

# All you need to know about resources, services & collaboration opportunities provided

## Bioinformatics infrastructure

### Platform provides computational and software resources for computational and wetlab biologists



**What:** shiny server for interactive R applications  
**For whom:** drylab (developing the app) + Wetlab (using the app)  
**Links:** <http://shiny.mdc-berlin.de>  
**How to get help:** <https://itsupport.mdc-berlin.net/>



**What:** In-house mirror of UCSC genome Browser  
**For whom:** drylab + Wetlab  
**Links:** <http://genome.mdc-berlin.de>  
**How to get help:** <https://itsupport.mdc-berlin.net/>



**What:** In-house instance of Galaxy Server  
**For whom:** all MDC users  
**Links:** <https://galaxy.mdc-berlin.net>  
**How to get help:** <https://itsupport.mdc-berlin.net/>



**What:** Reproducible scientific software for linux desktops & HPC  
**For whom:** People who need to install scientific software  
**Links:** <http://guix.mdc-berlin.de/documentation>  
**How to get help:** <https://itsupport.mdc-berlin.net/>



**What:** App servers, Workstations, all running a Linux Distribution (Ubuntu or RedHat based )  
**For whom:** BIMS users  
**How to get help:** <https://itsupport.mdc-berlin.net/>



**What:** Dedicated computational servers (assistance on deciding what to by → support until the server is retired)  
**For whom:** BIMS groups  
**How to get help:** [bimsb\\_itsupport@mdc-berlin.de](mailto:bimsb_itsupport@mdc-berlin.de)



**What:** Highly resources Virtual Machines for running scientific tools, webtools (with public access)  
**For whom:** All BIMS Users  
**How to get help:** [bimsb\\_itsupport@mdc-berlin.de](mailto:bimsb_itsupport@mdc-berlin.de)



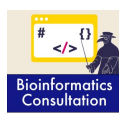
**What:** up to 25 training laptops preinstalled with the requested tools  
**For whom:** all training participants  
**How to get help:** [bimsb\\_itsupport@mdc-berlin.de](mailto:bimsb_itsupport@mdc-berlin.de)



**What:** data sharing over https  
**For whom:** All BIMS Users  
**Links:** <https://bimsbstatic.mdc-berlin.net/> / .de  
**How to get help:** [bimsb\\_itsupport@mdc-berlin.de](mailto:bimsb_itsupport@mdc-berlin.de)

## Consultation, Mentorship & Training

### Platform provides know-how for research and software projects in different formats



**What:** Consultation related to bioinformatics and data analysis related questions  
**For whom:** Researchers who are looking for advice on how to proceed on data analysis  
**How to get access:** <http://iris.mdc-berlin.de>



**What:** one-2-one mentorship to develop computational skills  
**For whom:** People who have some computational skills and want to improve them with a mentor  
**How to get access:** <http://iris.mdc-berlin.de>

#### Learning resources maintained by the platform:

- ✓ Linux+HPC tutorial: <http://bioinformatics.mdc-berlin.de/intro2UnixandSGE/>
- ✓ Computational Genomics with R book: <http://compgenomr.github.io/book>
- ✓ biostars forum for MDC (Q&A forum): <http://biostar.mdc-berlin.net/>
- ✓ PiGx pipeline manuals (Genomics pipelines): [http://bioinformatics.mdc-berlin.de/pigx\\_docs/](http://bioinformatics.mdc-berlin.de/pigx_docs/)

## Research Collaboration

### We provide multiple avenues for research collaboration

#### Large-Scale collaboration

**What:** These are projects have large-scale data and/or complex analysis requirements. Needs significant time investment.

**How to get access:** consultation request  
<http://iris.mdc-berlin.de>

#### Small-Scale collaboration

**What:** These are small-scale projects where someone in the lab has specialized expertise, set of existing scripts or established pipelines such as PiGx  
<https://bioinformatics.mdc-berlin.de/pigx> .

**How to get access:** consultation or small-scale project request <http://iris.mdc-berlin.de>