



What is it?

Data can be collected manually or automated in the laboratory/field or you can discover useful data through literature/repository search (see [Fact-Sheet 'Discover'](#)). GFBio offers workbenches and tools ([Diversity Workbench](#), [BExIS 2](#)) to make data collection easy for you. Data emerge through observation (during an experiment), interviews, counting, measuring with instruments, or simulation by algorithms. Researchers use different technologies like tape- or video-recorders, (portable) computers/instruments, hand-written data sheets, questionnaires, tools, workbenches, satellite or aerial images.

How to do it?

- 1) Before collecting data, set up a collection protocol/sampling strategy.
- 2) Define which data will be created/collected. (What?, Where?, When?, Who?, How?)
- 3) Consider how many populations/individuals/tissue you need to gain significant results.
- 4) Allow for separate columns/rows per individual variable (eschew compound variables).
- 5) Define the collection methodologies and standards (see [Fact-Sheet 'Describe'](#)) to ensure that your data will be compatible with standards. GFBio is currently creating a standardization-tool to convert data to the EML-metadata standard. Use our description-tool (that will soon be implemented within the [GFBio-Portal](#)) to ensure consistency and data quality.
- 6) Choose indicative, unique file names, reflecting the contents, place and time (keep it short: 20141104_Collect_Factsheet.xlsx). Document this consistent naming convention and coding that can be used by the whole research team.
- 7) Choose appropriate software and formats that are suited for long-term preservation and reuse.
- 8) Communicate your collection protocol to the involved team members so that everybody is on the same level.
- 9) Gain comprehensive knowledge about the item to be collected and its habitat/occurrence.
- 10) Organize the logistics, e.g. gain collection permits (if required).
- 11) Set up a backup plan and save your data on secure and geographically dispersed servers.

Who does it?

Currently every data producer integrating other data or creating own data within his/her research project or as partner in research programme (like ecologists, geo-scientists, geneticists etc.).

Key elements

- Use the GFBio workbenches and tools to collect data in a consistent, systematic manner throughout the study (reliability).
- Capture and create structured Metadata (EML, ABCD) by the aid of the GFBio-Description-Tool and the Standardization-Tool (both soon available).
- Experiment, observe, measure, simulate.

Useful links

<https://www.dataone.org/best-practices> (Best-Practices-Primer)

http://www.abctaxa.be/volumes/volume-8-manual-atbi/chapter-7/Chapter_7.pdf (tissue collection)

<http://www.dcc.ac.uk/training/train-the-trainer/dc-101-training-materials> (for a deeper understanding)

<http://www.diversitymobile.net> (tool which helps to record data in the field)

<https://www.youtube.com/watch?v=nNBiCcBlwRA> (nice video about what can go wrong – not only in medical science!)