



# MicroScope: an integrated platform for the annotation and exploration of microbial gene functions through genomic and metabolic comparative analysis

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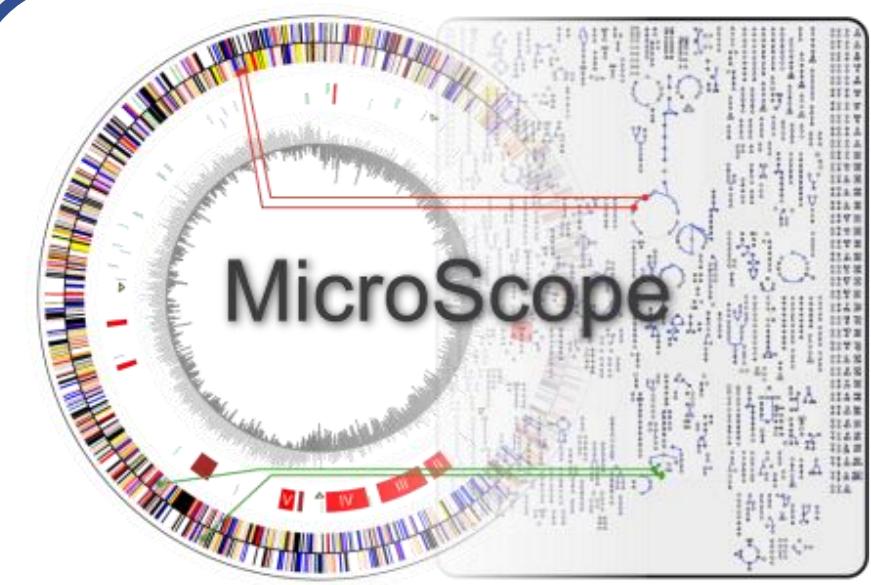
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# MicroScope<sup>\*1</sup>: an integrated platform for the annotation and exploration of microbial gene functions



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## MicroScope features

- High throughput automatic (re)-annotation of microbial genomes
- Expression and evolutionary studies by NGS data integration
- Function and biological process predictions
- Data exploration functionalities (comparative genomics, metabolism)
- Tools to assist curators in the evaluation of all available data
- Support to MicroScope users & regular user day (MicroScope Open Days)



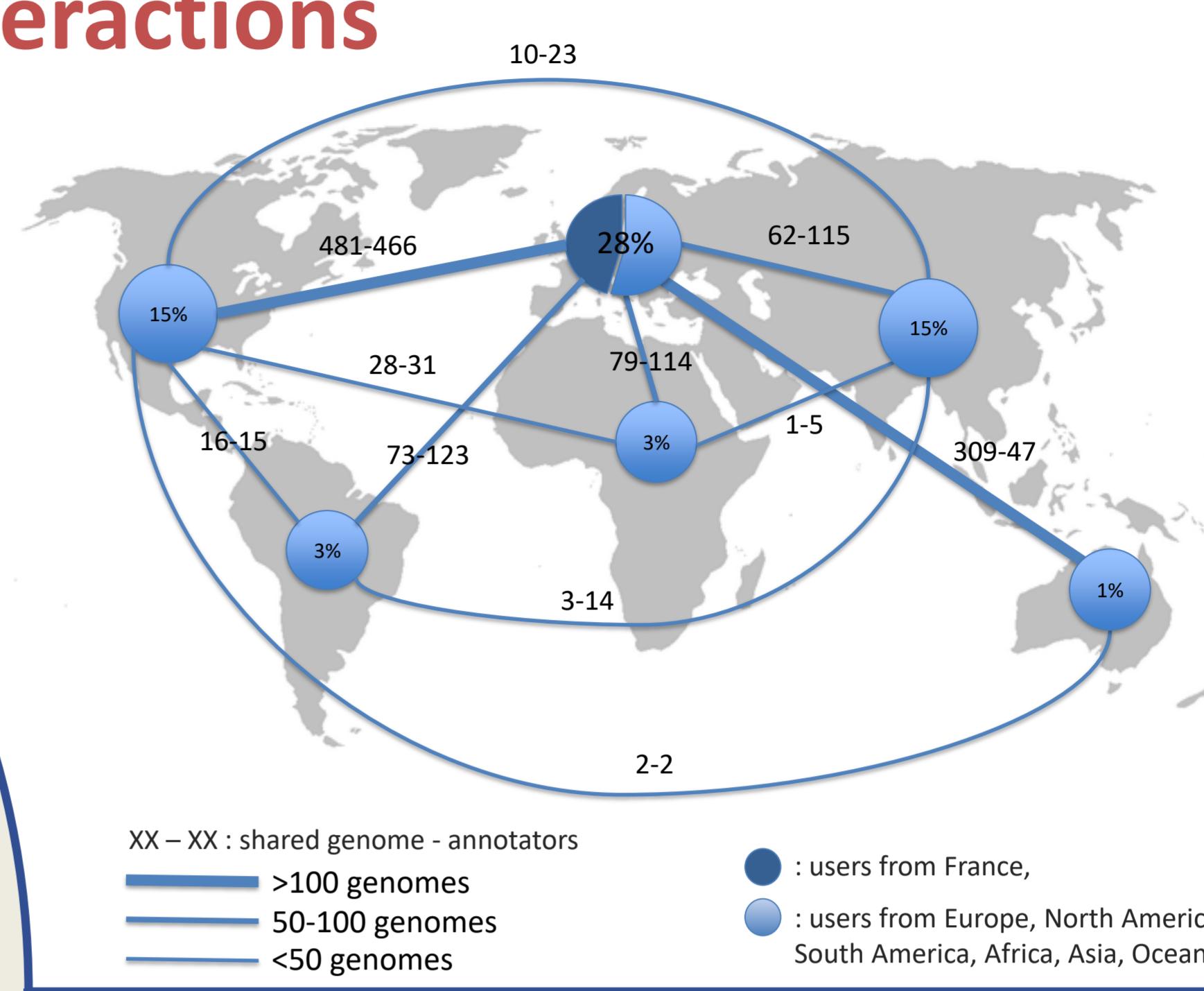
The Quality Management System of the LABGEM team has been certified according the ISO 9001:2015 and NF X50-900:2016 standards for its activities of research, developments, services and MicroScope trainings.

## Statistics

- 3 800 user accounts
- 375 active accounts per month
- 200 new genomes per month

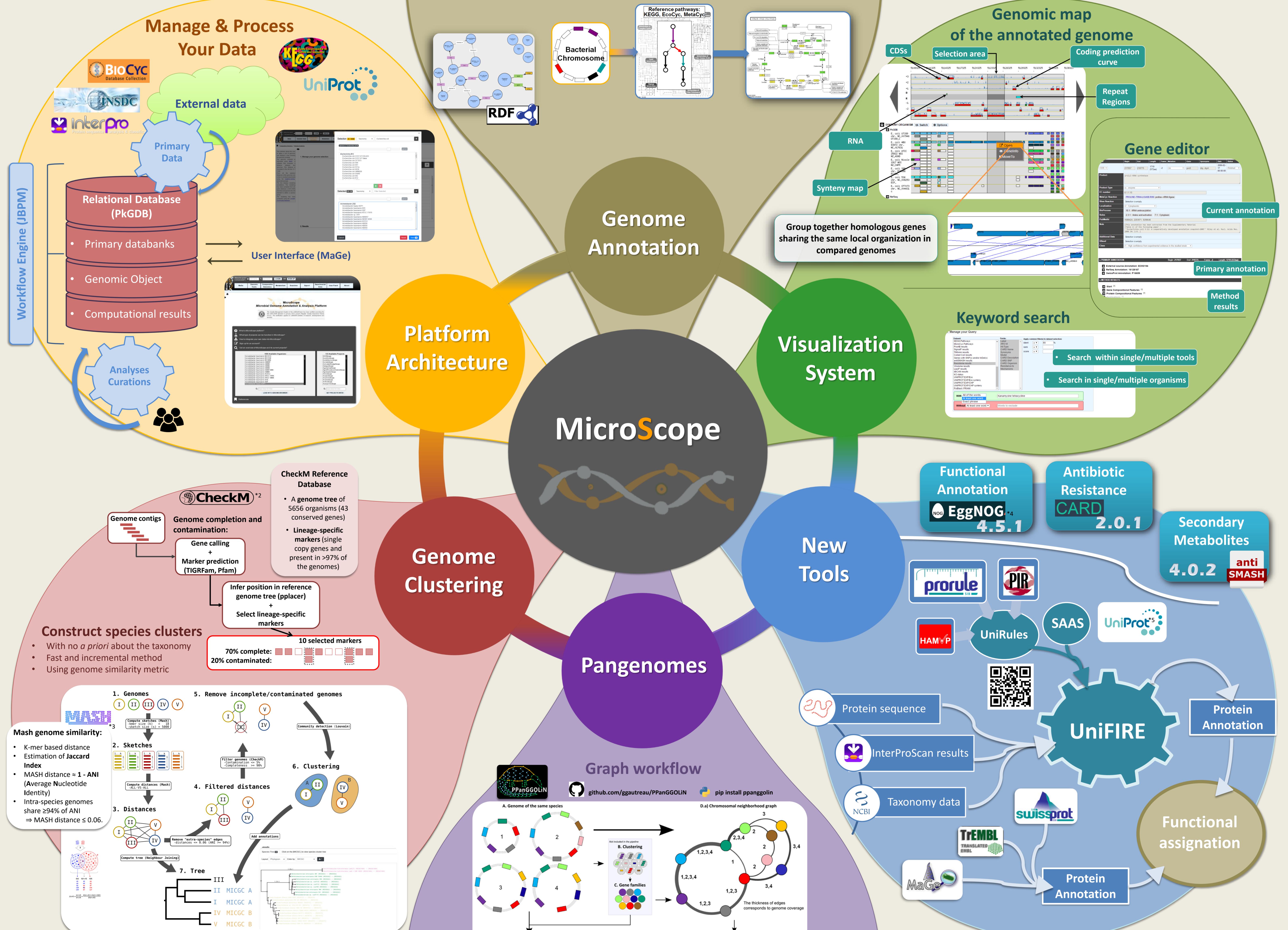
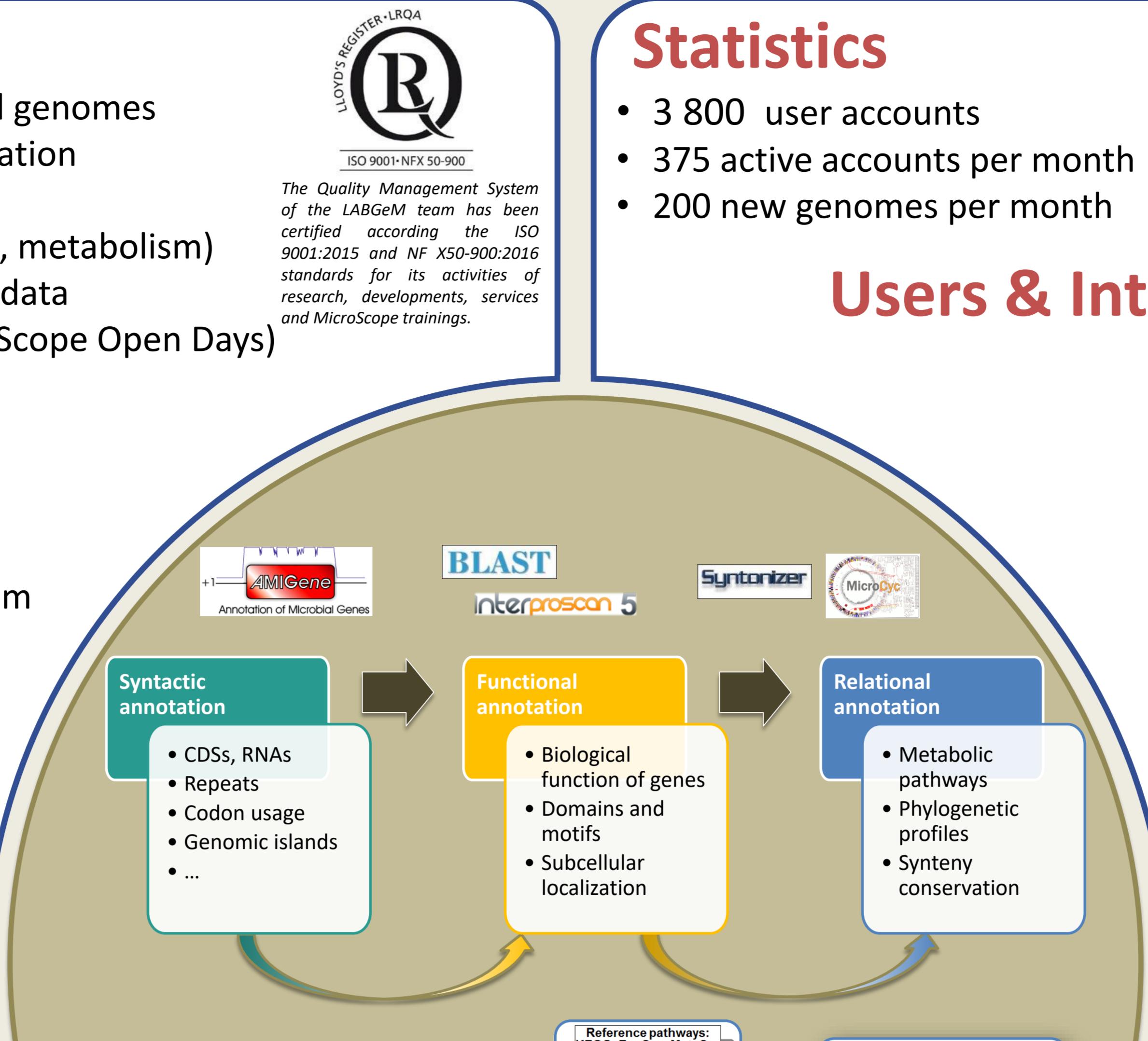
- About 10 000 genomes
- 3 000 human-expertized annotations a month
- 140 collaborative publications
- More than 800 citations

## Users & Interactions



## Professional Trainings

- Audience: researchers, engineers, students, industrials
- Dates:
  - General training on comparative genomics, metabolism exploration and expert annotation (4.5 days / November 2018)
  - Advanced course (2 days / December 2018)
- Location: at Evry University or external training session in France and abroad
- Since 2008 > 450 persons participated in our trainings
- Check our website!



## Collaborations & Services

- MicroScope is part of National infrastructures:
  - France Génomique ([www.france-genomique.com](http://www.france-genomique.com))
  - IFB ([www.france-bioinformatique.fr](http://www.france-bioinformatique.fr))
- Collaborations with industrials & Biotech companies
- ANR projects (Replay, BlueEnzyme)
- European project (IS-Elixir)



## References

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- \*3 B. Ondov *et al.* Genome Biology, 2016 doi: 10.1186/s13059-016-0997-x
- \*4 J. Huerta-Cepas *et al.* Nucl. Acids Res., 2016 doi: 10.1093/nar/gkv1248
- \*5 The UniProt Consortium. Nucl. Acids Res. 2017 doi: 10.1093/nar/gkw1099

[www.genoscope.cns.fr/agc/microscope](http://www.genoscope.cns.fr/agc/microscope)

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