

Wong Lim Soon

From: Hui Kam Man (NCCS) [cmrhkm@nccs.com.sg]
Sent: Tuesday, April 17, 2012 11:22 AM
To: wongls@comp.nus.edu.sg; ASCKKWOH@ntu.edu.sg
Subject: IT employment

Hi Limsoon and Chee Keong,
I would like to employ a IT M.Sc. graduate for my laboratory.
I have put together the job descriptions as follows.
I would appreciate it if you can recommend someone.
Thanks.
Kam

Job Descriptions:

The Bek Chai Heah Laboratory of Cancer Genomics is involved in the generation and analysis of large-scale genomic data generated from expression profiling studies and integration with clinical information to identify novel genes that are crucial for the development of human cancers. Applying genome-wide, microarray gene expression analysis to identify deregulated genes in different tumour types can provide potential gene candidates as diagnostic and prognostic tools and promising targets for drug development. In collaboration with different clinical departments, the laboratory has characterized tumour-specific genes for many human cancers that are common in Singapore. The group is investigating the potentials of some of these genes for the early diagnosis and treatment of these cancers.

We are looking for a Bioinformatician with a keen interest in medical bioinformatics. Your main responsibilities include microarray expression analysis, gene/protein pathway analysis and providing basic biostatistics support to the lab.

Requirements:

- Masters in bioinformatics/computational biology/computer science or higher
- A good team player who can work with a variety of different researchers
- Experience with data-mining platforms and statistical software programmes (such as R, Partek, IPA, SPSS, etc.) is preferred
- Experience in conducting research is a major advantage
- Strong work ethics and a keen interest to learn on the job is essential

Interested applicants, please email Ms Theresia Tan at cmrth@nccs.com.sg. Please address your application to Prof Hui Kam Man.

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