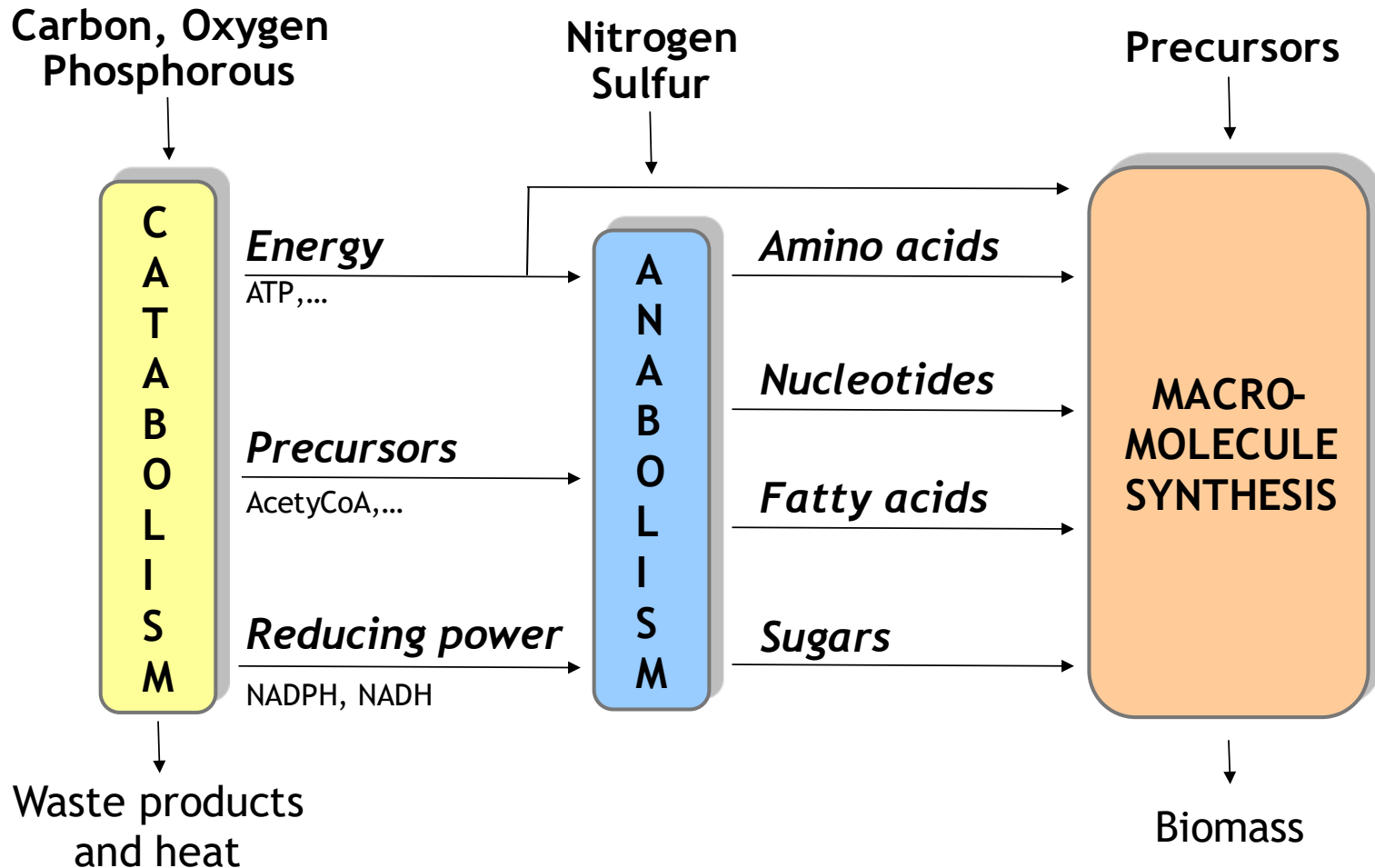


Introduction

1.1 – Batch Reactor (BSTR)

- ✓ 1.1.1 – Definitions
- ✓ 1.1.2 – Cell growth phases
- ✓ 1.1.3- Elementary Composition of the Biomass
- ✓ 1.1.4- Structured Cell Growth Models
- 1.1.5- Mass Balances to the Reactor
- 1.1.6- Relationship between Growth and Substrate Consumption
- 1.1.7- Effect of temperature and pH
- 1.1.8- Endogenous Respiration and Maintenance
- 1.1.9- Product Formation
- 1.1.10- Inhibition Models

Primary metabolism



1.1.1 – Definitions

- **Bioreactor**

system used for the development of cultures or biological processes

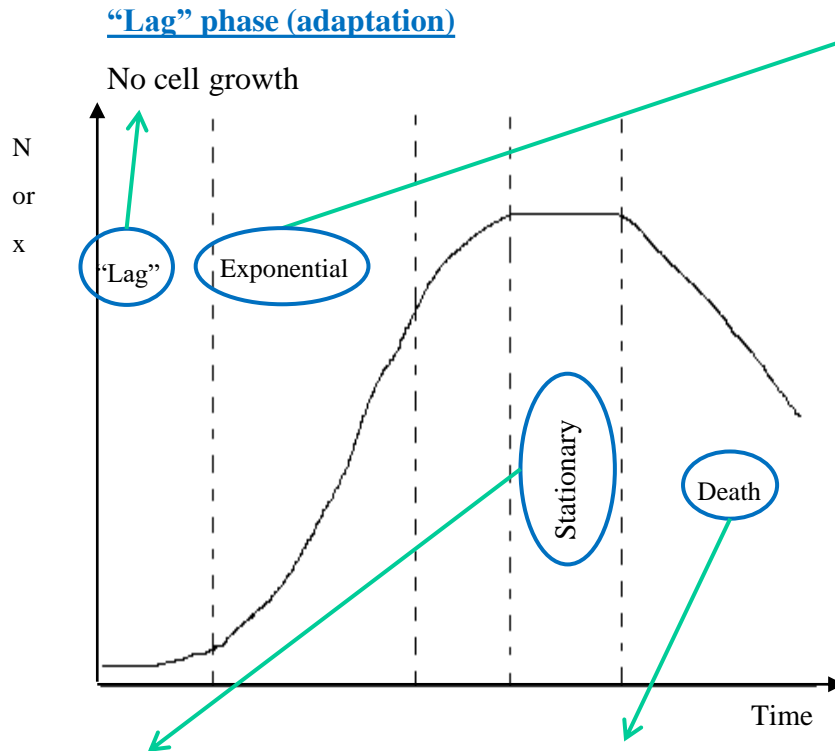
- **Batch Reactor**

all components are inserted into the bioreactor at the beginning of the process

- **Inoculum**

suspension of microorganisms of suitable concentration, used to start the fermentation process

1.1.2 – Cell growth phases



stationary phase

No cell growth

$$\frac{dx}{dt} = 0$$

Death phase

Decrease cell concentration

$$\frac{dx}{dt} = -k_d x$$

k_d – specific cell death rate (h^{-1})

Exponential phase

The cell growth rate is proportional to the cell concentration

x – cell concentration (mg/L)

μ – specific cell growth rate (t^{-1})

r_x – volumetric cell growth rate (mg cel/l.h)

$$\frac{dx}{dt} = \mu x = r_x$$

“Balanced growth”
Constant cell composition

$$\ln x = \ln x_0 + \mu t$$

$$\mu = \mu_{\max}$$

μ_{\max} – maximum specific cell growth rate (h^{-1})

1.1.3 – Elemental composition of the biomass

• Biomass = cells (bacteria, fungi, yeasts, microalgae, etc.) \rightarrow **X**

• Composition:

C **H** **O** **N** **S** **P** other elements

• Represented by chemical formulas:

$C_i H_j O_k N_l$ i, j, k, l – stoichiometric coefficients

1.1.3 – Elemental composition of the biomass

Important for estimating the microorganisms' nutrient requirements

For cell growth:

- C source
- N source

(- O₂ under aerobic conditions)

Chemical reaction for cell growth

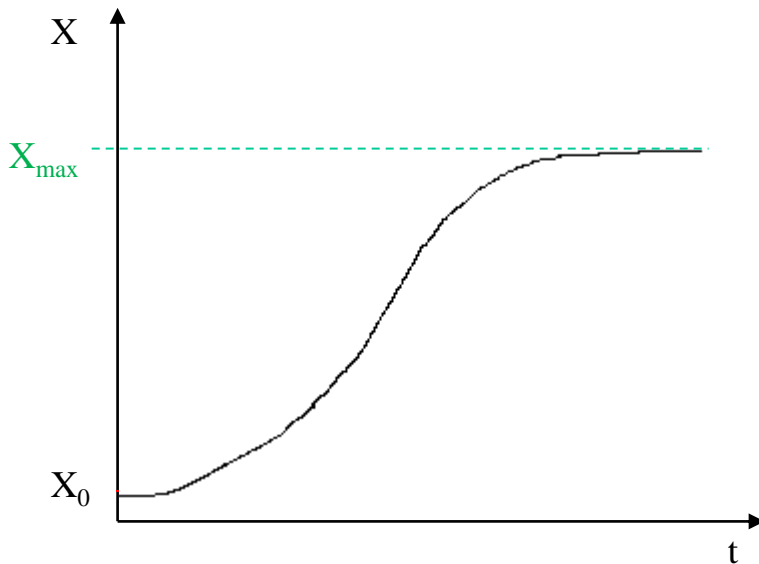


aerobiose

respiração
celular

1.1.4 – Non structured Models

- **Malthus Model:** $\frac{dx}{dt} = \mu x$ ou $r_x = \mu x$ Does not predict the appearance of the stationary phase



- **Verhulst Model:** Logistic model

$$\frac{dX}{dt} = k X (1 - \beta X)$$

$$X = \frac{X_{\max} X_0 e^{\mu_{\max} t}}{X_{\max} - X_0 (1 - e^{\mu_{\max} t})}$$

$$t = \frac{\ln\left(\frac{-(x \cdot x_{\max} - x_0 \cdot x)}{x \cdot x_0 - x_0 \cdot x_{\max}}\right)}{\mu_{\max}}$$

1.1.5 – Reactor Mass Balances

Cell growth $\frac{dx}{dt} = \mu x \quad (1)$

Cell death $\frac{dx}{dt} = -k_d x \quad (9)$

$$\frac{dx}{dt} = (\mu - k_d)x \quad (13)$$

For the exponential phase $\mu = \mu_{\max}$

$$\frac{dx}{dt} = (\mu_{\max} - k_d)x \quad (14)$$

$$\frac{dx}{x} = (\mu_{\max} - k_d)dt \Leftrightarrow \int_{x_0}^x \frac{dx}{x} = \int_0^t (\mu_{\max} - k_d)dt$$

$$\Leftrightarrow \ln \frac{x}{x_0} = (\mu_{\max} - k_d)t \Leftrightarrow x = x_0 e^{(\mu_{\max} - k_d)t} \quad (15)$$

1.1.5 – Reactor Mass Balances

$$\frac{dx}{dt} = (\mu_{\max} - k_d)x \quad (14)$$

$$\Leftrightarrow \int_{x_0}^{x_t} \frac{dx}{x} = \int_0^{t_b} (\mu_{\max} - k_d) dt$$

where t_b is the time needed to reach the maximum cell concentration ($x_t \Rightarrow x_{\max}$)

$$\Leftrightarrow \ln \frac{x_t}{x_0} = (\mu_{\max} - k_d)t_b \Leftrightarrow t_b = \frac{1}{\mu_{\max} - k_d} \ln \frac{x_t}{x_0} \quad (16)$$

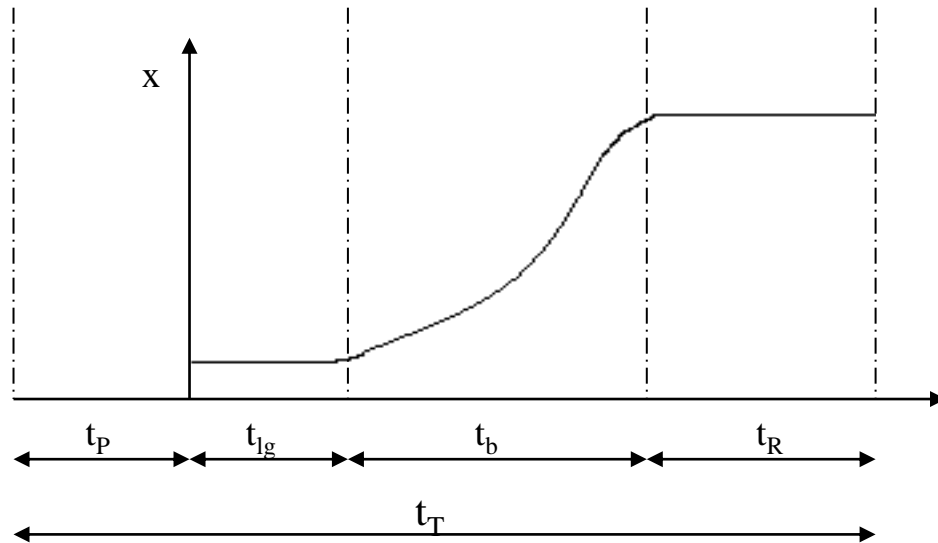
If death rate is negligible :

$$t_b = \frac{1}{\mu_{\max}} \ln \frac{x_t}{x_0} \quad (17)$$

1.1.5 – Reactor Mass Balances

Total operating time (t_T) of a Batch reactor:

$$t_T = \underbrace{t_p + t_r}_{\text{preparation}} + t_{\text{lag}} + t_b \quad (18)$$



t_p – reactor preparation time
(cleaning, sterilization, addition
of medium);

t_r – time to empty the reactor;

t_{lag} – “lag” phase time

1.1.6 – Relationship between Cell Growth and Substrate Consumption

Monod Model

$$\mu = \frac{\mu_{\max} S}{K_s + S} \quad (19)$$

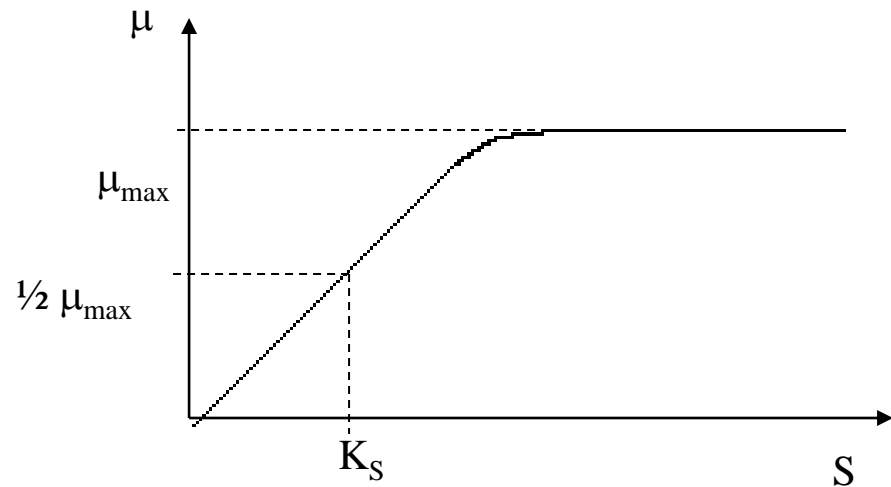
K_s - saturation constant or affinity constant (mgS/L)

S - limiting substrate concentration (mgS/L)

Assumes that only one nutrient limits growth
- **limiting substrate**

Graphic representation
of the Monod Model

$$\mu = f(s)$$



1.1.6 – Relationship between Cell Growth and Substrate Consumption

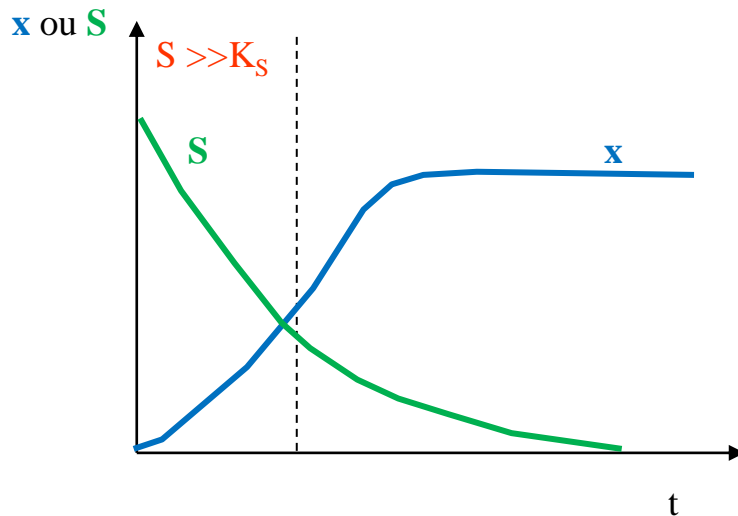
Typical values of μ_{\max} and K_s for various organisms and substrates
(at the optimum growth temperatures)

Organism (temperature)	Limiting nutrient	μ_{\max} (mg/L)	K_s (mg/L)
<i>Escherichia coli</i> (37 °C)	Glucose	0.8 – 1.4	2 – 4
	Glycerol	0.87	2
	Lactose	0.80	20
<i>Saccharomyces cerevisiae</i> (30 °C)	Glucose	0.5 – 0.6	25
<i>Candida tropicalis</i> (30 °C)	Glucose	0.5	25 - 75
<i>Candida</i> sp.	Oxygen	0.5	0.045 – 0.45
	Hexadecane	0.5	
<i>Klebsiella aerogenes</i>	Glycerol	0.85	9
<i>Aerobacter aerogenes</i>	Glucose	1.22	1 - 10

1.1.6 – Relationship between Cell Growth and Substrate Consumption

Representation of Model Monod X or $S = f(t)$

Two different zones: $S \gg K_s \Rightarrow K_s + S \approx S$



Note: The K_s value depends on the type of microorganism, and for each microorganism it depends on the type of substrate and conditions operating time of the reactor.

$$\mu = \frac{\mu_{max} S}{K_s + S} \Rightarrow \mu \approx \frac{\mu_{max} S}{S}$$

$$\Leftrightarrow \mu \approx \mu_{max}$$

$S \gg K_s \Rightarrow$ zero order rate $\Rightarrow \mu$ does not depend on S

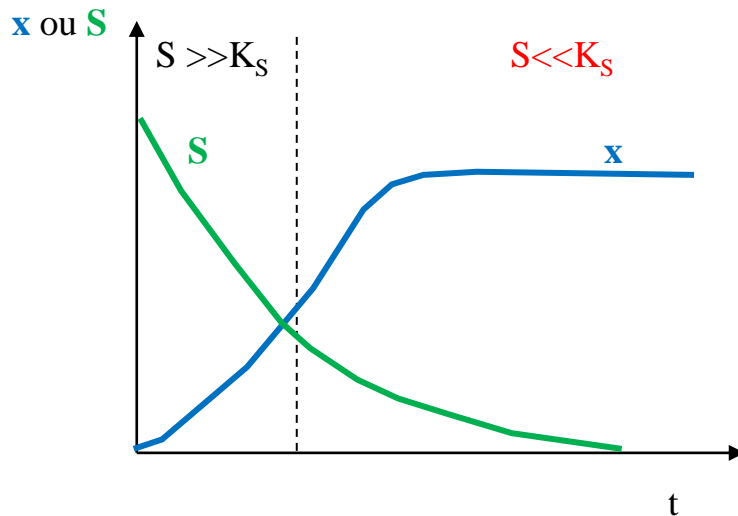
$\text{For } S \gg K_s \Rightarrow \mu = \mu_{max}$

 (21)

1.1.6 – Relationship between Cell Growth and Substrate Consumption

Representation of Model Monod X or S = f (t)

Two different zones: $S \ll K_s \Rightarrow K_s + S \approx K_s$



Note: The K_s value depends on the type of microorganism, and for each microorganism it depends on the type of substrate and conditions operating time of the reactor.

$$\mu = \frac{\mu_{max} S}{K_s + S} \Rightarrow \mu \approx \frac{\mu_{max} S}{K_s}$$

$S \ll K_s \Rightarrow$ first order rate $\Rightarrow \mu$ depends on S

$$\text{For } S \ll K_s \Rightarrow \mu = \frac{\mu_{max} S}{K_s} \quad (22)$$

1.1.6 – Relationship between Cell Growth and Substrate Consumption

Growth

$$\mu = \frac{\mu_{\max} S}{K_s + S} \quad (\text{eq. de Monod})$$

Substrate consumption

$$-\frac{ds}{dt} = r_s = \frac{v_{\max} S}{K_m + S} \quad (23) \quad (\text{eq. de Michaelis-Menten})$$

v_{\max} - maximum rate of substrate consumption (mg/l.h)

K_m – affinity constant (mgS/l)

r_s – volumetric rate of substrate consumption (mgS/l.h)

1.1.6 – Relationship between Cell Growth and Substrate Consumption

Growth yield $Y_{x/s}$

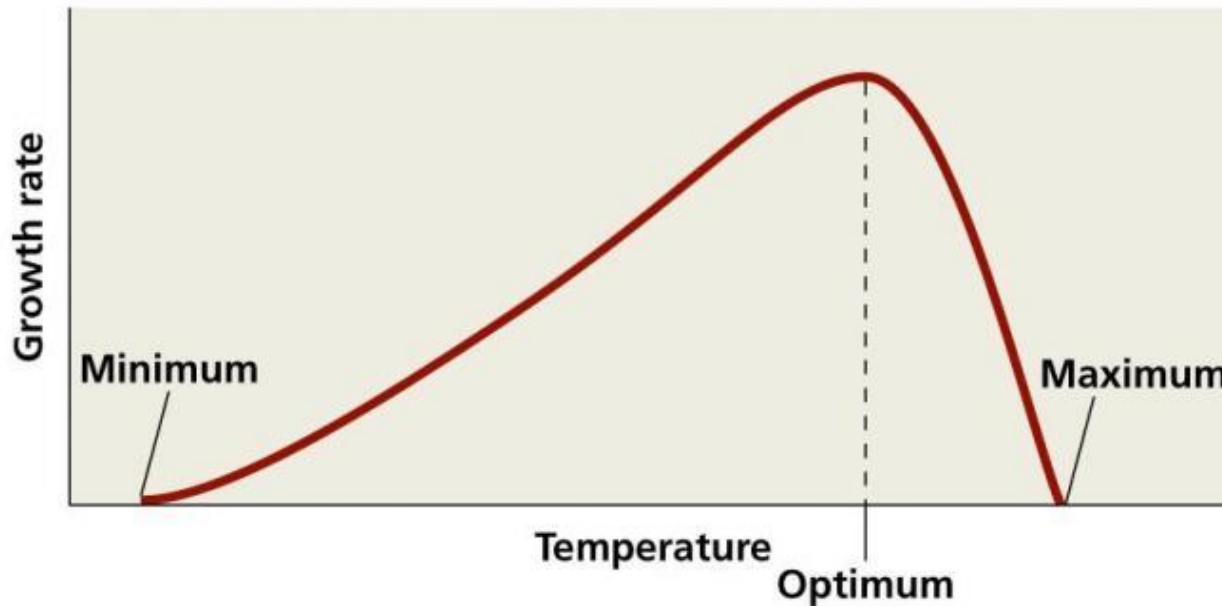
$$Y_{x/s} = \frac{x - x_0}{s_0 - s} \quad ou \quad Y_{x/s} = \frac{\Delta x}{\Delta S} \quad (24) \qquad Y_{x/s} = gX/gS$$

Note: Yield Coefficient may vary for the same medium and microorganism; may vary with μ

$$\text{if } Y_{x/s} \text{ is constant:} \qquad r_s = \frac{1}{Y_{x/s}} \mu x \quad (25) \quad ou \quad r_s = \frac{1}{Y_{x/s}} \frac{\mu_{\max} s}{K_s + s} x \quad (26)$$

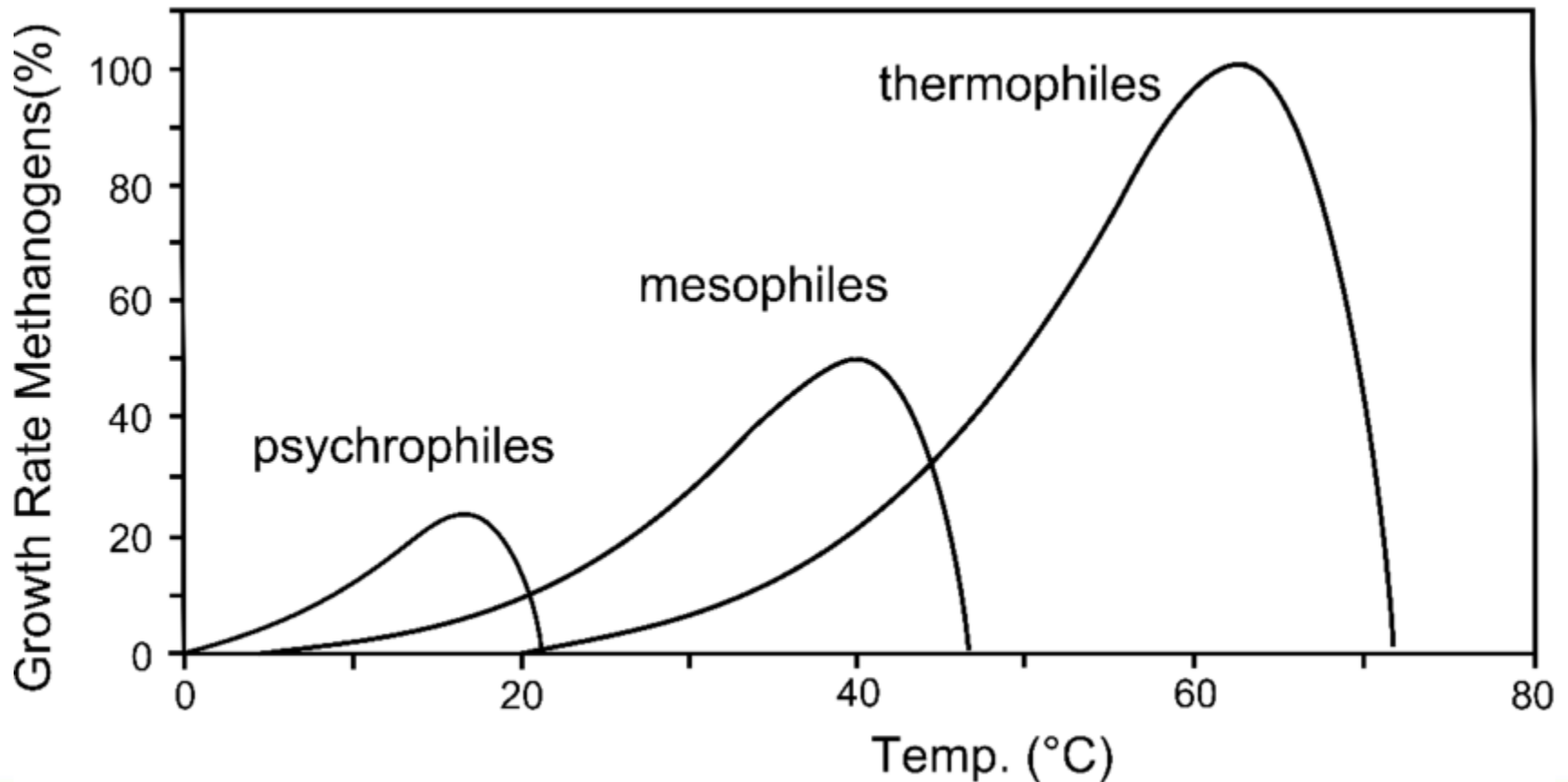
1.1.7 – Effect of Temperature and pH on growth

Effect of temperature



1.1.7 – Effect of Temperature and pH on growth

Effect of temperature



1.1.7 – Effect of Temperature and pH on growth

Effect of temperature

Classification of m.o. in function of temperature:

Group	Temperature (°C)		
	Minimum	Optimum	Maximum
Termophiles	40-45	55-75	60-80
Mesophiles	10-15	30-45	35-47
Psycrophiles			
Obligate	-5 a 5	15-18	19-22
Facultatives	-5 a 5	25-30	30-35

1.1.7 – Effect of Temperature and pH on growth

Effect of temperature

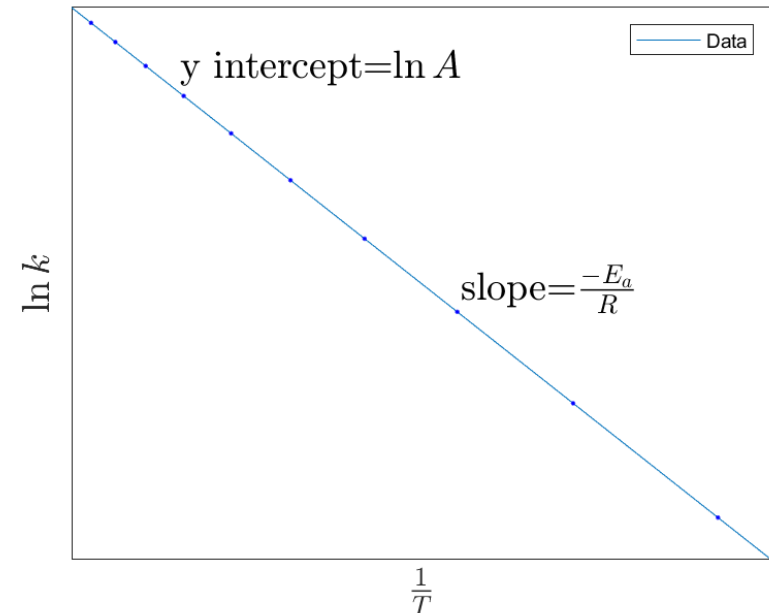
The effect of temperature on growth can be described by an equation that encompasses the Arrhenius equation and enzymatic deactivation

$$\mu = A e^{-\frac{E}{RT}}$$

$$\ln \mu = \ln A - \frac{E}{R} \frac{1}{T}$$

intercept slope

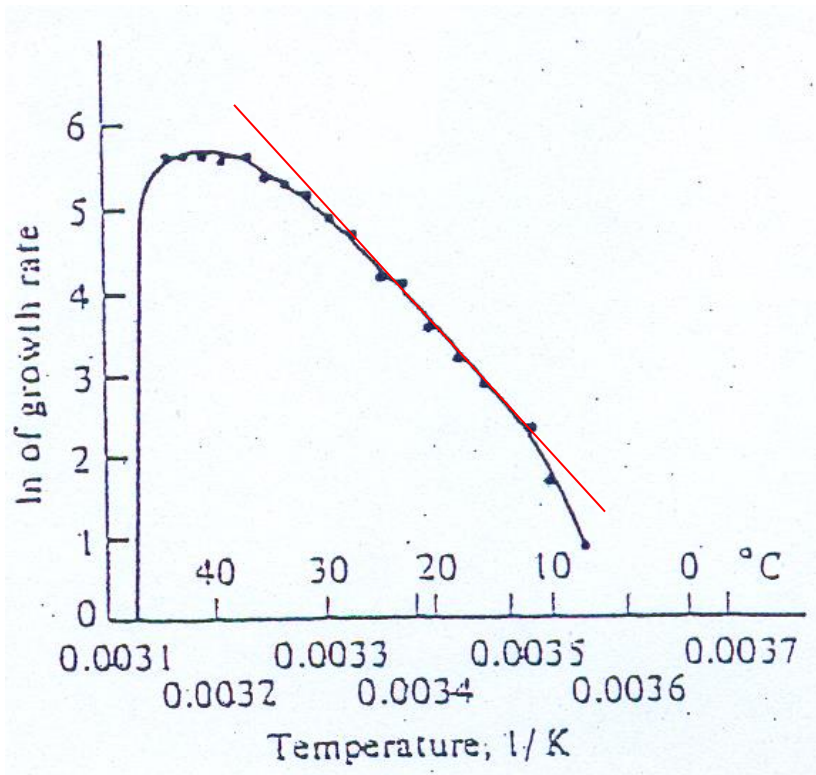
$\ln \mu$ vs $1/T$



1.1.7 – Effect of Temperature and pH on growth

Effect of temperature

The effect of temperature on growth can be described by an equation that encompasses the Arrhenius equation and enzymatic deactivation



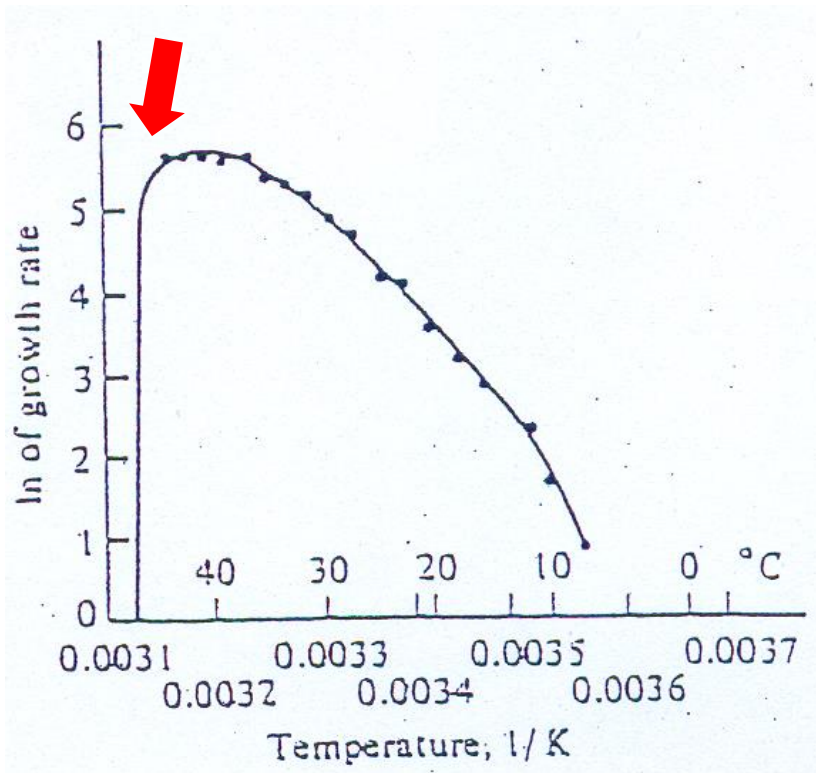
Two distinct zones:

- zone where μ increases linearly with the temperature at which the Arrhenius equation is valid:

1.1.7 – Effect of Temperature and pH on growth

Effect of temperature

The effect of temperature on growth can be described by an equation that encompasses the Arrhenius equation and enzymatic deactivation



Two distinct zones:

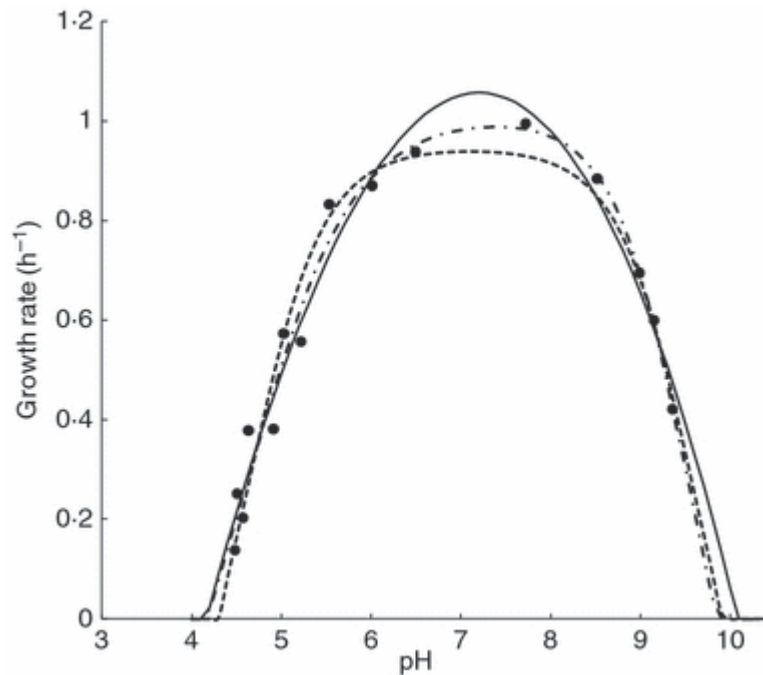
- zone where μ increases linearly with the temperature at which the Arrhenius equation is valid:
- zone in which μ decreases with the increase in temperature that corresponds to the enzymatic deactivation.

1.1.7 – Effect of Temperature and pH on growth

Effect of pH

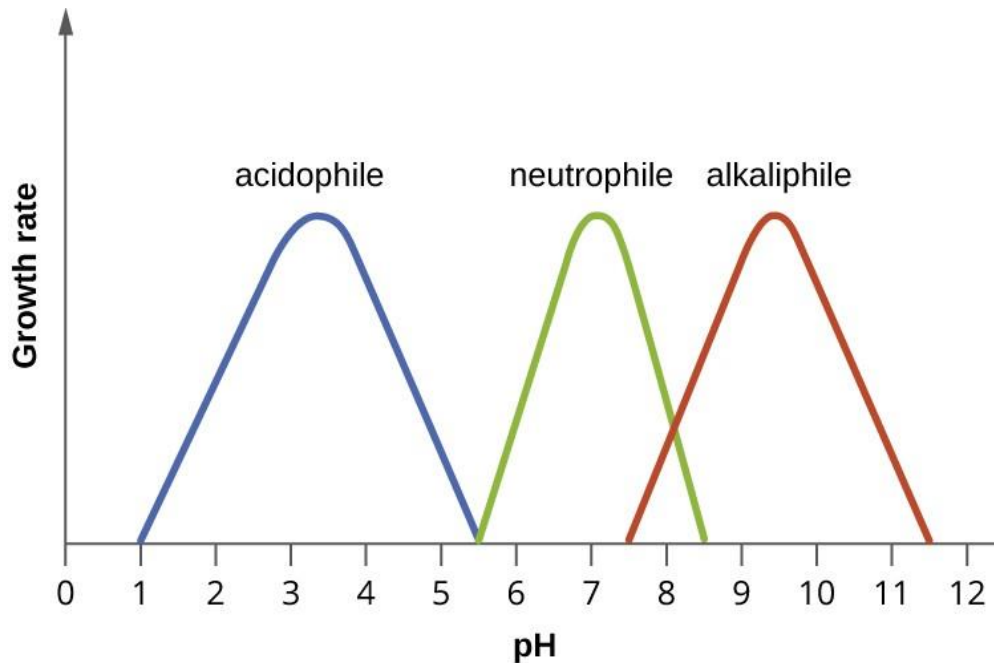
The pH influences:

- type of metabolism
- enzyme activity
- substrate or product inhibition
- biomass (cell wall) and morphological (fungi) composition



1.1.7 – Effect of Temperature and pH on growth

Effect of pH



- Most bacteria grow at pH 6.5-7.5
- Yeasts grow at pH 4-5
- Algae grow at pH=10 (contain cytoplasmic membranes that are not permeable to H^+ or OH^-)