



SYGNIS Bioscience GmbH & Co. KG is a wholly-owned subsidiary of SYGNIS AG, listed in the Prime Standard of the German Stock Exchange. It is specialized in the development of novel products in Molecular Biology focusing on polymerases with multiple applications in DNA- and RNA- amplification and NGS sequencing.

We have an immediate vacancy for a

### **Bioinformatician (m/f)**

#### Major Responsibilities:

- Conduct resequencing / de novo assemblies / transcriptome analyses
- Analyze characteristics of assemblies (coverage, biases, SNVs, CNVs etc.)
- Support planning of NGS-based experiments
- Responsible for maintenance of the Linux-based bioinformatic analyses tools (system / hardware)

#### Your Qualifications:

- Profound bioinformatics background and hands-on experience
- Junior level (e.g. after master's thesis) welcome to apply
- Good knowledge of Linux
- Programming skills in either Perl, Java, or C++
- Experience with common (command line) tools and platforms for sequence manipulation and analyses, able to adapt and tweak workflows for non-standard questions
- Some statistical knowledge is welcome, R is a plus

We offer an inspiring working atmosphere in a small and dedicated team developing novel NGS tools.

If you are interested please send your application or inquiries for the job by Email to:

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