



**HAL**  
open science

## Validation of analytical methods as an evaluation tool for research data reliability. Functional Ecology

Anne Jaulin, Elodie Ollivier, Nathalie Cheviron, Virginie Grondin, Amelie  
Trouve

► **To cite this version:**

Anne Jaulin, Elodie Ollivier, Nathalie Cheviron, Virginie Grondin, Amelie Trouve. Validation of analytical methods as an evaluation tool for research data reliability. Functional Ecology. Functional Ecology Conference - JEF AnaEE France, Dec 2018, Nancy, France. 2018. hal-02786058

**HAL Id: hal-02786058**

**<https://hal.inrae.fr/hal-02786058v1>**

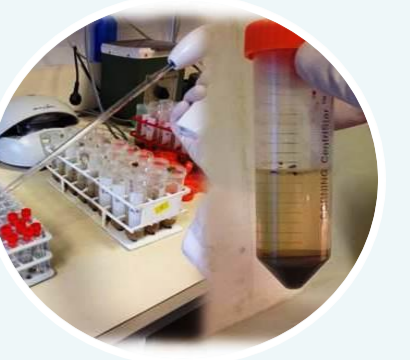
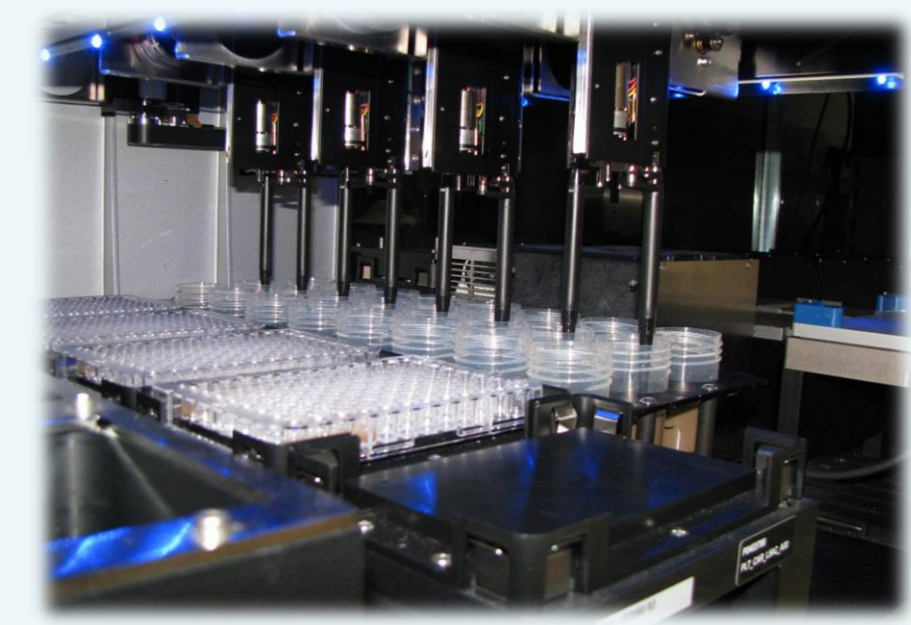
Submitted on 4 Jun 2020

**HAL** is a multi-disciplinary open access archive for the deposit and dissemination of scientific research documents, whether they are published or not. The documents may come from teaching and research institutions in France or abroad, or from public or private research centers.

L'archive ouverte pluridisciplinaire **HAL**, est destinée au dépôt et à la diffusion de documents scientifiques de niveau recherche, publiés ou non, émanant des établissements d'enseignement et de recherche français ou étrangers, des laboratoires publics ou privés.



Distributed under a Creative Commons Attribution - ShareAlike 4.0 International License



# VALIDATION OF ANALYTICAL METHODS AS AN EVALUATION TOOL FOR RESEARCH DATA RELIABILITY

Anne JAULIN\*, Elodie OLLIVIER, Nathalie CHEVIRON, Virginie GRONDIN, Amélie TROUVE

UMR ECOSYS INRA, AgroParisTech, Université Paris-Saclay, plateforme Biochem-Env, 78026, Versailles, France

\* [anne.jaulin@inra.fr](mailto:anne.jaulin@inra.fr)

[contact-biochemenv@inra.fr](mailto:contact-biochemenv@inra.fr)



## Context

The platform Biochem-Env:

- Was created in 2012 by INRA (French National Institute for Agricultural Research) with the support of the ANR program "Investissements d'avenir" as a service of the infrastructure ANAEE-France,
- For the **biochemical characterization of natural environments** (soils and sediments) and **associated macrofauna** in research projects,
- By developing and validating methods in order to **provide traceable analytical data with high level of confidence**.

For intra-laboratory validation of quantitative analytical methods, the INRA's Quality Guidelines for research and experimental units (2013) recommends "the accuracy profile" method according to the NF V03-110:2010 standard.

Could we use a same internally developed method to quantify proteins in various biological models ?

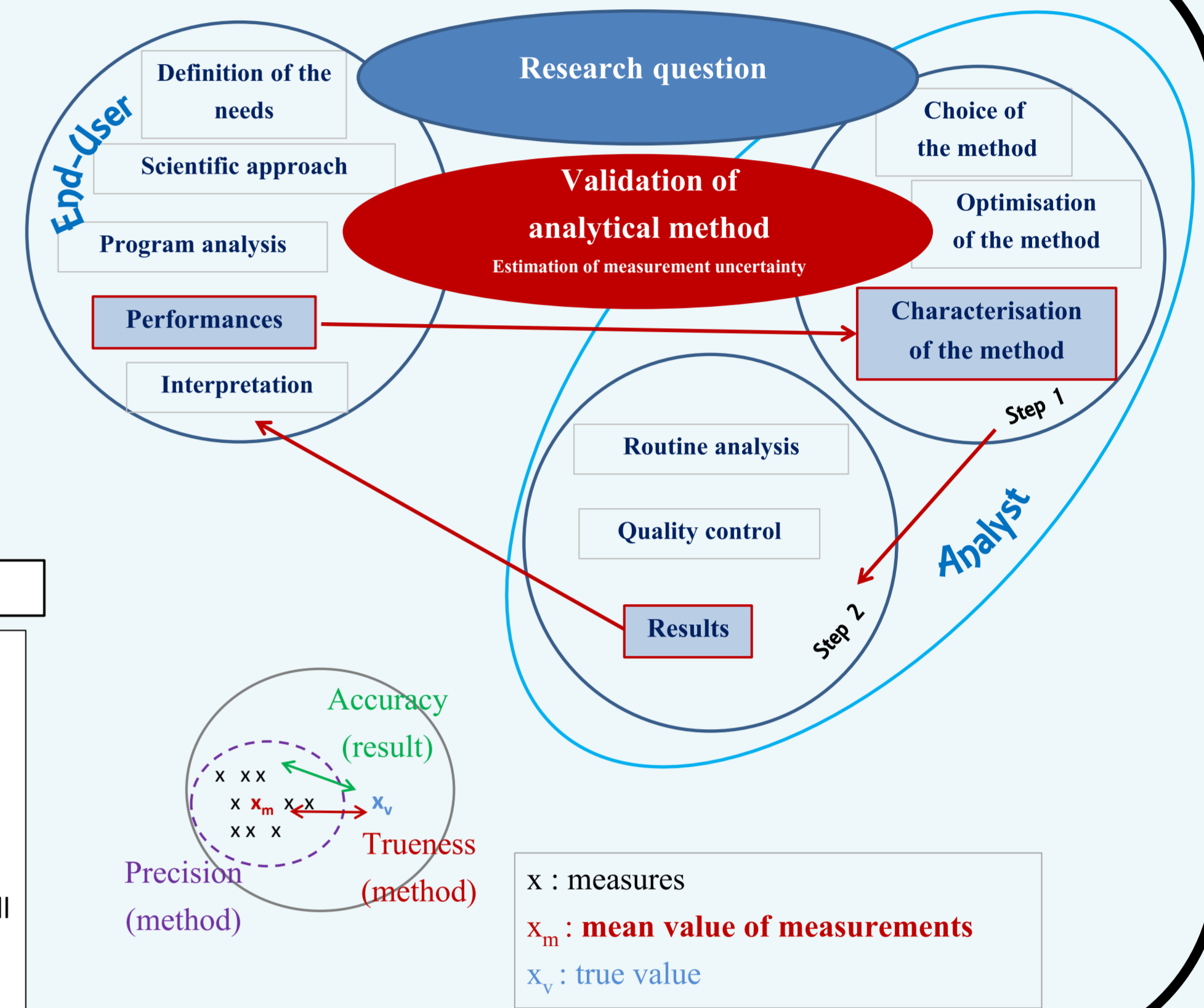
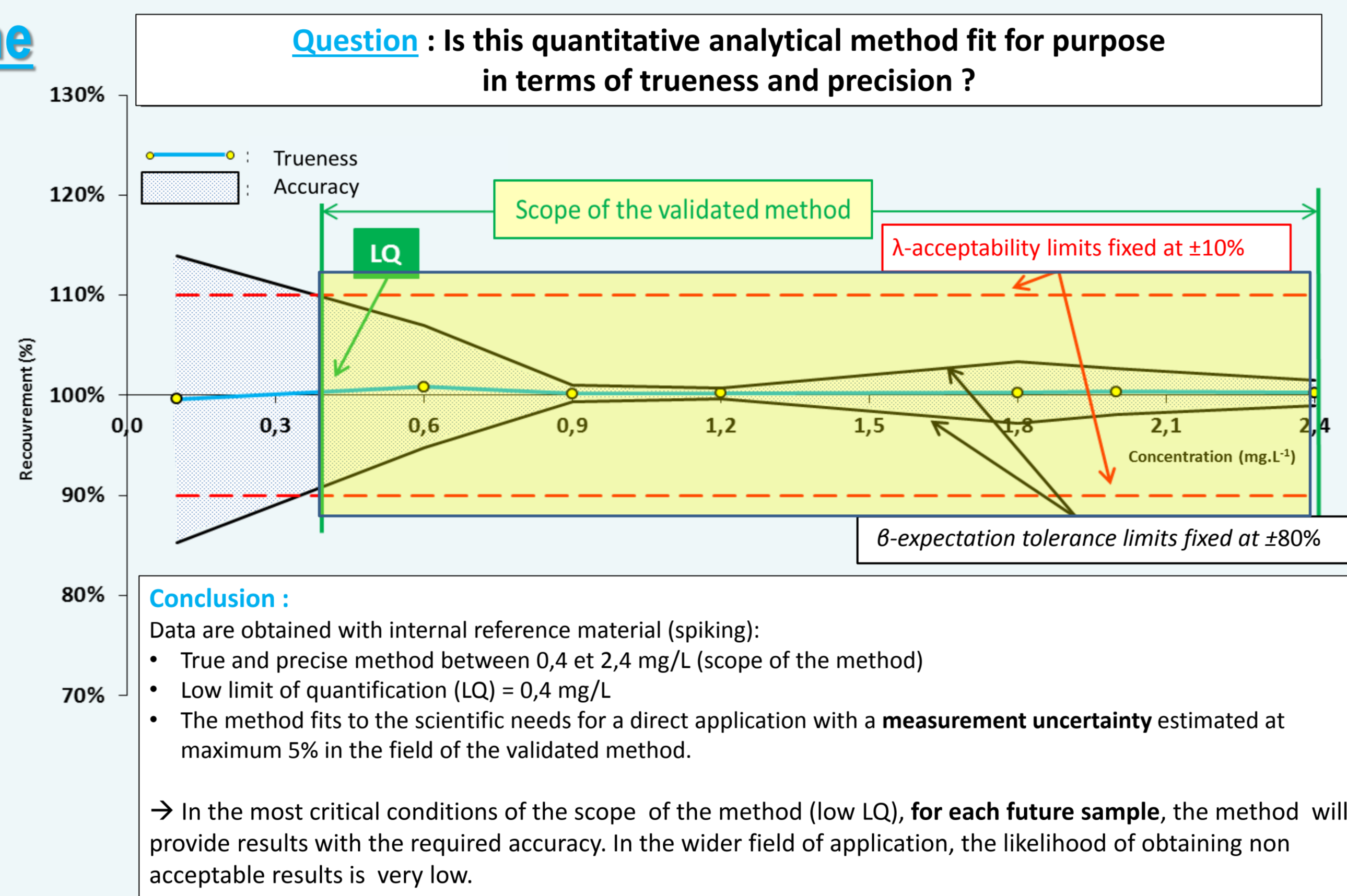
## Validation of analytical methods by the accuracy profile approach

### Purpose:

- To provide **guarantees on analytical results**, for the analyst and the end-user
- To demonstrate **analytical method fitting with the scientific objectives**
- To allow **laboratory recognition**
- To **improve analysts working practices**

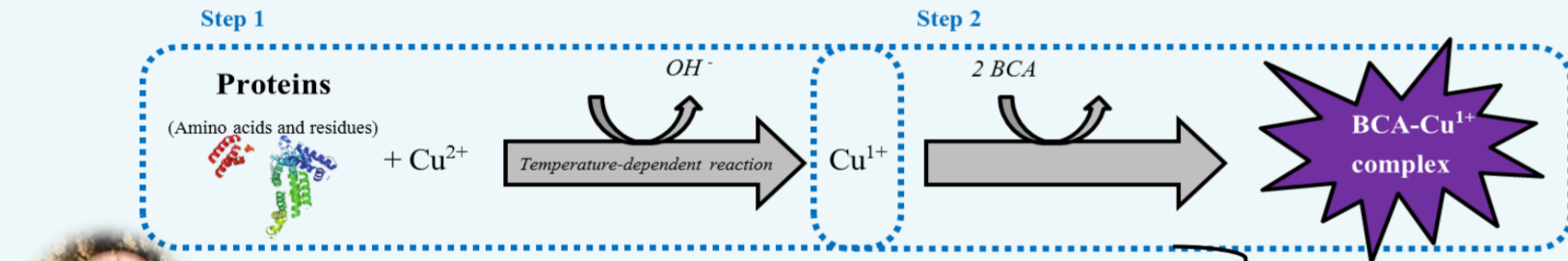
### Benefits of the accuracy profile approach:

- An **overall statistical method** combining **trueness and precision**
- A standardized approach : **NF V03-110:2010**
- A **simple and graphic interpretation** for a **rapid decision**
- The determination of the **scope of the method**
- The determination of **quantification limits**
- An estimation of **measurement uncertainty**



## Material and methods

Protein determination method by the Bicinchoninic acid (Ref. QPBCA-1KT, QuantiPro™ BCA Assay Kit, Sigma-Aldrich)

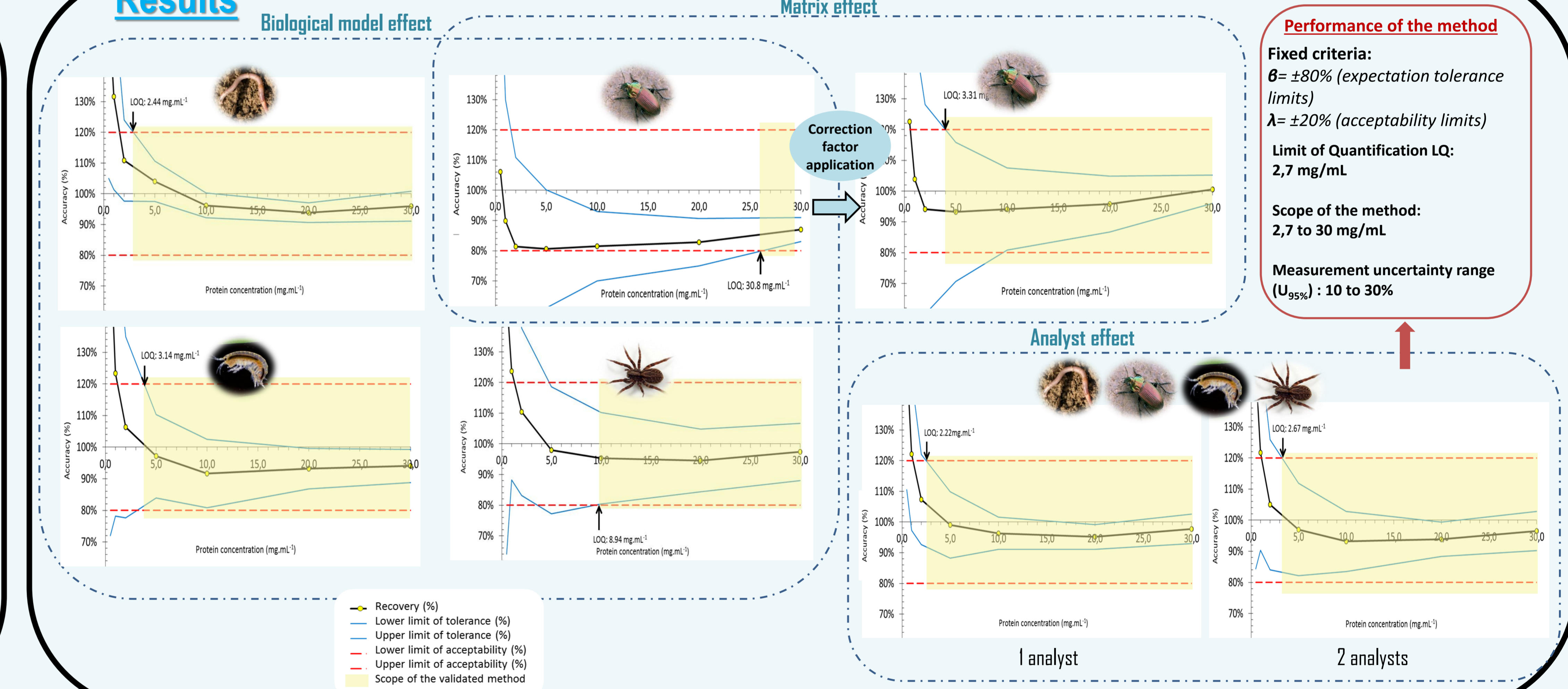


- 4 biological models
- Aporrectodea icterica* (earthworm)
  - Poecilus cupreus* (beetle)
  - Gammarus pulex* (gammarid)
  - Alopecosa pulverulenta* (spider)

Validation method according to the « accuracy profile » approach (NF V03-110:2010 standard):

- Selection of validation samples:**
  - Spiked reference material: Bovine Serum Albumine
  - 7 levels (0.5 ; 1 ; 2 ; 5 ; 10 ; 20 ; 30 mg.mL<sup>-1</sup>)
  - 3 replicates (repeatability)
  - 6 days of analysis (*intermediate precision*)
  - 2 analysts (*intermediate precision*)
- Calibration (indirect quantitative analysis):**
  - 6 levels (0 ; 2 ; 4 ; 8 ; 16 ; 20 ; 30 ; 40 mg.mL<sup>-1</sup>)
  - Regression model : polynomial

## Results



## Conclusions

### Accuracy profile approach:

- Global statistical combination of **trueness and precision** and pragmatic
- Simple graphic interpretation**, allowing a clear and easy comparison between method performance and intended use and, a **rapid decision**
- No limit in the choice of the calibration model => **large scope range**
- Methods with very low variability can be validated (not rejected by a H0)
- Diagnostic tool**, matrix effect taken into account
- Risks and guarantees managed for both end-users and laboratories
- Estimation of measurement uncertainty**

### The validation of this analytical method helped us to:

- Determine the performance of the method
- Improve the steps for sample preparation and analysis
- Assess the matrix effect
- Pointed out the importance of an experimented analyst

Decision on the fitness for purpose

- Analytical method for protein quantification in 4 biological models was validated with a good accuracy considering scientific specification and needs.**

### Perspectives :

- Adapt  $\lambda$ -acceptability values according to the concentration range
- Extend the method to other biological models and biomarkers (Lipid, glycogen...)

## References

- Validation des méthodes d'analyse quantitative par le profil d'exactitude. Numéro spécial du Cahier des Techniques de l'INRA, 2010.
- Labo-Stat : Guide de validation des méthodes d'analyse, Feinberg M. Retirage 2012.
- Jaulin A. et Deschamps M. La validation de méthodes d'analyse quantitative, c'est l'affaire de tous ! La démarche Qualité au service des activités de recherche. Journées de la mesure et de la métrologie J2M 2012 du 8-11/10/2012, Le Croisic.
- Feinberg M. et Laurentie M. A global approach to method validation and measurement uncertainty. 2006. Accred. Qual. Assu. 11, 3-9.
- Feinberg M. et al. New advances in method validation and measurement uncertainty aimed at improving the quality of chemical data. 2004. Anal. Bioanal. Chem. 380 : 502-514.