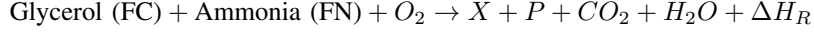


## SUPPLEMENTARY MATERIAL

### A. Detailed Mass Balances

#### General Reaction



#### General Mass Balance Formulation

$$\text{Accumulation} = \text{In} - \text{Out} + \text{Generation} - \text{Consumption}$$

**Batch Process:**  $F_{\text{in}} = F_{\text{out}} = 0$  and  $V = \text{constant}$ , thus:

$$\text{Accumulation} = \text{Generation} - \text{Consumption}$$

#### Biomass Balance

$$\begin{aligned} \frac{d}{dt}(X \cdot V) &= \mu \cdot X \cdot V \\ \frac{dX}{dt} \cdot V &= \mu \cdot X \cdot V \\ \frac{dX}{dt} &= \mu \cdot X \end{aligned}$$

Including the death rate constant  $k_d$ , the specific net growth rate becomes:

$$\frac{dX}{dt} = (\mu - k_d) \cdot X$$

#### Substrate and Product Balances

For any substrate or product  $S$ , assuming only microbial consumption or generation and constant volume  $V$ :

$$\frac{d}{dt}(S \cdot V) = -r_S \cdot V$$

The yield coefficient is defined as:

$$Y_{S/X} = \frac{r_X}{-r_S} = \frac{\mu X}{-r_S}$$

Therefore, the general expression becomes:

$$\frac{dS}{dt} = \pm \frac{\mu X}{Y_{S/X}}$$

where the sign depends on whether  $S$  is a substrate (negative) or a product (positive).

### B. Ionic Balance

$$[\text{OH}^-] = \frac{K_{a6}}{[H^+]} \quad (8)$$

$$[\text{KHPO}_4^-] = \frac{\text{KH}_2\text{PO}_4^{\text{tot}}}{\left(1 + \frac{[H^+]}{K_{a1}}\right)} \quad (9)$$

$$[\text{C}_6\text{H}_5\text{O}_7^{3-}] = \frac{\text{C}_6\text{H}_8\text{O}_7^{\text{tot}}}{\frac{[H^+]^3}{K_{a2}K_{a3}K_{a4}} + \frac{[H^+]^2}{K_{a3}K_{a4}} + \frac{[H^+]}{K_{a4}} + 1} \quad (10)$$

$$[\text{C}_6\text{H}_6\text{O}_7^{2-}] = \frac{[H^+] \cdot [\text{C}_6\text{H}_5\text{O}_7^{3-}]}{K_{a4}} \quad (11)$$

$$[\text{C}_6\text{H}_7\text{O}_7^-] = \frac{[H^+] \cdot [\text{C}_6\text{H}_6\text{O}_7^{2-}]}{K_{a3}} \quad (12)$$

$$[\text{HCO}_3^-] = \frac{\text{CO}_2^{\text{tot}}}{\left(\frac{[H^+]}{K_{a7}} + 1\right)} \quad (13)$$

$$[H^+] = [\text{OH}^-] + [\text{KHPO}_4^-] + 3[\text{C}_6\text{H}_5\text{O}_7^{3-}] + 2[\text{C}_6\text{H}_6\text{O}_7^{2-}] + [\text{C}_6\text{H}_7\text{O}_7^-] + [\text{HCO}_3^-] - [I_{\text{ext}}]$$

## C. Complementary Tables

TABLE II  
MODEL PARAMETERS USED IN THE DYNAMIC SIMULATION OF *M. smegmatis* GROWTH

Parameter	Description	Value
$t_{\text{lag}}$	Lag phase duration [h]	7
$k_C$	Monod constant for glycerol [g/L]	0.3828
$k_N$	Monod constant for ammonia [g/L]	$1.0027 \times 10^{-4}$
$Y_{X/C}$	Biomass yield on glycerol [g/g]	0.484
$Y_{X/N}$	Biomass yield on ammonia [g/g]	21.575
$Y_{X/CD}$	CO <sub>2</sub> yield [g/g]	0.3687
$Y_{X/O_2}$	O <sub>2</sub> yield [g/g]	1.352
$X_{\text{max}}$	Maximum biomass concentration [g/L]	1.4462
$\mu_{\text{max}}$	Maximum specific growth rate [1/h]	0.19
$k_d$	Death rate constant [1/h]	0.001
$\text{pH}_{\text{LL}}$	Lower limit of pH tolerance	3.4687
$\text{pH}_{\text{UL}}$	Upper limit of pH tolerance	7.4
$\text{pH}_{\text{alk}}$	Alkalinity value	7.2
$pK_{a1}$ to $pK_{a9}$	Acid dissociation constants [-]	See Table III
$[\text{KH}_2\text{PO}_4]$	Phosphate buffer (monobasic)	2.18
$[\text{C}_6\text{H}_8\text{O}_7]$	Citric acid (tricarboxylic acid)	2.00
$I_{\text{val}}$	pH inhibition shape parameter [-]	4.0413
$O_{2,\text{sat}}$	O <sub>2</sub> saturation concentration [g/L]	0.007267
$k_{L,a}$	Oxygen transfer coefficient [1/h]	86.26
$k_O$	Monod constant for oxygen [g/L]	0.0045

TABLE III  
ACID DISSOCIATION CONSTANTS ( $pK_a$ ) USED IN THE PH MODEL

Parameter	Chemical species or group	Value
$pK_{a1}$	KH <sub>2</sub> PO <sub>4</sub> (phosphate buffer)	6.86
$pK_{a2}$	Citric acid (C <sub>6</sub> H <sub>8</sub> O <sub>7</sub> )	3.13
$pK_{a3}$	Mono-deprotonated citric acid (C <sub>6</sub> H <sub>7</sub> O <sub>7</sub> ) <sup>-</sup>	4.76
$pK_{a4}$	Di-deprotonated citric acid (C <sub>6</sub> H <sub>7</sub> O <sub>7</sub> ) <sup>2-</sup>	6.40
$pK_{a5}$	Ammonium (NH <sub>3</sub> )	9.25
$pK_{a6}$	Glycerol (C <sub>3</sub> H <sub>8</sub> O <sub>3</sub> )	14.15
$pK_{a7}$	Carbon dioxide (CO <sub>2</sub> )	6.35
$pK_{a8}$	Bicarbonate (HCO <sub>3</sub> <sup>-</sup> )	10.33
$pK_{a9}$	Water (H <sub>2</sub> O)	14.00

TABLE IV  
INITIAL CONDITIONS USED IN THE SIMULATION OF *M. smegmatis* GROWTH

Variable	Description	Initial Value
$X_0$	Biomass concentration [g/L]	0.223
$C_0$	Glycerol concentration [g/L]	5.922
$N_0$	Ammonium concentration [g/L]	1.027
$CO_{2,0}$	Carbon dioxide concentration [g/L]	0.000439
$O_{2,0}$	Dissolved oxygen concentration [g/L]	0.65

TABLE V  
HIGHLY CORRELATED PARAMETER PAIRS ( $|\rho| > 0.95$ )

Parameter A	Parameter B	Correlation
$t_{\text{lag}}$	$k_C$	0.9712
$t_{\text{lag}}$	pH.UL	-0.9831
$Y_{X/CO_2}$	$k_d$	0.9877
$Y_{X/O_2}$	pH.LL	0.9988
$Y_{X/O_2}$	$I_{\text{val}}$	-0.9988
$Y_{X/O_2}$	$k_{\text{La}}$	-0.9998
pH.LL	$I_{\text{val}}$	-1.0000
pH.LL	$k_{\text{La}}$	-0.9995
$I_{\text{val}}$	$k_{\text{La}}$	0.9995
$O_{2,\text{sat}}$	$k_O$	0.9865

#### D. Complementary Figures

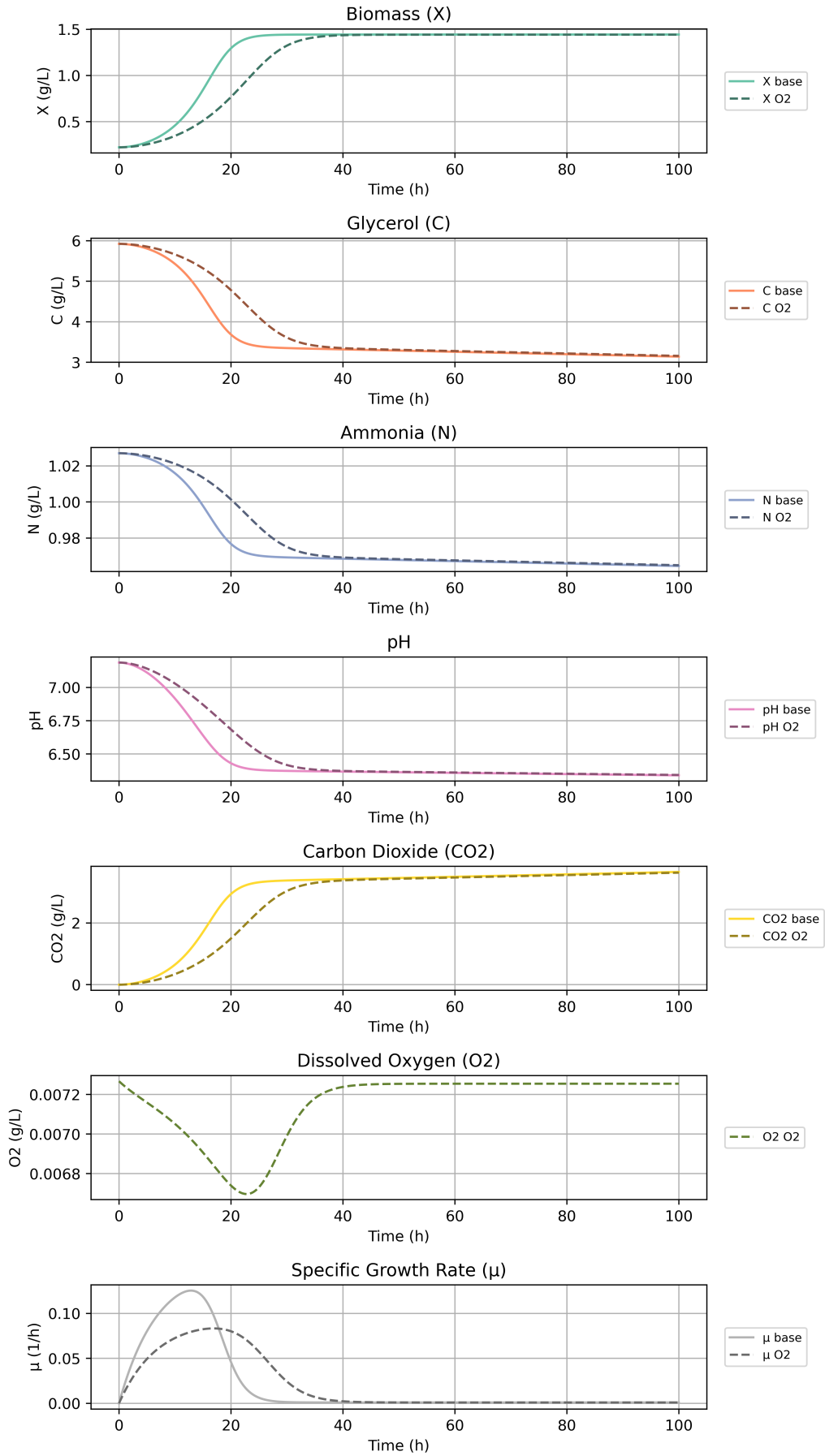


Fig. 1. Comparison between original model without oxygen, and the updated model with oxygen consumption on hypoxic conditions.

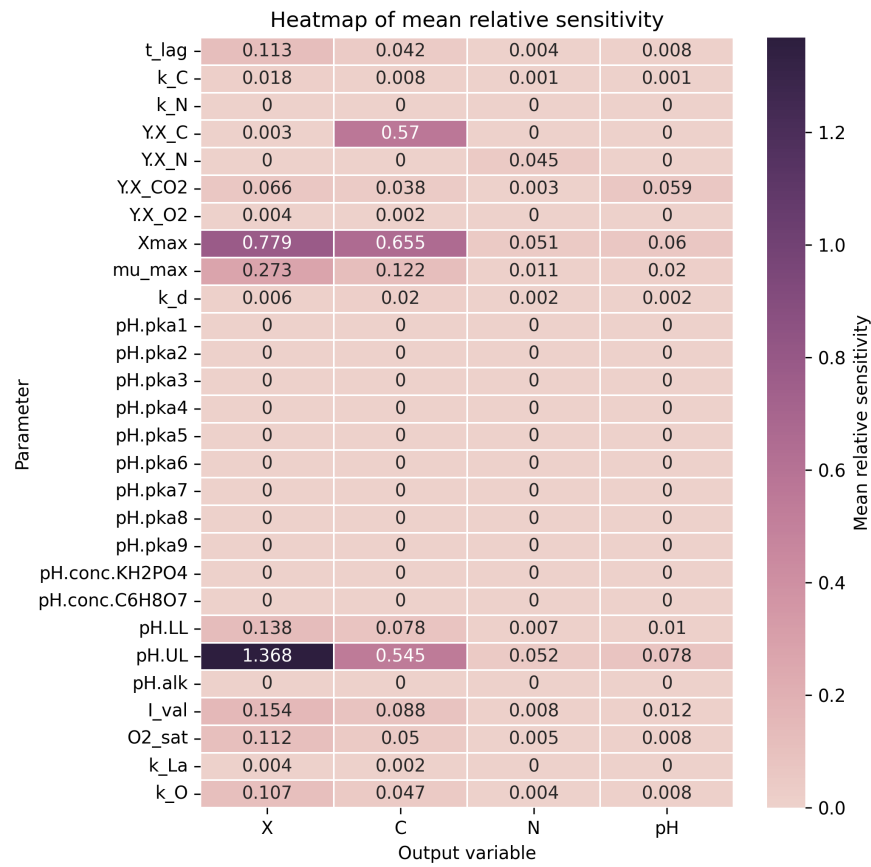


Fig. 2. Heatmap of the mean relative sensitivity of each parameter with respect to the model outputs.

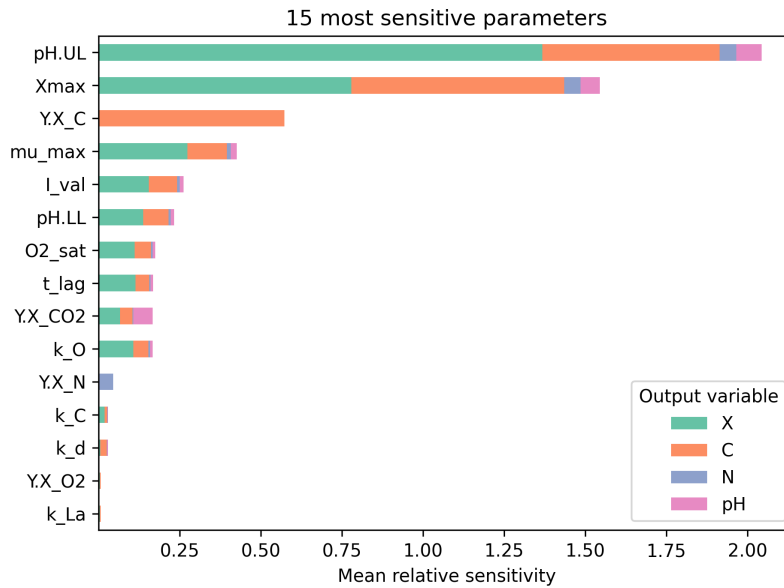


Fig. 3. Mean relative sensitivity of the 15 most influential parameters in the model, disaggregated by affected output variable. Each bar represents the cumulative sensitivity contribution of a parameter across all state variables.

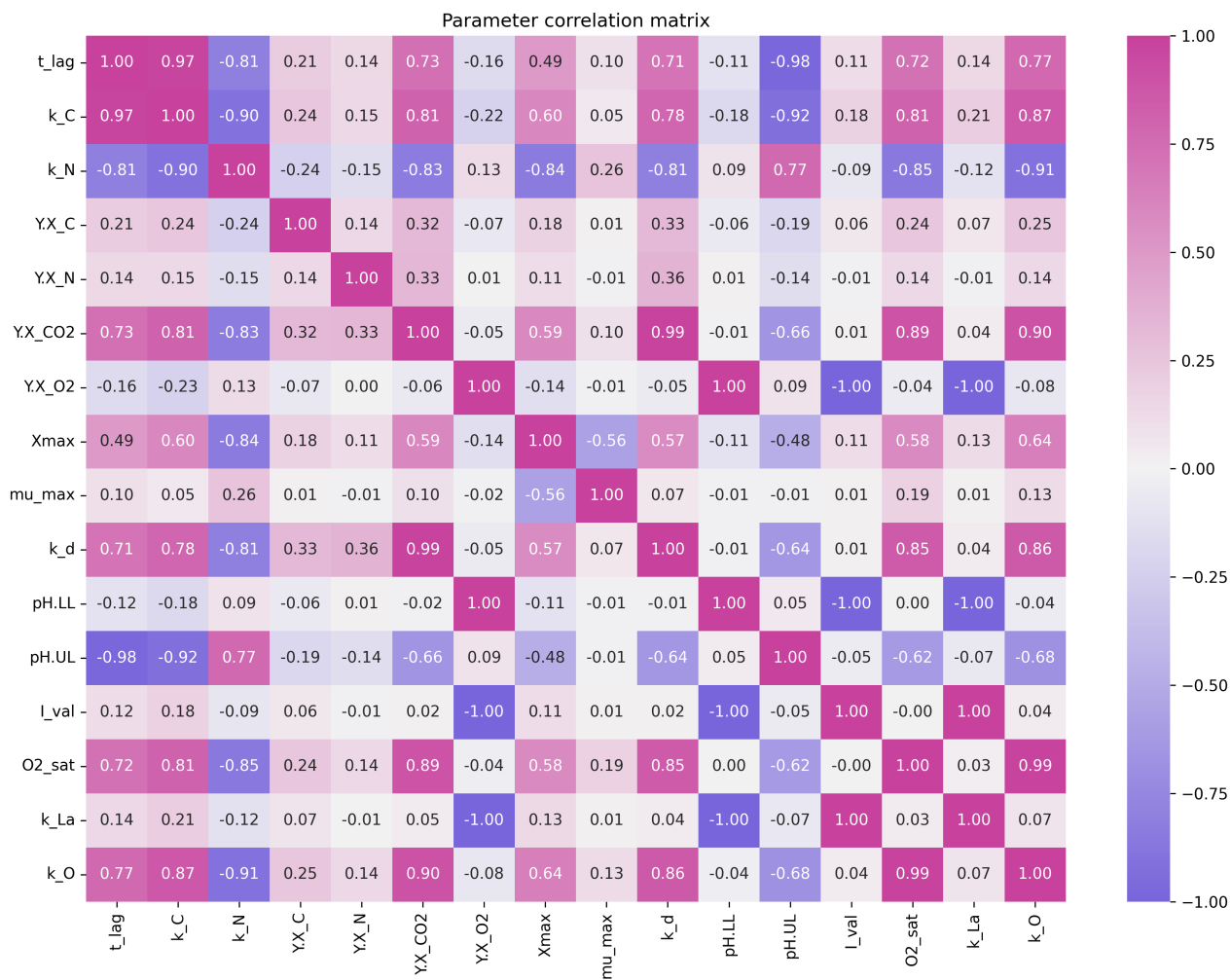


Fig. 4. Correlation matrix between the parameters retained after excluding non-identifiable pH-related terms.

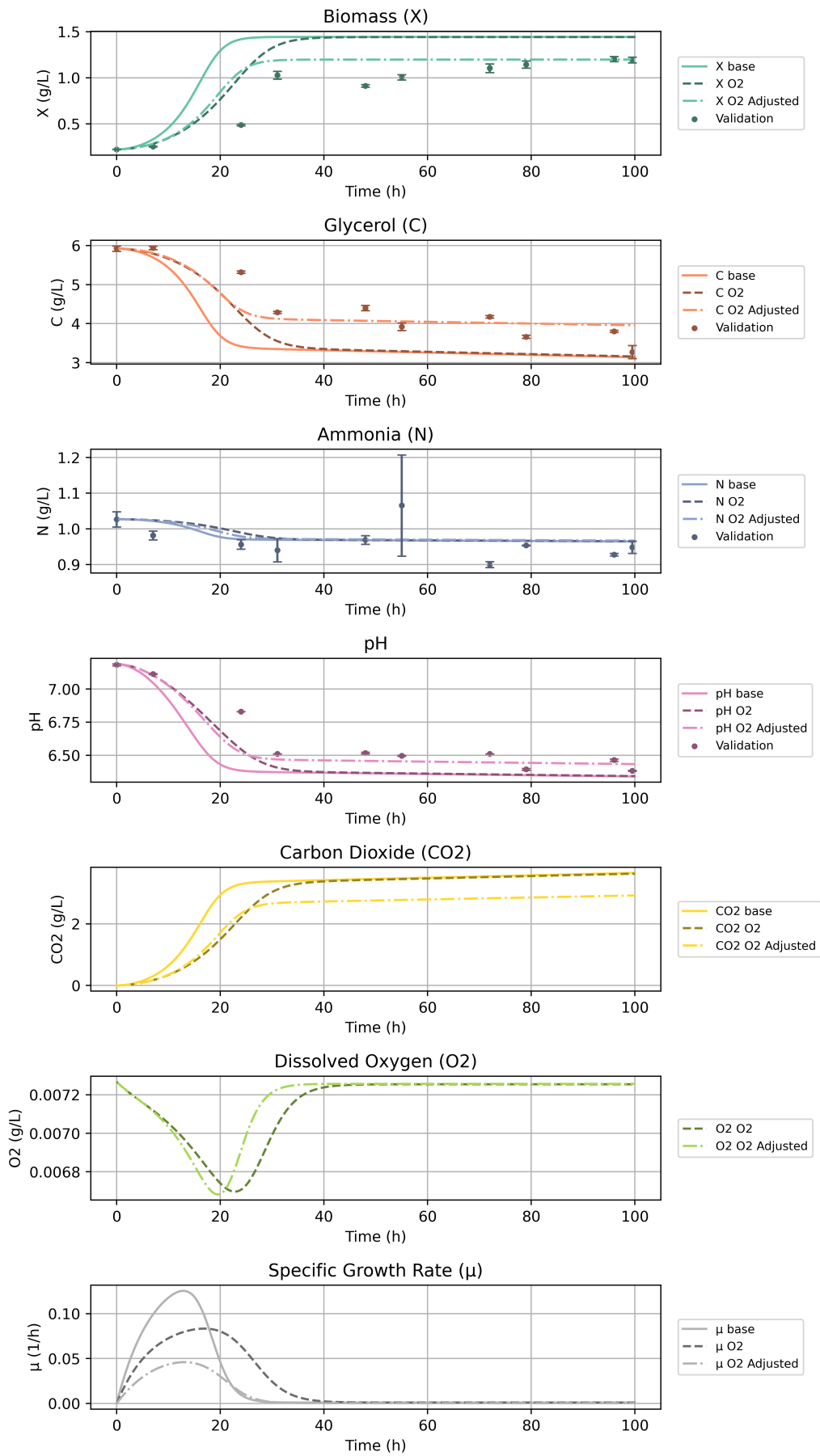


Fig. 5. Comparison between all the model tested, with the validation data. Error bars represent the relative deviation of experimental validation data.