Hoechst Marion Roussel

Hoechst Marion Roussel, the pharmaceutical company of Hoechst, discovers and develops novel therapies based on scientific innovations in fields such as genetic engineering and biotechnology and aims to bring those therapies to the global marketplaces in the form of branded prescription drugs.

HMR Global Bioinformatics is focused on the exploitation of high-throughput biological data from our genomics resources.

We are seeking a

Bioinformatics Scientist - Sequence analysis high throughput analysis pipelining

Members of this group are responsible for the enhancement and further development and maintenance of our tools for sequence analysis and the application of these tools for high throughput sequence analysis and annotation. Applicants for these positions must demonstrate knowledge of additional bioinformatics analysis tools (e.g. HMMer, Wise, PHYLIP, etc.) and good CGI programming skills.

We prefer highly skilled and motivated staff to fill positions at our research locations in Germany. Applicants for all positions should possess good UNIX and scripting skills, be familiar with standard bioinformatics tools and programs (GCG, BLAST, FASTA) and public sequence databases and demonstrate basic knowledge of programming concepts and
languages.

We offer a creative scientific and international environment which will allow you to develop your skills very quickly. Our state-of-the-art facilities give you the opportunity to fill this challenging position and prepare you for future career. In addition we offer a competitive salary and attractive bonus package which will suit your qualification..

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

HOECHST MARION ROUSSEL
DEUTSCHLAND GMBH
attn. Doris Madet
Human Resources
Building K 607
65926 Frankfurt
Germany

For more information on our company visit our web site at http://www.hmr.de
For job information, please contact Doris.Madet@hmrag.com

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