



Global sensitivity analysis in wastewater applications: A comprehensive comparison of different methods



Alida Cosenza^a, Giorgio Mannina^{a,*}, Peter A. Vanrolleghem^b, Marc B. Neumann^{b,c,d}

^aDipartimento di Ingegneria Civile, Ambientale, Aerospaziale, dei Materiali, Università di Palermo, Viale delle Scienze, 90128 Palermo, Italy

^bmodelEAU, Département de génie civil et de génie des eaux, Université Laval, 1065 av. de la Médecine, Québec, QC G1V 0A6, Canada

^cBasque Centre for Climate Change, Alameda Urquijo, 4-4°, 48008 Bilbao, Spain

^dIKERBASQUE, Basque Foundation for Science, 48011 Bilbao, Spain

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ABSTRACT

Three global sensitivity analysis (GSA) methods are applied and compared to assess the most relevant processes occurring in wastewater treatment systems. In particular, the Standardised Regression Coefficients, Morris Screening and Extended-FAST methods are applied to a complex integrated membrane bioreactor (MBR) model considering 21 model outputs and 79 model factors. The three methods are applied with numerical settings as suggested in literature. The main objective considered is to classify important factors (factors prioritisation) as well as non-influential factors (factors fixing). The performance is assessed by comparing the most reliable method (Extended-FAST), by means of proposed criteria, with the two other methods. In particular, similarity to results obtained from Extended-FAST is assessed for sensitivity indices, for the ranking of sensitivity indices, for the classification into important/non-influential factors and for the method's ability to detect interaction among factors and to provide results in a reasonable time.

It was found that the computationally less expensive SRC method was applied outside its range of applicability ($R^2 = (0.3-0.6) < 0.7$). Still, the SRC produced a ranking of important factors similar to Extended-FAST. For some variables significant interactions among the factors were revealed by computing the total effect indices S_{Ti} using Extended-FAST. This means that to obtain reliable variance decomposition and to detect and quantify interactions among the factors, the use of the Extended-FAST is recommended. Regarding the comparison between Morris screening and Extended-FAST a poor agreement was found. In particular, the Morris screening overestimated the number of both important and non-influential factors compared to Extended-FAST for the analysed case study.

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1. Introduction

In the field of mathematical modelling sensitivity analysis represents a very powerful tool as it provides information about how the variation in the outputs of the model can be apportioned to the variation of the model (input) factors (Saltelli, 2000). "Factors" is a term widely used in the sensitivity analysis literature and includes model parameters and model input variables. Saltelli (2000) singles out three main classes of sensitivity analysis methods: screening methods, local methods and global methods. Screening methods are economical and qualitative methods. Local methods provide a

* Corresponding author.

E-mail addresses: alida.cosenza@unipa.it (A. Cosenza), giorgio.mannina@unipa.it (G. Mannina), Peter.Vanrolleghem@gci.ulaval.ca (P.A. Vanrolleghem), marc.neumann@bc3research.org (M.B. Neumann).

measure of how the model output is affected by infinitesimal factor changes at a specific location in factor space. Global sensitivity analysis (GSA) provides information on how the model outputs are influenced by factor variation over the whole space of possible input factor values (Homma and Saltelli, 1996; Saltelli et al., 2004).

In the environmental modelling field the majority of sensitivity analysis applications are local. Moreover, often a one-at-a-time approach is used that does not allow identifying interacting factors. In recent years, several GSA techniques have been developed. Among them the most widely used are: (i) global screening methods such as the Morris screening method (Morris, 1991; Campolongo et al., 2007); (ii) variance decomposition methods such as Fourier Amplitude Sensitivity Testing (FAST), Extended-FAST and the Sobol indices method (Cukier et al., 1973; Schaibly and Shuler, 1973; Saltelli et al. 1999; Sobol, 2001); and (iii) regression-based methods such as the standardised regression coefficient (SRC) method (Saltelli et al., 2008). GSA may help the

List of symbols and abbreviations

MBR	Membrane BioReactor	K_{NO_3}	Half saturation parameter for S_{NO_3} for X_H
FAST	Fourier Amplitude Sensitivity Test	K_x	Half saturation parameter for X_S/X_H
SRC	Standardised Regression Coefficient	S_F	Fermentable organic matter
GSA	Global Sensitivity Analysis	S_A	Fermentation product (considered to be acetate)
ASM	Activated Sludge Models	X_{PAO}	Phosphorus accumulating organisms model variable
UCT	University of Cape Town	X_{PP}	Stored polyphosphates in PAOs
SMP	Soluble Microbial Product	X_{PHA}	Storage compound in PAOs
TSS	Total Suspended Solids	S_{ALK}	Alkalinity (HCO_3^-)
VSS	Volatile Suspended Solids	X_{AUT}	Autotrophic nitrifying organisms
COD	Chemical Oxygen Demand	S_{BAP}	Soluble biomass associated products
NH_4-N	Ammonia nitrogen	S_{UAP}	Soluble utilisation associated products
NO_2-N	Nitrite nitrogen	S_I	Soluble undegradable organics
NO_3-N	Nitrate nitrogen	X_I	Particulate undegradable organics
N_{TOT}	Total nitrogen	$K_{O,HYD}$	Half saturation/inhibition parameter for S_{O_2}
P_{TOT}	Total phosphorus	$K_{NO_3,HYD}$	Half saturation/inhibition parameter for S_{NO_3}
COD_{TOT}	Total COD model variable	μ_H	Maximum growth rate of X_H
S_{NH_4}	Ammonia nitrogen model variable	q_{FE}	Rate constant for fermentation/Maximum specific fermentation growth rate
S_{NO_3}	Nitrate nitrogen model variable	$\eta_{NO_3,H}$	Reduction factor for anoxic growth of X_H
S_{PO_4}	Soluble inorganic phosphorus model variable	b_H	Decay rate for X_H
MLSS	Mixed liquor suspended solid	K_F	Half saturation parameter for S_F
COD_{SOL}	Soluble modelled COD	K_{FE}	Half saturation parameter for fermentation of S_F
CTN	Total nitrogen model variable	K_A	Half saturation parameter for S_A
y	Model output	$K_{NH,H}$	Half saturation parameter for S_{NH_4} for X_H
x_i	i th model factor	K_P	Half saturation parameter for S_{PO_4} for X_H
b_i	Regression slopes	$K_{ALK,H}$	Half saturation parameter for S_{ALK} for X_H
ε	Random error of the regression model	q_{PHA}	Rate constant for S_A uptake rate
σ_{x_i}	i th factor standard deviation	q_{PP}	Rate constant for storage of polyphosphates
σ_y	Model output standard deviation	μ_{PAO}	Maximum growth rate of X_{PAO}
β_i	i th factor sensitivity index	$\eta_{NO_3,PAO}$	Reduction factor for anoxic growth of X_{PAO}
EE	Elementary Effect	b_{PAO}	Endogenous respiration rate of X_{PAO}
p	Sampling level of Morris screening method	b_{PP}	Rate constant for lysis of polyphosphates
Δ	Factor perturbation	b_{PHA}	Rate constant for respiration of X_{PHA}
μ	Mean of the EEs function	K_{PS}	Half saturation parameter for S_{PO_4} uptake
σ	Standard deviation of the EEs function	K_{PP}	Maximum ratio of X_{PP}/X_{PAO}
IF	Interaction among factors	K_{MAX}	Half saturation parameter for X_{PP}/X_{PAO}
μ^*	Mean of the absolute EEs function	K_{IPP}	Half inhibition parameter for X_{PP}/X_{PAO}
r	Sampling repetition for Morris screening method	K_{PHA}	Saturation constant for X_{PHA}/X_{PAO}
n	Model factors number	$K_{O,PAO}$	Half saturation parameter for S_{O_2} for X_{PAO}
$Var(Y)$	Total variance of the model output	$K_{NO_3,PAO}$	Half saturation parameter for S_{NO_3} for X_{PAO}
S_i	First order effect index of the i th factor	$K_{A,PAO}$	Half saturation parameter for S_A for X_{PAO}
S_{Ti}	Total effect index of the i th factor	$K_{NH,PAO}$	Half saturation parameter for S_{NH_4} for X_{PAO}
N_{MC}	Number of Monte Carlo simulations	$K_{P,PAO}$	Half saturation parameter for S_{PO_4} as nutrient (X_{PAO} growth)
S_{Ni}	Normalised interaction index	$K_{ALK,PAO}$	Half saturation parameter for S_{ALK} for X_{PAO}
ρ_s	Spearman's rank correlation index	μ_{AUT}	Maximum growth rate of X_{AUT}
r_p	Pearson correlation index	b_{AUT}	Decay rate for X_{AUT}
PF	Position Factor	$K_{O,A}$	Half saturation parameter for S_{O_2} for X_{AUT}
<i>Rel</i>	Relevance	$K_{NH,A}$	Half saturation parameter for S_{NH_4} for X_{AUT}
NS	Number of simulations	$K_{ALK,A}$	Half saturation parameter for S_{ALK} for X_{AUT}
PAOs	Phosphorus Accumulating Organisms	$K_{P,A}$	Half saturation parameter for S_{PO_4} for X_{PAO}
$Rel_{IMPORTANT}$	Relevance of important factors	$k_{H,BAP}$	Hydrolysis rate coefficient for S_{BAP}
$Rel_{NON-INFLUENTIAL}$	Relevance of non-influential factors	$k_{H,UAP}$	Hydrolysis rate coefficient for S_{UAP}
k_H	Maximum specific hydrolysis rate	$k_{LaT,3}$	Overall oxygen transfer coefficient aerobic tank
$\eta_{NO_3,HYD}$	Correction factor for hydrolysis under anoxic conditions	$k_{LaT,4}$	Overall oxygen transfer coefficient MBR tank
η_{FE}	Correction factor for hydrolysis under anaerobic conditions	Y_H	Yield for X_H growth
K_O	Half saturation parameter for S_{O_2} for X_H	f_{X_I}	Fraction of X_I generated in biomass decay
S_{O_2}	Dissolved oxygen	Y_{PAO}	Yield for X_{PAO} growth
X_H	Ordinary heterotrophic organisms	Y_{PO_4}	Yield for X_{PP} requirement per X_{PHA} stored
X_S	Particulate biodegradable organics	Y_{PHA}	Yield for X_{PP} storage per X_{PHA} utilised
X_H	Ordinary heterotrophic organisms	Y_A	Yield of X_{AUT} growth per S_{NO_3}
		f_{BAP}	Fraction of S_{BAP} generated in biomass decay
		f_{UAP}	Fraction of S_{UAP} generated in biomass decay

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