

RBC: Fkz 031A538C RNA Bioinformatics Center: BIMSB The Berlin Institute for Medical Systems Biology

Bora Uyar, Vedran Franke, Ricardo Wurmus, Dan Munteanu, Wolfgang Kopp, Nikolaus Rajewsky, Uwe Ohler, Altuna Akalin The Berlin Institute for Medical Systems Biology, Max-Delbrueck Center for Molecular Medicine, Berlin, Germany

Short description of the project

We offer tools, services and training for the analysis of RNA-binding proteins and post-transcriptional regulation.

de.NBI services

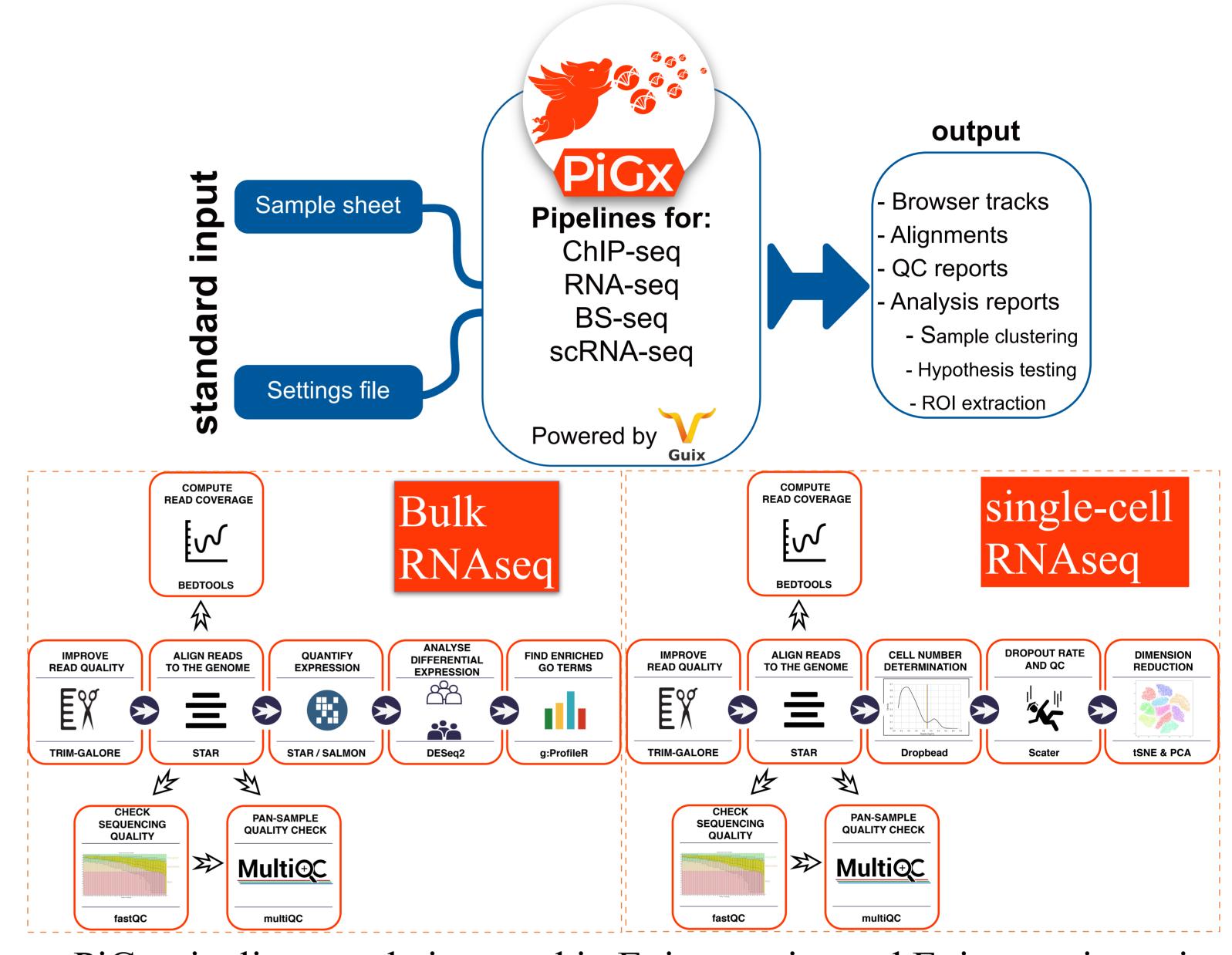
| | Service | Description |
|-----------------|------------------|---|
| Data- bases | Galaxy Server | Galaxy server at MDC |
| | doRiNA | A database of RNA interactions |
| | circBase | A database to explore and discover circular RNAs |
| RNA (Seq) Tools | PiGx | RNAseq and single-cell RNA-seq pipelines |
| | RCAS | RNA-centric annotation system |
| | NASTIseq | identifies cis-Natural Antisense Transcripts from RNA-seq data |
| | RiboTaper | Ribo-Seq analysis pipeline |
| miRNA Tools | PIPmiR | identifies novel plant miRNA genes from deep sequencing data |
| | PicTar | Web resource for microRNA targets |
| | miRDeep | microRNA detection tool in deep sequencing data |
| | miReduce | discovers motifs in mRNAs and correlates with gene expression changes |
| RBP Tools | PARalyzer | PAR-CLIP sequencing analysis |
| | micro- MUMMIE | microRNA target-site prediction in PAR- CLIP data |
| | cERMIT | finds optimal motifs in high-throughput ChIP or RIP datasets |

General information on the project

- No. of staff paid from de.NBI grant: 2
- Other staff involved: 3

Progress report

Developed and Maintained Software



PiGx pipelines are being used in Epigenomics and Epitranscriptomics QC group (EpiQC) as part of the FDA's SEQC2 Study.

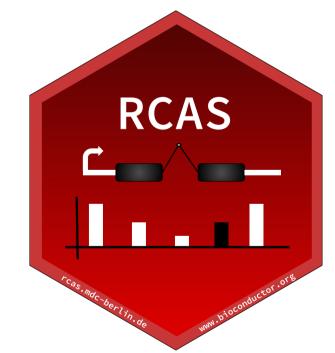
Local Galaxy Server at MDC

> 100 Total Users



RCAS: RNA-Centric Annotation System

- ~ 120 monthly downloads on Bioconductor
- Packaged using Conda, Guix, and available on the local Galaxy Server



doRiNA database of RNA interactions doRiNA



de.NBI Training and education

Provided one-on-one mentorships and consultations

Selected Publications

Wyler, Emanuel, **Vedran Franke** et al. 2019. "Single-Cell RNA-Sequencing of Herpes Simplex Virus 1-Infected Cells Connects NRF2 Activation to an Antiviral Program." Nature Communications 10 (1): 4878.

Müthel, Stefanie, **Bora Uyar** et al. 2019. "The Conserved Histone Chaperone LIN-53 Is Required for Normal Lifespan and Maintenance of Muscle Integrity in Caenorhabditis Elegans." Aging *Cell* 18 (6): e13012.





