

Defining training requirements for biomolecular researchers with high computational needs

BioExcel Educational Webinar Series #9

Presenters: Cath Brooksbank

Host: Adam Carter

23 November, 2016

Partners



Funding



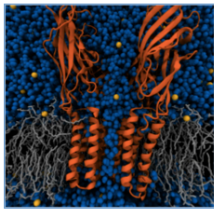


This webinar is being recorded

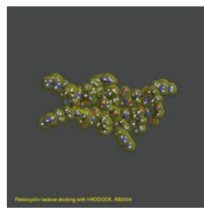
BioExcel Overview

- **Excellence in Biomolecular Software**

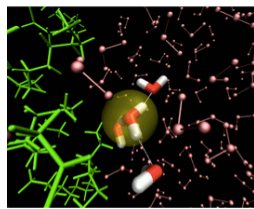
- Improve the performance, efficiency and scalability of key codes



MD simulations
/GROMACS/



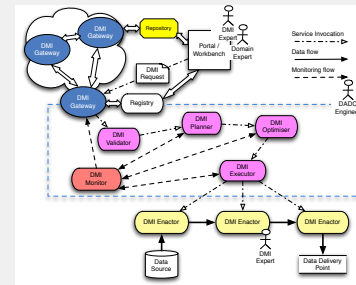
Docking
/HADDOCK/



QM/MM
/CPMD/

- **Excellence in Usability**

- Devise efficient workflow environments with associated data integration



Key Workflows
and Platforms



- **Excellence in Consultancy and Training**

- Promote best practices and train end users



Interest Groups

- Integrative Modeling IG
- Free Energy Calculations IG
- Best practices for performance tuning IG
- Hybrid methods for biomolecular systems IG
- Biomolecular simulations entry level users IG
- Practical applications for industry IG
- **Training**
- Workflows

Support platforms

<http://bioexcel.eu/contact>



Forums



Code Repositories



Chat channel

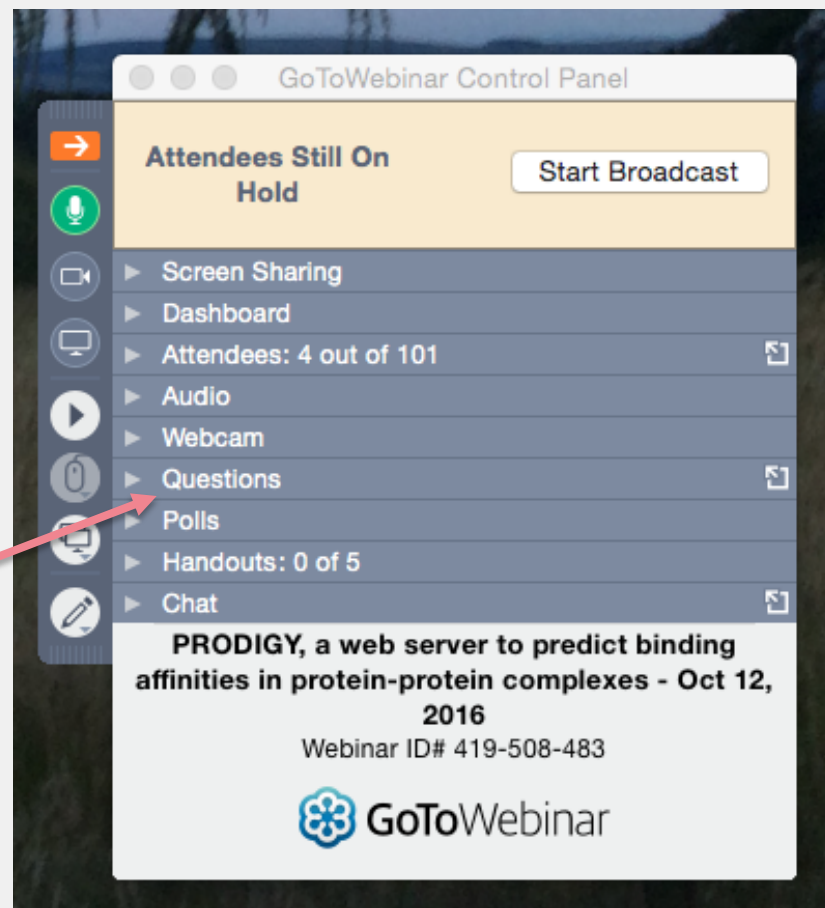


Video channel

Audience Q&A session

Please use the **Questions** function in GoToWebinar application

Any other questions or points to discuss after the live webinar? Join the discussion the discussion at <http://ask.bioexcel.eu>.



Today's Presenter



Cath Brooksbank

<presenter's slides here>

Objectives for this webinar

- Gain an overview of BioExcel and its goals
- Appreciate how to perform a simple training needs analysis as a precursor to developing a new training programme
- Discover how you can provide input into BioExcel's training programme
- Interact with others who have an interest in training for molecular life scientists, to learn from them and share your experience with them

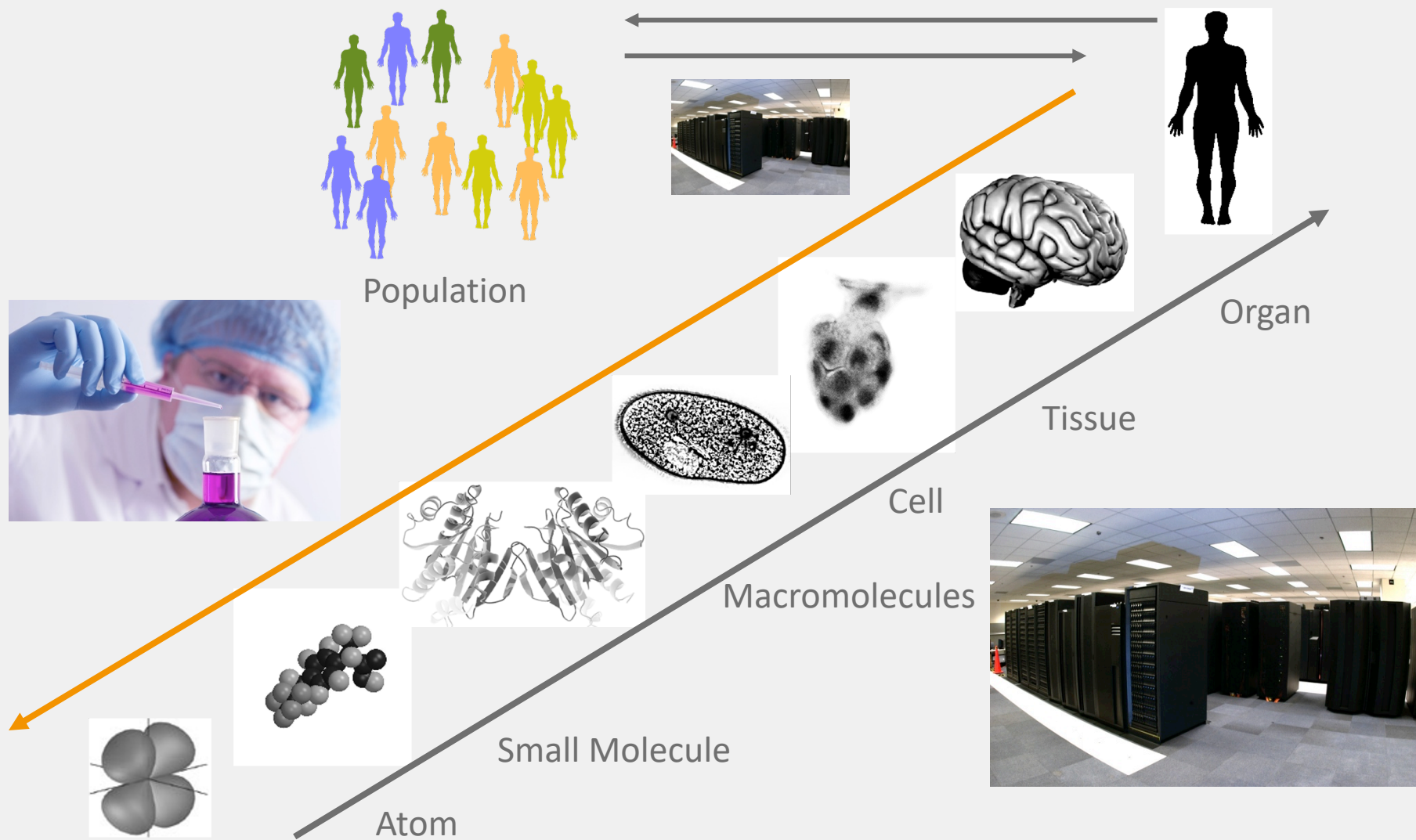
Poll 1

What is your motivation for joining this webinar?

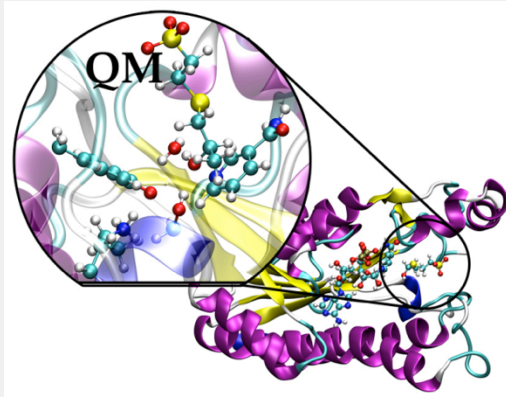
- To learn about BioExcel in general
- To learn about the BioExcel Training Interest Group
- To learn about our approach to performing a training needs analysis
- Other (please tell us using the chat function)

What is BioExcel?

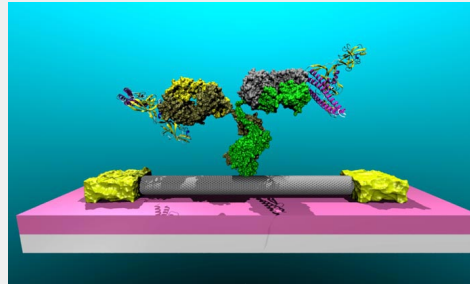
Life science is becoming computationally intensive



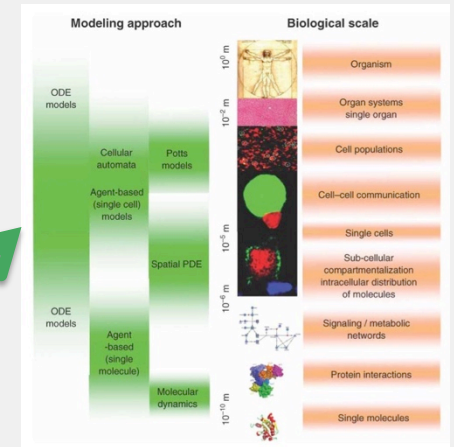
Electronic structure



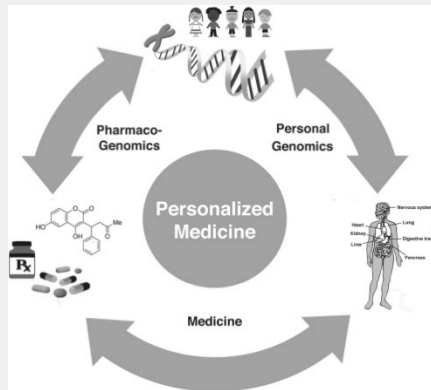
Biomarkers design



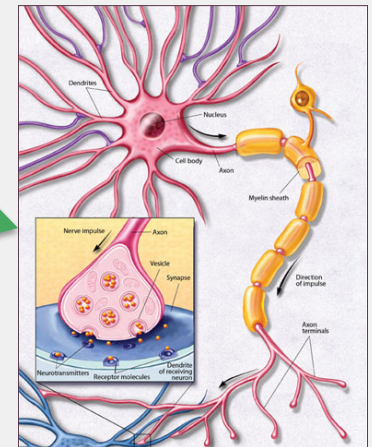
Physiology



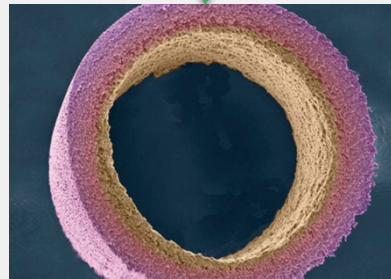
Biomolecular Modelling and Simulations



Personalized medicine



Neuroinformatics



Biomaterials science and nanotechnology

BioExcel Center of Excellence

SOLUTIONS

SOFTWARE

Widely used, fast and scalable codes for integrative modelling and molecular simulations

CORE DEVELOPERS

The scientists who wrote the code and know it best work with us!

WORKFLOWS

User-friendly and efficient systems for workflow executions and data processing



SERVICES

TRAINING

Webinars, "ask-me-anything" sessions, hands-on workshops for everyone from newbies to advanced users

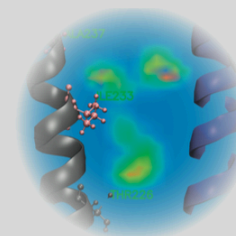
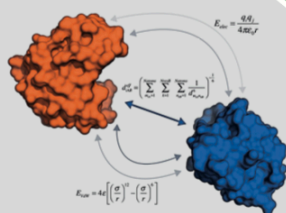
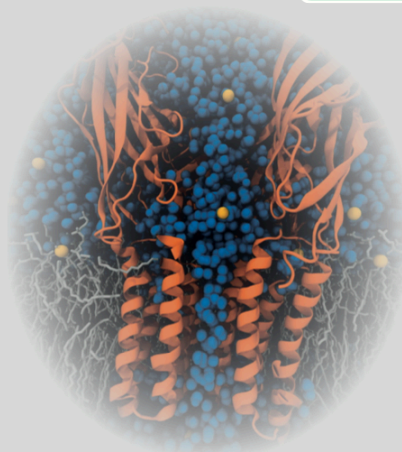
CUSTOMISATION

Tailored solutions adapted to your needs

CONSULTANCY

Personalized support with software usage, tuning and scientific aspects of the research

ACADEMIA INDUSTRY



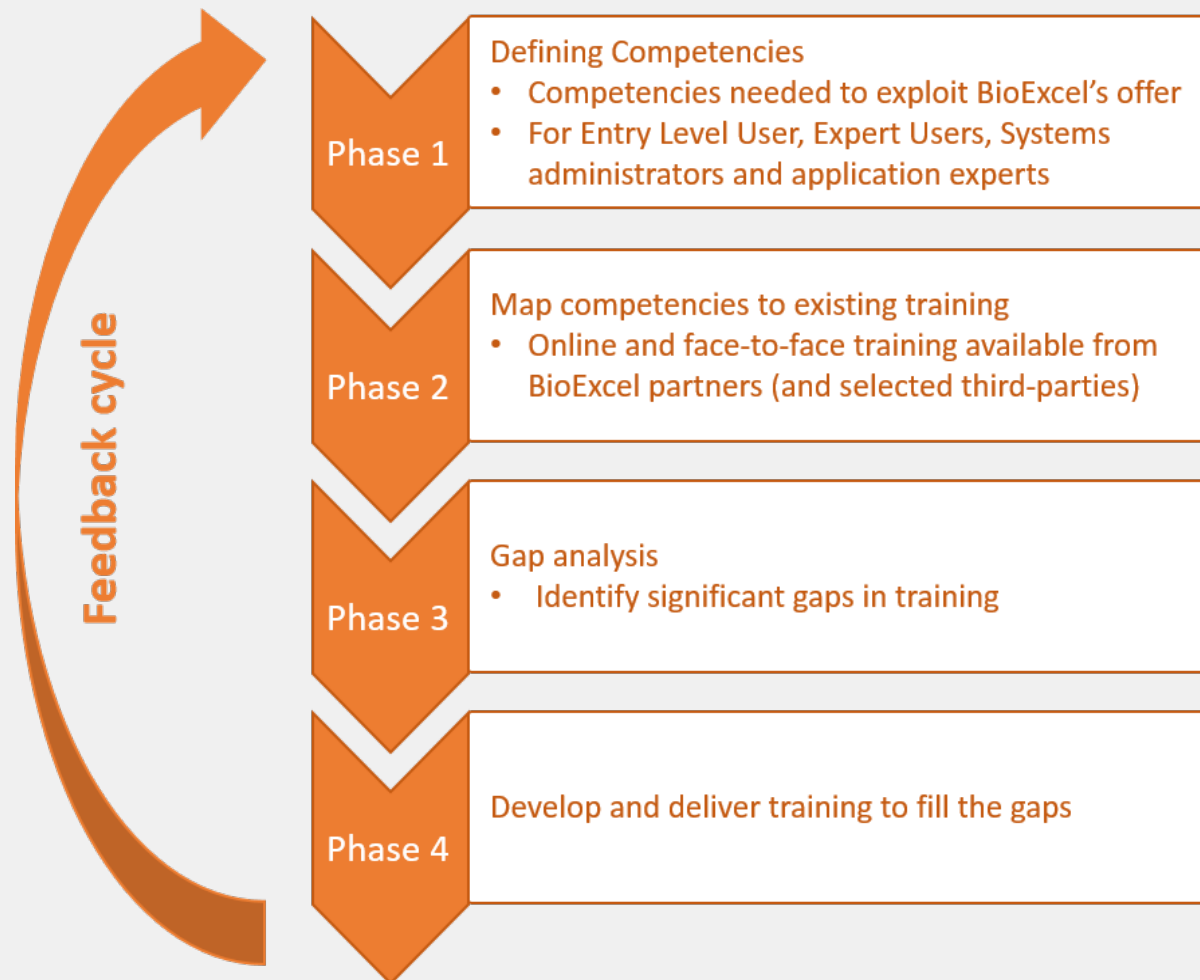
How are we defining BioExcel's training requirements?

Poll 2

What's your professional background?

- Life sciences
- Computer science
- Learning and development
- Other (please tell us using the chat function)

BioExcel Training Programme



What is a Competency?

Competency is ‘an observable ability of any professional, integrating multiple components such as knowledge, skills, values and attitudes’.

Acquisition can be validated objectively.

Shared ‘currency’ applicable to learning of all types and at all career stages

Competency profile

Defines the competencies required to fulfil a particular role

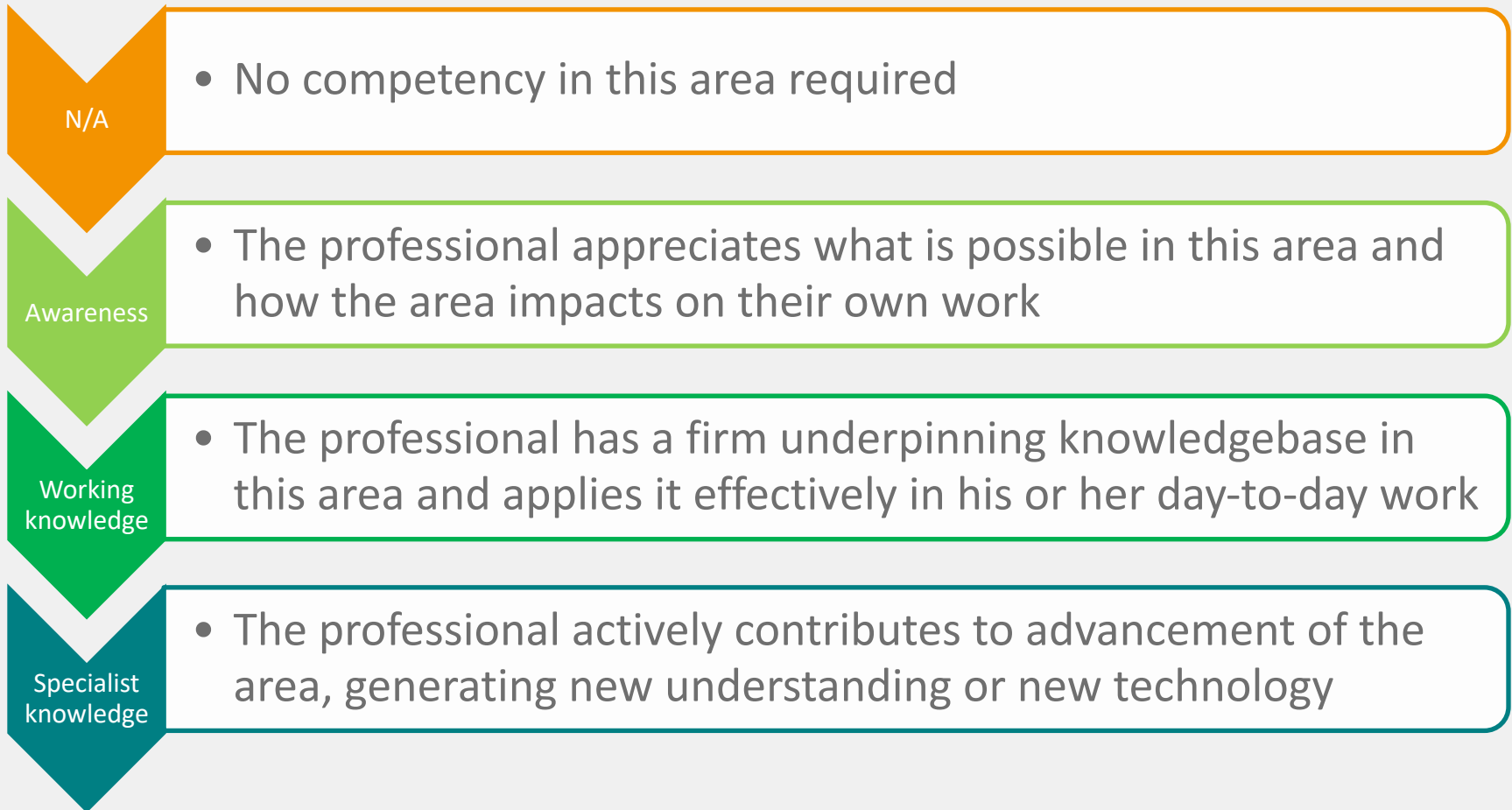
Typically defined by professional bodies / learned societies in collaboration with employers

An example...

"Write his/her own scripts to perform tasks in context of biomolecular research"

Knowledge	Skills	Behaviour
Knowledge of existing commands/libraries to re-use	Is able to automate the process of executing processes remotely	Uses appropriate scripting languages
Judges when a task should be automated	Write & debug scripts	

What level of competency is required?



How should their competency develop as they progress through their career?

Apprentice

- has the prerequisite knowledge but hasn't gained experience of applying it in the workplace

Journeyman

- has some experience of applying knowledge and is in the process of gaining further experience

Master

- Has sufficient mastery of the role to be able to coach an apprentice to journeyman level

The process



BioExcel competency profile

Group related competencies together

- Generic competencies (5)
- Scientific competencies (13)
- Generic computing competencies (8)
- Parallel computing competencies (5)

Different types of users

- Entry-level user
- Specialist user
- Systems administrator / applications expert

→ You might be entry level for some competencies and specialist for others

Method and competency terminology have been aligned with similar international initiatives

Competency-based projects



Generic competencies (5)

Highlighted the most important ones for this group

Comprehend and comply with professional, ethical, legal, security and social issues and responsibilities

Function effectively in a team

Communication

Continuing Professional Development (CPD)

User needs

Scientific competencies (13)

Distributed data management & data management planning

Impact of HPC/HTC

Computing expertise

Expertise in formal & natural sciences

Licensing policy

Scientific process

Use computer-based systems

Evaluate computer-based systems

Find, read & assess literature

Data-driven science

Identify and compile datasets

Present your results

Data management

Generic Computing Competencies (8)

Operating system

Write/adapt programs

Good programming practice

Install biomolecular software

Identify & define computing requirements

SYSTEMS MONITORING

Write scripts

Deploy & test non-commercial software

Parallel computing competencies (5)

Workflow systems

Batch systems

Performance profiling

Parallel programming

**Advantages & limitations for
deploying, executing and
optimising computations in
cloud/grid environment**

Want a closer look?

Competency survey

goo.gl/dEvs8N

Full profile

goo.gl/8OqKYK

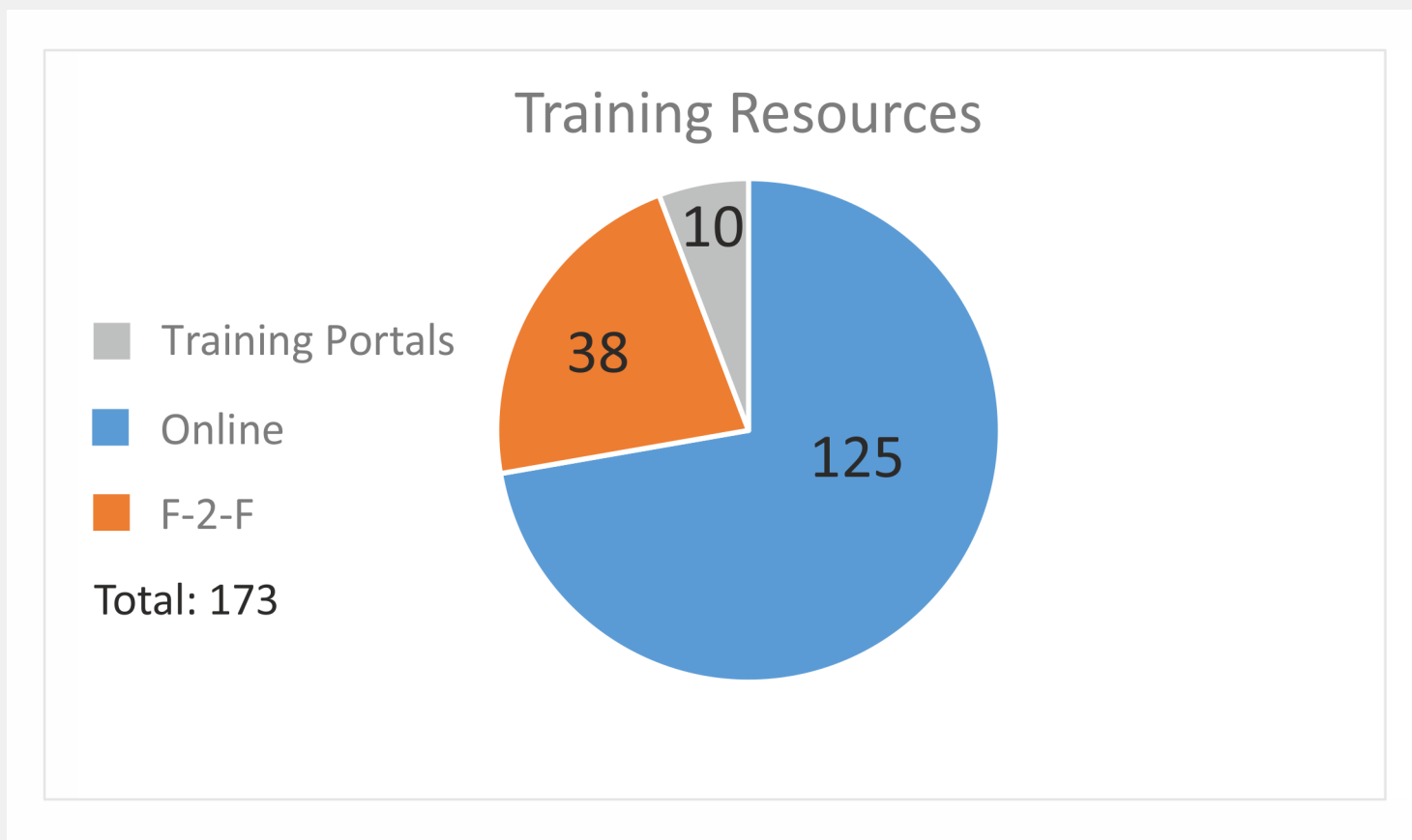
Let us know what you think!

What will this will be used for?

- **To develop high-value training materials** and adapt existing ones
- As components of **VM-based pre-packaged bundles** of applications and inputs for user-friendly exercises
- **To incorporate materials into workshops** at major Infrastructure events (e.g. PRACE, EGI, ELIXIR; INSTRUCT) or life science conferences (e.g. ECCB)
- **To collaborate with major related training activities** of PRACE, ELIXIR, CECAM, Pistoia Alliance and CCP-BioSim for reaching out to larger user communities

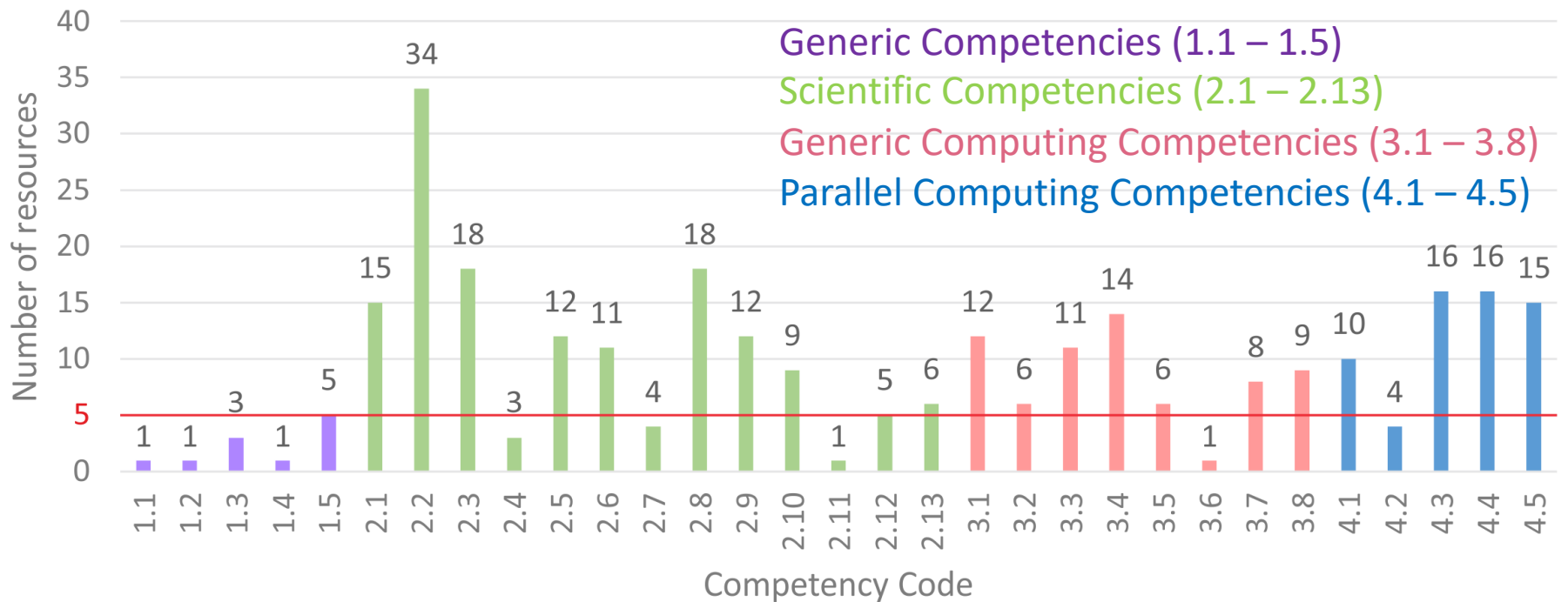
What kinds of training do we need to develop?

Identifying training resources



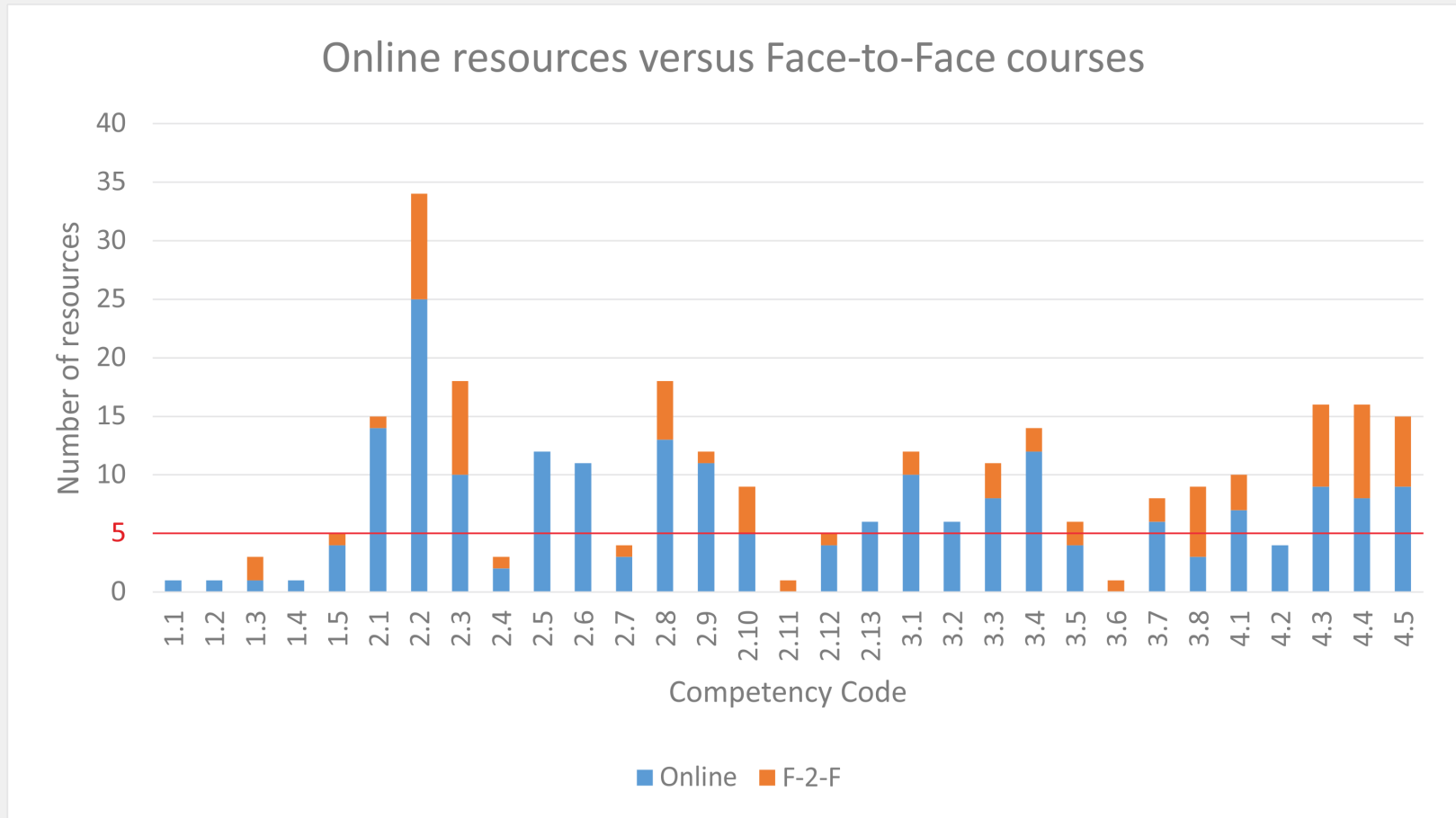
Mapping

Number of resources mapped to each Competency



5 (total training resources) picked as a minimum viable number for coverage

Online versus Face-to-Face



Gap Analysis

<p>Key</p>	<p>Insufficient training resources identified but outside BioExcel scope</p>	<p>BioExcel or External training resources identified</p>	<p>Insufficient training resources identified</p>
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Side notes on Mapping & Gap Analysis

- Falsely optimistic view due to **partial mapping** of resources to competencies
- **Suitability** of the training resources for the BioExcel User Groups (entry-level users, expert users, and systems administrators)
 - Likely only internal or project associated resources

Gap Analysis – insufficient coverage

2. Scientific Competencies

2.4 Comprehension of, and compliance with, licensing policy.

2.7 Comprehension of the local and global impact of high-performance computing (HPC) and high-throughput computing (HTC) on individuals, organizations, and society.

2.11 Comprehension of, and compliance with, best practice in distributed data management and data management planning.

2.13 Presenting your results to the community (writing papers, conference presentation, YouTube).

Gap Analysis – insufficient coverage

3. Generic Computational Competencies

3.2 Analyse a problem and identify and define the computing requirements appropriate to its solution (e.g., define algorithmic time and space complexities and hardware resources required to solve a problem).

3.3 Apply knowledge of the operating system.

3.6 Install biomolecular simulation software on his/her computer.

3.7 Deploy and test non-commercial software, including software that is built collaboratively and on a volunteer basis.

3.8 Apply knowledge of systems monitoring (e.g. queue monitoring, systems availability and optimisation, storage used; scheduling maintenance at appropriate times and communicating this to users).

Gap Analysis – insufficient coverage

4. Parallel Computing Competencies

4.2 Apply knowledge of batch system

BioExcel Training Programme

- **Entry Level Users:** Focus on bridging the gap between life-science and HPC/HTC computing (users transitioning from a bench-based molecular life science background).
- **Expert Users:** training needs for this group will probably be quite specific for each use case (e.g. specific software packages, optimisation). We anticipate input from the Interest Groups to inform highest priority training needs
- **Systems Administrators & Application experts:** Goals include Enabling communication between this group and users in addition to technical training

Poll 3

Which (if any) BioExcel user group do you identify most closely with?

Entry level user

Expert user

Sys admin/application expert

None (Please use chat to specify an alternative group)

BioExcel Training Programme

Face-to-Face training events

- 3-4th May 2016 - BioExcel: addressing training needs for advanced simulations in biomolecular research, EMBL-EBI
- 20-21st October 2016 - BioExcel: Workflow Training for Computational Biomolecular Research, BSC/IRB
- **10-13th April 2017** - PRACE & BioExcel Spring School – HPC in the Life Sciences (NAMD, AMBER, VMD, and GROMACS), KTH
- **3-7th July 2017** - BioExcel Summer School – Foundation skills for HPC in computational biomolecular research @EMBL-EBI
- **Autumn 2017/Early 2018** - Introduction to HPC for Life Scientists (with PRACE) @EPCC and/or BSC
- **Summer 2018** – Summer School 2018

BioExcel Training Programme

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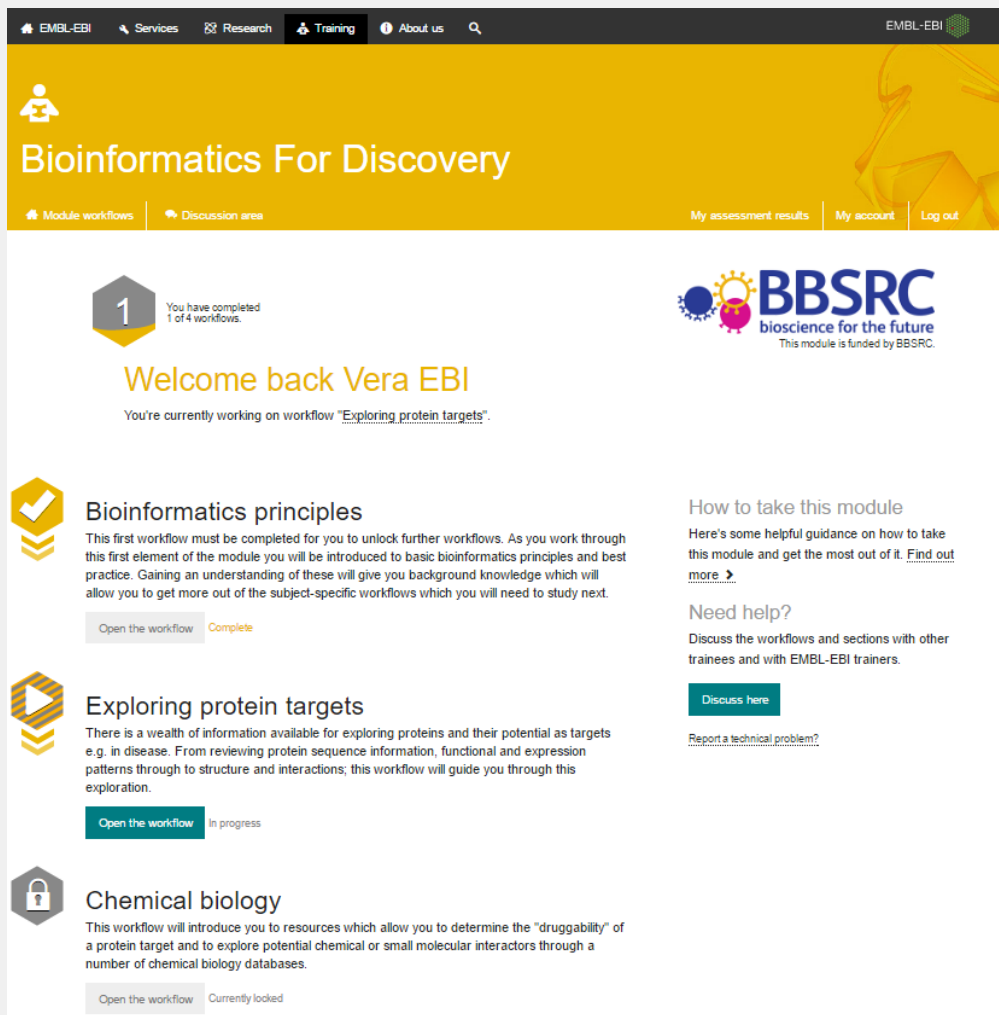
Webinars (in preparation)

- Computational resources: local computer vs Cloud / HPC
- Quality control of structures / models
- Entry level webinars (Biomolecular modelling, Molecular Dynamics etc.)
 - Aimed at general science background
 - Turn into e-learning

E-Learning (workflow-based training)

- Repurpose course material from face-to-face
- Webinars
- Bespoke online courses
- Knowledge base

Workflow-based training



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Bioinformatics For Discovery

Module workflows Discussion area My assessment results My account Log out

1 You have completed 1 of 4 workflows.

Welcome back Vera EBI
You're currently working on workflow "[Exploring protein targets](#)".

Bioinformatics principles
This first workflow must be completed for you to unlock further workflows. As you work through this first element of the module you will be introduced to basic bioinformatics principles and best practice. Gaining an understanding of these will give you background knowledge which will allow you to get more out of the subject-specific workflows which you will need to study next.
[Open the workflow](#) Complete

Exploring protein targets
There is a wealth of information available for exploring proteins and their potential as targets e.g. in disease. From reviewing protein sequence information, functional and expression patterns through to structure and interactions; this workflow will guide you through this exploration.
[Open the workflow](#) In progress

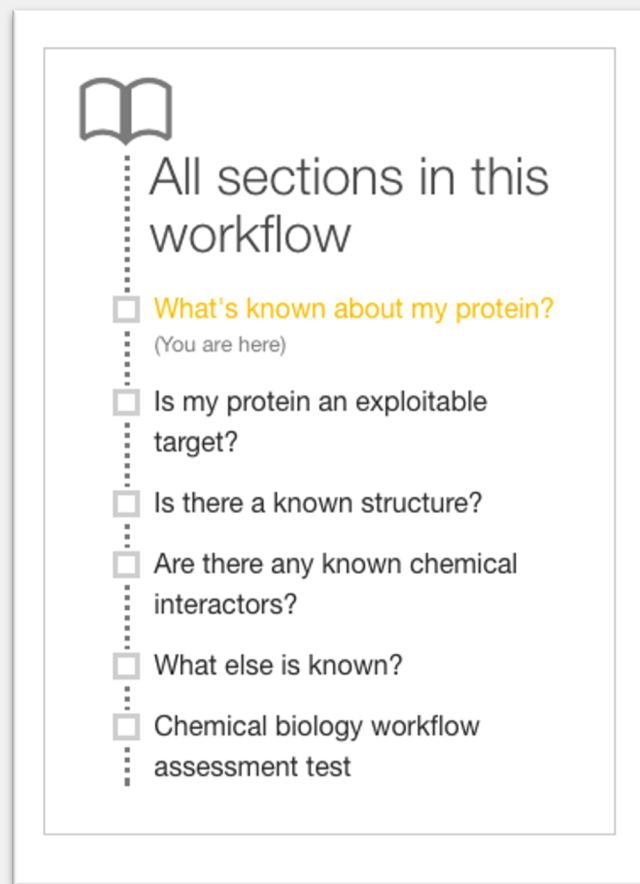
Chemical biology
This workflow will introduce you to resources which allow you to determine the "druggability" of a protein target and to explore potential chemical or small molecular interactors through a number of chemical biology databases.
[Open the workflow](#) Currently locked

BBSRC
bioscience for the future
This module is funded by BBSRC.

How to take this module
Here's some helpful guidance on how to take this module and get the most out of it. [Find out more >](#)

Need help?
Discuss the workflows and sections with other trainees and with EMBL-EBI trainers.
[Discuss here](#)

[Report a technical problem?](#)



www.ebi.ac.uk/training/events/2017/bioinformatics-discovery-0

Collaborations (Training)

Ritrain (<http://ritrain.eu/>) - mission is to improve and professionalize the training of managerial and leadership staff in research infrastructures (RIs).

CORBEL (<http://www.corbel-project.eu/>) - is an initiative of eleven new biological and medical research infrastructures (BMS RIs), includes competency-based training Work Package aimed at technical operators of BMS RIs.

ELIXIR (www.elixir-europe.org) - unites Europe's leading life science organisations in managing and safeguarding the increasing volume of data being generated by publicly funded research.

PRACE Advanced Training Centres (<http://www.prace-ri.eu/>) - The PRACE Advanced Training Centres (PATCs) provide top-class education and training opportunities for computational scientists in Europe and are the primary source for PRACE training portal materials. The BioExcel partners BSC and EPCC are PATCs and can act as liaisons to PRACE.

CompBioMed (CoE) – (<http://www.compbioimed.eu/>) A Centre of Excellence in Computational Biomedicine

Training Interest Group (TIG)

Aims:

- Improve visibility of training initiatives and individual courses/resources. Members of the Training IG will be able to act as **liaisons** to other training initiatives
- Facilitate **collaboration** between projects to promote best practice and efficient use of resources
- Improve **communication** between “computational trainers” and “life science trainers”
- Promote the importance of **high quality training**

TIG is for you if you are responsible for, or have an interest in, training within your institute or project or are a training professional (trainer, coordinator, lecturer) in a life sciences and/or computational field

TIG - plans

Webinars

- 23.11.2016: Defining training requirements for biomolecular researchers with high computational needs (IG kick-off)
- Early 2017: Engaging the hard to engage – encouraging biomedical scientists to embrace high-end computing

Face-to-face

- Feb 2017: Using containers and VMs in training
- Jul 2017: ISMB/ECCB competency workshop – BioExcel use case
- 5–7 Dec 2017: Lifelong Learning in the Biomedical sciences (EMBL Conference)

**What other activities would
you like to see in the TIG?**

Audience Q&A session

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