

27. Mai 1999

The Novartis Forschungsinstitut is a Vienna-based pharmaceutical research centre representing an important element in the international drug discovery effort of the Novartis group.

For our

Genetics Unit

we are seeking a

Scientific Coworker Bioinformatics

with a thorough working knowledge of a Unix operating system environment. Good programming skills (preferably C, Perl, shell scripts, and ideally also Java) as well as experience in installing and integrating external software are essential. A solid background in the molecular biology of gene expression and experience with sequence databanks, ideally also with methods for EST clustering, is highly desirable. The successful candidate will become part of a team working on a large-scale gene discovery effort. The post will provide the opportunity to participate in employing and developing methods and algorithms for high-throughput EST sequence processing including automated annotation, efficient data mining, and linking to gene expression data. Further enquiries can be made by E-mail to Mr. H. Lapp (hilmar.lapp@pharma.novartis.com).

We are seeking a highly motivated individual with good teamwork, communication, and problem solving skills. In return, we offer a creative scientific and international environment, state-of-the-art facilities and equipment and competitive salary. The post is available immediately, for 3 years.

To apply, please send a full curriculum vitae including names of references to:

Novartis Forschungsinstitut GmbH
attn. Ms. Marion Kala, Human Resources
Brunner Straße 59, A-1235 Vienna, Austria

For more information on our Institute visit our web site at <http://www.at.novartis.com>

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