### Target audience
Professionals from the biotechnology sector: project leaders, researchers, programmers, students, teachers, …

### Prerequisite
- Good knowledge of biostatistics and RSoftware
- STAT-1.1: L’analyse statistique appliquée aux sciences du vivant
- STAT-2.1: Initiation au logiciel d’analyse statistique libre R

### Objectives
People with no knowledge of data mining will learn to:
- Recognize benefits and understanding the complexities of data mining;
- Understand how to extract added value from their most valuable medical and genomics data;
- Decide which data-mining technique is the most appropriate in given situations;
- Study real applications of data mining in life sciences (bioinformatics, genomics);

### Content
- Theory: data mining methods, supervised, unsupervised
- Advanced R tutorial; Case studies making use of TCGA data using R: survival analysis, gene differential expression analysis, etc using different data types (integromics)

### Methodology
1 day lecture, 1 day training

### Experts
- Gianluca Bontempi (Interuniversity Institute of Bioinformatics in Brussels (IB), Prof., Director)
- Catharina Olsen (ULB, Computer Science Department)
- Antonio Colaprico (ULB, Computer Science Department)

### Coordinator
Biopark Formation - Erika Baus, formatrice (erika.baus@ulb.ac.be)

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### In practice
**Référence:** DATA-3.1
**Location:** Biopark Charleroi
**Duration:** 2 days
**Schedule:** 9:15 to 17:00

### REGISTRATIONS AND FEES
Biophare: [http://www.biophare.eu](http://www.biophare.eu)

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