MicroScope satisfaction survey 2012

#### Microscope global survey 2012

- Measure user satisfaction of MicroScope users
- Assess needs, particularly for prokaryotic metagenomics
- Tool: Lime survey
- Sent to all registered MicroScope users
- Anonymous

Started the 2<sup>nd</sup> of april 2012
2 recalls 17th april, 3rd may
Results analysed the 5th of june 2012

### Participation

• 1120 accounts / 300 « active » accounts

#### o 200 answers

- 17,9% participation / total account number
- 67% participation / « active » accounts
- Many comments and suggestions

#### Who are you ?

### Research field / area



### Current position



#### How did you hear about LABGeM's MicroScope platform ?



### How long have you been using the MicroScope platform ?



### How often do you use the platform ?



### Which type of project(s) are you working on ?



#### Easiness of the MicroScope Tools use



#### Tool offer

### Suitability of our tools for your needs

- (1) very adapted 53,55%
- (2) Rather adapted to my needs, but not completely 41,53%
- 2% Not really adapted, I generally use other resources

## Select the tool(s) you most frequently use

- o(1) Search
- o (2) BLAST
- o (3) Genome Browser / synteny
- (4) annotation editor
- o (5) Metabolic pathway
- o (6) Phyloprofile
- o (7) Gene cart
- o (8) Metabolic profile
- o (9) RGP finder
- o (10) RNA seq tools

Do you think an extension to MicroScope dedicated to the study of complex metagenomes would be useful ?



### Trainings

Do you plan to attend the 'Annotation and analysis of prokaryotic genomes using the MicroScope platform' training in 2013 ?



# Would you be interested in one (or several) of the following specialized training ?



### Ergonomics

How would you rate our website in terms of clarity (presentation and navigation) ?



How do you find the online documentation related to the MicroScope tools?



## Are you satisfied with the speed of the page display ?

