

WP5 MEETING REPORT, PARIS, 13 JUNE 2007

Were here :

Marc Brugman (ETI Bioinformatics, Netherlands, involved in Key to life, IdentifyLife, Encyclopedia Of Life, GBIF...)

Elise Kuntzelmann (LIS/UPMC – MNHN, France)

Karol Marhold (IBSAS, Slovakia)

Régine Vignes (LIS/UPMC – MNHN, France)

Comment on D5.47 and chosen tools :

Keep all the chosen tools (Delta-Access, Delta-Intkey, IKBS, Lucid, Linnaeus, Xper², Navikey) but check if EFG2 is really accessible to taxonomists.

The online editor FRIDA should be added to the list in D5.47 : allowed to create online new applications.

Discussion on draft version of Milestone « Criteria for descriptive tools » :

We have completed the list of criteria and it appeared to us that the different systems involved are probably quite equivalent and that the most important criteria is data exchange (import/export data formats).

List of taxonomists that could be interviewed for the project :

Karol Marhold will play an important role to organize these task (to contact taxonomists, to prepare the guidelines of the interviews, to interpret the feedback of taxonomists).

How to sample ?

We propose to interview three taxonomists in each town concerned.

It could be important to interview taxonomists in different kind of institutes.

We should interview someone of each EDIT institution.

People to contact :

David Taylor (biologist, Kew) already interviewed taxonomists for EDIT to analyze how they work.

Karol suggests that Paris could ask him about the content of such an interview.

Marc Brugman will join taxonomists in Netherlands.

Paris have already taken contact with taxonomists at the museum in Paris and made some interviews.

We should ask tools developers if we could contact some known users.

Taxonomists interviews should also take place during workshops concerning taxonomists : TDWG in September, GBIF in October...

Guidelines for the taxonomists interviews :

When do they use a taxonomic tool during their work ?

Interview of taxonomists that are using these tools and some that don't use.

Why do they use or don't use these tools ?

Why do they stop or continue using a tool ?

If they use a tool what is missing ?

What could be done to make them use a tool ?

Ask taxonomists about what they expect from descriptive tools

Do you use a tool for several knowledge base ?

Do you use the software alone or in community ?

How do you work with your colleagues on a knowledge base ?

For the taxonomists that don't use taxonomic tools

We could propose a document, a demo on principles, concepts, functions but not a notice of the tools.

We should prepare all the tools with the same knowledge base (maybe a zoology data set and a botany data set) in order to present it to the taxonomists interviewed.

Plan a workshop with taxonomists : try before or after the TDWG meeting in Bratislava (16 - 23 September) or beginning 2008.