

Computational Biologist / Bioinformatician / NGS Data Analyst



The Stark group at the Research Institute of Molecular Pathology (IMP) is looking for a Computational Biologist / Bioinformatician / NGS (Next Generation Sequencing) Data Analyst.

We are an internationally leading group in genomics and the regulation of gene expression, combining genome-wide experiments (e.g. STARR-seq, ChIP-seq, RNA-seq, PRO-seq), deep sequencing with on-site Illumina machines, and bioinformatics analyses.

To support our projects, we are looking for a highly motivated individual with a Masters or PhD degree in bioinformatics or related areas and a strong expertise in Unix, R, statistics, NGS data analyses, version-control, and Slurm-based compute clusters or similar. If you are passionate and excited to improve your skills and apply them to exciting biological questions in an outstanding environment, you're the one!

The IMP is a world-class research institute that provides outstanding support, competitive salaries, and excellent social benefits in Vienna, Austria – the number one city in the Mercer 'quality of living' ranking.

Applications with a cover letter, CV, and the names of your referees should be sent to manuela.steurer@imp.ac.at by September 1st 2020. The position is available immediately.

Selected publications

Haberle et al., Nature 2019; Arnold et al., Nature Biotech. 2017; Stampfel et al., Nature 2015; Zabidi et al., Nature 2015; Yanez-Cuna et al., Genome Res. 2014; Arnold et al., Science 2013

Additional information

<http://www.starklab.org>

<http://www.imp.ac.at>

