



(https://lbbe.univ-lyon1.fr/sites/default/files/styles/img_1280x768_image_scale_crop_main/public/media/images/arton6036.jpg?itok=MCdmkdvC)

Bioinformatic engineer position on transcriptomic/epigenomic analysis available at the LBBE in Lyon

Publié le 31 août 2016

A 2 years position is available at the Laboratory of Biometry and Evolutionary Biology (LBBE) located near the city of Lyon, France. We are looking for a highly motivated person with strong expertise in bioinformatics and statistics, to work on the analysis of genomic, RNAseq and ChIPseq data from *Drosophila* wild-type strains.

The successful candidate will be involved in several research projects dedicated to the analysis of abiotic and biotic stress on *Drosophila*, with a particular focus to the repeated sequences.

The ideal candidate should have expertise/experience with differential expression analysis and ChIP seq data coming from multiple samples, and with programming. An expertise in statistics is also necessary. She/he also should appreciate to work in a collaborative system and on different projects.

Degrees: we expect to recruit a master level engineer (BAC+ 3 to BAC +5). The choice of the candidate will depend on the experience and motivation.

Environment: our team is located in the Laboratory of Biometry and Evolutionary Biology at the University Lyon1 in the Doua campus. The LBBE provides a very rich scientific environment from genetics to genome evolution, ecology, modeling and bioinformatics. The Doua campus is close to the center of Lyon, which is a very attractive and cosmopolitan city.

For more information about the job and to apply, candidates should send a brief summary of research experience, a CV, and names and e-mail addresses of three references to :

cristina.vieira@univ-lyon1.fr

website: <https://lbbe.univ-lyon1.fr/>