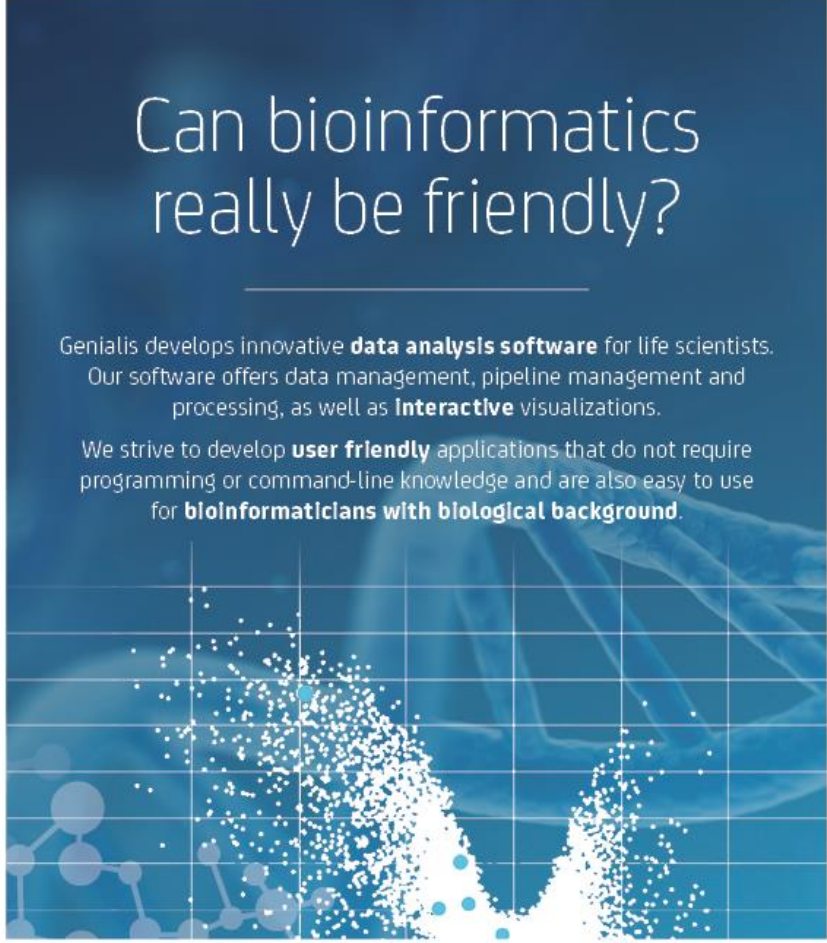


On November 27th **GENIALIS** will present a proprietary software for microarray and NGS sequencing data analysis

Come at 11:00 in room 6203 (PRC building) !



Can bioinformatics
really be friendly?

Genialis develops innovative **data analysis software** for life scientists. Our software offers data management, pipeline management and processing, as well as **Interactive** visualizations.

We strive to develop **user friendly** applications that do not require programming or command-line knowledge and are also easy to use for **bioinformaticians with biological background**.

Come and make your life easier!

November 27th at 11:00 in room 6203 (PRC BUILDING)

Genialis
www.genialis.com | info@genialis.com | +386 59 054 580

Genialis is a bioinformatics spin-out from Laboratory of Bioinformatics, Faculty of Computer and Information Science, University of Ljubljana. Genialis' mission is to offer friendly bioinformatics. Genialis has a team of life scientists, bioinformaticians and computer scientists that completely understands problems in bioinformatics.

Genialis develops innovative **data analysis software** for analyzing Next Generation Sequencing and Microarray data. Software offers support for open source tools for genomic data analysis and management, pipeline management and processing. It provides experienced bioinformaticians with full control over analysis pipelines. Also, Genialis' software supports **interactive visual browsing** through data, to get biological answers faster and easier. We strive to develop **user friendly** web applications that do not require deep programming or command-line skills and are also **easy to use for bioinformaticians with biology background**. Genialis has already established commercial and research collaborations with Garvan Institute of Medical Research (Sydney) and Baylor College of Medicine (Houston).