Bioinformatician in cancer research

We are looking for a talented **bio/informatician** to join our research team in the group of Prof. Ondřej Slabý at CEITEC MU, Brno, CZ (https://www.ceitec.eu; http://slabylab.ceitec.cz/).

We're interested in the **analysis of molecular changes in various solid malignancies using different Next-generation sequencing methods** (Whole Exome/Genome Sequencing, RNA sequencing, methylome sequencing etc.).

The focus of the position will be:

- **Computational analysis of genomic data**
- **Implementation of analytical pipelines**
- **Development of software tools**

Skills in scripting languages (R, Python) and good spoken and written English are essential. **Candidates of any level of education are encouraged to apply.** Candidates with experience in high-throughput sequencing data analysis, machine learning and statistics, as well as in biology and medicine, and first-author publications will be favored.

The working capacity is preferably full-time, but a half-time contract is negotiable. The salary will be set based on candidate’s experience with a possible increase after the probation period.

Please **apply until the 31.1.2020** to be fully considered. Send us a **short CV** including contact details of at least one referee to Hana Nosková, Ph.D. at hana.noskova@ceitec.muni.cz and Karolína Trachtová, M.Sc. at karolina.trachtova@ceitec.muni.cz. Questions are welcome. Shortlisted candidates will be invited for an interview (on-site / Skype).