

Bioinformatics @ Stark Group IMP Vienna



Bioinformatician / NGS Data Analyst

The Stark group at the Research Institute of Molecular Pathology (IMP) is looking for a 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), deep sequencing with on-site Illumina machines, and bioinformatics analyses. 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.

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 & interest in Unix, programming, R, statistics, and NGS data analyses. If you are passionate, hard working, and excited to learn and improve your skills in an outstanding research environment, you’re the one!

Applications with a cover letter, CV, and the names of your referees should be sent to manuela.steurer@imp.ac.at by August 31st 2017. Interviews will follow shortly after this date.

Selected publications

Arnold & Zabidi et al., *Nature Biotechnology* 2017; Stampfel et al., *Nature* 2015; Zabidi & Arnold et al., *Nature* 2015; Arnold & Gerlach et al., *Nature Genetics* 2014; Kvon et al., *Nature* 2014; Shlyueva et al., *Molecular Cell* 2014; Zabidi & Arnold et al., *Nature* 2014; Yanez-Cuna et al., *Genome Research* 2014; Arnold et al., *Science* 2013

Additional information

<http://www.starklab.org>

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

<https://www.wien.gv.at/english>

