

The what, how and who in the **Bio-IT** project

"Community initiative which aims to build, support, and promote computational biology activity at EMBL"

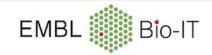
Why Computational Biology

- 1. **Integral to experimental design**: support experimental design planning and data processing.
- 2. **Necessary to scale up**: to process and store big data, to compare data in different context, even from small-scale experiments.
- 3. **Data (and science) doesn't end at publication**: ensure scientific FAIRness by properly storing your data and processes and making them findable and available.
- 4. **Data-driven research**: advanced computational techniques applied to biology define a new scientific paradigm.



Why Computational Biology at EMBL

- 1. EMBL is committed to **Open Science** policies.
- 2. A **dynamic research context** such as EMBL needs FAIR data and processes to ensure usability.
- 3. In a **collaborative environment** researchers can easily take advantage of each others' skills: design experiments to spare time and money and achieve better science.
- 4. Even more because of EMBL's **internationality**: wide network of collaborations and data sources.
- 5. A specific interest of EMBL scientists is populating and maintaining biological **databases and tools**.





"Community initiative which aims to build, support, and promote computational biology activity at EMBL."



Training and consulting

- Courses and seminars
- Course materials
- Consulting sessions



Infrastructure and resources

- GitLab
- Coding platforms and tools



Community building and networking

- Clubs
- Grassroots
- Community blog



Knowledge aggregation and dissemination

- EMBL centers / IT services info
- WikiHows
- Newcomers guide



Bio-IT community

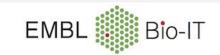
We will present the Bio-IT community through personas.



Personas disclaimer: "all persons fictitious"

In this context, personas are fictional characters used to present different way to interact with the community.





The Observer

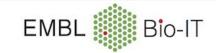


The Observer

- Mostly applies **wet lab biology** techniques in experiments
- Sometimes **uses computational tools** through their graphical interfaces to make some predictions
- Got interested in learning how to query them programmatically and tried writing some code

To do so, ended up in Bio-IT website





The Observer

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Consulted the list of Bio-IT resources to choose the right platform Consulted the list of Bio-IT resources to choose the right platform

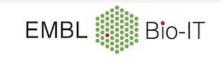
Uses the chat for consulting



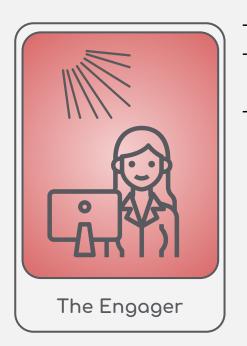
@ Bio-IT
@ Bio-IT announce
@ Coding Club
@ Python
@ R_and_statistics

Attends external and internal courses



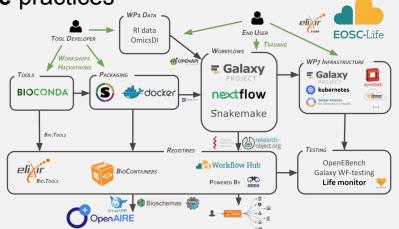


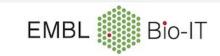
The Engager



Strong computational skills

- Developer of computational **tools** for biological data analysis
- Promotes open science practices
 - 🦊 EMBL GitLab
 - 💶 EMBL Galaxy





The Engager

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Chairs and presents in clubs



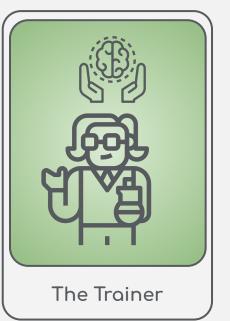


- Shares coding stories and resources in chat channels
- Proposes and leads themed seminars
- Joins hackathons
- Sometimes joins courses as trainer





The Trainer

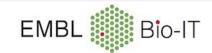


Leads a research group

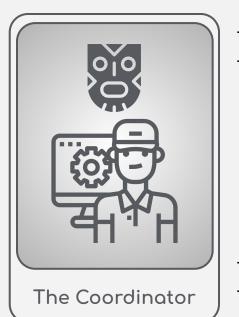
- Has wide experience in computational biology
- Uses computational tools from **command line** for exploratory analysis
- Has little time to join Bio-IT activities, but keeps updated
- Is listed in the Grassrots consulting table
- Follows and interacts with the Twitter profile @EMBL_BioIT

Trainer in external and internal courses

EMBL Bio-IT Project @EMBL_BioIT Building and supporting the #bioinformatics/#compbio community at @embl. Training; networking; resources; information; community.



The Coordinator



- Enjoys coding and testing new tools and resources
- Maintains Bio-IT resources, admin of GitLab



- Maintains Bio-IT website
- Chair of meetings and clubs, organiser
- Available for consulting via chat or in the Bio-IT drop-in
- Organises small groups sessions for discussion



Thank you! Thank you! @paladin #engage Thank you! #stay informed @ralves Thank you! Thank you! chat.embl.org Thank you! bio-it.embl.de @ Bio-IT Thank you! **@** Bio-IT announce Thank you! - Read the blog @ Coding Club Thank you! - Check upcoming courses / activities * **@** Python Thank you! Thank you! - Check the news about the resources **@** R and statistics @EMBL_BioIT Thank you! Thank you! Thank you! Thank you! MONDAY TUESDAY WEDNESDAY THURSDAY FRIDAY Thank you! Thank you! BIO-IT drop-in Thank you! BIO-IT 10-12 AM Thank you! taskforce Thank you! #join Thank you! 2-3 PM Coding Club Thank you! (R or Python) Thank you! Thank you! 4-5 PM * EPUG community discussion (28 September) Thank you! * Git Week (15-19 November) Thank you! bio-it@embl.de Thank you! * Bio-IT Hackathon (1-2 December) Thank you!