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Baobab is a French research team of the Laboratoire de Biométrie et Biologie Évolutive, and at the same time represents the core of a European research team of

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called

[Erable](#) ↗

. Besides the members of Baobab, Erable has thus members in three institutions in Italy (Sapienza University of Rome, Luiss University, and University of Pisa) and two institutions in the Netherlands (CWI and Free University of Amsterdam).

Baobab has two main sets of research goals that currently cover four axes:

URL of the page: <https://lbbe.univ-lyon1.fr/en/baobab-group>

➤ Goals:

- The first is related to the original areas of expertise of the team, namely combinatorial and statistical modelling and algorithms, although more recently the team has also been joined by members that come from biology including experimental.
- The second set of goals concern its main Life Science interest which is to better understand interactions between living systems and their environment. This includes close and often persistent interactions between two living systems (symbiosis), interactions between living systems and viruses, and interactions between living systems and chemical compounds.

➤ Axes:

- (pan)genomics and transcriptomics in general,
- metabolism and (post)transcriptional regulation,
- (co)evolution
- health in general, of living systems and environmental.

A longer objective of the team is to become able in some cases to suggest the means of controlling for or of re-establishing equilibrium in an interacting community by acting on its environment or on its players, how they play and who plays.

Two major steps are constantly involved in the research done by the team: a first one of modelling (*i.e.* translating) a Life Science problem into a mathematical one, and a second of algorithm analysis and design. The algorithms developed are then applied to the questions of interest in Life Science using data from the literature or from collaborators. More recently, thanks to the recruitment of young researchers (PhD students and postdocs) in biology, the team has become able to start doing experiments and producing data or validating some of the results obtained on its own.

From a methodological point of view, the main characteristic of the team is to consider that, once a model is selected, the algorithms to explore such model should, *whenever possible*, be *exact* in the answer provided as well as *exhaustive* when more than one exists for a more accurate interpretation of the results. More recently, the team has become interested in exploring the interface between exact algorithms on one hand, and probabilistic or statistical ones on the other such as used in machine learning approaches. More in particular, the team is interested in investigating an area of research called “interpretable machine learning” that has been developing more recently and its potential relations with exact, combinatorial approaches.

Besides being at the core of a European team, Baobab has a number of other collaborations at the international level.

Baobab is also strongly involved in teaching at the University of Lyon and Insa-Lyon, well as in other research institutions in Europe, directly or through the members of Erable that are not in France.

For more information, you may also visit the site of the Inria team Erable here:

<http://team.inria.fr/erable/en/> 

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