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 - Expression Methods

Members of this group are responsible for the enhancement and further development and implementation of our methods for expression (DNA and protein) analysis. Applicants for these positions must understand the statistics underlying these methods and be able to link results to sequence analysis, physical and functional classifications and metabolic/signaling pathway data systems. Additional skills include statistics, solid programming skills (e.g. C/C++ or Java) and an working knowledge of expression analysis technologies

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:

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For more information on our company visit our website at http:// www.hmr.de
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