Study on sensitive analysis criteria for activated sludge models combined with settling model

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Abstract

The purpose of this paper was to propose proper sensitivity analysis criteria which can minimize calibration efforts and produce reliable effluent quality predictions. ASM1 and ASM3 parameter sensitivity (in combination with a one-dimensional settling model) was analyzed by various criteria based on the step variation of single parameters (SVM) or random variations of all parameters (RVM), using the *IWA Simulation Benchmark* (Copp et al. 2002). SVM was not significantly affected by the analysis conditions and it produced reliable Δ EQ values in every case. Moreover, it was the easiest and simplest methodology. Once selected the parameters were estimated with a genetic algorithm. It was concluded that SVM was the best sensitivity analysis criteria for both ASM1 and ASM3 in this case.

Key Words: Activated sludge model, genetic algorithm, sensitivity analysis, Simulation Benchmark

INTRODUCTION

The complicated and nonlinear characteristics of the activated sludge models (ASMs) make it very difficult to identify the system behavior. Parameter estimation is essential for process modeling, but it generally requires lots of time and effort. It has been reported that the success and failure of a model application are strongly related to the cost for stoichiometric and kinetic parameter estimation (Sollfrank and Gujer, 1991; Ko *et al.*, 2001).

Sensitivity analysis is an essential procedure for selecting significant parameters which can have serious effects on process behavior, *prior* to numerical parameter optimization. The aim of this study was to understand the sensitivity of ASM1 and ASM3 parameters and to suggest a criteria for selecting sensitive parameters. Even though a sedimentation process might have an impact on the biological process, the settling model has been commonly ignored during previous sensitivity analysis research. In this paper, the sensitivity of settling model parameters also has been analyzed.

METHODS

Target process and models

The target process was the denitrifying layout of the IWA Simulation Benchmark (Copp et al., 2002). Simulation Benchmark adopts ASM1 and a one-dimensional settling model (Takacs et al., 1991) for biological reactors and the clarifier, respectively. In this study, ASM3 was also included and sensitivity analysis of the parameters was performed.

Effluent Quality (EQ) index for sensitivity indexes and objective function calculation

Selection of sensitive parameters might be influenced by the sensitivity index (SI) calculation method (Saltelli *et al.*, 2000). In this study, eleven SI calculation methods were examined (Fig. 1). Sensitivity indices were based on Effluent Quality (EQ) as defined by the IWA Task Group. Slight modifications had to be made for calculating the ASM3 EQ (Kim *et al.* 2004).

The difference in EQ (Δ EQ) was calculated as below;

$$\Delta EQ = EQ_{\text{Ref}} - EQ_{\text{Var}}$$
 (Eq. 1)
where
$$EQ_{\text{Ref}} = EQ \text{ calculated with reference parameters values}$$

where $EQ_{Ref} = EQ$ calculated with reference parameters value $EQ_{Var} = EQ$ calculated with varied parameters values

After selecting the parameters, they were estimated with a genetic algorithm (GA) aimed at minimizing ΔEQ .

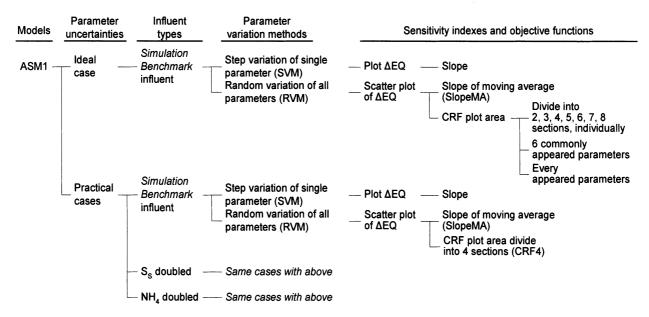
Sensitive parameter selection criteria

The criteria for selecting sensitive parameters is shown in Fig. 1. Each parameter was changed from 50 to 200% of its

reference value. The parameters were changed by 10% for the step variation method (SVM). For the random variation method (RVM), 2,000 simulations were conducted with randomly selected sets of parameters.

Tab. 1. Values of insensitive parameters according to the type of model.

	ASM1		ASM3				
	Reference simulation	Estimation step	Reference simulation	Estimation step			
Ideal case	Simulation Benchmark defaults (Copp et al., 2002)	Simulation Benchmark defaults	ASM3 defaults (Henze et al., 2000)	ASM3 defaults			
Practical cases	ASM1 defaults (Henze et al., 2000)	Simulation Benchmark defaults	Tuned for Bio-P Module (Rieger et al., 2001)	ASM3 defaults			



ASM3 — Everye cases done with ASM1

Fig. 1. Tested sensitivity analysis criteria.

Parameter estimation

After parameters were classified into sensitive or insensitive, only the most sensitive parameters were optimized with GA (Kim et al., 2002). Carroll's GA was applied (Kim et al. 2004, Yang et al., 1998) and the applied genetic operator was a two-member tournament selection, uniform crossover, flip and creep mutation, elitism and niching (Goldberg, 1989).

RESULTS AND DISCUSSTION

Sensitivity analysis

Step variation of single parameter. The impact on EQ was plotted against the parameter value and the greater the impact, the higher the sensitivity. The most sensitive five parameters of ASM1 and ASM3 selected by SVM were as following:

- ASM1 : Y_H , b_H , $\mu_{max,A}$, $K_{O,A}$, b_A
- ASM3 : $Y_{STO,NO}$, $Y_{H,NO}$, $\mu_{max,A}$, η_{NO} , b_A

Random variation of all parameters. Calculated ΔEQs versus the varied parameters were presented as a scatter plot. If a parameter was insensitive, dots distribution inside each vertical section was uniform, implying that the effect from the variation of that parameter was not significant and could be compensated by the variation in other parameters (Fig.2a) The sensitive parameters resulted in a different distribution of dots (Fig.2b). For quantifying the sensitivity, two different methods were used; 1) slope of moving average in each vertical section as shown in Fig. 2a and 2b, 2) area of cumulative relative frequency (CRF) (data not shown). K_S and $\mu_{max,A}$ were selected as typical examples of insensitive and sensitive parameters, respectively.

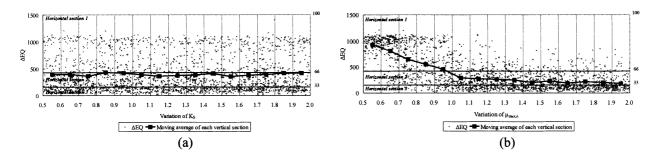


Fig. 2. Sensitivity parameter selection criteria based on RVM; (a), (b) scatter plot and slope of moving averaged K_S and $\mu_{max.A}$ in ASM1, respectively

Selected sensitive parameters and simulation results in the ideal case

<u>Sensitive parameters of ASM1</u>. SVM identified Y_H , b_H , $\mu_{max,A}$, b_A and $K_{O,A}$, as being sensitive and RVM identified Y_H , b_H , $\mu_{max,A}$, b_A , k_h (for ASM1) and v_0 and r_h (for the one-dimensional settling model). Y_H , $\mu_{max,A}$, and r_h were identified by RVM in every case, and v_0 was identified in all cases except SlopeMA. RVM identified one or two settling parameters each time, but those did not show high sensitivity with SVM (Tab. 2).

Sensitive parameters of ASM3. SVM identified $Y_{STO,NO}$, $Y_{H,NO}$, $\mu_{max,A}$, b_A and η_{NO} as being sensitive, while RVM identified $Y_{H,O2}$, $i_{N,Xs}$, $\mu_{max,A}$, b_A , $K_{O,A}$, r_h and v_0 . Just like in ASM1, one or two settling parameters were identified by RVM in each case except SlopeMA and CRF8 both of which did not identify any settling parameters. The nitrogen content of X_S ($i_{N,Xs}$), was identified by RVM and it was reasoned that this parameter had a large impact on the nitrogen concentration to be nitrified. No criteria identified b_H and k_h as sensitive which is in contrast to the ASM1 results.

Tah '	2 Selected	sensitive nara	meters and sim	ulation recul	ts at ideal case

ASM1						ASM1																					
Analys	is methods		Hi	ighly sensit	live parame	ters		Δ	Q	Abso	lute er	тог (mg	(L)	Analysis	methods		1	lighly se	nsitive par	ameters					Abso	lute er	rror (mg L)
Variation method	Sensitivity Index	Stoichiometric		Kinetic			Settling	Settling	•	COD NI	NЩ	NO ₃	, TSS	Sensitivity Index	Influent type	Stoichiometric		Kinetis	Kinetic			Settling			COD	NH.	NO ₃ TS
SVM		Y _H ,		nex.A ba,				(.57	0.45	0.01	0.02	0.33	SVM	S. B. default	Y _H .	$b_{H},\\$	μ _{max.A} ,	b _A , K _O ,					2.09	0.13	0.02	0.02 0.6
RVM	SlopeMA	Y _H ,	μα	b _A .	Ko.a.	rh		(.91	0.09	0.02	0.02	0.07		S _s doubled	Y _H ,	b _H ,	μ _{max,A} ,	ba, Ko.					2.52	1.48	0.03	0.02 0.0
	CRF2	Y _B ,	μπ	b _A .		r _h ,	. v ₀	3	.69	0.01	0.18	0.01	0.00		NH4 doubled	Yн.	b _H ,	μ _{max.A} ,	b _A , K _O ,					11.62	0.31	0.07	0.45 0
	CRF3	Y _H ,	μα	b _A .		r _h ,	. V ₀	3	.69	0.01	0.18	0.01	0.00	RVM-													
	CRF4	Y _H .	μπ	sex.A	k	h. rh,	. V ₀	1	.08	0.12	0.01	0.02	0.18	SlopeMA	S. B. default			μ _{max,A} ,	ba, Koa		r _h			3.03	0.78	0.01	0.05 0.5
	CRF5	Y _H ,	b _н , ^{µп}	nex_A		r _h ,	. v ₀	4	.01	0.50	0.10	0.04	0.29	SiopciviA	S _s doubled	Y _H ,	К _{о.н} .	μ _{max.A} ,	K _{0,4}				V ₀	1.13	0.37	0.01	0.01 0.1
	CRF6	Y _H ,	μα	ък.А	k	h, rh,	. V ₀	1	.08	0.12	0.01	0.02	0.18		NH₄ doubled	Y _H ,		µ _{mex.A} ,	ba, Koa			V0_		8.57	0.92	0.12	0.22 0.4
	CRF7	Y _H ,	μα	ux.A	k	h. rh,	. v ₀	1	.08	0.12	0.01	0.02	0.18	RVM-CRF4	S. B. default	Y _H ,		μ _{max.A} ,		\mathbf{k}_{h}	r _h ,	V ₀		6.13	0.26	0.23	0.01 0.9
	CRF8	Y _H ,		nex.A	k	h. rh,	. v ₀	1	.08	0.12	0.01	0.02	0.18		S _s doubled	Y _H ,	b _H ,	μ _{max.A} .			r _h ,	V_0		17.75	0.58	0.03	0.37 8.6
	Common6	Y _H ,	μα	b _A .	k	h, r h,	. V ₀	11	.33	2.79	0.18	0.00	2.15		NH doubled	v				k _h ,	_						
	Every7	Y _н ,	b _Η , μ _σ	b _A ,	k	h, rh,	. V ₀		.72	0.45	0.15	0.02	0.42		NH4 doubled	1 H,		μ _{mex.A} .		Kh.	r _h ,	V ₀		11.16	0.30	0.09	0.33 1.2
ASM	3													ASM3													
		Stoichio	metric		Kine	tic	Settling	g								Stoich	iometric		Kii	etic			Settling				
SVM		Y _{STO.NO} ,	YHA	νο, μ	max.A. ba.		η_{NO}	2	.66	0.02	0.12	0.02	0.01	SVM	S. B. default	$Y_{STO,NO}$	$Y_{\text{H,NO}},$		$\mu_{max,A},$	\mathbf{b}_{A}		η_{NO}		0.53	0.02	0.02	0.01 0.0
RVM	SlopeMA	Y _{H.02}	,	i _{N.Xs} . μ _r	mex.A. b _A .	K _{O,A}		(.76	0.04	0.02	0.01	0.03		S ₃ doubled	$Y_{STO,O2}, Y_{H,O2},$	Y _{H.NO} ,		μ _{max,A} ,				$\mathbf{r}_{\mathbf{h}}$	2.40	0.35	0.06	0.01 0.3
	CRF2	Y _{H.02}	,	i _{N.Xs} , μ _r	пых.А.		r _h ,	V ₀ 1	.76	0.05	0.02	0.06	0.10		NH4 doubled	Y _{STO,NO} , Y _{H,O2} ,	Y _{H.NO} ,		μ _{mex.A} .	b _A				0.47	0.00	0.00	0.01 0.0
	CRF3	Y _{H.02}		i _{N.Xs} , μ _r	nex.A. b _A ,			V ₀	.81	0.21	0.01	0.02	0.08	RVM-	S. B. default	Y _{H.02} ,					 K						
	CRF4			i _{N.Xs} . μ _r	nex.A. bA.		r _h ,	v ₀ i	.58	0.26	0.02	0.04	0.22	SlopeMA				S.	μ _{max.Λ} .	b _A ,	N.	D.A		5.41	0.86	0.25	0.15 0.7
	CRF5			$i_{N,Xs},\ \mu_{e}$	_{nax.A} , b _A ,		\mathbf{r}_{h}	Vo I	.58	0.26	0.02	0.04	0.22	Siopeivira	S _s doubled	Y _{STO,O2} , Y _{H.O2} ,	i _{N,N}	s. μ _{max}	H. Hmax.A					0.91	0.33	0.00	0.00 0.2
	CRF6	Y _{H.02}		$i_{N,Xs},\ \mu_r$	nax_A. b _A .			V ₀ (81	0.21	0.01	0.02	0.08		NH4 doubled	Y _{H.02} ,	f _{Xi}		μ _{max,A} ,	b _A .	ηνο	END		2.83	0.03	0.10	0.04 0.0
	CRF7				max.A. ba,			V ₀ 2	22	0.14	0.05	0.03	0.11	RVM-CRF4	S. B. default		i _{N,X}	is.	μ _{max.A} .	b _A .			r _h	V ₀ 19.82	0.02	0.65	0.34 0.0
	CRF8	Y _{H.02}		$i_{N,Xs}, \mu_r$	mex.A. b _A ,	Koa		(87	0.03	0.03	0.01	0.02		S. doubled	Y _{STO,O2} , Y _{H,O2} ,			μ _{max.A} ,		Ko		r _h	2 19	0.56	0.00	0.03 0.3
	Common6	Y _{H,O2} .		i _{N.Xs} , μ _r	nex.A. ba.		r _h ,	V ₀ I	14	0.25	0.02	0.00	0.20		NIU doubled	- 010,011 - 11,011			PHINCH				-	- 66.03			

Selected sensitive parameters and simulation results in the practical cases

<u>Sensitive parameters of ASM1</u>. In the practical cases, a different set of insensitive parameter values was used for the reference simulation. Three criteria were examined at practical cases; SVM, SlopeMA and CRF4. In the practical case, SVM identified precisely the same parameters as identified in the ideal case. This suggests that SVM is not significantly affected by influent

type and the insensitive parameter values and further suggests that this methodology is valid for ASM1. This contrasts the SlopeMA and CRF4 results which identified different parameters depending on the simulation setup (Tab. 3).

<u>Sensitive parameters of ASM3</u>. The various simulation conditions identified a total of 15 sensitive parameters as compared to 9 for the ASM1 simulations. In this case, different parameters were selected and no criteria exhibited consistent results.

Parameter estimation with GA and validation

With the sensitive parameters identified, a genetic algorithm was then used to estimate the values of the sensitive parameters to see if the genetic algorithm could find the solution and minimize ΔEQ . Previous work (Kim *et al.* 2002) suggested that the use of a genetic algorithm was a suitable choice for finding optimal parameter values. However in this case, finding the solution is complicated by the fact that the IWA Simulation Benchmark quantity, EQ, is a composite variable composed of carbon and nitrogen state variables meaning that it could be possible to minimize ΔEQ (or even find a ΔEQ of zero) with a different combination of nitrogen and carbon species. A list of sensitive parameters, ΔEQ and absolute errors of effluent COD, NH_4 , NO_3 and TSS according to sensitivity analysis criteria are shown in Tab. 2 and 3 under ideal and practical cases, respectively.

The highest Δ EQs (ignoring Common6 and Every7) were 4.01 and 2.66 for ASM1 and ASM3 respectively in the ideal case. These relatively high values were attributed to an NH₄⁺-N and NO₃.-N prediction error less than 0.2 mg/L. For practical purposes an error of this magnitude is negligible, but it is amplified in this case by the IWA weighting factor of 20 applied to nitrogenous species. It was therefore concluded that SVM combined with a well-tuned GA is a suitable approach for sensitivity analysis and parameter estimation.

CONCLUSIONS

Various sensitivity analysis criteria were examined in an effort to minimize calibration efforts and produce reliable prediction results. For ASM1, the step variation of single parameters (SVM) proved to be any acceptable approach and provided similar results regardless of the simulation conditions. In previous research, $K_{O,A}$ and b_A were regarded as insensitive, but in this case those parameters showed high sensitivity. Other criteria based on random variation of all parameters (RVM) exhibited different sets of sensitive parameters in each case demonstrating that SVM was the better approach. For ASM3, no criteria exhibited consistent results and no conclusion could be reached about which ASM3 parameters were the most sensitive. But, it should be noted that SVM presented reliable Δ EQ values at every influent condition. Moreover, it was the simplest methodology. Therefore, it is concluded that SVM is a reasonable approach for both ASM1 and ASM3.

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