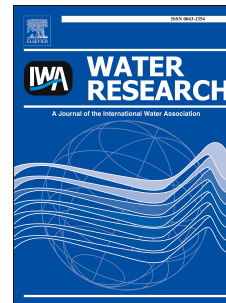


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New systematic methodology for incorporating dynamic heat transfer modelling in multi-phase biochemical reactors

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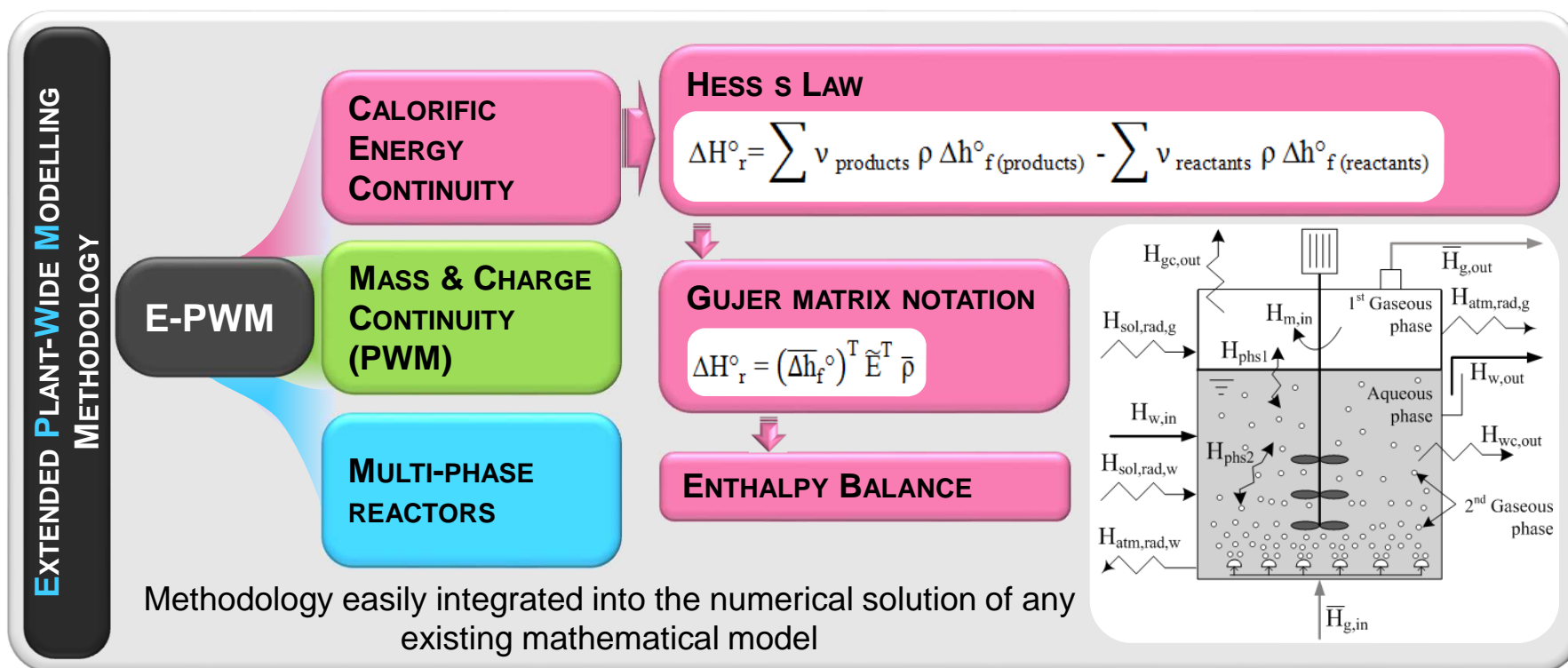
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1 **New systematic methodology for incorporating dynamic heat transfer modelling**
2 **in multi-phase biochemical reactors**

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9
10 **Abstract**

11 This paper presents a new modelling methodology for dynamically predicting the heat
12 produced or consumed in the transformations of any biological reactor using Hess's law.
13 Starting from a complete description of model components stoichiometry and formation
14 enthalpies, the proposed modelling methodology has integrated successfully the
15 simultaneous calculation of both the conventional mass balances and the enthalpy change
16 of reaction in an expandable multi-phase matrix structure, which facilitates a detailed
17 prediction of the main heat fluxes in the biochemical reactors. The methodology has been
18 implemented in a plant-wide modelling methodology in order to facilitate the dynamic
19 description of mass and heat throughout the plant. After validation with literature data, as
20 illustrative examples of the capability of the methodology, two case studies have been
21 described. In the first one, a predenitrification-nitrification dynamic process has been
22 analysed, with the aim of demonstrating the easy integration of the methodology in any
23 system. In the second case study, the simulation of a thermal model for an ATAD has
24 shown the potential of the proposed methodology for analysing the effect of ventilation and
25 influent characterization.

27 **Keywords**

28 Heat balance; Hess's law; Mathematical modelling; Multi-phase models; Plant-Wide Modelling

29 methodology

30

31 **NOMENCLATURE**

A_{bubbles}	Contact area between aqueous phase and 2 nd gas phase (m ²)
A_{g}	1 st gas phase surrounding area (m ²)
A_{tank}	Contact area between aqueous phase and 1 st gas phase (m ²)
A_{w}	Aqueous phase surrounding area (m ²)
\overline{Cp}_i	Specific heat capacity of i gaseous phase components at constant pressure (kJ gE ⁻¹ K ⁻¹)
$Cp_{\text{w,H}_2\text{O}}$	Specific heat capacity of water at constant pressure (kJ gH ₂ O ⁻¹ K ⁻¹)
\tilde{E}_{ij}	i phase stoichiometric matrix for the transformations between i and j phase
g	Gravitational acceleration (m s ⁻²)
Gr	Grashof number
$\bar{h}_{i,\text{ref}}$	Reference enthalpy of the i gaseous phase components (kJ gE ⁻¹)
$h_{\text{w,ref,H}_2\text{O}}$	Reference enthalpy of water (kJ gH ₂ O ⁻¹)
H_{Act}	Enthalpy transmitted from the actuators (kJ d ⁻¹)
$H_{\text{atm_rad}}$	Longwave atmospheric radiation flux (kJ d ⁻¹)
$H_{\text{Cond/Conv}}$	Surface conduction and convection enthalpy flux (kJ d ⁻¹)
\bar{H}_{in}	Input enthalpy (kJ d ⁻¹)
\bar{H}_{out}	Output enthalpy (kJ d ⁻¹)
$H_{\text{sol_rad}}$	Shortwave solar radiation flux (kJ d ⁻¹)
H_{T}	Net heat exchange of the analysed phase (kJ d ⁻¹)
H_{trans}	Heat released or absorbed by the transformations in the analysed phase (kJ d ⁻¹)
$k_{\text{sol,rad}}$	Total energy incident on the surface (kJ d ⁻¹ m ⁻²)
k_{wls}	Heat conduction through walls and pipes coefficient (kJ d ⁻¹ K ⁻¹ m ⁻²)
$\bar{m}_{i,\text{in}}$	Inlet i phase mass flux (gE d ⁻¹)
$\bar{m}_{i,j}$	Mass flux transport between i and j phases (gE d ⁻¹)
$\bar{m}_{i,\text{out}}$	Outlet i phase mass flux (gE d ⁻¹)
\bar{M}_i	Mass vector for the components present in the i phase (gE)

$n_{\text{phs},i}$	Constant exponent of the Pr and Gr numbers (aqueous and i gaseous phases conduction/convection)
NC	No. of components in the i phase
NT	No. of transformations in the i phase
P_m	Power supplied by the engines and aeration systems (kJ d^{-1})
Pr	Prandtl number
T_{atm}	Atmospheric temperature (K)
T_i	i phase temperature (K)
$T_{i,\text{in}}$	i phase inflow temperature (K)
$\bar{T}_{i,\text{ref}}$	Temperature corresponding to the reference enthalpy of the i gaseous phase components (K)
$T_{\text{w,ref,H2O}}$	Temperature corresponding to the reference enthalpy of water (K)
\bar{X}_i	Mass fraction of the gaseous phase components ($\text{gE}_i \text{gE}_{\text{phase}}^{-1}$)

Greek Symbols

$\alpha_{\text{phs},i}$	Correction factor for surfaces facing up (aqueous and i gaseous phases conduction/convection)
α_{rad}	Solar absorptivity
β_{air}	Atmospheric radiation factor
β_{phs}	Coefficient of volume expansion (K^{-1})
δ	Characteristic length (m)
Δh_f	Specific formation enthalpy (kJ gE^{-1})
Δh_r	Specific reaction enthalpy (kJ gE^{-1})
ΔH_r	Net reaction enthalpy (kJ d^{-1})
ε_{atm}	Surface emissivity
η_m	Efficiency of engines/aeration systems
$\bar{\lambda}_i$	Thermal conductivity of the i gaseous phase components ($\text{kJ d}^{-1} \text{K}^{-1} \text{m}^{-1}$)
λ_{atm}	Surface reflectivity
μ_{gl}	Dynamic viscosity of the gaseous phase components ($\text{kgE m}^{-1} \text{s}^{-1}$)
$v_{i,j}$	Rate coefficient for component i on process j ($\text{gE}_i \text{gE}_{\text{reference component}}^{-1}$)
$\bar{p}_{i,j}$	Kinetic rate for the transformations between i and j phases ($\text{gE}_{\text{removed component}} \text{d}^{-1}$)
σ	Stefan Boltzman constant ($\text{kJ d}^{-1} \text{m}^{-2} \text{K}^{-4}$)
ν_{gl}	Kinematic viscosity of the gaseous phase components ($\text{m}^2 \text{s}^{-1}$)

Superscripts

*	Absolute temperature (°C)
°	Standard state values (25°C)

Subscripts

g	gaseous phase
g ₁	1 st gaseous phase
g ₂	2 nd gaseous phase
in	Input
k	No. of state variables in the water phase
m	No. of state variables in the 1 st gaseous phase
n	No. of transformations in the water phase
out	Output
s	solid phase
z	No. of state variables in the 2 nd gaseous phase
w	Aqueous phase

32

33 **1. INTRODUCTION**

34 Operating temperature plays an important role in wastewater treatment efficiency (Metcalf and
35 Eddy, 2004). Temperature dynamics affect microbial activity as well as physicochemical properties,
36 such as dissolved oxygen saturation concentration, diffusivity, viscosity, density and the settling
37 velocity (Sedory and Stenstrom 1995). An increased temperature of a few degrees might stimulate
38 the metabolic activity of the bacteria, while a substantial reduction of several degrees would result
39 in reduced process stability, albeit temporarily, and a possible shift in the population of the reactor
40 (Gallert and Winter, 2005). The causes of these temperature variations are often due to a
41 combination of different factors. The most representative are conduction/convection phenomena,
42 short-wave and long-wave radiations, the heat transmitted by the actuators and, to a greater or lesser
43 degree, the heat produced or consumed by biochemical, chemical or physico-chemical
44 transformations. The influence of each term in the system is varied and depends largely on the
45 analysed process and climate of the place. In the activated sludge units, a typical diurnal

46 temperature difference between water inlet and outlet fluxes only varies between 0.5 and 1 degree
47 (Makinia *et al.*, 2005). However, in some parts of the world, treatment systems are subjected to
48 significant winter cooling and summer heating. In some nontemperate zones for instance, the
49 atmospheric diurnal air temperature variation can be considerable, ranging from 2 degrees during
50 the early morning to over 25 degrees during mid-afternoon (Paul, 2013), affecting considerably to
51 the water temperature. In membrane bioreactors, the temperature rise in the tank may be quite
52 significant comparing with the activated sludge reactors because of the higher biological heat
53 production. These reactors have a higher biomass concentration and bacterial activity, which
54 implies a higher biological heat production due to the exothermicity of the oxidation, nitrification
55 and denitrification reactions (Sethi *et al.*, 2011). In sludge line processes, the effect of temperature
56 has greater importance and practically all units work above atmospheric temperature. The anaerobic
57 digestion for example, is often an endothermic process which operates around 35 or 55 degrees,
58 thus requiring a heat supply to maintain the digester temperature and support the microbial activity
59 (Inoije *et al.*, 1996), and in the Autothermal Thermophilic Aerobic Digestion (ATAD) the organic
60 matter is oxidized under aerobic conditions with a concomitant biological heat release that is able to
61 maintain thermophilic temperatures. Despite the clear knowledge that exists to date on the
62 phenomena involved in the transfer of heat, the exact influence of temperature is difficult to
63 determine because of its interaction with mass transfer, chemical equilibrium and growth rate (Van
64 Hulle *et al.*, 2010). It is for this reason that the joint modelling of biological transformations and
65 accurate heat transfer models are becoming increasingly practical and necessary, not only to predict
66 the temperature of the system, but also to identify the diverse heat flows, to analyse the contribution
67 of these in the heat transfers, to relate the temperature variations with the chemical, biochemical and
68 physico-chemical transformations and, ultimately, to understand the process better. An accurate
69 temperature model allows determining a precise microbial activity, which is extremely important
70 aspect in the analysis of new operational strategies as well as new configurations

71 (www.novedar.com). Nowadays, thermal processes are increasingly being incorporated into
72 treatment plants (Daigger, 2011), making temperature prediction models essential for a proper use
73 and recovery of the heat.

74

75 Nonetheless, most of the existing mathematical models of activated sludge units and anaerobic
76 digestion models focus mainly on microbiology and often assume constant temperature (Makinia *et*
77 *al.*, 2005). There are several papers which have included temperature prediction in the general
78 model, the studies carried out by Novotny and Krenkel (1973), Argaman and Adams (1977),
79 Sedony and Stenstrom (1995), la Cour Jansen *et al.* (1992), Gillot and Vanrolleghem (2003),
80 Makinia *et al.* (2005) and Lippi *et al.* (2009) being some of the most representative works in water
81 line modelling. In the field of municipal sludge digestion, almost all heat transfer models published
82 in literature focus on the representation of the ATAD (Vismara, 1985; Messenger *et al.*, 1992 &
83 1993; Pitt *et al.*, 1994; Lapara *et al.*, 1999; Gómez *et al.*, 2007; and de Gracia *et al.*, 2009), focusing
84 less on anaerobic digestion. All these publications propose a detailed and accurate heat transfer
85 model for a precise prediction of the temperature. However, most of them are very specific for the
86 system under study, with a limited capacity for being expanded or adapted to include additional
87 transformations in water or gas phases. Moreover, the description of heat generated or consumed in
88 biochemical transformations are normally only based on COD removal. Therefore, a rigorous and
89 systematic methodology for constructing more detailed and flexible heat transfer models in
90 biochemical reactors is demanded.

91 Observing the low-detail used in the definition of the reaction heat, the first goal of this work has
92 been the development of a systematic, generic and rigorous methodology for the dynamic prediction
93 of the heat produced and consumed in all transformations of any biological reactor. And, the second
94 goal has been the restructuration of the mass and heat balances to allow the modeller to construct
95 mathematical models as complex as required in a systematic and modular way.

96

97 The complexity of today configurations and processes with recirculations and interrelations among
98 the different units makes it necessary to consider a plant-wide perspective in order to establish an
99 optimum solution for the design or operation of entire plants (Copp *et al.*, 2003; Grau *et al.* 2007;
100 Jeppsson *et al.*, 2007). The PWM methodology (Grau *et al.*, 2007) was proposed by the CEIT as a
101 systematic methodology for constructing mathematical models able to describe the whole plant as
102 complex as required in each case study. Along these years, the premises in which this methodology
103 was based (elemental mass characterisation) have been demonstrated as the most suitable
104 framework for developing mathematical models able to face with current needs, such as the
105 technologies where liquid, gaseous and solids phases coexist with energetic aspects. With this
106 frame, an update of the Plant-Wide Modelling (PWM; Grau *et al.*, 2007) methodology has been
107 carried out in order to describe technologies in which not only the water phase is considered but
108 also the gas phase in contact with it and the transfers among them and to incorporate heat balances
109 that allow a better estimation of process temperature and a proper use and recovery of the heat.

110

111 **2. EXTENDED PLANT-WIDE MODELLING METHODOLOGY (E-PWM)**

112 Plant-Wide Modelling (PWM) methodology (Grau *et al.*, 2007) was proposed for the systematic
113 construction of compatible unit-process models for describing the dynamic behaviour of the water
114 and sludge lines in an integrated way. This methodology is based on selecting the set of process
115 transformations required to model all unit-process elements incorporated into each specific WWTP.
116 For the unification and standardisation of this set of biochemical, chemical and physico-chemical
117 transformations, the incorporation of activation/inhibition terms in the kinetics must be accurate, in
118 order to reproduce the appropriate bacterial activity under aerobic, anoxic and anaerobic conditions.
119 Additionally, this unification permits the definition of a unique component vector for the whole
120 plant, without the need to develop specific transformers to interface the unit-process models,

121 making it a flexible methodology for both the definition of new reactions or processes and defining
122 new units. The accurate definition of the stoichiometry ensures the elemental mass (in terms of C,
123 N, O, H, P or other additional elements) and charge continuity in all the transformations. In order to
124 achieve this, some components must act as source-sink terms (or mass “compensation terms”) for
125 the correct adjustment of the C, N, O, H, P and charge balances (Reichert *et al.*, 2001; de Gracia *et*
126 *al.*, 2006). More details of this methodology can be found in Grau *et al.* (2007).

127

128 In this paper an update of PWM called Extended Plant-Wide Modelling methodology (E-PWM) is
129 proposed. This update is focused on facilitating the modelling of heat transfer phenomena in multi-
130 phase biochemical reactors. For this purpose, the methodology has been based mainly on two
131 aspects: (1) the multi-phase structure of the mathematical model and (2) the incorporation of
132 generic heat transfer models based on the automatic calculation of the enthalpy change of reaction
133 associated with all transformations of the system.

134

135 **2.1. Multi-phase model construction in the Extended Plant-Wide Modelling methodology**

136 The description of ASM models (Henze *et al.*, 2000) is based on the Gujer matrix notation that
137 presents the process kinetics and stoichiometry in a single product of matrices ($\tilde{E} \cdot \tilde{p}$). These models
138 only considered the water phase reactions, but in current technologies different aqueous, gaseous or
139 solid phases can be considered in a single unit-process multiplying the equations required to models
140 definition. With the purpose to develop flexible and easily understood models, the E-PWM
141 methodology proposes a general procedure for multi-phase model construction in which mass
142 balances are constructed systematically considering as many type of phases (gas, liquid, solid) and
143 number (1 liquid + 1 gas, 1 liquid + 2 gas phases, etc.) as required in each case study. The first step
144 of the procedure consists on the selection of the phases and transformations to model the WWTP
145 under study. Each phase (aqueous, gaseous or solid) as an independent system with its

146 corresponding mass vector (\bar{M}) will be considered. Accordingly, each phase will have an
 147 independent matrix ($\tilde{E} \cdot \bar{\rho}$) to define the transformations that take place in the phase (such as, for
 148 example, bacterial growth, or chemical equilibria reactions in the aqueous phase) and another set of
 149 matrices ($\tilde{E} \cdot \bar{\rho}$) to define the transformations between different phases (such as
 150 evaporation/condensation, stripping/absorption, precipitation/dissolution or deposition/sublimation).
 151 Thus, the model will be constructed using as many matrices as phases and interactions between
 152 phases. This modular construction can be visualized by means of the generic example depicted in
 153 Figure 1. And the second step consists on the definition of the mass balances of the unit processes.
 154 The definition of the set of matrices allows a systematic mass balance description based on the
 155 structure shown in equation 1.

$$\left(\frac{d\bar{M}}{dt}\right)_i = \tilde{E}_{i,i}^T \bar{\rho}_{i,i} + \sum_{j=1}^{\text{No. adj. phase}} \tilde{E}_{i,j}^T \bar{\rho}_{i,j} + \sum_{\text{in}} \bar{m}_{\text{phase}} - \sum_{\text{out}} \bar{m}_{\text{phase}} \quad (1)$$

156 $i = 1, 2 \dots \text{No. phases (number of phases)}$
 157 $j = 1, 2 \dots \text{No. adj. phase (number of adjacent phases to phase } i)$

158 Thus, the mass balance in each phase will be constructed as follows: the matrix ($\tilde{E}_{i,i} \cdot \bar{\rho}_{i,i}$) with the
 159 transformations that take place in the phase, plus the matrix with the transformations between
 160 different phases ($\tilde{E}_{i,j} \bar{\rho}_{i,j}$), plus the mass inputs and outputs of the system.

161
 162 This way of distinguishing the different phases and applying mass balances in each one, enables the
 163 modeller to construct systematically mathematical models as complex as required, considering
 164 different aqueous, gaseous or solid phases in a single unit-process. For example, in the case of
 165 layered models for settlers or biofilm systems in which biological reactions need to be included,
 166 each layer is considered to be an aqueous phase and mass transfers among them due to diffusion,
 167 convection or gravity effects are described by means of a stoichiometric matrix and kinetic vector.
 168 To understand better this methodology, an example is shown in section 3.

169

170 **2.2. New methodology for temperature prediction based on formation enthalpies**

171 As previously mentioned, the PWM methodology is based on the components characterization in
 172 their elemental mass and charge density. This quality is essential to avoid redundancies in
 173 component definition and to guarantee elemental mass and charge continuity throughout the whole
 174 plant. This detailed components characterization is also the one that enables the estimation of
 175 formation enthalpies for each model component and makes possible a systematic calculation of the
 176 heat released or absorbed by each transformation, guaranteeing heat energy continuity at any point
 177 in the plant. Thus, the specific enthalpy change of reaction (Δh_r° in $\text{kJ g}_{\text{stoich. unit}}^{-1}$) due to
 178 biochemical, physico-chemical or chemical transformations can be defined as the difference
 179 between the enthalpy of formation of the products and the enthalpy of formation of the reactants
 180 (applying Hess's law).

$$\Delta h_r^\circ = \sum v_{\text{products}} \Delta h_{\text{f(products)}}^\circ - \sum v_{\text{reactants}} \Delta h_{\text{f(reactants)}}^\circ \quad (2)$$

181 To calculate the net enthalpy change of reaction (ΔH_r° in KJ d^{-1}) instead, multiplying this specific
 182 enthalpy change of reaction by its corresponding kinetic and stoichiometry is sufficient:

$$\Delta H_r^\circ = \sum v_{\text{products}} \rho \Delta h_{\text{f(products)}}^\circ - \sum v_{\text{reactants}} \rho \Delta h_{\text{f(reactants)}}^\circ \quad (3)$$

183 Therefore, the net enthalpy change (see Equation 5) can be expressed as a function of the well-
 184 known Gujer transformation matrices (Equation 4; Henze *et al*, 2000) and the enthalpy of formation
 185 vector ($\overline{\Delta h_f^\circ}$).

$$\dot{x} = \tilde{E}^T \bar{\rho} \quad (4)$$

$$\Delta H_r^\circ = (\overline{\Delta h_f^\circ})^T \dot{x} \quad (5)$$

186 where,

$$\overline{\Delta h_f^\circ} = \begin{pmatrix} \Delta h_{f1} \\ \Delta h_{f2} \\ \vdots \\ \Delta h_{fj} \\ \vdots \\ \Delta h_{fk} \end{pmatrix}; \quad \tilde{E} = \begin{pmatrix} v_{11} & v_{12} & \dots & v_{1j} & \dots & v_{1k} \\ v_{21} & v_{22} & \dots & v_{2j} & \dots & v_{2k} \\ \vdots & \vdots & & \vdots & & \vdots \\ v_{i1} & v_{i2} & \dots & v_{ij} & \dots & v_{ik} \\ \vdots & \vdots & & \vdots & & \vdots \\ v_{n1} & v_{n2} & \dots & v_{nj} & \dots & v_{nk} \end{pmatrix}; \quad \bar{\rho} = \begin{pmatrix} \rho_1 \\ \rho_2 \\ \vdots \\ \rho_i \\ \vdots \\ \rho_n \end{pmatrix};$$

187 This close relationship maintaining mass and heat balances makes it a systematic methodology that
 188 can be easily integrated into the numerical solution of any existing mathematical model. The only
 189 requisite is the definition of the enthalpies of formation of the compounds present in these
 190 transformations. For this paper a set of transformations have been selected from the E-PWM library
 191 which gather all the components and transformations able to describe aerobic, anoxic and anaerobic
 192 COD biodegradation; nitrogen removal in two steps ($\text{NO}_3^- \rightarrow \text{NO}_2^- \rightarrow \text{N}_2$); anaerobic ammonia
 193 oxidation (Anammox process); aerobic or anaerobic sludge digestion; as well as, sludge
 194 pretreatments. Enthalpies of formation needed for the definition of these transformations are
 195 summarized in Table 1 and Table 2.

196

197 Most of these enthalpies of formation of of inorganic compounds, gaseous components and some
 198 organic compounds have been found in chemical handbooks and in specialized databases. To
 199 estimate the rest of enthalpies of formation by contrast, different methods and approaches have been
 200 used, such as the molecular group contribution approach (Constantinou and Gani, 1994; Hukkerikar
 201 *et al.*, 2013) or the strengths of bonds approach (Nelson *et al.*, 2005).

202

203 3. IMPLEMENTATION OF THE E-PWM METHODOLOGY IN AN AERATED 204 COMPLETELY STIRRED CLOSED TANK REACTOR

205 One illustrative example of the usefulness of the presented E-PWM methodology can be shown in
 206 aerated biological closed reactors. In these, an aqueous phase coexists with two gaseous phases; one
 207 corresponds to the total volume occupied by the bubbles submerged in the mixed liquor (gas hold-
 208 up) and the other one to the volume of the gas in contact with the free surface of the mixed liquor

209 (off-gas). The distinction of these volumes represents an interesting advantage over other simpler
 210 models, both for the oxygen transfer estimation, as for the correct prediction of the temperature in
 211 each phase.

212

213 3.1. Mass balance for an aerated biological closed reactor

214 For the analysis of three phases, three mass balances are necessary, as shown in equations 6, 7, and

215 8.

$$\left(\frac{d\bar{M}_w}{dt}\right) = \tilde{E}_{w,w}^T \bar{\rho}_{w,w} + \tilde{E}_{w,g1}^T \bar{\rho}_{w,g1} + \tilde{E}_{w,g2}^T \bar{\rho}_{w,g2} + \bar{m}_{w,in} - \bar{m}_{w,out} \quad (6)$$

$$\left(\frac{d\bar{M}_{g1}}{dt}\right) = \tilde{E}_{g1,w}^T \bar{\rho}_{g1,w} + \bar{m}_{g2,g1} - \bar{m}_{g,out} \quad (7)$$

$$\left(\frac{d\bar{M}_{g2}}{dt}\right) = \tilde{E}_{g2,w}^T \bar{\rho}_{g2,w} + \bar{m}_{g,in} - \bar{m}_{g2,g1} \quad (8)$$

216 For this specific case, the assumptions listed below have been made: (1) perfect and continuous
 217 mixing is assumed; (2) reactions in the gaseous phases have not been considered, only the aqueous
 218 phase reactions; (3) the transition from the first gaseous phase (gas hold-up) to the second gaseous
 219 phase (off-gas) has been defined as a mass transfer, and not as reaction. Thus, the first term of the
 220 aqueous mass balance ($\tilde{E}_{w,w}^T \bar{\rho}_{w,w}$) gathers all transformations that can take place in this phase
 221 (such as biochemical reactions or chemical equilibria among others) and the remaining
 222 transformations correspond to liquid-gas transfers ($\tilde{E}_{i,j}^T \bar{\rho}_{i,j}$) such as evaporation/condensation or
 223 stripping/absorption reactions (see Figure 2).

224

225 3.2. Heat balance for an aerated biological closed reactor

226 The completely mixing assumption commonly used in wastewater treatment plant process
 227 modelling, permits thermal uniformity in the reactor, avoiding temperature gradients. Thus, based
 228 on the energy conservation principle, the general one-dimensional dynamic heat transfer model

229 includes biochemical, chemical or physico-chemical heat transfers (H_{trans}), conduction and
 230 convection fluxes ($H_{\text{Cond/Conv}}$), heat energy fluxes transmitted by the actuators (H_{Act}) and short-
 231 wave (solar) and long-wave (atmospheric) radiations ($H_{\text{sol_rad}}$, $H_{\text{atm_rad}}$), as can be shown in
 232 equation 9.

$$\left(\frac{dH_T}{dt}\right)_{\text{phase}} = \sum H_{\text{trans}} + \sum H_{\text{Cond/Conv}} + \sum H_{\text{Act}} + \sum H_{\text{sol_rad}} + \sum H_{\text{atm_rad}} + \sum \bar{H}_{\text{in}} + \sum \bar{H}_{\text{out}} \quad (9)$$

233 The formulation for the time-derivative of enthalpy-state-variables has also followed the multi-
 234 phase principle proposed. Thus, equation 9 is transformed into the following equations:

$$\begin{aligned} \left(\frac{dH_w}{dt}\right) = & -(\overline{\Delta h_{fw}})^T \left(\tilde{E}_{w,w}^T \bar{p}_{w,w}\right) - (\overline{\Delta h_{fw}})^T \left(\tilde{E}_{w,g1}^T \bar{p}_{w,g1}\right) - (\overline{\Delta h_{fg1}})^T \left(\tilde{E}_{g1,w}^T \bar{p}_{g1,w}\right) \\ & - (\overline{\Delta h_{fw}})^T \left(\tilde{E}_{w,g2}^T \bar{p}_{w,g2}\right) - (\overline{\Delta h_{fg2}})^T \left(\tilde{E}_{g2,w}^T \bar{p}_{g2,w}\right) + H_{w,g1} + H_{w,g2} + H_{w,\text{in}} - H_{w,\text{out}} \\ & - H_{\text{phs1}} - H_{\text{phs2}} - H_{\text{wc,out}} + H_{\text{m,in}} - H_{\text{atm,rad,w}} + H_{\text{sol,rad,w}} \end{aligned} \quad (10)$$

$$\left(\frac{dH_{g1}}{dt}\right) = \sum_{i=1}^m (\bar{H}_{g1,g2}) + \sum_{i=1}^m (\bar{H}_{g1,w}) - \sum_{i=1}^m (\bar{H}_{g,\text{out}}) + H_{\text{phs1}} - H_{\text{gc,out}} - H_{\text{atm,rad,g}} + H_{\text{sol,rad,g}} \quad (11)$$

$$\left(\frac{dH_{g2}}{dt}\right) = \sum_{i=1}^z (\bar{H}_{g,\text{in}}) + \sum_{i=1}^z (\bar{H}_{g2,w}) + \sum_{i=1}^z (\bar{H}_{g2,g1}) + H_{\text{phs2}} \quad (12)$$

235 These expressions in Figure 3 have been schematically represented and the definition of each term
 236 in Table 3. Thus, the first term of the water heat balance $\left((\overline{\Delta h_{fw}})^T \left(\tilde{E}_{w,w}^T \bar{p}_{w,w}\right)\right)$ refers to the heat
 237 associated to the transformations that take place in the aqueous phase, namely, it encompasses the
 238 heat of all transformations that occur in this phase. The second term $\left((\overline{\Delta h_{fw}})^T \left(\tilde{E}_{w,g1}^T \bar{p}_{w,g1}\right)\right)$
 239 provides the stoichiometric enthalpies of formation of the components present in the aqueous phase,
 240 and the third term $\left((\overline{\Delta h_{fg1}})^T \left(\tilde{E}_{g1,w}^T \bar{p}_{g1,w}\right)\right)$ the stoichiometric enthalpies of formation of the
 241 components present in the first gaseous phase. Applying the first law of thermodynamics, the sum

242 of both terms refers to the water-gas/gas-water convective transferences, namely, the heat
243 transferred due to the stripping/absorption and evaporation/condensation transferences. In such a
244 transfer, besides heat production or consumption, an enthalpy associated to the matter is transferred
245 between the phases, producing an increase in the net heat of the target phase, though not of the
246 specific heat. In the model these terms have been defined as enthalpy inputs and outputs due to
247 transformations, as can be seen in Table 3.

248

249 By way of simplification, for the aqueous flow heat estimation, only water enthalpy has been
250 considered, ignoring the enthalpy of dissolved compounds that make up the aqueous phase. In the
251 modelling of convective phenomenon, natural convection heat transfer on the phase surface
252 depends on the contact surface, the temperature difference between the phases and also on the
253 thermophysical properties of the phases involved (see Table 3). One of the most noticeable
254 differences to be considered in open tank modelling is precisely this term. In closed tank modelling,
255 natural convection is considered, whereas in open tank modelling with forced convection, this term
256 can be highly influenced by wind velocity and temperature differences between the phases and the
257 atmosphere.

258

259 **4. VALIDATION OF THE METHODOLOGY FOR ESTIMATING**

260 **TRANSFORMATION HEAT (ΔH_r°)**

261 First validation of the proposed methodology has been focused on comparing the theoretical model-
262 based heat values obtained with the E-PWM with other studies. Table 4 shows a comparison
263 between the transformation heat values calculated with the proposed methodology and a set of
264 values presented in bibliography, previously estimated both experimentally (E) or theoretically (T).
265 In order to validate the methodology as rigorously as possible, some model reactions had to be
266 modified (for example, some bibliographic reactions don't consider the biomass growth) to make
267 more validation cases available.

268

269 It is interesting to note that the deviations with almost all reactions have been as low as 5%, with the
270 exception of the enthalpy change of reaction estimated theoretically for the conversion of acetic
271 acid to methane (reaction 7), where the discrepancies may be due to differences in the enthalpies of
272 formation used for the products and/or reactants. Therefore, these comparative results confirm that
273 the enthalpy change of reaction estimated by the Extended Plant-Wide modelling methodology is in
274 accordance with the values traditionally estimated both theoretically and experimentally.

275

276 Despite numerous experimental studies published, there is no consensus on the determination of the
277 specific heat yields (see Table 4), although it has been estimated an approximate range of 12–18 kJ
278 gCOD⁻¹ removed. These discrepancies can be attributed to the heterogeneity in the samples'
279 composition due to the diversity in the organic components present (Cooney *et al.*, 1968). It is also
280 important to note that, although the enthalpy change of reaction values proposed in literature are
281 attributed to the oxidation of organic materials, these experimental estimations implicitly include
282 other smaller transformation heats like, for example, those associated with hydrolysis or CO₂
283 stripping. In order to show the additional information offered by the proposed E-PWM
284 methodology, Table 5 first row presents the theoretical specific heat yields for the seven
285 carbonaceous substrates included in the model under study, showing the small differences
286 associated with substrate composition, but within the range proposed in literature. Additionally, the
287 second row illustrates the expected changes in the theoretical heats when substrate oxidation and
288 CO₂ stripping transformations are both considered in the calculations, demonstrating the relevance
289 of taking into account all the simultaneous transformations. This fact can explain some of the
290 discrepancies found in the experimental results presented in the literature.

291

292 Once the appropriate calculation of the specific heat yields has been validated, the applicability of

293 the proposed E-PWM methodology for the dynamic simulation of heat fluxes and temperature
294 prediction has been analysed. For this, two illustrative examples have been constructed and
295 implemented in the WEST simulation platform (www.mikebydhi.com), for its subsequent analysis.

296

297 **4.1. Example I: Analysis of dynamic heat exchanges in a WWTP biological reactor**

298 This first case study shows the usefulness of the E-PWM methodology for a detailed prediction of
299 the dynamic evolution of the heat exchanges generated in the biological reactors of WWTPs. The
300 simulation example corresponds to the water line of the BSM2 layout (Jeppsson *et al.*, 2007),
301 maintaining its physical attributes and using the 365-day dynamic influent data file. The data of
302 atmospheric temperature, solar radiation and wind velocity have been extracted from the Spanish
303 National Meteorological Agency (<http://www.aemet.es>) corresponding to measurements made in
304 northern Spain in 2012. Regarding the influent water temperature, the profile has been estimated
305 from atmospheric data and previous year's tendencies in that area (every day of the year, the input
306 water temperature is higher than the temperature of the atmosphere).

307

308 Figure 4a shows the predicted contribution of the most significant heat fluxes in the overall balance.
309 The mild temperatures and abundant rainfall that characterized the oceanic climate means that the
310 solar radiation flux (ΔH radiation) and the conduction and convection fluxes (ΔH cond/conv) do not
311 have the significance that can be expected in other areas. Although surprised that the term of heat
312 transfer due to aeration hasn't appeared in the figure 4a, this term in the ΔH cond/conv term has
313 been included along with the transfer between the atmosphere and the water phase. The aeration
314 provide heat to the system but this heat is offset by losses due to contact between the atmosphere
315 and the water.

316 Figure 4b shows a detailed description of the main heat fluxes within the reactors, differentiating
317 between the heat supplied by the biological transformations and the heat consumed by evaporation

318 and stripping. Simulations results clearly show how the most significant heat fluxes are due to COD
319 removal (denitrification in anoxic reactors and heterotrophic growth in aerobic reactors) and
320 nitrification reactions. To a lesser extent, but also relevant is the stripping of CO₂ and evaporation.

321

322 Simulation results (not presented) have shown that the expected overall increase in the water
323 temperature from influent to effluent ranges between 0.8 and 1.5 degrees depending on the
324 radiation term, coinciding with the ranges previously published by la Cour Jansen *et al.* (1992) and
325 Makinia *et al* (2005). The proposed E-PWM methodology facilitates a detailed description of the
326 contribution of different transformations in the heat flows and temperature variations. For example,
327 at this particular case-study, simulation results show that the nitrification and heterotrophic growth
328 reactions increase the tank temperature 0.6-0.8 degrees while denitrification only by 0.23 degrees.

329 Evaporation and stripping reactions are widely influenced by atmospheric temperature, and
330 decrease the temperature by 0.05-0.2 degrees. As expected, the effect of transformation heats in
331 process temperature is not very relevant in conventional activated sludge reactors. However, it can
332 be very significant in other processes that exhibit higher biological activity and lower heat transport
333 with the environment like, for example, the Autothermal Thermophilic Aerobic Digesters (ATAD).

334

335 **4.2. Example II: Analysis of the heat exchanges generated in an ATAD reactor**

336 The ATAD digesters exhibit a high organic matter oxidation, being able to maintain thermophilic
337 conditions without the external addition of heat (except recirculations/mixing energy). To analyse
338 the predictive capacity of the methodology applied to an ATAD reactor, the first step has been the
339 comparison between the simulation results and the experimental data obtained in the full scale
340 ATAD at the Tudela WWTP (de Garcia *et al.*, 2009; Gomez *et al.*, 2007). This ATAD digester is a
341 completely stirred tank reactor with a total effective volume of 350 m³. In the analysed 14-day
342 period, the digester operated at 10 days hydraulic retention time with discontinuous charges and

343 discharges of the reactor every 24h. A period with abrupt changes in the organic loading rate (OLR)
344 was selected for the validation of the thermal model under both oxygen limiting and substrate
345 limiting conditions. To achieve these two conditions, the aeration flow rate of $9,200 \text{ m}^3 \text{ d}^{-1}$
346 remained constant, gradually increasing the OLR (see Table 6). A detailed description of the
347 stoichiometric, kinetic and physico-chemical parameters can be found in de Gracia *et al.* (2009).

348

349 The dynamic results obtained in the simulation are shown in Figure 5a, where the experimental and
350 simulated aqueous phase temperatures obtained in the selected period are compared, obtaining a
351 proper fit both in terms of substrate limiting conditions (days 0 to 9) and oxygen limiting conditions
352 (days 9 to 14). The evolution of temperature in oxygen limiting conditions makes possible the
353 calibration of the oxygen transfer coefficient (k_{La}), whereas temperature data under substrate
354 limiting conditions supplies the information required for calibrating the heat flux associated with
355 aeration.

356

357 By analysing the thermal flux of the system (Figure 5b), it can be seen that biological processes
358 (ΔH_r water) are the most important heat flux contributors, representing 60-70% of the heat in the
359 process. The contour of this flux area shows the evolution of organic matter oxidation, clearly
360 differentiating substrate limiting and oxygen limiting areas. The heat dissipated due to the liquid
361 and gas phase transformations (ΔH_r water-gas) is the second most important heat flux, representing
362 15-30% of the heat in the process. Of this percentage, 10% corresponds to CO_2 stripping, and the
363 remaining 90% to the heat lost by evaporation. In this example, atmospheric and solar radiation
364 have not been graphed since they were negligible compared to other terms.

365

366 The proposed E-PWM methodology also makes possible a detailed dynamic analysis of the
367 different terms in heat generation and transport as a function of the influent load or operational

368 strategies. Next examples briefly show the effect of ventilation and the substrate fed in the ATAD
369 temperature.

370

371 **4.2.1. Model-based exploration of the effect of air flow in an ATAD digester**

372 Aeration is a key factor in aerobic digestion from the point of view of biological reaction and heat
373 balance. If the process is working with oxygen concentrations under the stoichiometric value
374 (oxygen limiting conditions), the oxygen transfer will mark the substrate consumption rate of the
375 system and therefore the sludge heating (Messenger *et al.*, 1990). Thus, the digester temperature can
376 be controlled with the air flow, until the substrate is exhausted or the stoichiometric oxygen value is
377 reached. Conversely, if the system is working with oxygen concentrations over the stoichiometric
378 (substrate limiting conditions), the heating degree will depend on the amount of substrate fed. In
379 contrast to oxygen limiting conditions, an excess of aeration can cause digester cooling due to the
380 temperature gradient of the bubbles and the liquid phase.

381

382 The model's ability to identify these ventilation effects is essential for an appropriate selection of
383 the operational conditions. Still, it is important to mention that for a correct prediction of the heat of
384 reaction, it is necessary to estimate experimentally the respiration quotient, especially important
385 when the system is working under substrate limiting conditions. Figure 6 shows an exploration of
386 the expected reactor temperature of the Tudela ATAD for the same OLR of $4.0 \text{ Kg COD}\cdot\text{m}^{-3} \text{ d}^{-1}$
387 under different aeration flows. It can be clearly observed how excess air can cause a cooling of the
388 aqueous phase, decreasing the temperature to values obtained with oxygen limiting conditions, with
389 the disadvantage of having higher costs. Model-based exploration can facilitate the selection of the
390 most appropriate aeration flow. The proposed E-PWM methodology is already prepared for dealing
391 with different gas phases, making possible the simulation of pure oxygen supply.

392

393 4.2.2. Model-based exploration of the effect of influent characterization in an ATAD digester

394 Different studies (Heijnen, 1999; Heidrich *et al.*, 2011; Hill *et al.*, 2012) have shown that oxidation
395 heat is closely related to the composition of the feed. In this section, the effect of a change in the
396 composition of the sludge fed in the liquid phase has been explored by simulation, using the E-
397 PWM methodology. Table 7 shows the three influent characterisations used for the model-based
398 exploration.

399
400 Figure 7a shows the predicted evolution of process temperature for the three load characterisations.
401 The rise in the concentration of carbohydrates has led to an increase in liquid temperature of 0.4 °C
402 for this specific OLR (a decrease of 0.4 °C for Characterisation 2). This temperature variation
403 would have been greater using a higher OLR or energy crops such as maize grains, artichoke or
404 wheat, among others.

405
406 The enthalpy change of reaction of each characterisation is plotted in Figure 7b showing the
407 capacity of E-PWM methodology for exploring the effect of influent characterisation in
408 temperature. As expected the greatest heat are obtained for the influent rich in carbohydrates. It is
409 interesting to note that the global transformations heat and, consequently the specific heat yield, is
410 varying in time, depending on the evolution of substrate fractioning within the reactor.

411 412 5. CONCLUSIONS

413 The paper presents a new methodology, based on the well-known Hess's law, for the rigorous and
414 systematic calculation of heat balances in biochemical reactors. Starting from a complete
415 description of model components stoichiometry and enthalpies of formation, the proposed
416 modelling methodology, called Extended Plant-Wide Modelling (E-PWM), has integrated
417 successfully the simultaneous calculation of both the conventional mass balances and the heat

418 balances in an expandable multi-phase matrix structure, which facilitates a detailed prediction of the
419 main heat fluxes in the biochemical reactors. Without wanting to focus on the mere calculation of
420 heat exchanges in a single reactor, the heat balance has been also integrated into the CEIT Plant-
421 Wide Modelling methodology (Grau *et al*, 2007) in order to conduct a comprehensive analysis of
422 entire WWTPs.

423

424 The transformation heats and specific heat yields estimated with the proposed methodology have
425 been successfully contrasted with other experimental or theoretical studies previously presented in
426 the literature. Additionally, the E-PWM offers very comprehensive information about the different
427 heat fluxes associated with each particular reaction or transport, facilitating, for example, an
428 exhaustive analysis of the effect of operational conditions or influent fractioning in temperature
429 evolution. Finally, it is interesting to remark that the proposed heat transfer modelling methodology
430 can be easily incorporated to the existing WWTPs simulators that are based on a detailed
431 description of model components and transformations like, for example, those based on the PWM
432 methodology.

433

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437

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Table 1 - Gas phase formation enthalpies

i	Name	Formula	Description	Δh_f° (kJ mol ⁻¹)	Δh_f° (kJ g _{stoich. unit} ⁻¹)	Reference
1	G _{CO2}	CO ₂	Carbon dioxide	-393.51	-32.79 kJ g _C ⁻¹	Perry & Green (1999)
2	G _{H2}	H ₂	Hydrogen	0.00	0.00 kJ g _{COD} ⁻¹	Perry & Green (1999)
3	G _{CH4}	CH ₄	Methane	-74.80	-1.17 kJ g _{COD} ⁻¹	Perry & Green (1999)
4	G _{NH3}	NH ₃	Ammonia	-45.90	-3.28 kJ g _N ⁻¹	Perry & Green (1999)
5	G _{N2}	N ₂	Nitrogen	0.00	0.00 kJ g _N ⁻¹	Perry & Green (1999)
6	G _{O2}	O ₂	Oxygen	0.00	0.00 kJ g _{O2} ⁻¹	Perry & Green (1999)
7	G _{H2O}	H ₂ O	Water steam	-241.81	-13.43 kJ g _{H2O} ⁻¹	Perry & Green (1999)

Table 2 - Water phase formation enthalpies

i	Name	Formula	Description	Δh_f° (kJ mol ⁻¹)	Δh_f° (kJ g _{stoich} ⁻¹)	Reference
1	S _{H2O}	H ₂ O	Water steam	-285.84	-15.88 kJ g _{H2O} ⁻¹	Perry & Green (1999)
2	S _{O2}	O ₂	Dissolved Oxygen	0.00	0.00 kJ g _{O2} ⁻¹	Perry & Green (1999)
3	S _{H⁺}	H ⁺	Protons	0.00	0.00 kJ g _H ⁻¹	Brown <i>et al.</i> (1993)
4	S _{OH⁻}	OH ⁻	Hydroxide ions	-230.00	-230.00 kJ g _H ⁻¹	Brown <i>et al.</i> (1993)
5	S _{HPO4=}	(HPO ₄) ⁼	Hydroxy phosphate	-1298.70	-41.89 kJ g _P ⁻¹	Chang (1999)
6	S _{H2PO4-}	(H ₂ PO ₄) ⁻	Dihydroxy	-1302.48	-42.01 kJ g _P ⁻¹	Chang (1999)
7	S _{NH4+}	(NH ₄) ⁺	Ammonium	-132.50	-9.46 kJ g _N ⁻¹	Brown <i>et al.</i> (1993)
8	S _{NH3}	NH ₃	Ammonia	-80.29	-5.73 kJ g _N ⁻¹	Brown <i>et al.</i> (1993)
9	S _{CO2}	CO ₂	Dis. Carbon dioxide	-412.90	-34.41 kJ g _C ⁻¹	Chang (1999)
10	S _{HCO3-}	(HCO ₃) ⁻	Bicarbonate	-691.10	-57.59 kJ g _C ⁻¹	Chang (1999)
11	S _{SU}	C ₆ H ₁₂ O ₆	Monosaccharides	-1268.20	-6.61 kJ g _{COD} ⁻¹	Reger <i>et al.</i> (2010)
12	S _{AA}	C ₄ H _{6.10} O _{1.2N}	Amino acids	-306.10	-2.29 kJ g _{COD} ⁻¹	NIST
13	S _{FA}	C ₁₆ O ₂ H ₃₂	LCFAs	-848.40	-1.15 kJ g _{COD} ⁻¹	Lebedeva (1964)
14	S _{HVA}	C ₅ H ₁₀ O ₂	Valeric acid	-558.90	-2.69 kJ g _{COD} ⁻¹	Adriaanse <i>et al.</i> (1965)
15	S _{VA-}	C ₅ H ₉ O ₂ ⁻	Valerate	-501.07	-2.4128 kJ g _{COD} ⁻¹	Constantinou <i>et al.</i> (1994) Hukkerikar <i>et al.</i> , 2013
16	S _{HBU}	C ₄ H ₈ O ₂	Butyric acid	-533.92	-3.34 kJ g _{COD} ⁻¹	Lebedeva (1964)
17	S _{BU-}	C ₄ H ₇ O ₂ ⁻	Butyrate	-519.20	-3.25 kJ g _{COD} ⁻¹	Everett <i>et al.</i> (1939)
18	S _{HPRO}	C ₃ H ₆ O ₂	Propionic acid	-510.80	-4.56 kJ g _{COD} ⁻¹	Lebedeva (1964)
19	S _{PRO-}	C ₃ H ₅ O ₂ ⁻	Propionate	-507.79	-4.53 kJ g _{COD} ⁻¹	Everett <i>et al.</i> (1939)
20	S _{HAC}	C ₂ H ₄ O ₂	Acetic acid	-483.52	-7.56 kJ g _{COD} ⁻¹	Steele <i>et al.</i> (1997)
21	S _{AC-}	C ₂ H ₃ O ₂ ⁻	Acetate	-482.09	-7.53 kJ g _{COD} ⁻¹	Everett <i>et al.</i> (1939)
22	S _{H2}	H ₂	Hydrogen	0.00	0.00 kJ g _{COD} ⁻¹	Perry & Green (1999)
23	S _{CH4}	CH ₄	Dis. Methane	-82.97	-1.30 kJ g _{COD} ⁻¹	NIST
24	S _{N2}	N ₂	Dis. Nitrogen	0.00	0.00 kJ g _N ⁻¹	Perry & Green (1999)
25	S _{NO2-}	NO ₂ ⁻	Nitrites	-104.60	-7.47 kJ g _N ⁻¹	Masterton & Hurley (2003)
26	S _{HNO2}	HNO ₂	Nitrous acid	-116.00	-8.29 kJ g _N ⁻¹	Guillaumont <i>et al.</i> (2003)
27	S _{NO3-}	(NO ₃) ⁻	Nitrates	-206.57	-14.76 kJ g _N ⁻¹	Chang (1999)
28	S _I	C ₇ H _{9.1} O _{2.65} NP _{0.05}	Soluble Inerts	-495.00	-1.34 kJ g _{COD} ⁻¹	Need estimation
29	S _P	C ₇ H _{9.1} O _{2.65} NP _{0.05}	Lysis sol. Product	-495.00	-1.34 kJ g _{COD} ⁻¹	Need estimation
30	X _{C1}	C _{13.7} H ₂₄ O _{3.8} N _{0.5} P _{0.035}	Composites	-555.90	-1.94 kJ g _{COD} ⁻¹	Need estimation
31	X _{C2}	C ₅ H _{6.9} O ₂ NP _{0.1}	Decay complex	-414.02	-2.54 kJ g _{COD} ⁻¹	Prochazka <i>et al.</i> (1973)
32	X _{CH}	C ₆ H _{9.95} O ₅ P _{0.05}	Carbohydrates	-979.00	-5.06 kJ g _{COD} ⁻¹	Brown (2011)
33	X _{PR}	(C ₄ H _{6.1} O _{1.2} N) _x	Proteins	-284.10	-2.13 kJ g _{COD} ⁻¹	Nelson <i>et al.</i> , (2005)
34	X _{LI}	C ₅₁ H _{97.9} O ₆ P _{0.1}	Lipids	-2474.17	-1.06 kJ g _{COD} ⁻¹	Freedman <i>et al.</i> (1989)
35	X _H	C ₅ H _{6.9} O ₂ NP _{0.1}	Heterotrophic bac.	-414.02	-2.537 kJ g _{COD} ⁻¹	Prochazka <i>et al.</i> (1973)
36	X _{AOB}	C ₅ H _{6.9} O ₂ NP _{0.1}	Nitrosomona bac.	-414.02	-2.537 kJ g _{COD} ⁻¹	Prochazka <i>et al.</i> (1973)
37	X _{NOB}	C ₅ H _{6.9} O ₂ NP _{0.1}	Nitrobacter bac.	-414.02	-2.537 kJ g _{COD} ⁻¹	Prochazka <i>et al.</i> (1973)
38	X _{SU}	C ₅ H _{6.9} O ₂ NP _{0.1}	Sugar degrader bac.	-414.02	-2.537 kJ g _{COD} ⁻¹	Prochazka <i>et al.</i> (1973)
39	X _{AA}	C ₅ H _{6.9} O ₂ NP _{0.1}	Amino-acid	-414.02	-2.537 kJ g _{COD} ⁻¹	Prochazka <i>et al.</i> (1973)
40	X _{FA}	C ₅ H _{6.9} O ₂ NP _{0.1}	LCFA degrader bac.	-414.02	-2.537 kJ g _{COD} ⁻¹	Prochazka <i>et al.</i> (1973)
41	X _{C4}	C ₅ H _{6.9} O ₂ NP _{0.1}	Val/but degrader	-414.02	-2.537 kJ g _{COD} ⁻¹	Prochazka <i>et al.</i> (1973)
42	X _{PRO}	C ₅ H _{6.9} O ₂ NP _{0.1}	Propionate degrader	-414.02	-2.537 kJ g _{COD} ⁻¹	Prochazka <i>et al.</i> (1973)
43	X _{AC}	C ₅ H _{6.9} O ₂ NP _{0.1}	Acetate degrader	-414.02	-2.537 kJ g _{COD} ⁻¹	Prochazka <i>et al.</i> (1973)
44	X _{H2}	C ₅ H _{6.9} O ₂ NP _{0.1}	Hydrogen degrader	-414.02	-2.537 kJ g _{COD} ⁻¹	Prochazka <i>et al.</i> (1973)
45	X _{AN}	C ₅ H _{6.9} O ₂ NP _{0.1}	Anammox bac.	-414.02	-2.537 kJ g _{COD} ⁻¹	Prochazka <i>et al.</i> (1973)
46	X _I	C ₇ H _{9.1} O _{2.65} NP _{0.05}	Particulate inert	-718.41	-3.11 kJ g _{COD} ⁻¹	Need estimation
47	X _P	C ₇ H _{9.1} O _{2.65} NP _{0.05}	Lysis particulate	-718.41	-3.11 kJ g _{COD} ⁻¹	Need estimation

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48	X _{II}	X	Inorganic inert	-	-	-
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Table 3 - Variables, parameters and expressions used in the energy balance of a C-CSTR

	Term	Description	Equations
Input / Output due to Transformations	$\bar{H}_{g1,g2}$	Enthalpy lost/gained by the 1 st gaseous phase due to transformation (from/to the 2 nd gaseous phase)	$(\bar{E}_{g1,g2}^T \bar{p}_{g1,g2})_i [(Cp_{g1})_i (T_{g1} - (T_{g1,ref})_i) + (h_{g1,ref})_i]$
	$\bar{H}_{g1,w}$	Enthalpy lost/gained by the 1 st gaseous phase due to transformation (from/to the aqueous phase)	$(\bar{E}_{g1,w}^T \bar{p}_{g1,w})_i [(Cp_{g1})_i (T_{g1} - (T_{g1,ref})_i) + (h_{g1,ref})_i]$
	$\bar{H}_{g2,g1}$	Enthalpy lost/gained by the 2 nd gaseous phase due to transformation (from/to the 1 st gaseous phase)	$(\bar{E}_{g2,g1}^T \bar{p}_{g2,g1})_i [(Cp_{g2})_i (T_{g2} - (T_{g2,ref})_i) + (h_{g2,ref})_i]$
	$\bar{H}_{g2,w}$	Enthalpy lost/gained by the 2 nd gaseous phase due to transformation (from/to the aqueous phase)	$(\bar{E}_{g2,w}^T \bar{p}_{g2,w})_i [(Cp_{g2})_i (T_{g2} - (T_{g2,ref})_i) + (h_{g2,ref})_i]$
Input	$H_{w,g1}$	Enthalpy lost/gained by the aqueous phase due to transformation (from/to the 1 st gaseous phase)	$(\bar{E}_{w,g1}^T \bar{p}_{w,g1})_{H_2O} [Cp_{w,H_2O} (T_w - T_{w,ref,H_2O}) + h_{w,ref,H_2O}]$
	$H_{w,g2}$	Enthalpy lost/gained by the aqueous phase due to transformation (from/to the 2 nd gaseous phase)	$(\bar{E}_{w,g2}^T \bar{p}_{w,g2})_{H_2O} [Cp_{w,H_2O} (T_w - T_{w,ref,H_2O}) + h_{w,ref,H_2O}]$
Output	$H_{w,in}$	Enthalpy associated with the aqueous influent	$[Cp_{w,H_2O} (T_{w,in} - T_{w,ref,H_2O}) + h_{w,ref,H_2O}] (\dot{m}_{w,in})_{H_2O}$
	$\bar{H}_{g,in}$	Enthalpy associated with the gaseous influent	$[(Cp_{g2})_i (T_{g2,in} - (T_{g2,ref})_i) + (h_{g2,ref})_i] (\dot{m}_{g,in})_i$
	$H_{w,out}$	Enthalpy associated with the aqueous effluent	$[Cp_{w,H_2O} (T_w - T_{w,ref,H_2O}) + h_{w,ref,H_2O}] (\dot{m}_{w,out})_{H_2O}$
	$\bar{H}_{g,out}$	Enthalpy associated with the gaseous effluent	$[(Cp_{g1})_i (T_{g1} - (T_{g1,ref})_i) + (h_{g1,ref})_i] (\dot{m}_{g,out})_i$
Conduction / Convection	H_{phs1}	Enthalpy flow transfer between the water and the 1 st gaseous phase	$\frac{\alpha_{phs,g1} (Gr Pr)^{n_{phs,g1}} (\bar{k}_{g1} \bar{X}_{g1}^T) (T_w - T_{g1}) A_{tank}}{\delta}$
	H_{phs2}	Enthalpy flow transfer between the water and the 2 nd gaseous phase	$\frac{\alpha_{phs,g2} (Gr Pr)^{n_{phs,g2}} (\bar{k}_{g2} \bar{X}_{g2}^T) (T_w - T_{g2}) A_{bubbles}}{\delta}$
	$H_{gc,out}$	Enthalpy lost/gained through walls and pipes by conduction (gaseous phase)	$k_{wls} (T_w - T_{atm}) A_w$
	$H_{wc,out}$	Enthalpy lost/gained through walls and pipes by conduction (aqueous phase)	$k_{wls} (T_{g1} - T_{atm}) A_g$
Radiation	$H_{atm,rad,g}$	Longwave atmospheric radiation flux (gaseous phase)	$k_{SB} A_g [\varepsilon_{atm} (T_{g1}^*)^4 - (1 - \lambda_{atm}) \beta_{air} (T_{atm}^*)^4]$
	$H_{atm,rad,w}$	Longwave atmospheric radiation flux (aqueous phase)	$k_{SB} A_w [\varepsilon_{atm} (T_w^*)^4 - (1 - \lambda_{atm}) \beta_{air} (T_{atm}^*)^4]$
	$H_{sol,rad,g}$	Shortwave solar radiation flux (gaseous phase)	$\alpha_{rad} k_{sol,rad} A_g$
	$H_{sol,rad,w}$	Shortwave solar radiation flux (aqueous phase)	$\alpha_{rad} k_{sol,rad} A_w$
Actuator	$H_{m,in}$	Enthalpy transmitted from the actuators (mechanical)	$P_m (1 - \eta_m)$

where,

$$Gr = \frac{g \beta_{phs} (T_w - T_{med}) \delta^3}{(\bar{u}_{med} \bar{X}_{med}^T)^2}$$

$$Pr = \frac{(\bar{\mu}_{med} \bar{X}_{med}^T) (\bar{Cp}_{med} \bar{X}_{med}^T)}{(\bar{k}_{med} \bar{X}_{med}^T)}$$

Table 4 – Comparison of transformation heat estimated in modelling with experimental and theoretical literature data

Reaction	Δh_r°		Reference
	Model	Literature	
1 Substrate (aq.) + xO ₂ (g) → yH ₂ O(l) + zCO ₂ (g)+Biomass		12.28 – 16.46 kJ/gCOD _{rem.} ^(E) 13.32 – 15.12 kJ/gCOD _{rem.} ^(E) 14.30 kJ/gCOD _{rem.} ^(E) 13.5 kJ/gCOD _{rem.} ^(E) 12.50 – 16.66 kJ/gCOD _{rem.} ^(E) 13.6 kJ/gCOD _{rem.} ^(E) 13.62 – 14.62 kJ/gCOD _{rem.} 12.1 – 13.7 kJ/gCOD _{rem.} ^(E) 12.8 kJ/gCOD _{rem.} ^(E) 15.68 kJ/gCOD _{rem.} ^(E) 12.4 kJ/gCOD _{rem.} ^(E) 13.9 kJ/gCOD _{rem.} ^(E) 17.8 kJ/gCOD _{rem.} ^(E)	Cooney <i>et al.</i> , 1968 Andrews and Kambhu, 1969 McCarty, 1972 Zanoni and Mueller, 1982 Jewell, 1982 Heinritz <i>et al.</i> , 1990 Messenger <i>et al.</i> , 1992 Pitt <i>et al.</i> , 1995 Riley and Forster, 2002 Shizas and Bagley, 2004 Gomez <i>et al.</i> , 2007 Heidrich <i>et al.</i> , 2011
2 C ₆ H ₁₂ O ₆ (aq.)+3O ₂ (g) →3H ₂ O(l) +3CO ₂ (g)+Biomass	-893.1 kJ/mol _{SSU rem.}	-890.0 kJ/mol _{SSU rem.} ^(T)	Gallert <i>et al.</i> , 2005
3 NH ₄ ⁺ (aq.)+1.5 O ₂ (g) → 2H ⁺ (aq.) + NO ₂ ⁻ (aq.)+H ₂ O (l)	-258.0 kJ/mol _{N rem.}	-259.0 kJ/mol _{N rem.} ^(T)	Daverio <i>et al.</i> , 2003
4 NO ₂ ⁻ (aq.)+0.5 O ₂ (g) → NO ₃ ⁻ (aq.)	-102.0 kJ/mol _{N rem.}	-99.4 kJ/mol _{N rem.} ^(T)	Daverio <i>et al.</i> , 2003
5 NH ₄ ⁺ (aq.)+NO ₂ ⁻ (aq.) → N ₂ (g)+ 2H ₂ O (l)	-334.6 kJ/mol _{N rem.}	-333.0 kJ/mol _{N rem.} ^(T)	Buttiglieri <i>et al.</i> , 2010
6 C ₆ H ₁₂ O ₆ (aq.)→ 2.85CH ₄ (g) + 2.85CO ₂ (g)	-132.8 kJ/mol _{SSU rem.}	-131.0 kJ/mol _{SSU rem.} ^(T)	Gallert <i>et al.</i> , 2005
7 CH ₃ COOH(aq.)→ CH ₄ (aq.) + CO ₂ (aq.)	-12.4 kJ/mol _{SHAC rem.}	-15.3 kJ/mol _{SHAC rem.} ^(T)	Oh <i>et al.</i> , 2007
8 CH ₃ COOH(aq.)→ CH ₄ (g) + CO ₂ (aq.)	-4.8 kJ/mol _{SHAC rem.}	$\Delta h_r^\circ < 0$ ^(T)	Oh <i>et al.</i> , 2007
9 CH ₃ COOH(aq.)→ CH ₄ (g) + CO ₂ (g)	15.2 kJ/mol _{SHAC rem.}	$\Delta h_r^\circ > 0$ ^(T)	Oh <i>et al.</i> , 2007
10 HAC digestion (aq.) (methanogenesis)	-9.8 kJ/mol _{SHAC rem.}	$\Delta h_r^\circ < 0$ ^(T)	Oh <i>et al.</i> , 2010
11 HPRO digestion (aq.) (acetogenesis & methanogenesis)	0.6 kJ/mol _{SHPRO rem.}	$\Delta h_r^\circ > 0$ $\Delta h_{r,11}^\circ > \Delta h_{r,10}^\circ$ ^(T)	Oh <i>et al.</i> , 2010
12 HBU digestion (aq.) (acetogenesis & methanogenesis)	6.8 kJ/mol _{SHBU rem.}	$\Delta h_r^\circ > 0$ $\Delta h_{r,12}^\circ > \Delta h_{r,11}^\circ$ ^(T)	Oh <i>et al.</i> , 2010
13 HVA digestion (aq.) (acetogenesis & methanogenesis)	16.9 kJ/mol _{SHVA rem.}	$\Delta h_r^\circ > 0$ ^(T) $\Delta h_{r,13}^\circ > \Delta h_{r,12}^\circ$ ^(T)	Oh <i>et al.</i> , 2010

Theoretical ^(T) and Experimental Data ^(E)

Table 5 – Specific heat yields (or energy content) estimated for the different substrates used in the model

	S_{SU}	S_{AA}	S_{FA}	S_{HVA}	S_{HBU}	S_{HPRO}	S_{HAC}
Oxidation heats (kJ/gCOD_{rem.})	-15.23	-14.38	-14.64	-14.11	-14.13	-14.16	-14.28
Oxidation + CO₂ stripping heats (kJ/gCOD_{rem.})	-14.62	-13.80	-13.62	-13.64	-13.65	-13.64	-13.67

Table 6 – Evolution of the organic loading rate (OLR) applied to the ATAD reactor

Time (d)	0.2	1.2	2.2	3.2	4.0	5.0	6.0	7.0	8.0	9.0	10.0	11.0	12.0	13.0
OLR (Kg COD/m ³ d)	2.95	3.00	3.03	3.35	3.81	3.51	3.53	4.05	0.00	4.90	5.56	5.24	5.57	4.42

Table 7 – Characterization of the influent

	Fraction of lipids	Fraction of proteins	Fraction of carbohydrates	Fraction of inert
Baseline Characterisation	19%	25%	19%	37%
Characterisation 1	15%	20%	28%	37%
Characterisation 2	22%	31%	10%	37%

Figure 1 - Schematic representation of the matrix restructuration

Figure 2 - Schematic representation of the mass balance in an aerated biological closed reactor

Figure 3 - Schematic representation of the enthalpy balance in a C-CSTR

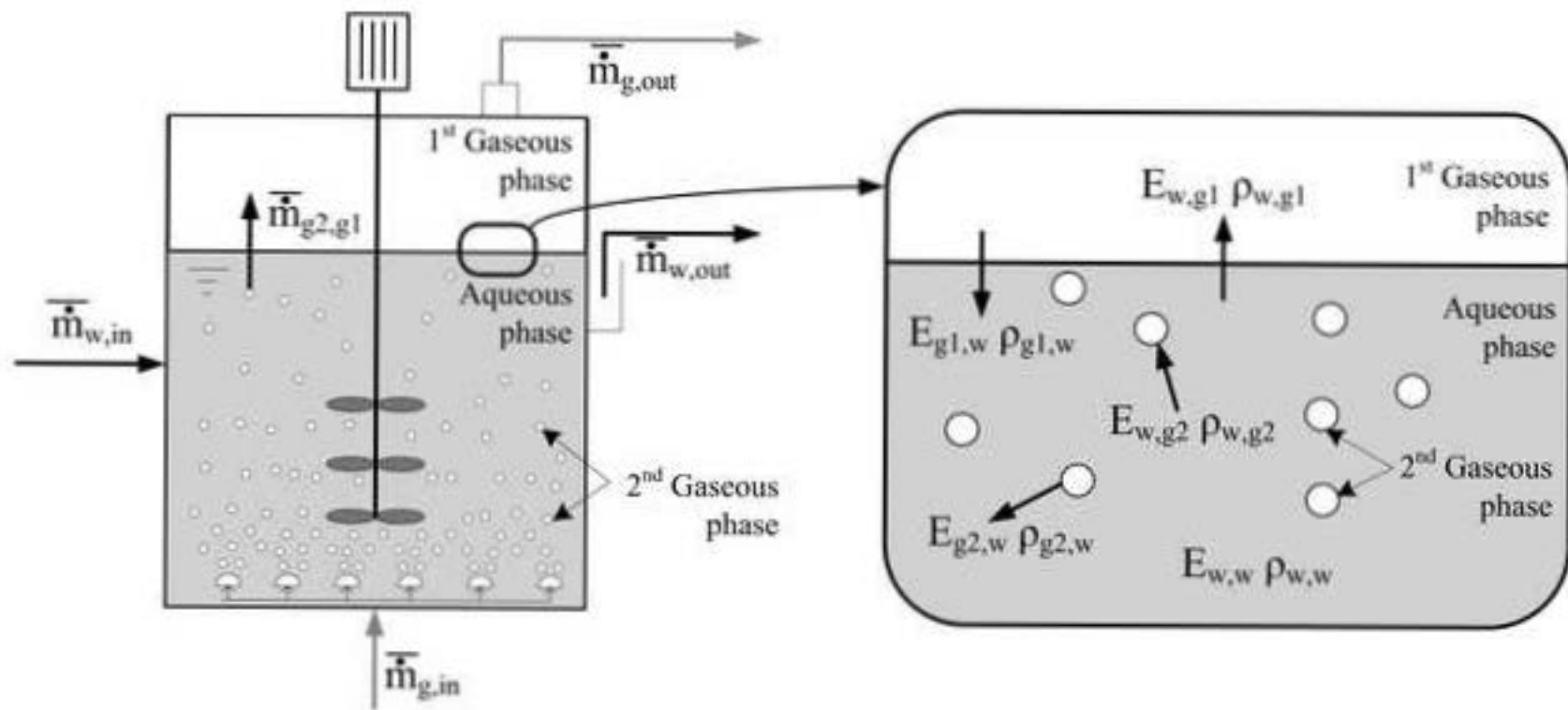
Figure 4 – (a) Total enthalpy produced and consumed in the system with the contribution of each thermal flux, (b) Transformation heat fluxes

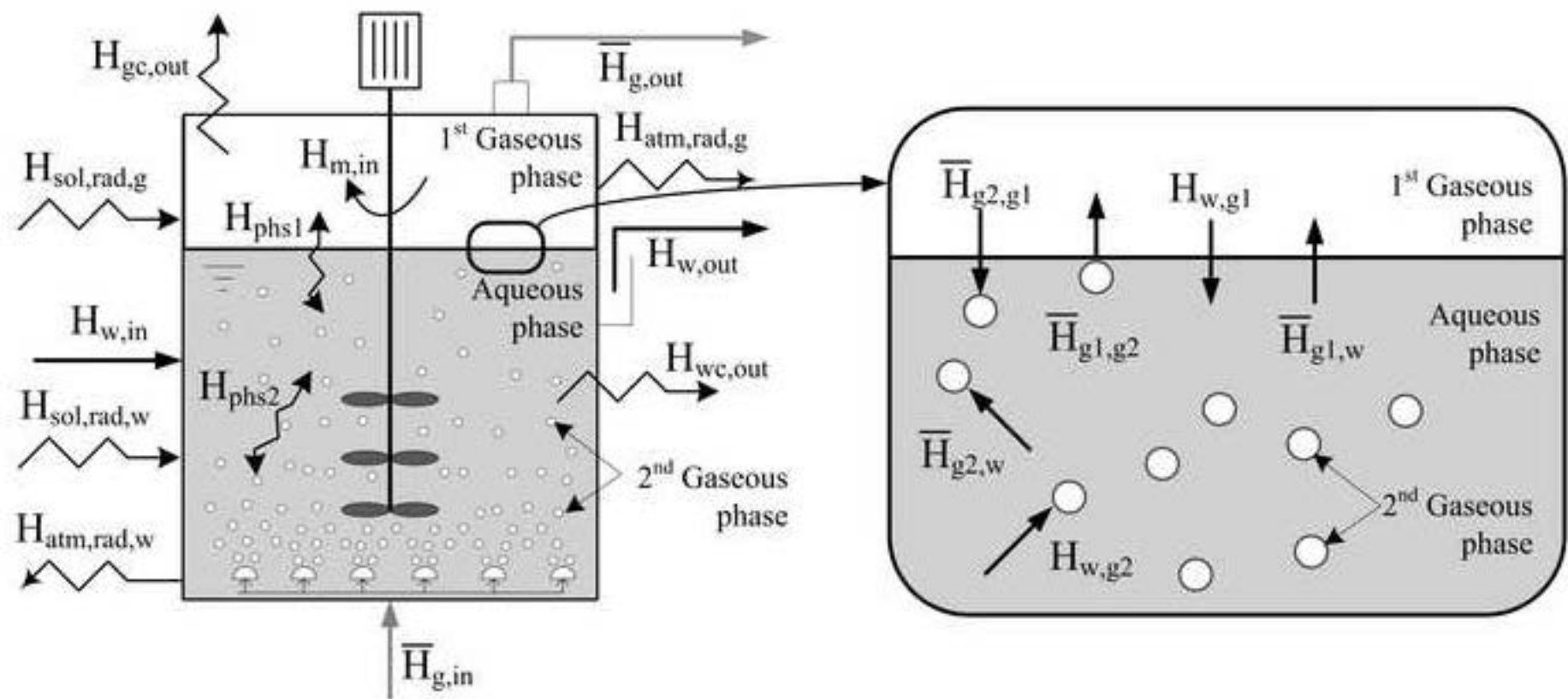
Figure 5 – (a) Experimental and simulation results for the full scale ATAD, (b) Total enthalpy produced and consumed in the system with the contribution of each thermal flux

Figure 6 – Behaviour of the temperature in a full scale ATAD for constant OLR and variable aeration flows

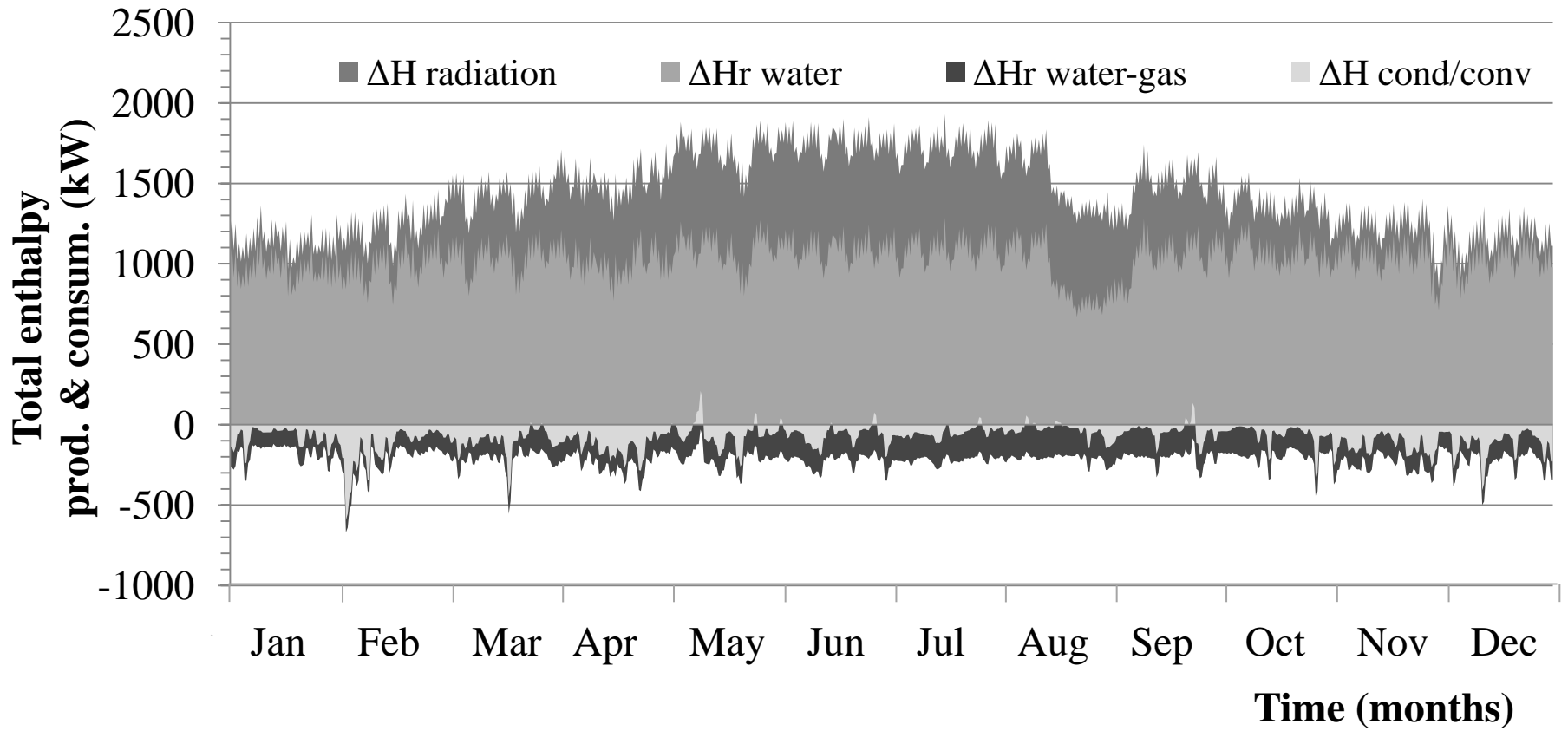
Figure 7 – (a) Temperature evolution for different influent characterizations (b) Aqueous transformation heats (specific heat yield) for different influent characterizations

	AQUEOUS $\bar{M}_w (NC_w \times I)$	GASEOUS $\bar{M}_g (NC_g \times I)$	SOLID $\bar{M}_s (NC_s \times I)$	
$\bar{p}_{w,w}$	$\tilde{E}_{w,w} (NT_w \times NC_w)$			Transformations in the aqueous phase
$\bar{p}_{g,g}$		$\tilde{E}_{g,g} (NT_g \times NC_g)$		Transformations in the gas phase
$\bar{p}_{w,g} = \bar{p}_{g,w}$	$\tilde{E}_{w,g} (NT_{wg} \times NC_w)$	$\tilde{E}_{g,w} (NT_{wg} \times NC_g)$		Liquid-Gas / Gas-Liquid transfers: Evaporation / Condensation & Stripping / Absorption
$\bar{p}_{w,s} = \bar{p}_{s,w}$	$\tilde{E}_{w,s} (NT_{ws} \times NC_w)$		$\tilde{E}_{s,w} (NT_{ws} \times NC_s)$	Liquid-Solid / Solid-Liquid transfers: Precipitation / Dissolution
$\bar{p}_{g,s} = \bar{p}_{s,g}$		$\tilde{E}_{g,s} (NT_{gs} \times NC_g)$	$\tilde{E}_{s,g} (NT_{gs} \times NC_s)$	Gas-Solid / Solid-Gas transfers: Deposition / Sublimation
	

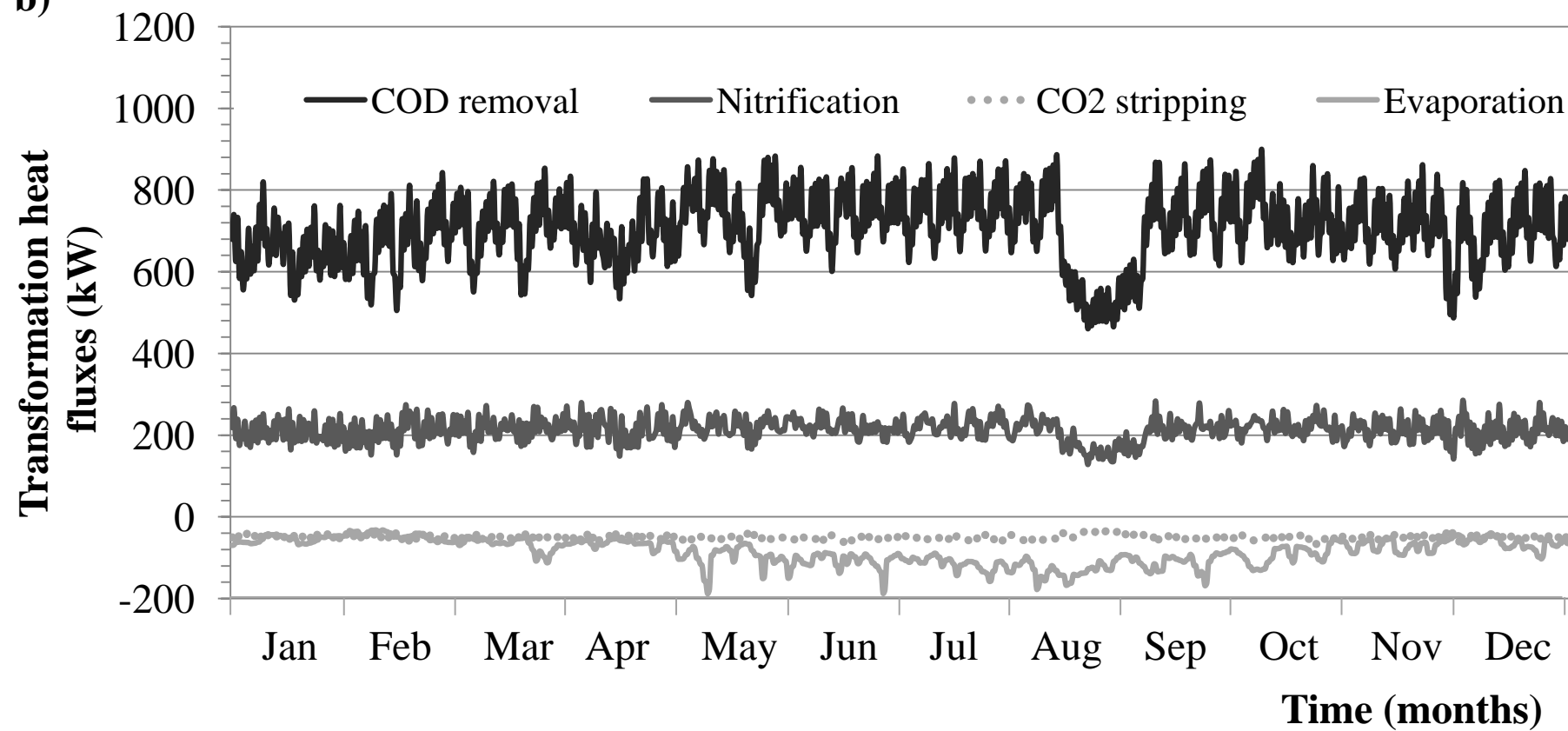




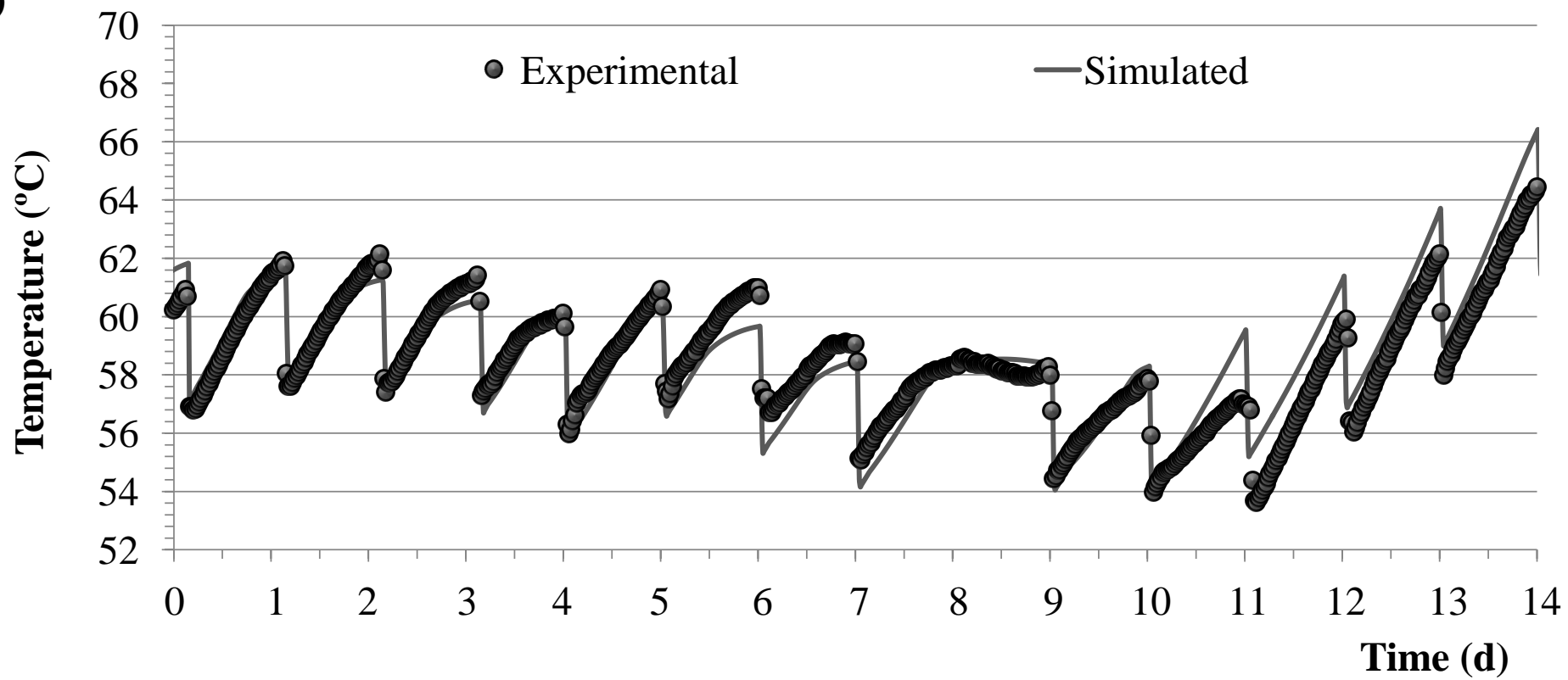
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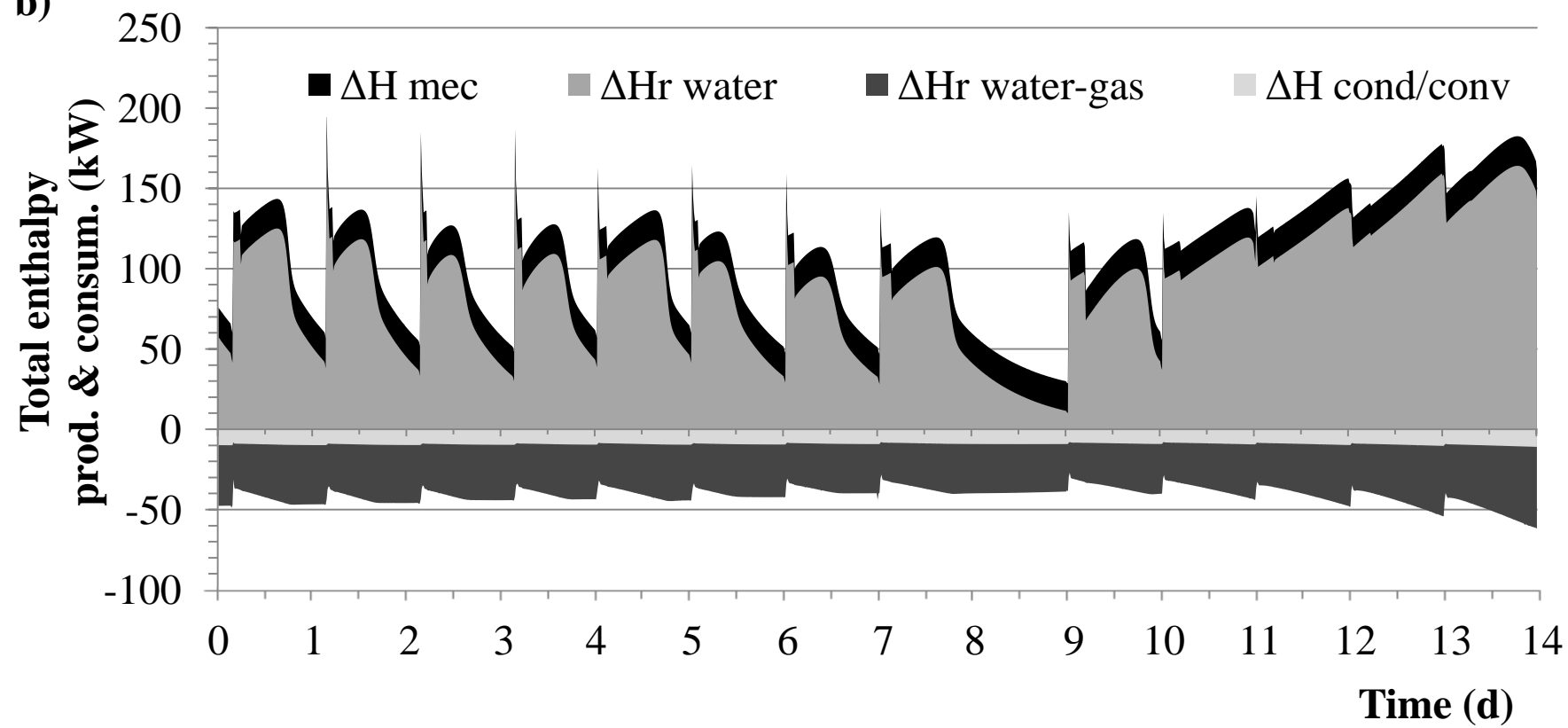
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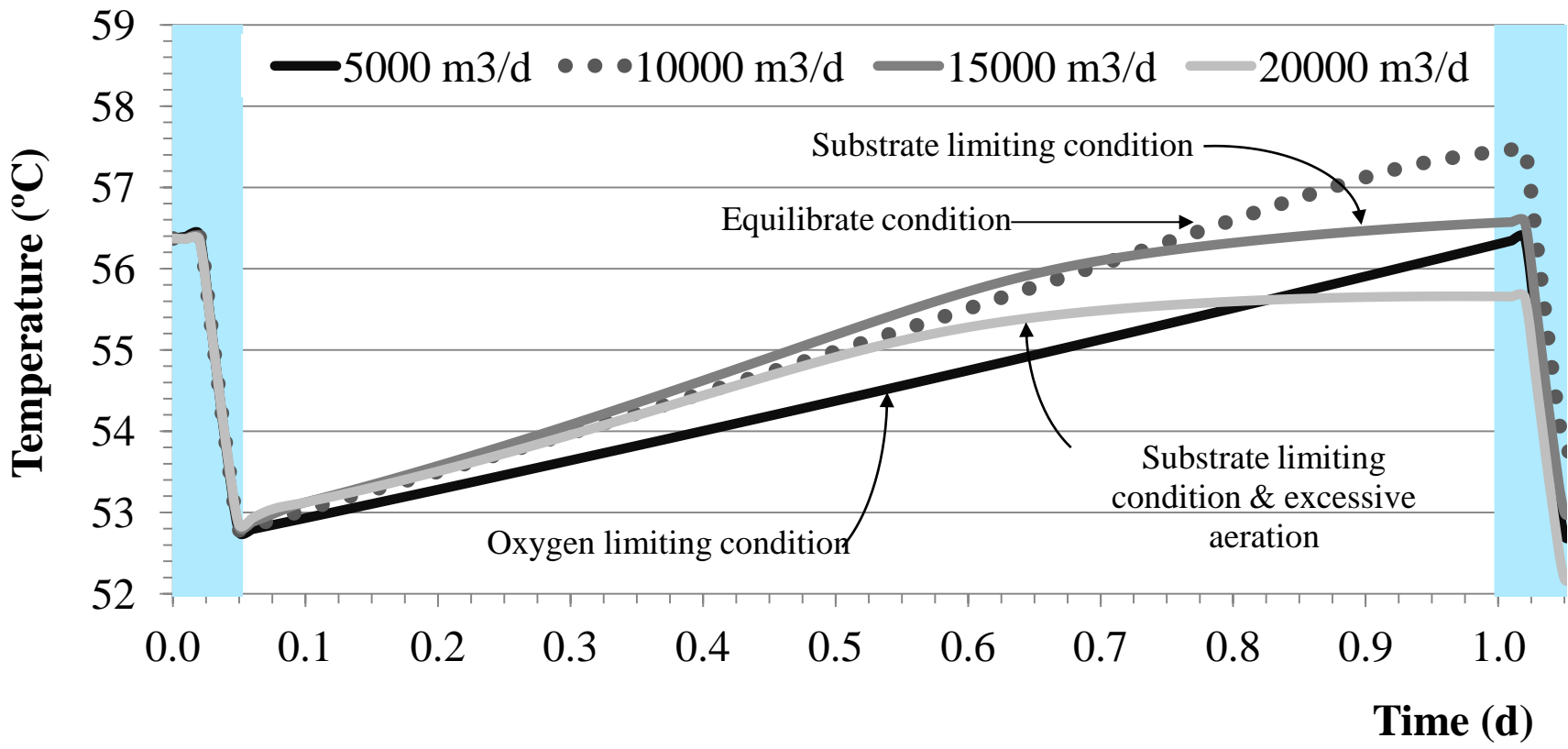


a)

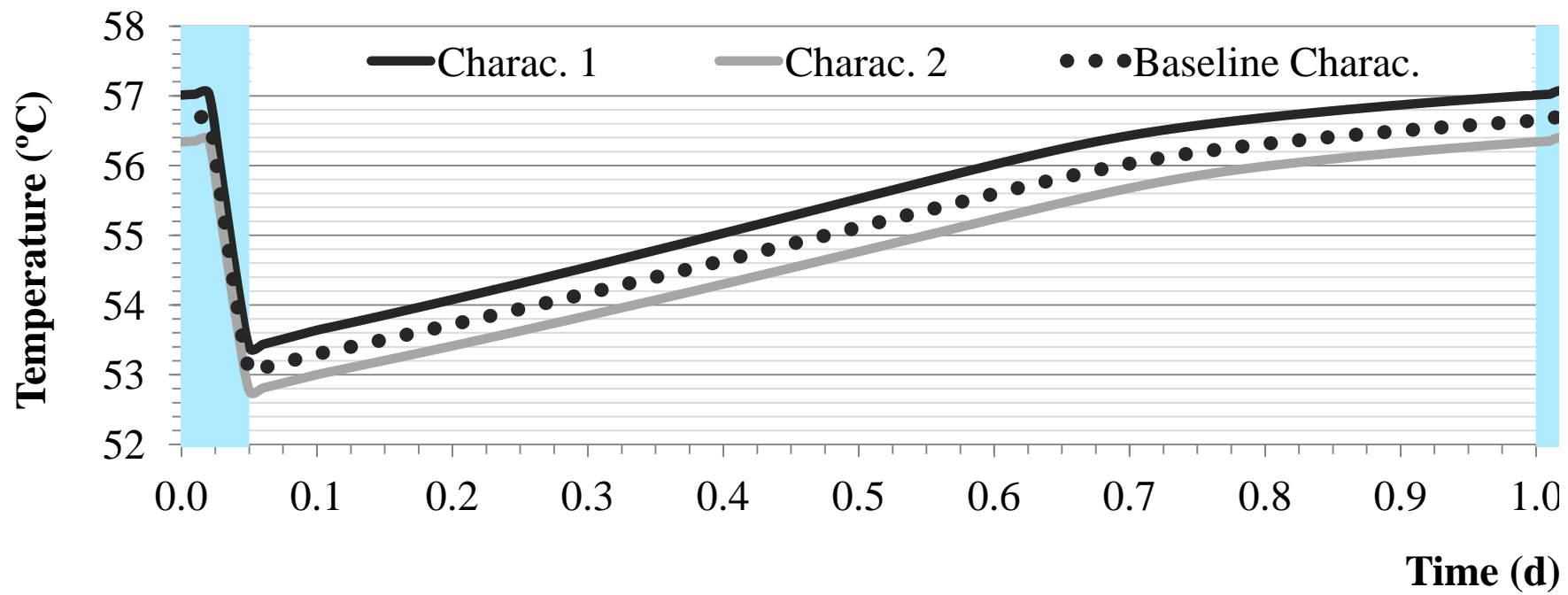


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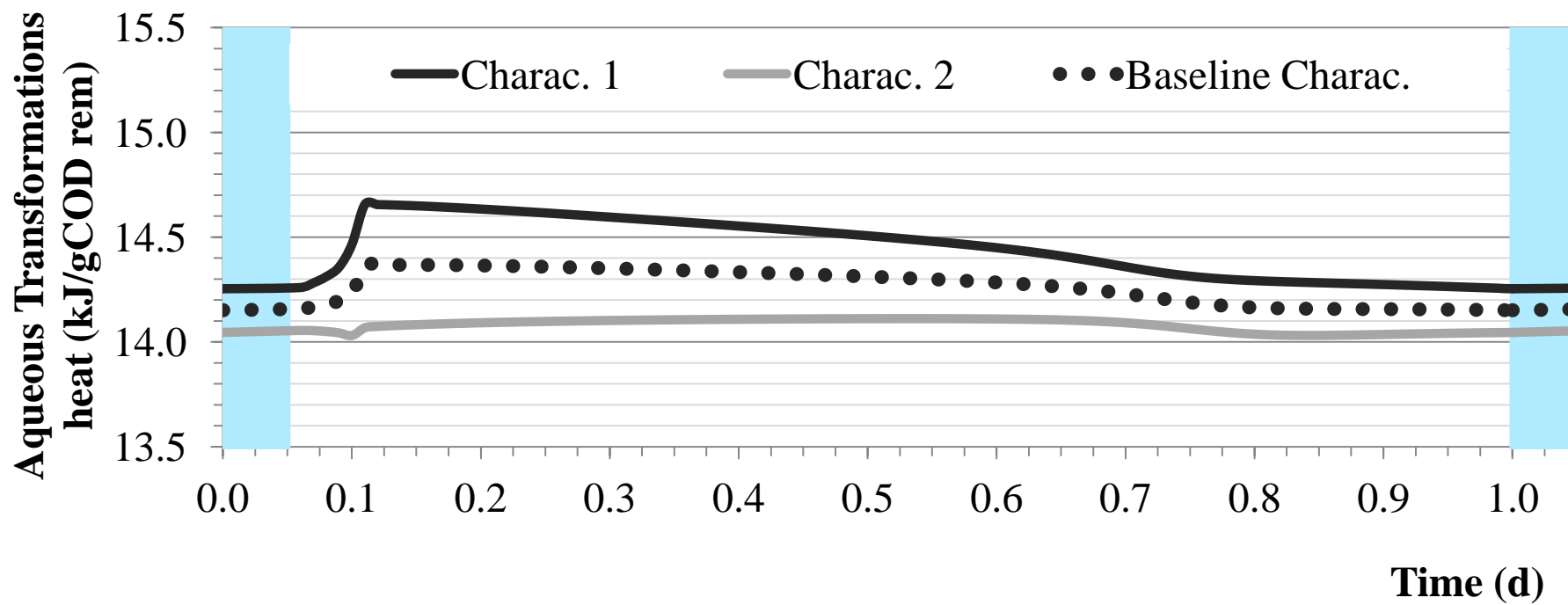




a)



b)



- New methodology for dynamic ΔH_r prediction in biological reactors using Hess's law
- Methodology implemented in a plant-wide modelling methodology
- Simultaneous calculation of the conventional mass balances and the enthalpy balance
- Multi-phase matrix structure to facilitate the prediction of mass and heat fluxes
- Methodology easily integrated into the numerical solution of existing models