



## Implementing a Text Mining Service Offer on the Migale Bioinformatics Platform

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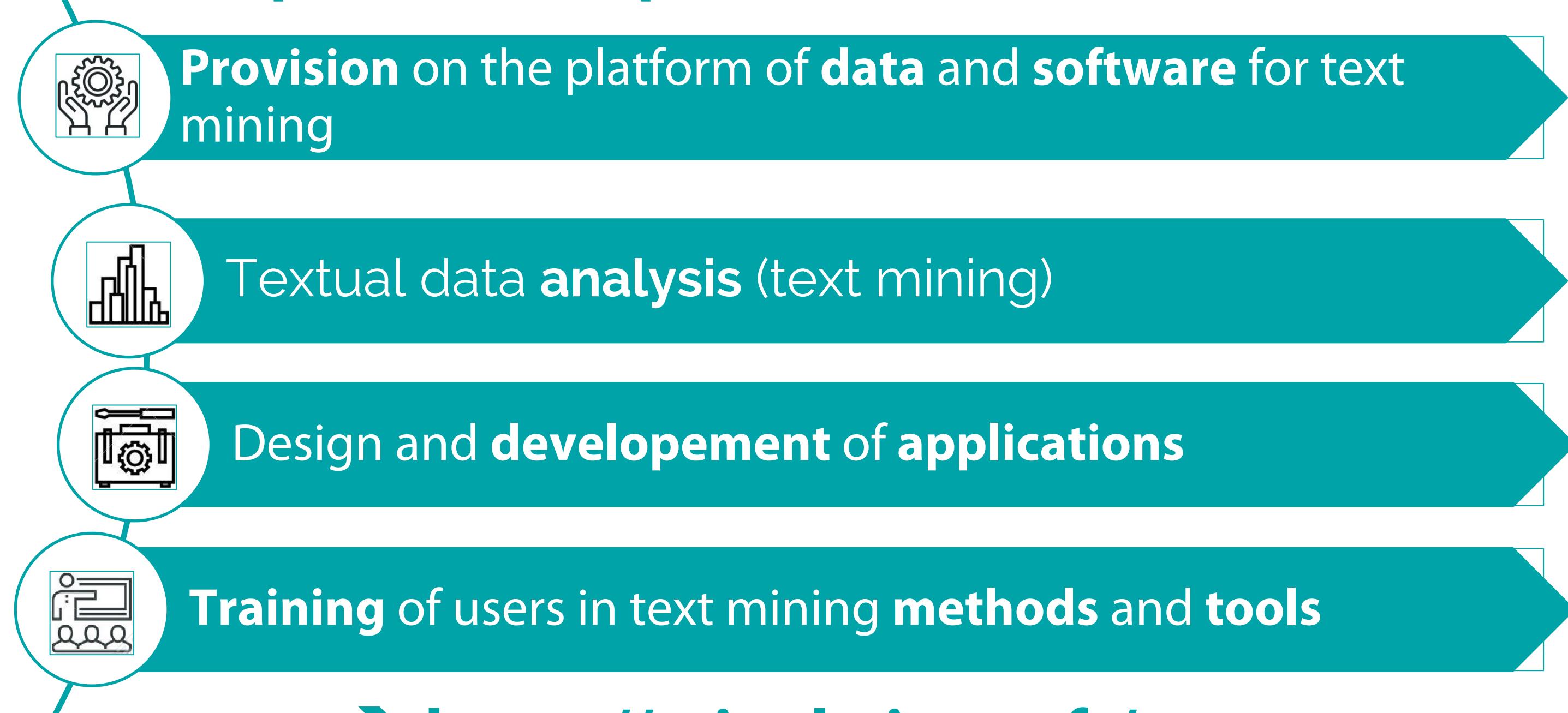
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## Implementing a Text Mining Service Offer on the Migale Bioinformatics Platform

Migale, as a collective scientific infrastructure of INRAE, is building a text mining service offer to enable the bioinformatics community to more easily exploit and extract the information contained in the scientific literature.

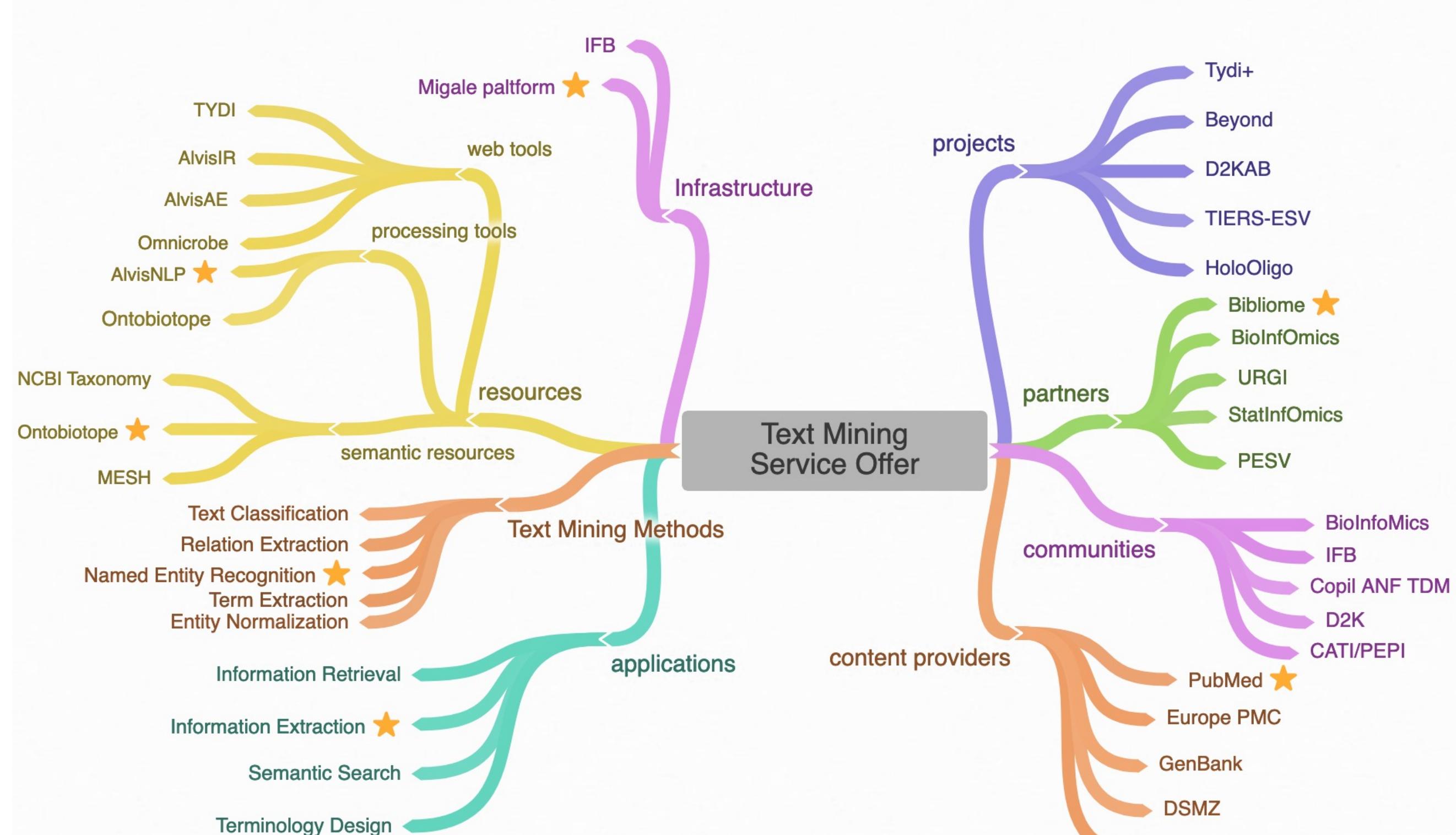


### As part of the platform missions



→ <https://migale.inrae.fr/>

### Reinforce the integration and use of text mining in bioinformatics



### Data and Tools

- Provision** of ~ 15 instances of text mining tools developed by partners
  - AlvisIR: generic semantic search engine
  - Tydi: terminology editor
  - AlvisAE: text annotation editor
- Regular updating of PubMed local databank**
- Packaging, deployment and management** of text mining **tools**



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### Data Analysis

#### Expertise Area:

- Text and data mining

#### Applications:

- Creation of thematic corpora
- Extraction of named entities
- Classification of texts

#### Application Domain:

- Microbiologie

#### Example of projet « TDM4AnimalPhysiology »

- Extraction of genes, pesticides, polyphenols using thematic corpora collected from PubMed
- Study of male and female fertility and energy metabolism of different organisms (mammals, fish, c elegans, drosophila, plants, etc.)

### Design and Development

- Text Mining APIs:** development of APIs wrapping text mining processes based on specialized uses
- Tydi+:** deployment and management of instances of a new application for terminology edition
- Omnicrobe:** management and evolution of the information extraction workflow that implements the text mining process of the Omnicrobe Information System

### Trainings

- A text mining module** in the « Bioinformatics by practicing » cycle of Migale
- Entitled: « **Introduction au text mining avec AlvisNLP** »
- Target audience: (Bio-)informaticians
- Contents:
  - Techniques for Named Entity Recognition (NER)
  - Use cases in biology (recognition of genes, proteins, habitats of bacteria, etc.)
  - Practice with **AlvisNLP**, a corpus processing engine developed by the Bibliome team at INRAE

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