain insights in proteomics and metabolomics within cancer datasets. We are equipped with state of the art facilities on which to develop your data-science and high performance computing skillset. The project has access to Cyfronet Prometheus (~55, 000 cores) and CI TASK Tryton (~38, 000 cores) clusters, which are consistently represented among the top 500 super computers in the world. As the work is international in nature, we are constantly integrating with sister-groups

around the world. There will be ample training opportunities for those who join the team and we offer an internationally competitive salary competitive with your experience.

How to apply:

You are encouraged to contact Dr. Ted Hupp (Ted.Hupp@ed.ac.uk) and Dr. Javier Alfaro (Javier.Alfaro@proteogenomics.ca) before submitting an application.

Responsibilities:

A post for a Computational Biologist in cancer vaccine science is currently available within the International Center for Cancer Vaccine Science led by Dr. Javier Alfaro. You will be (or become) an expert in a diversity of proteomics and metabolomics analysis pipelines involved in analyzing including single cell analysis, bulk-tumour analysis and blood biopsy analysis. You will support and develop grant applications within the team and provide guidance to enable best practice for experimental design. Within the bioinformatics group, you will work within a cohesive interdisciplinary team of data scientists that are located internationally in a variety of sites including France, Canada, UK and Poland. You will enable programs and pipelines for analysis, establish approaches for the integration of the analysis with multiple genome-wide '-omics datasets including processed DNA, RNA and proteomics datasets. Your results will be used to explore questions in cancer vaccine science.

Responsibilities may also include:

- Provide consulting on biological data analysis for ICCVS researchers conducting cutting-edge molecular biology research
- Advise and help researchers with their experimental design, data analysis and interpretation of results
- Provide training (2-5 days courses) in the area of machine learning and data science in biology
- Mentorship of junior staff and co-op students
- Collaborate and interact with other scientists at ICCVS and partner networks in

an international, interdisciplinary, and highly collaborative work environment

Preferred Qualifications:

- Bachelor's degree (MSc preferred) or further training in Machine Learning or Bioinformatics
- Experience with version control systems (SVN, Git, Mercurial, etc)
- Experience with some of the following: R, Python, C, C++, Java
- Knowledge of UNIX/Linux environments

Beneficial skills include:

- Experience in analyzing single cell 'omics, imaging, -omics, proteomics, or microfluidics data sets
- Experience in teaching
- Experience in R as a package developer or maintainer
- Knowledge of Python or other scripting languages
- Knowledge of scientific authoring frameworks (LaTeX, Rmarkdown) and R-Shiny
- Experience with version control, method benchmarking, or data management technologies
- experience with Amazon Web Services, Google Cloud Engine or other clouds

biochemistry, vaccine technology and computational biology.

- Employees will have the opportunity to develop skills in machine learning and high performance computing. The center has access to Cyfronet Prometheus (~55, 000 cores) and CI TASK Tryton (~38, 000 cores) clusters, which are consistently represented among the top 500 super computers in the world.
- Our software engineers, statisticians and computer-scientists are encouraged to further develop their skills in data-science through the attendance of workshops, conferences etc.
- The ICCVS provides an exceptional opportunity for motivated data scientists to work in an international multidisciplinary training environment to tackle major challenges at the interface between basic cancer discovery science and translational medicine.
- As we are an international center our infrastructure is remote-work compatible. A benefit enjoyed by excellent communicators and team players.

Benefits

- The ICCVS is housed in state-of-the-art facilities at the UG with facilities for mass spectrometry, virology, protein

