

Software sensors for bioprocesses

Jérôme Harmand, Alain Rapaport

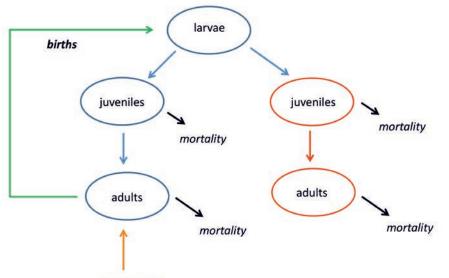
▶ To cite this version:

Jérôme Harmand, Alain Rapaport. Software sensors for bioprocesses. Complex Systems: from biology to landscapes, Agropolis International Editions, 2019. hal-02785901

HAL Id: hal-02785901 https://hal.inrae.fr/hal-02785901

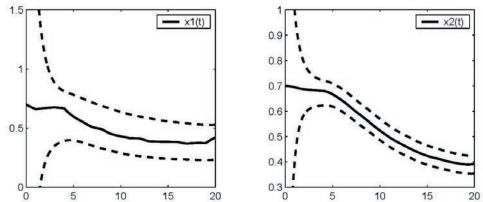
Submitted on 4 Jun 2020

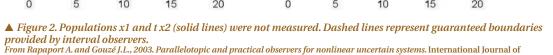
HAL is a multi-disciplinary open access archive for the deposit and dissemination of scientific research documents, whether they are published or not. The documents may come from teaching and research institutions in France or abroad, or from public or private research centers. L'archive ouverte pluridisciplinaire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



immigration

▲ Figure 1. Simply by observing infertile adult populations (red), all other populations can be reconstructed over time. From De Dreuzy J.R. and Rapaport A., 2013. Mieux gérer les ressources naturelles, Les mathématiques de la Terre, Textes et Documents pour la Classe (TDC). 1062: 20-23.





Control. 76(3): 237-251. https://doi.org/10.1080/0020717031000067457

Software sensors for bioprocesses

Sensors available for biological monitoring often do not provide continuous measurement of all variables describing reaction progress (or they are unreliable or too expensive). Through a mathematical model, these unmeasured variables can nevertheless be reconstituted over time as a function of other available measurements using software sensors (e.g. Kalman filters). Specific mathematical conditions are required for building these software sensors, especially the observability property (i.e. the possibility of reconstructing the state of a system from observation data). Not everything is possible of course. Studies on this property allow sensor selection (from among those available) to enable the reconstruction of unmeasured variables (See Fig. 1 above).

The sensor choice is not always intuitive for large or complex systems. Some terms of the biological model are at times poorly known or uncertain (e.g. growth rate depending on the climatic conditions). When the statistical data are not sufficiently abundant to support probabilistic hypotheses on uncertain situations, but the poorly known terms are functionally bounded, 'interval observers' can be implemented. Instead of means and variances, guaranteed lower and upper values can be determined for each unmeasured variable over a time course. A pair of software sensors is thus obtained rather than a single one (see Fig. 2 above). This guaranteed approach is well adapted to transient bioprocesses (tanks containing microorganisms that transform matter, e.g. fermenters, digesters and water purification bioreactors), where there is a risk of biomass washout when their concentrations are too low, which has to be detected as early as possible. Mathematically, this technique is based on the cooperativity property of dynamical systems, which is not always verified by models but can be applied more conservatively to combined variables.

Contacts: J. Harmand (UPR LBE), jerome.harmand@inra.fr, A. Rapaport (UMR MISTEA), alain.rapaport@inra.fr