

# Análise de modos elementares

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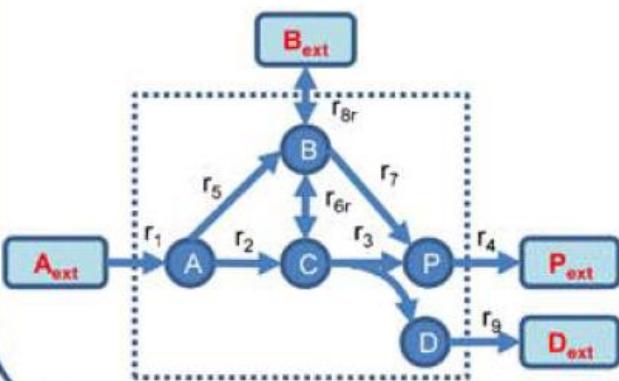


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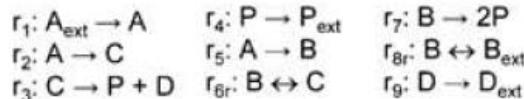
# analysis of cellular metabolism

## Problem statement

Network



## Stoichiometric reactions



## Equations to solve

A

$$\underline{S} \cdot \underline{r} = 0$$

Thermodynamic constraints:

$$r_{1,5,7,9} \geq 0$$

## Stoichiometric matrix

	$r_1$	$r_2$	$r_3$	$r_4$	$r_5$	$r_{6r}$	$r_7$	$r_{8r}$	$r_9$
A	1	-1	0	0	-1	0	0	0	0
B	0	0	0	0	1	-1	-1	-1	0
C	0	1	-1	0	0	1	0	0	0
D	0	0	1	0	0	0	0	0	-1
P	0	0	1	-1	0	0	2	0	0

$\underline{S} = [r_1 \ r_2 \ r_3 \ r_4 \ r_5 \ r_{6r} \ r_7 \ r_{8r} \ r_9]^T$

$$\frac{d}{dt} \underline{C} = \underline{S} \times \underline{r} - \mu \times \underline{C}, \quad \mu \cdot C \text{ (negligible)} \quad dC/dt = 0 \text{ (steady state)}$$

$$\underline{S} \cdot \underline{r} = 0 \text{ (Eq 2)}$$

$$r_i \geq 0 \text{ (Eq 3)}$$

Tools for analysis of cellular metabolism can be grouped into three categories, all of them developed from the same mathematical model:

- (1) Metabolic flux analysis,
- (2) Flux balance analysis and
- (3) Metabolic pathway analysis (Elementary mode analysis).

### Metabolic Flux Analysis

$$\underline{S}_{\text{u}} = \begin{bmatrix} r_3 & r_4 & r_5 & r_{6r} & r_7 \\ A & 0 & 0 & -1 & 0 & 0 \\ B & 0 & 0 & 1 & -1 & -1 \\ C & -1 & 0 & 0 & 1 & 0 \\ D & 1 & 0 & 0 & 0 & 0 \\ P & 1 & -1 & 0 & 0 & 2 \end{bmatrix}$$

$$\underline{S}_{\text{m}} = \begin{bmatrix} r_1 & r_2 & r_{8r} & r_9 \\ A & 1 & -1 & 0 & 0 \\ B & 0 & 0 & -1 & 0 \\ C & 0 & 1 & 0 & 0 \\ D & 0 & 0 & 0 & -1 \\ P & 0 & 0 & 0 & 0 \end{bmatrix}$$

### Measured fluxes

$$\underline{r}_m = \begin{bmatrix} r_1 \\ r_2 \\ r_{8r} \\ r_9 \end{bmatrix} = \begin{bmatrix} 1 \\ 0.3 \\ 0 \\ 0.75 \end{bmatrix}$$

### Equations to solve

$$\begin{aligned} \underline{S}_{\text{u}} \cdot \underline{r} &= 0 \\ \left[ \underline{S}_{\text{u}} \quad \underline{S}_{\text{m}} \right] \begin{bmatrix} \underline{r}_u \\ \underline{r}_m \end{bmatrix} &= 0 \end{aligned}$$

$$\underline{r}_u = -\underline{S}_{\text{u}}^{-1} \cdot \underline{S}_{\text{m}} \cdot \underline{r}_m$$

### Solution

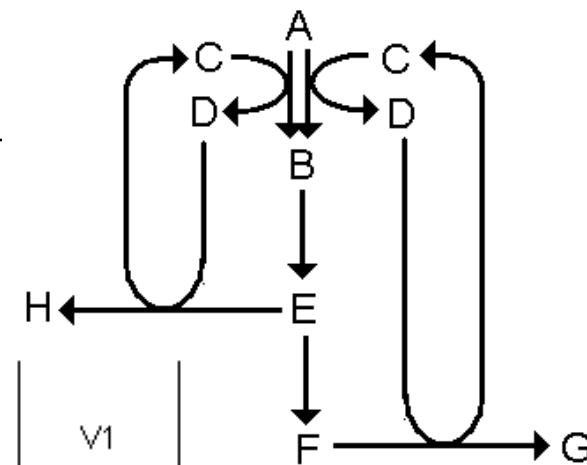
**B**

$$\begin{bmatrix} r_3 \\ r_4 \\ r_5 \\ r_{6r} \\ r_7 \end{bmatrix} = \begin{bmatrix} 0.75 \\ 1.25 \\ 0.7 \\ 0.45 \\ 0.25 \end{bmatrix}$$



A      B      E      F      C      D      G      H

$$\begin{bmatrix} -1 & 1 & 0 & 0 & -1 & 1 & 0 & 0 \\ 0 & -1 & 1 & 0 & 0 & 0 & 0 & 0 \\ 0 & 0 & -1 & 1 & 0 & 0 & 0 & 0 \\ 0 & 0 & 0 & -1 & 1 & -1 & 1 & 0 \\ 0 & 0 & -1 & 0 & 1 & -1 & 0 & 1 \end{bmatrix}$$



V1  
V2  
V3  
V4  
V5

$$F = J - K$$

C

## Flux Balance Analysis

Obj:  $\max r_4$

s.t.:  $\underline{S} \cdot \underline{r} = \underline{0}$

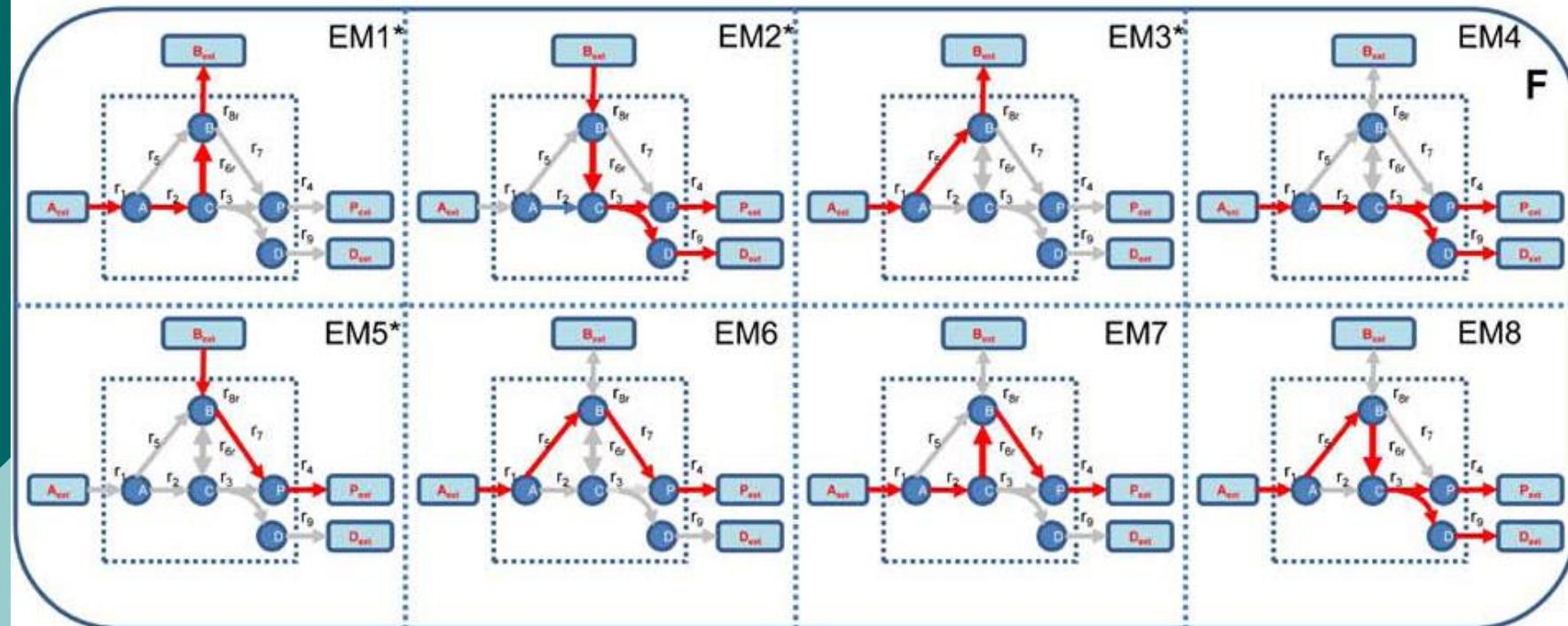
$$r_1 = 1$$

$$r_{8r} = 0$$

$$r_{2-5,7,9} \geq 0$$

$$\underline{r} = \begin{bmatrix} r_1 \\ r_2 \\ r_3 \\ r_4 \\ r_5 \\ r_{6r} \\ r_7 \\ r_{8r} \\ r_9 \end{bmatrix} = \begin{bmatrix} 1 \\ 0.35 \\ 0 \\ 2 \\ 0.65 \\ -0.35 \\ 1 \\ 0 \\ 0 \end{bmatrix}$$

# Metabolic pathway analysis (Elementary (flux) analysis)

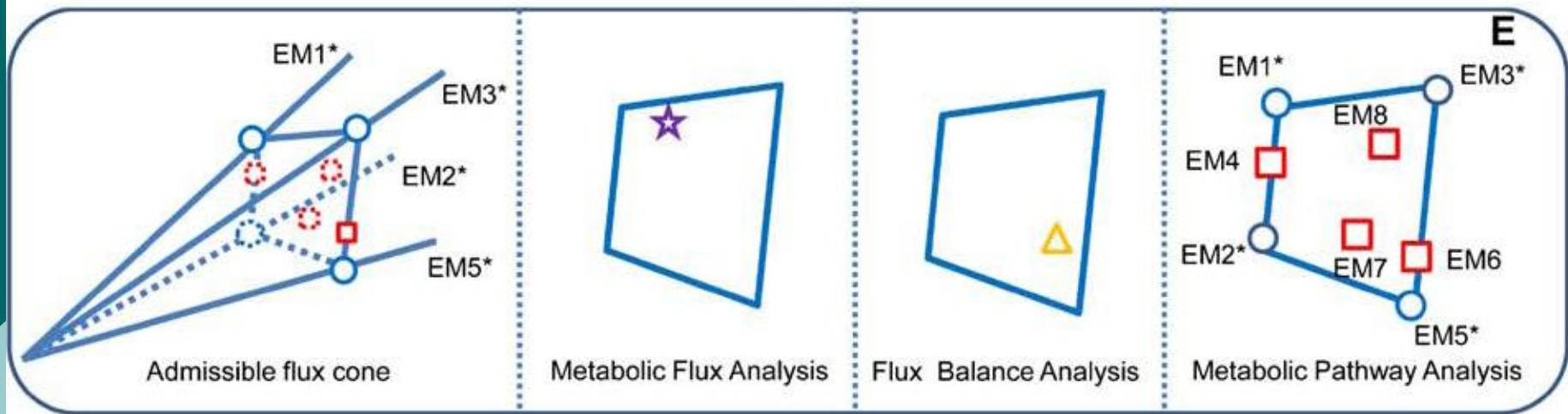


$$S.r = 0 \text{ (Eq 2)}$$

$$r_i \geq 0 \text{ (Eq 3)}$$

Elementary mode analysis calculates all solutions in the admissible flux space by solving Eq 2 in conjunction with the thermodynamic constraint (3) and additional non-decomposability and systematic independence constraints. Each solution (re)presents an elementary (flux) mode.

# Interpretação Geométrica



- ✓ O cone de fluxos admissíveis representa todas as possíveis vias que podem existir.
- ✓ Alguns modos elementares ficam na face ou na base do cone.
- ✓ AFM identifica somente uma via que se localiza em qualquer local do cone. ABF representa somente uma via em qualquer local do cone e satisfaz a função objetivo definida.
- ✓ AVM identifica todas as vias geneticamente independentes, com vias extremas em azul e modos elementares em vermelho.

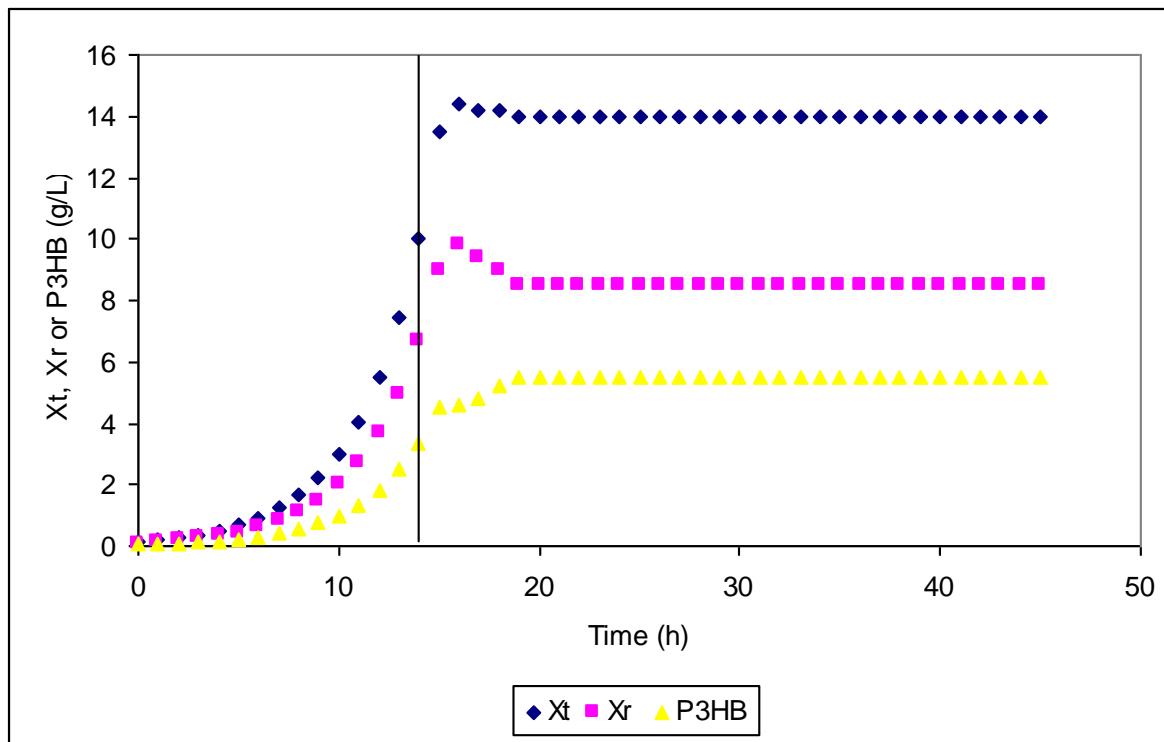
# Steady state



# Steady state

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Continuous culture



# Steady state

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Continuous culture

$$\frac{dX}{dt} = \mu = \text{constante}$$

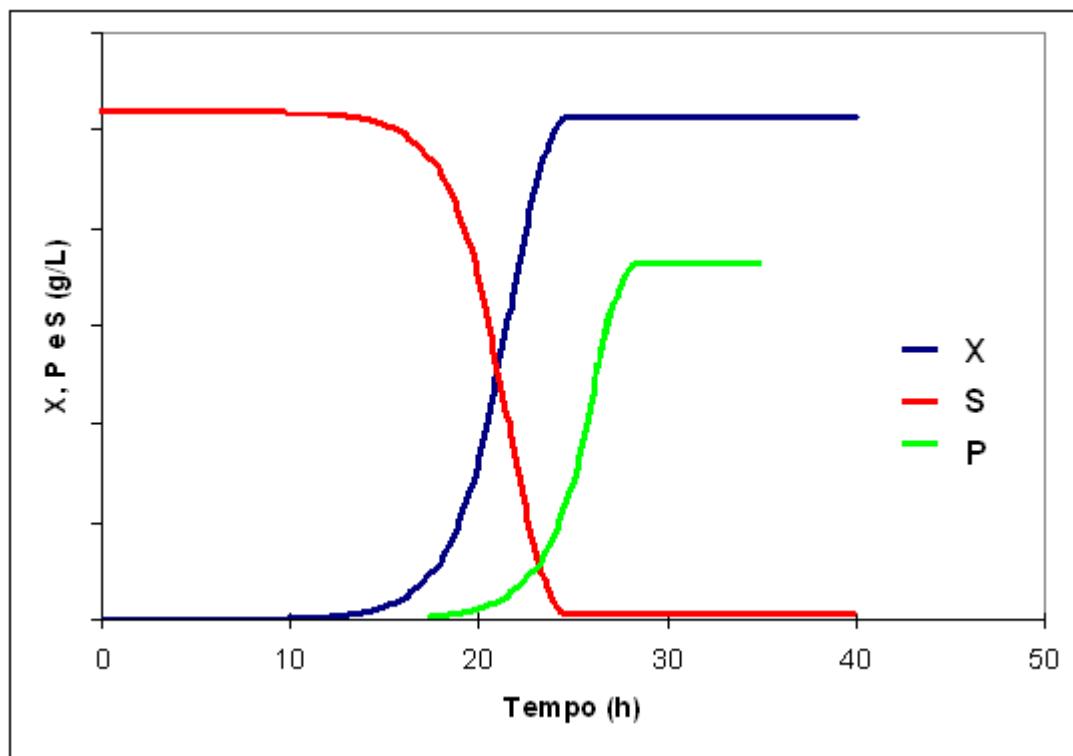
$$\frac{dP}{dt} = q_P = \text{constante}$$

$$\frac{dS}{dt} = q_S = \text{constante}$$

Y são constantes?

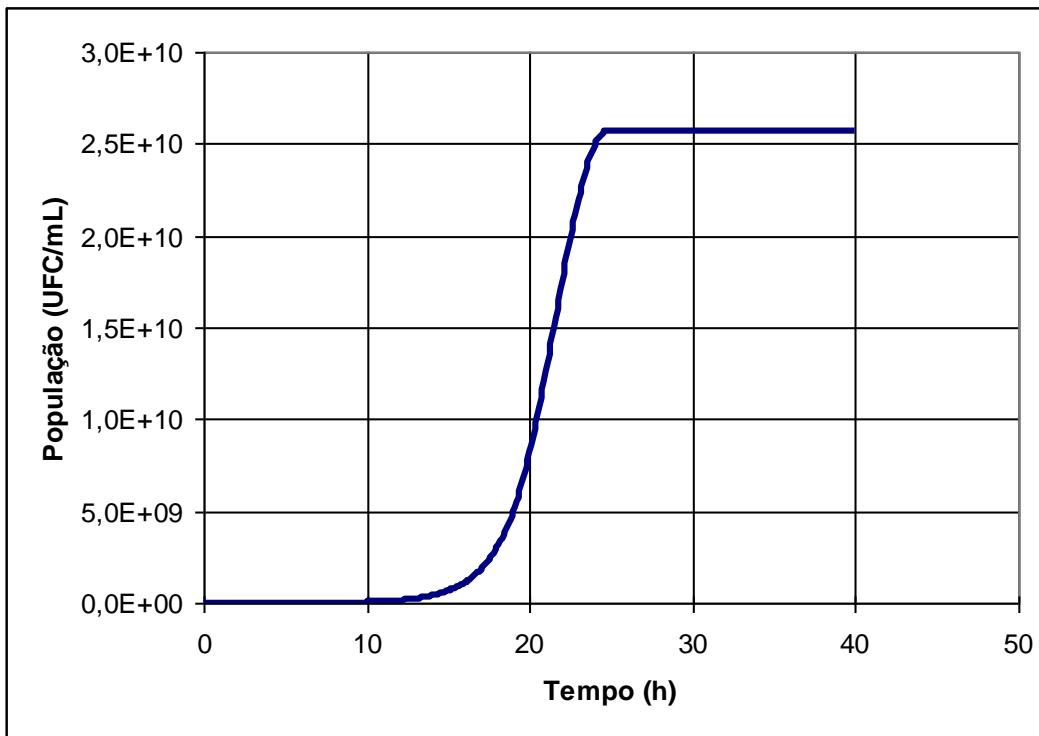
# Batch

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# Fases de crescimento

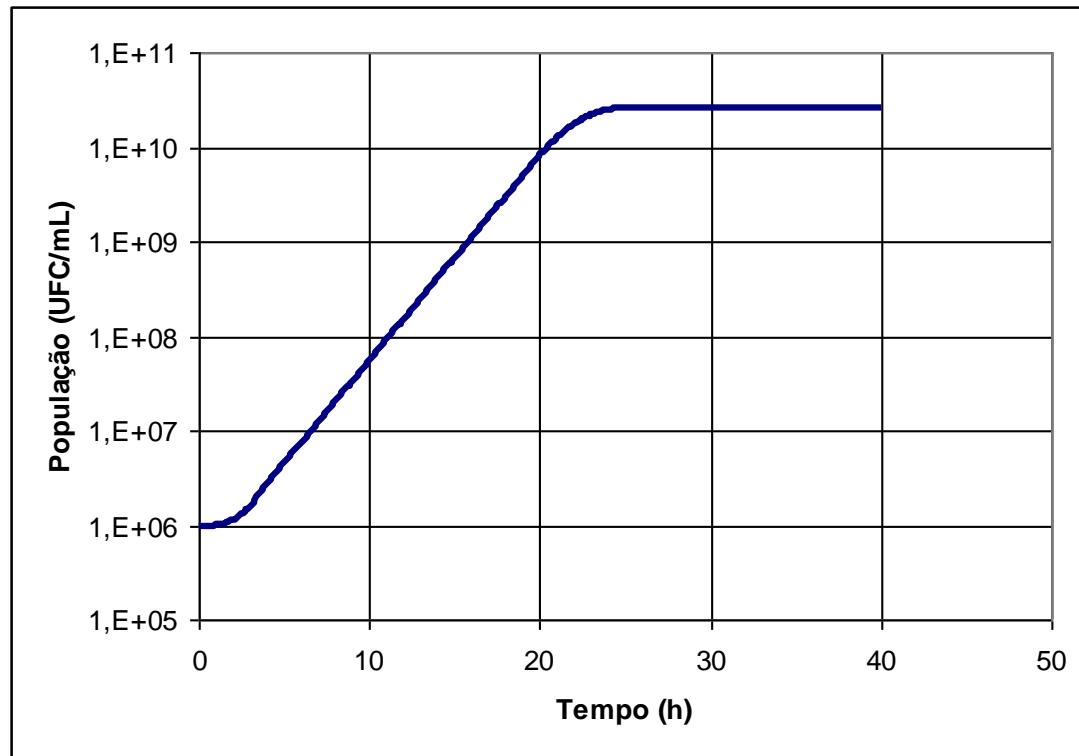
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Lag  
Exponencial  
Estacionária

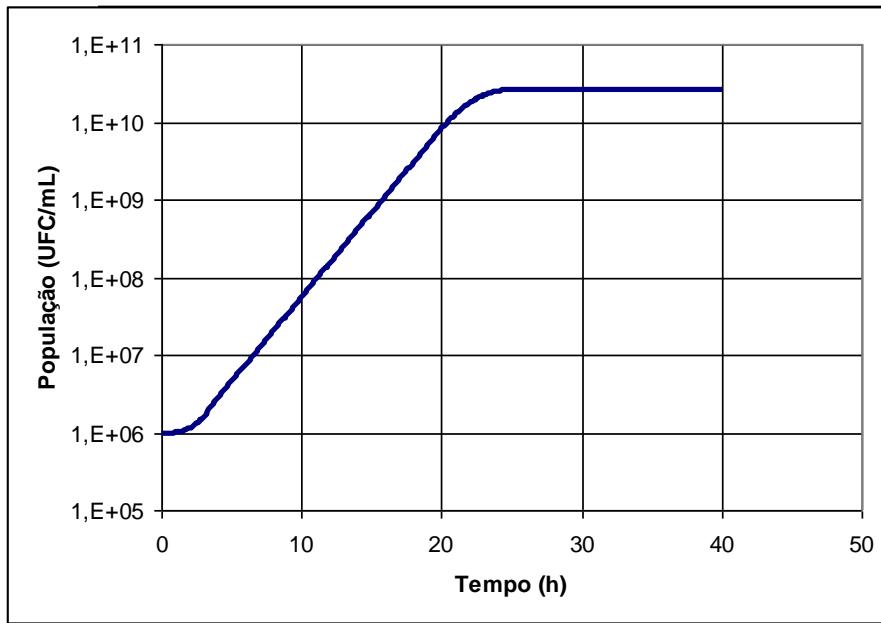
# Fases de crescimento

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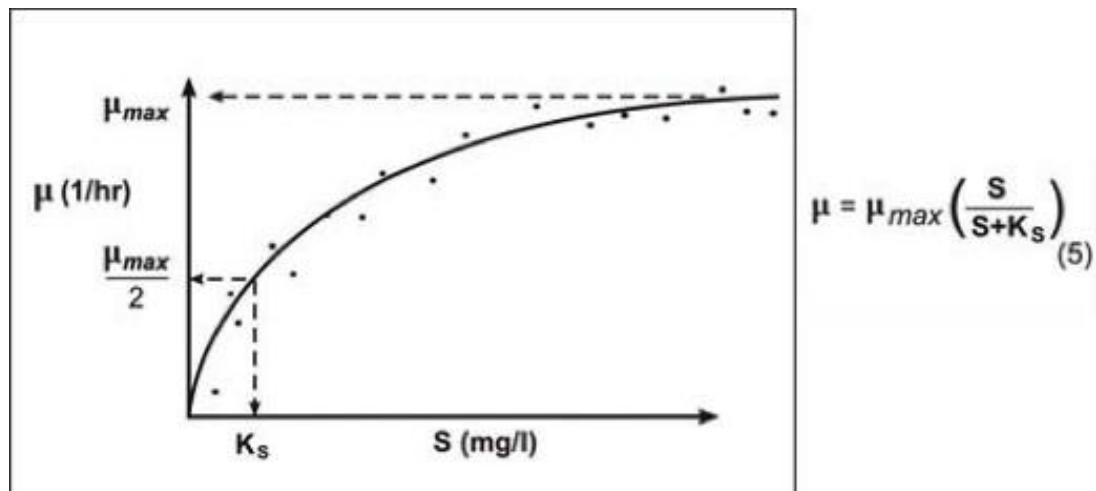


Lag  
Exponencial  
Estacionária

# Fases de crescimento



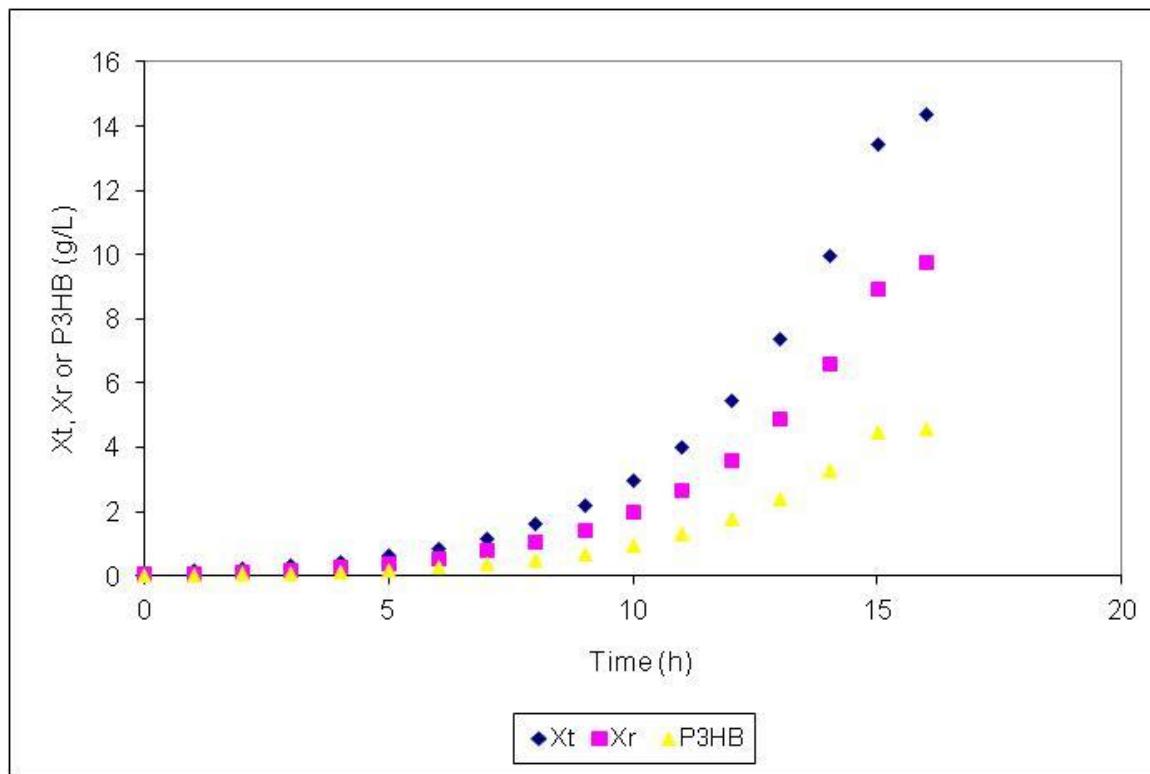
Lag  
Aceleração  
Exponencial  
Desaceleração  
Estacionária



# (Pseudo or Quase)Steady state

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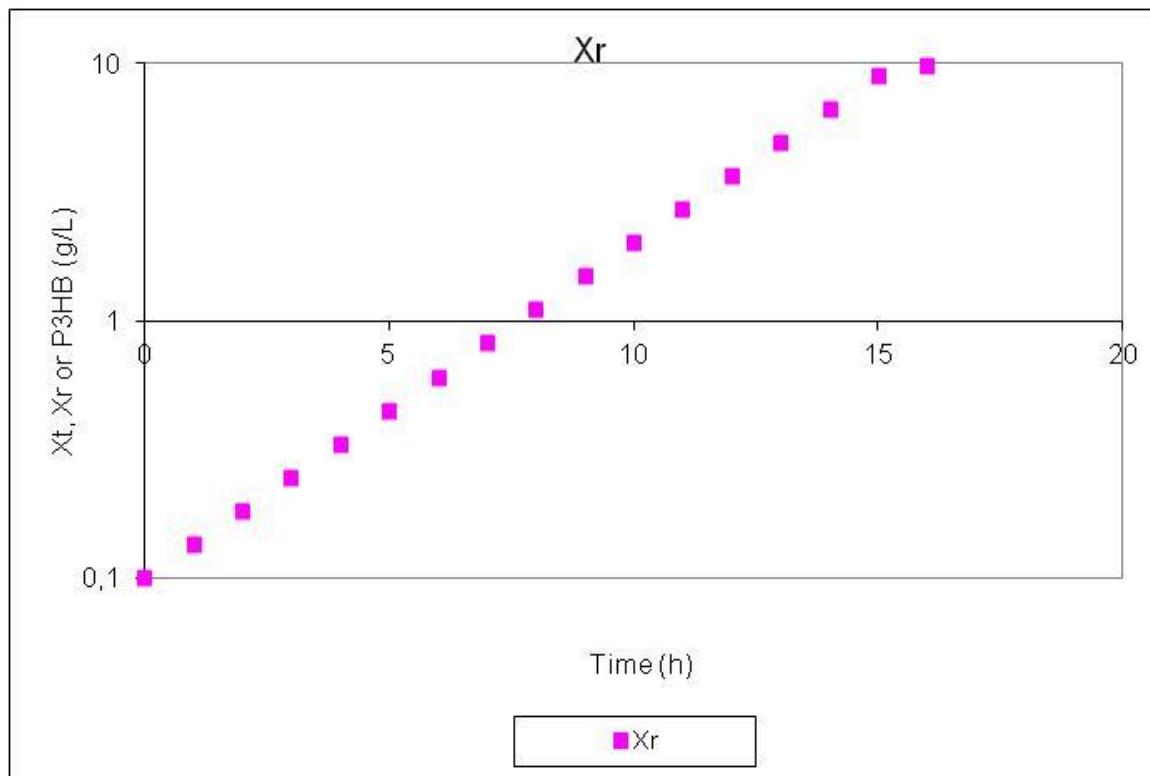
Batch – Exponential Growth phase



# (Pseudo or Quase)Steady state

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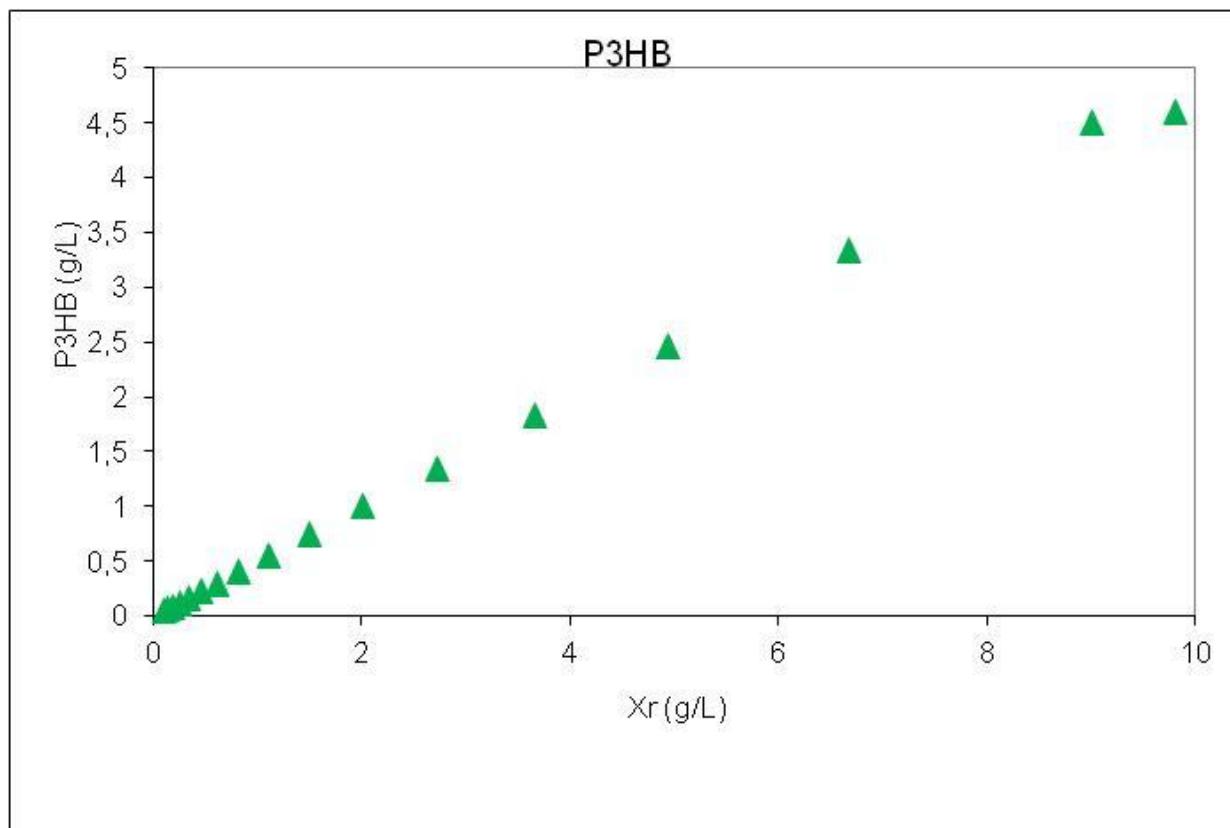
Batch – Exponential Growth phase



# (Pseudo or Quase)Steady state

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Batch – Exponential Growth phase



# (Pseudo or Quase)Steady state

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Batch – Exponential Growth phase

$$\frac{dX}{dt} = \mu = \text{constante}$$

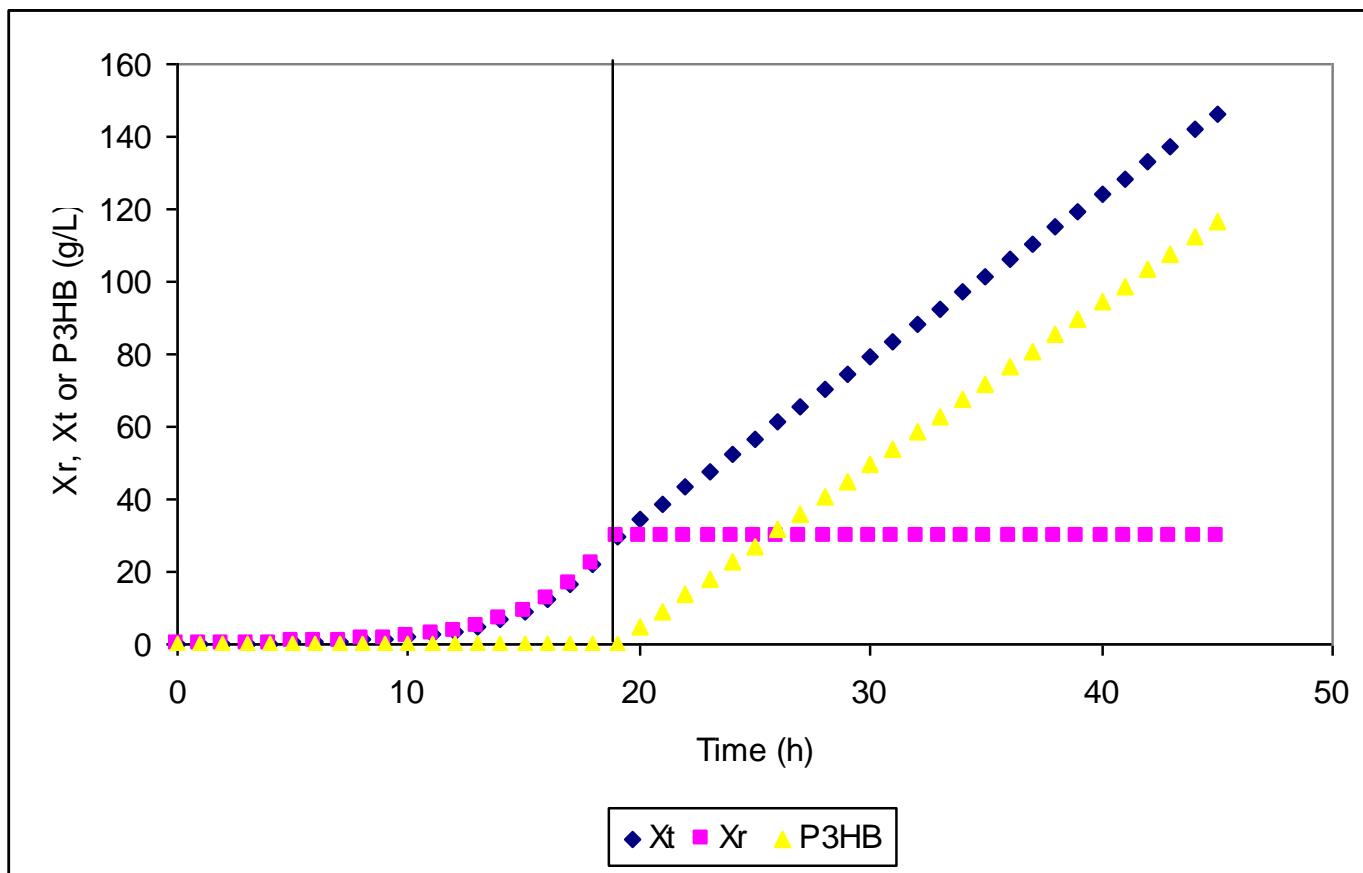
$$\frac{dP}{dt} = q_P = \text{constante}$$

$$\frac{dS}{dt} = q_S = \text{constante}$$

Y são constantes?

# (Pseudo or Quasi)Steady state

Batch or Fed-batch



# (Pseudo or Quasi)Steady state

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Batch or Fed-batch  
(stationary phase)

$$\frac{dX}{dt} = \mu = \text{constante} = 0$$

$$\frac{dP}{dt} = q_P = \text{constante}$$

$$\frac{dS}{dt} = q_S = \text{constante}$$

$$\frac{dX}{dt} = \text{constante} = 0$$

$$\frac{dP}{dt} = \text{constante}$$

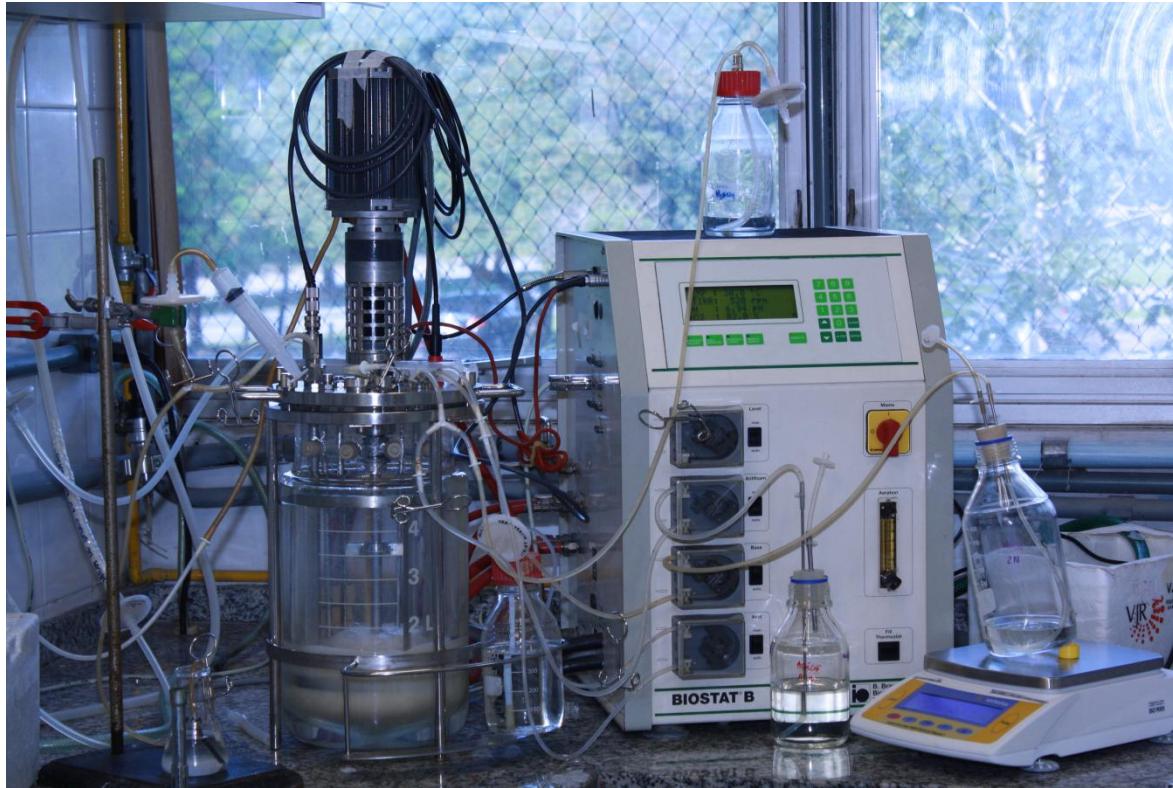
$$\frac{dS}{dt} = \text{constante}$$

Y são constantes?

# (Pseudo or Quasi)-Steady state

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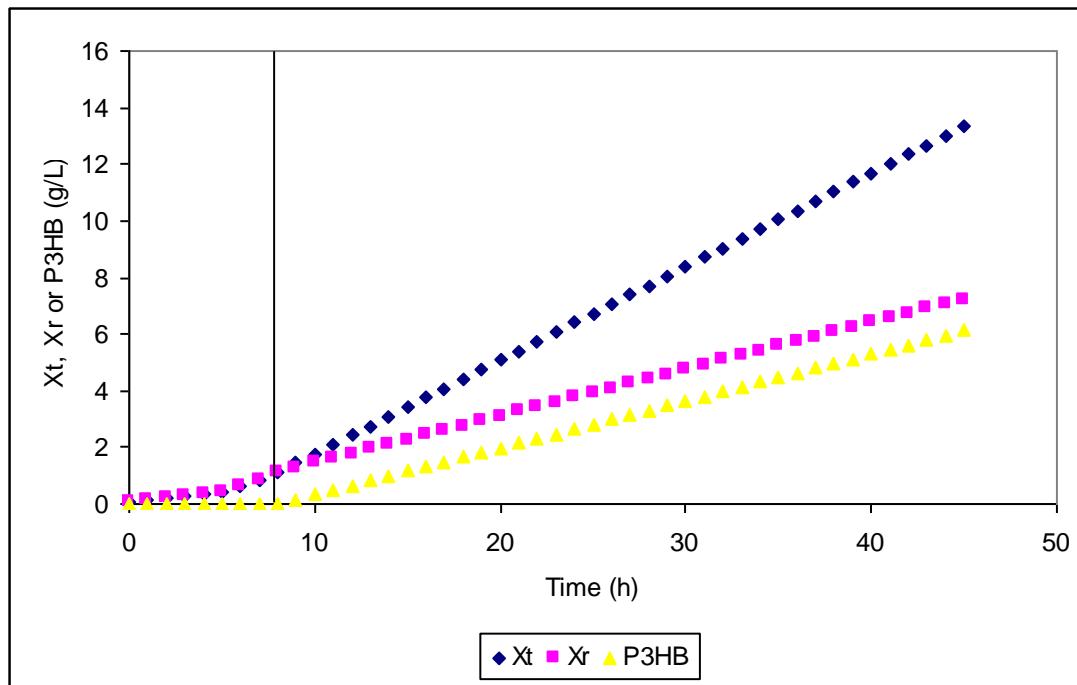
Fed-batch  
Continuous feeding



# (Pseudo or Quasi)-Steady state

Fed-batch

Continuous feeding



# (Pseudo or Quasi)-Steady state

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Fed-batch

Continuous feeding

$$\frac{dX}{dt} = \mu \text{ (Não constante)}$$

$$\frac{dP}{dt} = q_P \text{ (Não constante)}$$

$$\frac{dS}{dt} = q_S \text{ (Não constante)}$$

$$\frac{dX}{dt} = \text{constante}$$

$$\frac{dP}{dt} = \text{constante}$$

$$\frac{dS}{dt} = \text{constante}$$

Y são constantes?