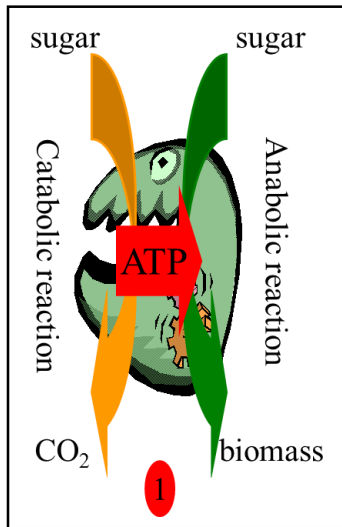


Herbert-Pirt (Extended)

Including product formation

Growth



Herbert-Pirt formulation without non-catabolic product

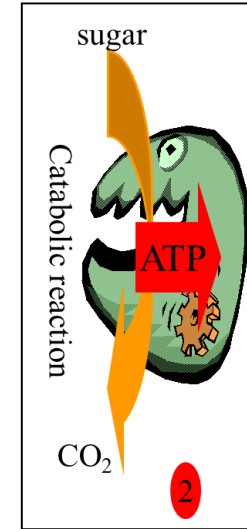
$$q_S = \frac{1}{Y_{SX}^{\max}} \mu + m_S$$

$$q_S = q_S^{\max} \frac{C_S}{K_S + C_S}$$

4 parameters: Y_{SX}^{\max} , m_S , q_S^{\max} , K_S

- q_P rate is a linear function of μ only!!!
- All q_i rates are governed by μ rate

Maintenance



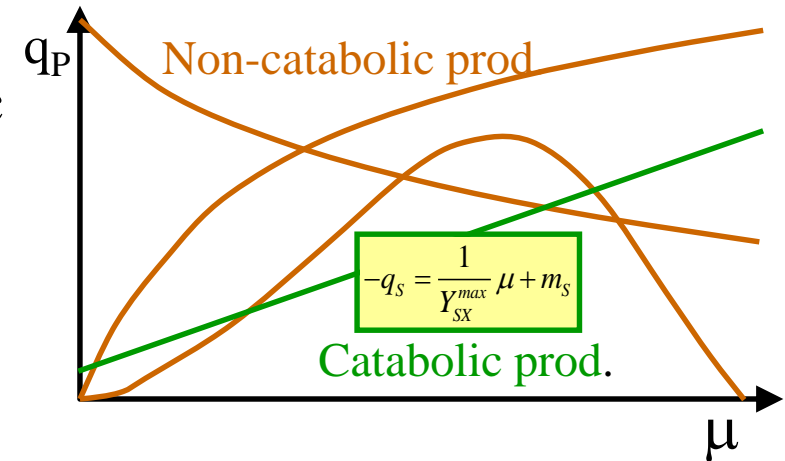
For growth with non-catabolic product ?

- Extension of Herbert-Pirt Eq. is needed
- An additional kinetic function for q_P is needed
- A modified $\mu = f(C_S)$ relation
- Other Herbert-Pirt relations

Growth with non-catabolic product ?

Various $q_P = f(\mu)$ functions

- Many different possibilities and case specific
- For catabolic products $q_P = f(\mu)$ is linear
- $q_P = f(\mu)$ function can easily be estimated in chemostat or fed batch, like $q_S = f(\mu)$



Kinetic functions for q_P

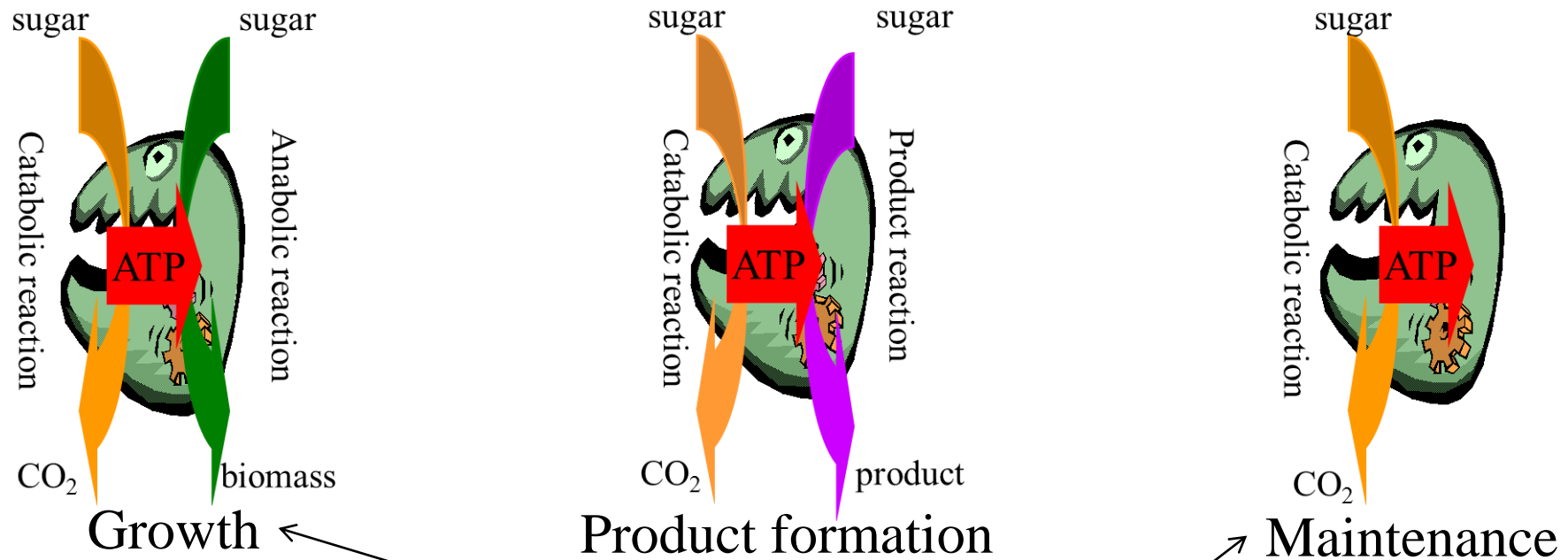
Assuming: single substrate limited conditions

Then $q_P = f(C_S)$. But C_S is difficult to measure

However for a given, μ , C_S is also fixed as $C_S = f(\mu)$
by eliminating $C_S \rightarrow q_P = f(\mu)$

This is the $q_P = f(\mu)$ concept \rightarrow Extended Herbert-Pirt Eq.

Extended Herbert-Pirt Eq.



$$-q_S = \frac{1}{Y_{SX}^{\max}} \cdot \mu + \frac{1}{Y_{SP}^{\max}} \cdot q_P + m_S$$

Substrate consumption:

1. Herbert-Pirt (not-extended) rate for **growth** and **catabolic** products
2. Extended Herbert-Pirt rate for **non-catabolic** products

All other q_i rates

All other rates follow from stoichiometric growth-catabolic coupling & substrate Herbert-Pirt equation. From known:

1. $q_S = f(C_S)$ hyperbolic
2. $q_P = f(\mu)$
3. Growth stoichiometry
4. Extended Herbert-Pirt Eq. for substrate

$$-q_S = \frac{1}{Y_{SX}^{\max}} \cdot \mu + \frac{1}{Y_{SP}^{\max}} \cdot q_P + m_S$$

$$q_{CO_2} = \dots$$

$$q_P = \dots$$

$$q_{O_2} = \dots$$

$$q_{\text{heat}} = \dots$$

$$q_{H_2O} = \dots$$

etc...

Ex: $-q_S = 2 \mu + 1.5 q_P + 0.01$ [kgS.kgX⁻¹.h⁻¹]; $q_P = 0.049$ [kgP.kgX⁻¹.h⁻¹] (independent of μ)
 $\mu = 0.02$ [h⁻¹]

Thus $Y_{SX}^{\max} = 0.5$ [kgX.kgS⁻¹]; $Y_{SP}^{\max} = 0.66$ [kgP.kgS⁻¹]; $m_S = 0.01$ [kgS.kgX⁻¹.h⁻¹] ...
 $\rightarrow Y_{SX} = 0.16$ [kgX.kgS⁻¹]; $Y_{SP} = 0.39$ [kgP.kgS⁻¹] (← Why & How???)

And substrate is consumed for:

- 32% \rightarrow biomass production: $(2 \cdot 0.02) / [(2 \cdot 0.02) + (1.5 \cdot 0.049) + 0.01]$
- 60% \rightarrow product formation: $(1.5 \cdot 0.049) / [0.1235]$
- 8% \rightarrow maintenance: $0.01 / [(2 \cdot 0.02) + (1.5 \cdot 0.049) + 0.01]$

Extended Herbert-Pirt (non-catabolic product) q_i rates

Lysine ($C_6H_{15}N_2O_2^+$) fermentation from glucose ($C_6H_{12}O_6$) and ammonia (NH_4^+).

Available data: $1/Y_{SX}^{\max} = 0.333$ [$kgC_moleX.moleS^{-1}$]; $1/Y_{SP}^{\max} = 1.5$ [$moleP.moleS^{-1}$];
 $m_S = 0.025$ [$moleS.C_moleX^{-1}.h^{-1}$]

Catabolic reaction: $-1 C_6H_{12}O_6 - 6 O_2 + 6 CO_2 + 6 H_2O$

Global growth reaction (which needs energy):

$-0.333 C_6H_{12}O_6 - 0.2 NH_4^+ - 0.95 O_2 + 1 C_1H_{1.8}O_{0.5}N_{0.2} + 0.2 H^+ + 1 CO_2 + 1.4 H_2O$

Catabolism for growth: $0.95/6 * (-1 C_6H_{12}O_6 - 6 O_2 + 6 CO_2 + 6 H_2O)$

$\rightarrow -0.158 C_6H_{12}O_6 - 0.95 O_2 + 0.95 CO_2 + 0.95 H_2O$ Substrate for anabolism $0.333 - 0.158 = 0.175$

Anabolism: $-0.175 C_6H_{12}O_6 - 0.2 NH_4^+ + 1 C_1H_{1.8}O_{0.5}N_{0.2} + 0.2 H^+ + 0.05 CO_2 + 0.45 H_2O$

Product reaction (global which needs energy):

$-1.5 C_6H_{12}O_6 - 2 NH_4^+ - 2 O_2 + 1 C_6H_{15}N_2O_2^+ + 3 CO_2 + 2 H^+ + 5 H_2O$

Catabolism for product production: $2/6 * (-1 C_6H_{12}O_6 - 6 O_2 + 6 CO_2 + 6 H_2O)$

$\rightarrow -0.33 C_6H_{12}O_6 - 2 O_2 + 2 CO_2 + 2 H_2O$ Substrate for Product synthesis $1.5 - 0.33 = 1.17$

Product synthesis: $-1.17 C_6H_{12}O_6 - 2 NH_4^+ + 1 C_6H_{15}N_2O_2^+ + 1.02 CO_2 + 1.08 H^+ + 2.98 H_2O$

Extended Herbert-Pirt (non-catabolic product) q_i rates

Lysine ($C_6H_{15}N_2O_2^+$) fermentation from glucose ($C_6H_{12}O_6$) and ammonia (NH_4^+).

Available data: $1/Y_{SX}^{\max} = 0.333$ [moleS.kgC_moleX⁻¹]; $1/Y_{SP}^{\max} = 1.5$ [moleS.moleP⁻¹];
 $m_S = 0.025$ [moleS.C_moleX⁻¹.h⁻¹]

Substrate rate Herbert-Pirt (Extended) Eq.: $-q_S = 0.333 \mu + 1.5 q_P + 0.025$

Catabolic reaction $- 1 C_6H_{12}O_6 - 6 O_2 + 6 CO_2 + 6 H_2O$

Global growth reaction (which needs energy) at μ rate

$- 0.333 C_6H_{12}O_6 - 0.2 NH_4^+ - 0.95 O_2 + 1 C_1H_{1.8}O_{0.5}N_{0.2} + 0.2 H^+ + 1 CO_2 + 1.4 H_2O$

Product reaction (global which needs energy) at q_P rate

$- 1.5 C_6H_{12}O_6 - 2 NH_4^+ - 2 O_2 + 1 C_6H_{15}N_2O_2^+ + 3 CO_2 + 2 H^+ + 5 H_2O$

Thus all q_i rates are given by Herbert-Pirt linear expressions: $q_i = f(\mu, q_P)$

$-q_S = 0.333 \mu + 1.5 q_P + 0.025$ with $q_P = \text{function}(\mu)$

$-q_{O_2} = 0.95 \mu + 2 q_P + 6 \cdot 0.025$

$+q_{CO_2} = 1 \mu + 3 q_P + 6 \cdot 0.025$

$+q_{H^+} = 0.2 \mu + 2 q_P$

$-q_N = 0.2 \mu + 2 q_P$

$+q_{H_2O} = 1.4 \mu + 5 q_P + 6 \cdot 0.025$

→ Hence all q_i and Y_{ij} are complex functions of μ , q_P [or only μ , if ($q_P = f(\mu)$)]

Extended Herbert-Pirt (non-catabolic product) Y_{ij} yields

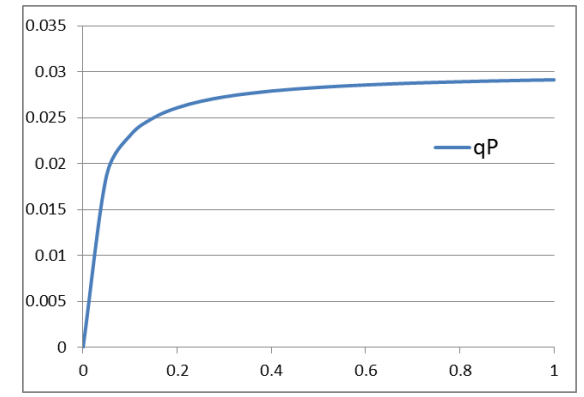
From q_i rates are given by **Herbert-Pirt** linear expressions: $q_i = f(\mu, q_P)$

- $q_s = 0.333 \mu + 1.5 q_P + 0.025$

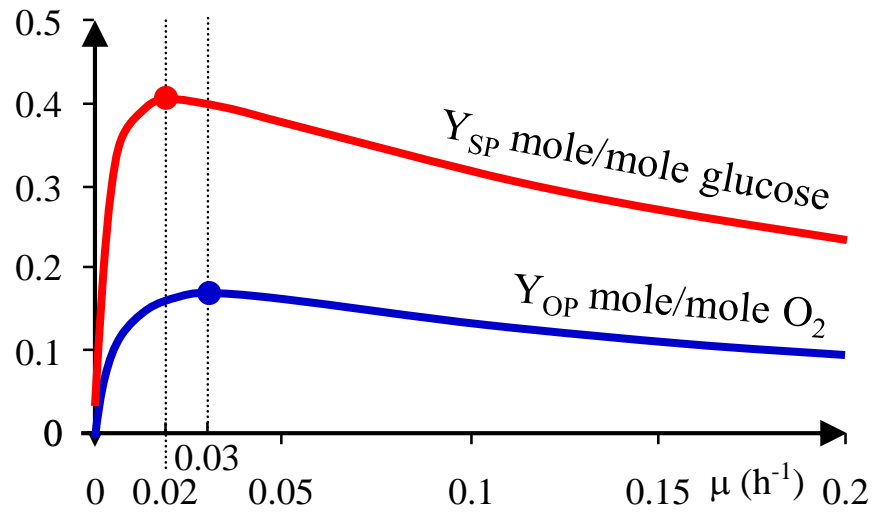
- $q_{O_2} = 0.95 \mu + 2 q_P + 6 \cdot 0.025$

If from measurements $q_P = f(\mu)$ is given:

$q_P = (0.03 \cdot \mu) / (0.03 + \mu)$ [moleP.C_moleX⁻¹.h⁻¹]



Thus: $Y_{SP} = \frac{q_P}{q_s} = \frac{0.03\mu / (0.03 + \mu)}{0.333\mu + 1.5 \cdot 0.03\mu / (0.03 + \mu) + 0.025}$ $Y_{OP} = \frac{q_P}{q_{O_2}} = \frac{0.03\mu / (0.03 + \mu)}{0.95\mu + 2 \cdot 0.03\mu / (0.03 + \mu) + 0.15}$



Which optimal μ to choose, 0.02 or 0.03 h⁻¹?

1. For best productivity on substrate criterion, μ should be 0.02 h⁻¹
2. But often, oxygen mass transfer are limited in bioreactor, in this case, the best lysine production of Lysine per consumed O_2 should be $\mu = 0.03$ h⁻¹

Black box kinetics

“Beyond single substrate limited growth”

Up to now, one assumption: Only one limited substrate
(all others non limiting)

What about:

➤ T, pH

What is the kinetic effect of temperature and pH?

➤ Multiple substrates

What about kinetic effects of other compounds than substrate?

➤ Death/inhibition/activation

Micro-organisms death?

What about the “corpses”?

➤ Mixed substrates

What if the micro-organism “sees” more than one substrate?

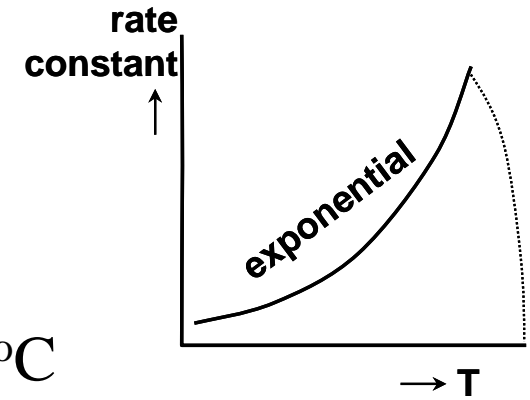
Temperature effect on kinetic parameters

- Affinity constants (K_S , etc.) are **weakly dependent** on T
- Rate constants (q_S^{\max} , m_S , q_P^{\max} , etc.) depend on T

→ **Arrhenius Eq.**

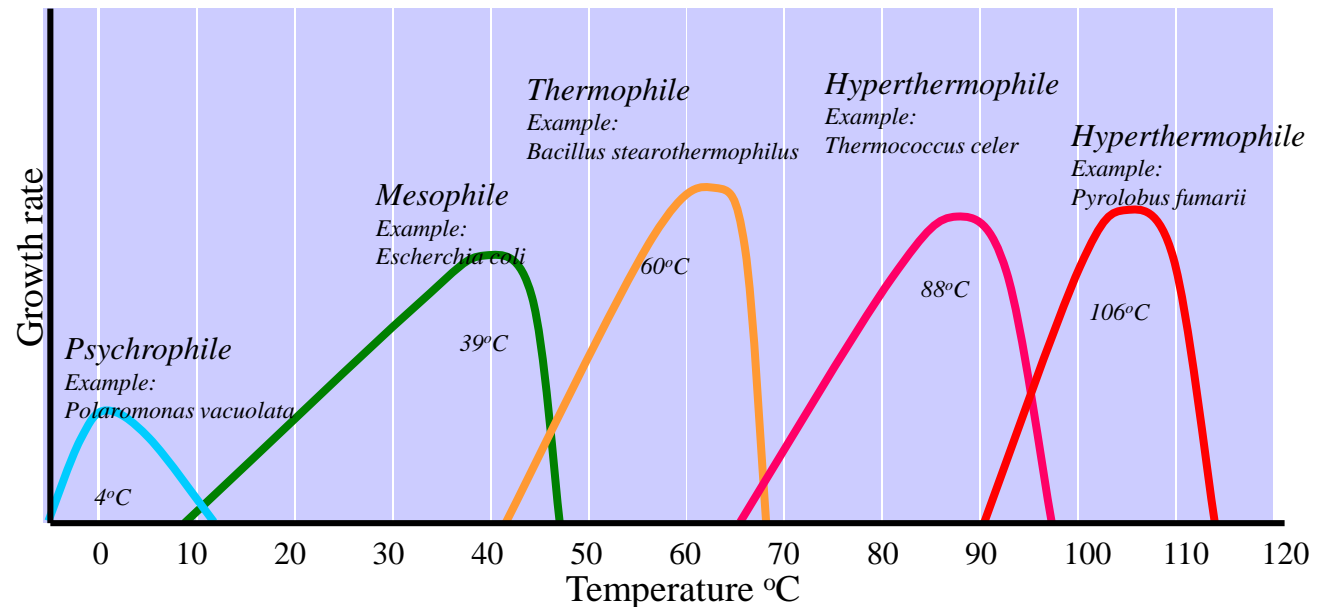
$$\text{Rate_Cst}(T) = \text{Rate_Cst}_{298K} * \exp \left[\frac{69000}{R} \left(\frac{1}{298} - \frac{1}{T} \right) \right]$$

with $R = 8.314 \text{ J/mol K}$



Increase with T: rate constants double for each $\Delta T \approx 10 \text{ }^\circ\text{C}$

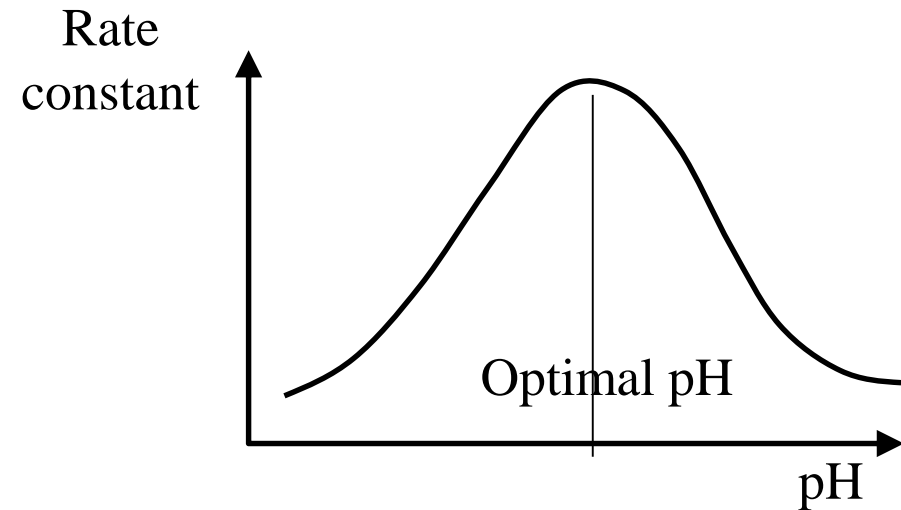
And collapse at high T



pH effect on kinetic parameters

Often empirical functions

→ see also enzymatic kinetics



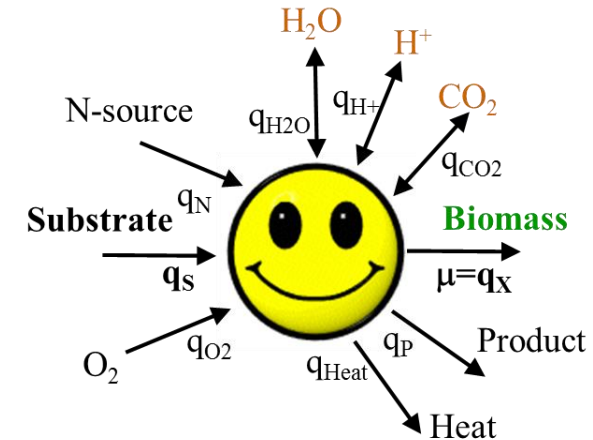
Cell Growth at pH Extremes

- Most natural environments have **pH's of 5-9**
- Most organisms have a growth **pH range of 2-3** units
- Organisms that grow at low pH are called **acidophilic**
- Organisms that grow at high pH are **alkaliphilic**
- Very few organisms grow below **pH=2** or above **pH=10**
- **Neutrophils** [pH= 7 (pH= 6-8)]

Multiple substrate limitations

What about the effect of other compound concentrations than C_S , on kinetics (C_{O_2} , $C_{NH_4^+}$, ...)

Growth requires substrates, O_2 , NH_4^+ , etc...
If one is missing, $C_{O_2} = 0$ or $C_{NH_4^+} = 0$, etc,
Rates stop! But the decrease is continuous...



The hyperbolic **Switching Function** $C_i/(K_i+C_i)$ can often be used.
(Already seen in ASM modeling):

$C_i = 0$ Switching function = 0

$C_i \gg K_i$ Switching function ≈ 1

$$q_S = q_S^{\max} \frac{C_S}{K_S + C_S} \frac{C_O}{K_O + C_O} \frac{C_N}{K_N + C_N}$$

Substrate
O₂
Ammonia
term
term
term

Multiple substrates: Inhibition

Irreversible inhibition

By reactive/toxic chemicals

→ **Death** of biomass

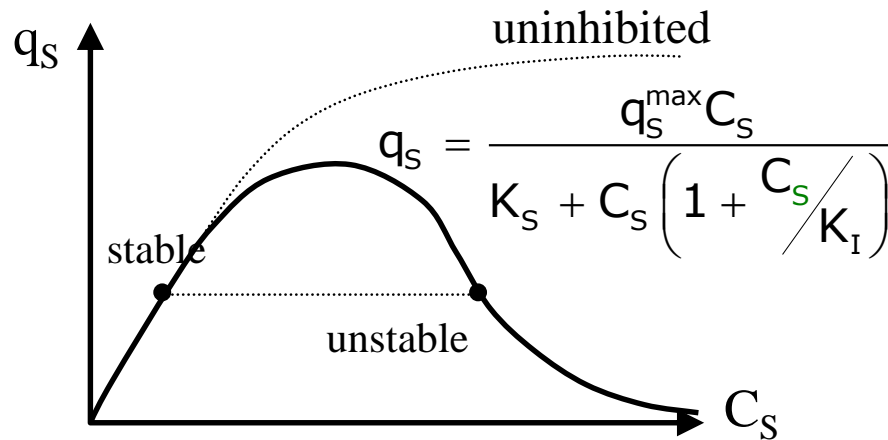
Use of simple empirical equations!

Reversible inhibition

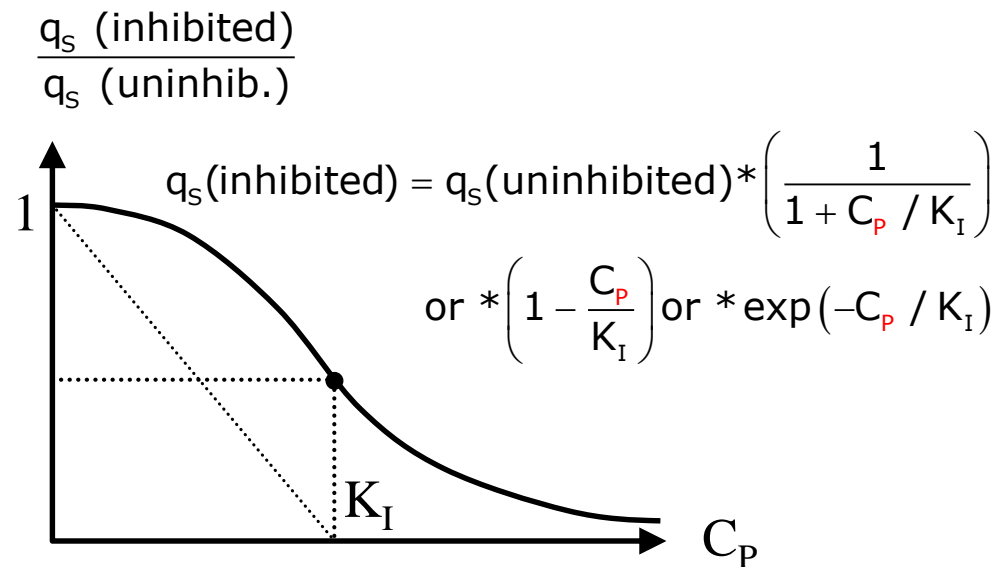
Due to high concentrations of :

- Substrates (phenol)
- Products (lactic acid, ethanol)
- Unspecific (salts, O₂ for denitrifiers)

→ **Activity** of micro-organisms drops



(S)ubstrate inhibition
(Haldane kinetics)



(P)roduct inhibition
(on substrate consumption)

Inert Organic Matter production /Active Biomass

(Microbial death - For more see ASM 1 Process#4 (p. 9))

Particularly relevant for : - Biological waste water treatment (high SRT)
- High cell density fermentation

Mechanism (Concerning C_X Active Biomass)

Growth: $C_S \rightarrow C_X$ (μ rate)

Death: $C_X \rightarrow C_S + C_{inert}$ (decay rate)

with $C_S = (1 - f)C_X$ Re-consumed
and $C_{inert} = f C_X$ Cell wall +
polymeric residues

As decay rate $k_d \approx cst$ [$kgX_{(dead)} \cdot kgX^{-1} \cdot hr^{-1}$]

and $f = C_{Inert} / X_{Dead} \approx cst$ [$kgC_{Inert} \cdot kgX_{(dead)}^{-1}$]

$1-f = C_S / X_{Dead} \approx cst$ [$kgC_S \cdot kgX_{(dead)}^{-1}$]

Biomass lysis produces C_S and C_{inert} at cst rates:

C_S : rate $q_{S,I} = k_d \cdot (1-f) = k_{S,I}$ [$kgC_S \cdot kgX^{-1} \cdot hr^{-1}$]

C_{Inert} : rate $q_I = k_d \cdot f = k_I$ [$kgC_{Inert} \cdot kg X^{-1} \cdot hr^{-1}$]

Inerts from biomass lysis can be seen as a “non-catabolic” product, at rates:

q_S Ext. Herbert-Pirt kinetic
[$kgC_S \cdot kgX^{-1} \cdot hr^{-1}$]

q_I ($=q_P$) kinetic (cst and independent of μ)
[$kgC_{Inert} \cdot kg X^{-1} \cdot hr^{-1}$]

$$-q_S = \frac{1}{Y_{SX}^{max}} \mu - k_{S,I} + m_S$$

$$q_I = (q_P) = k_I$$

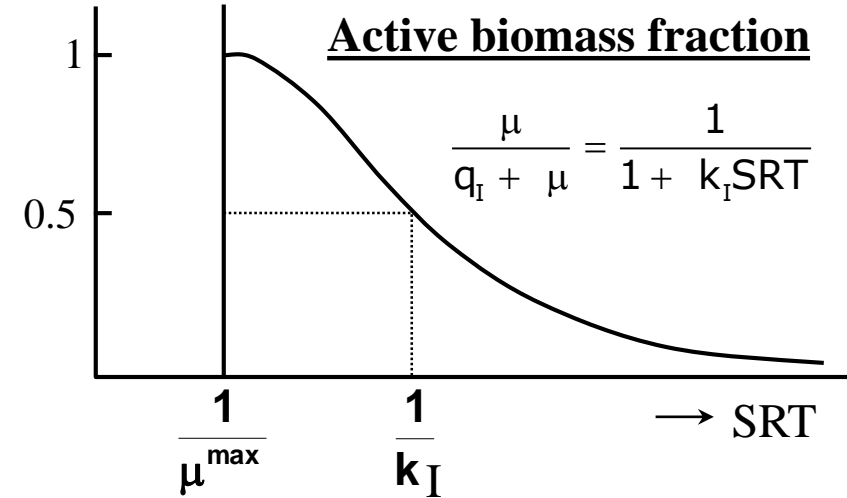
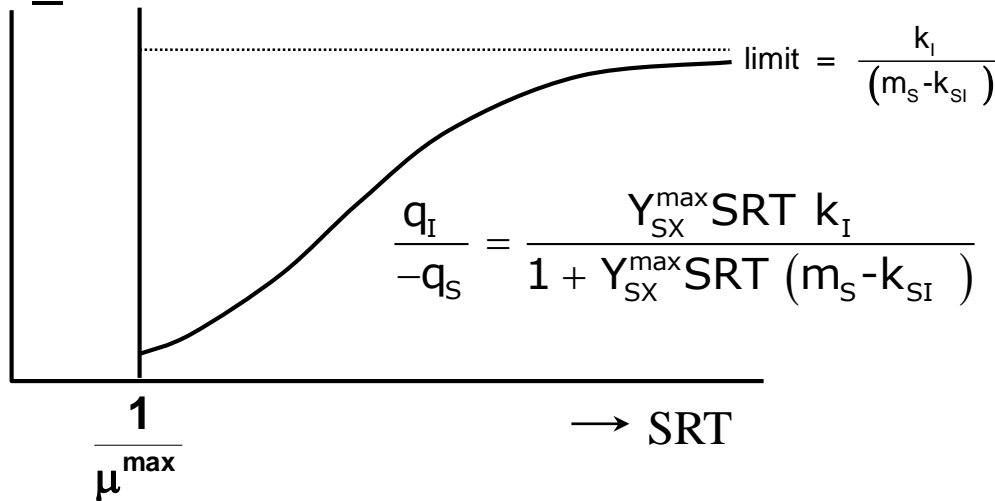
Inert Organic Matter production / Active Biomass (Microbial death - For more see ASM 1 Process#4)

$$-q_S = \frac{1}{Y_{SX}^{max}} \mu - k_{SI} + m_S$$

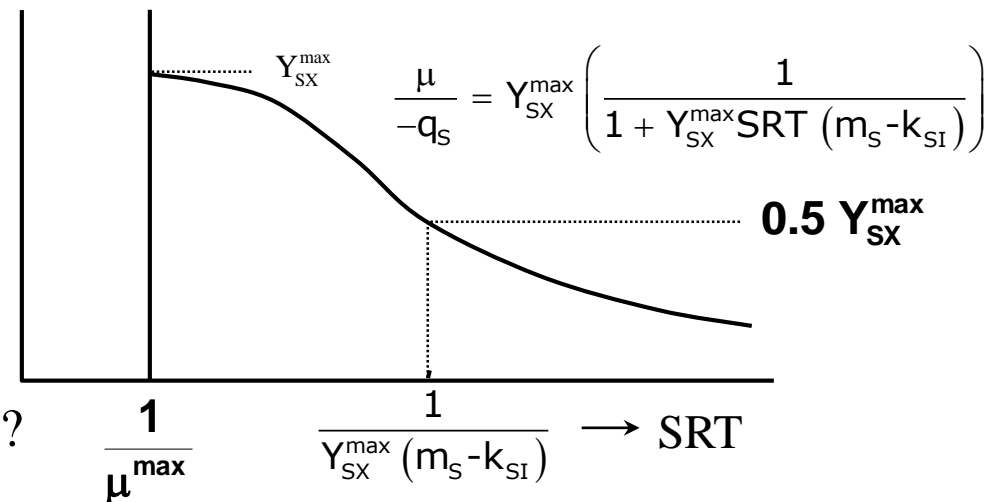
$$q_I = (q_P) = k_I$$

$$SRT = \frac{1}{\mu}$$

Y_{SI} yield of Inerts I on Substrate



Yield of active biomass on substrate

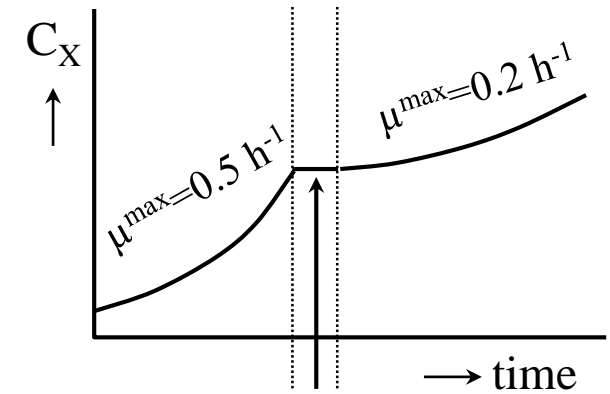
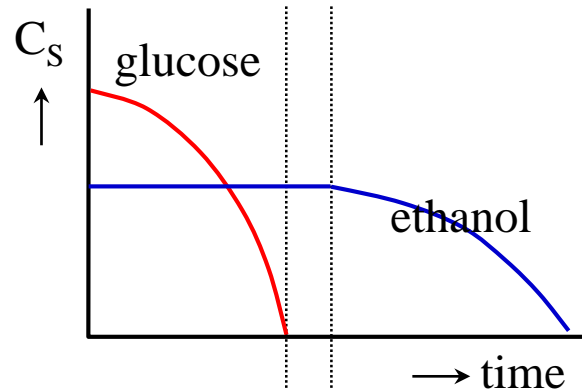


Conclusion: At high SRT

- active biomass fraction decreases strongly
- inert fraction increases strongly

? What about mineral suspended solids in WWTP ?

Mixed substrates / diauxy



time period needed to induce ethanol processing enzymes

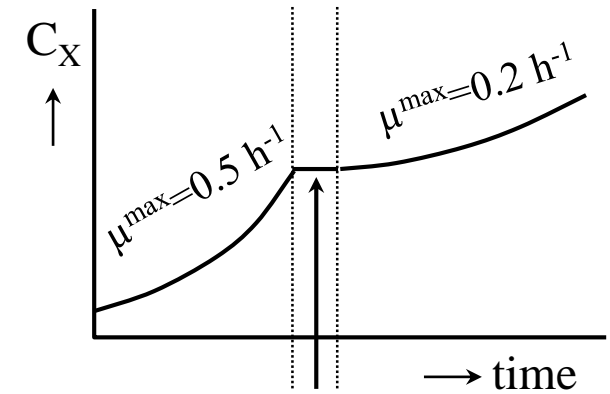
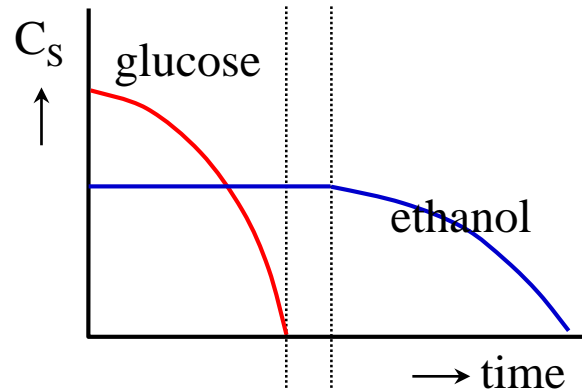
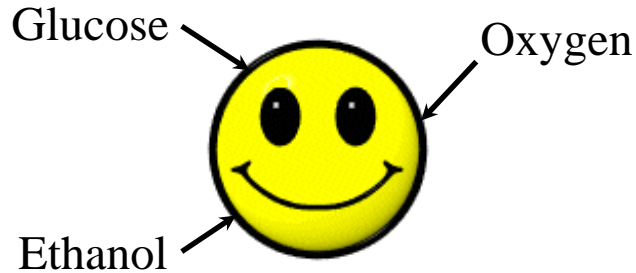
Growth observation in batch reactor, high glucose & high ethanol:

1. Substrate with highest μ^{\max} is preferred, repressing the enzymes required for the 2nd substrate (which are inhibited by 1st substrate).
2. At low glucose concentration ethanol consumption is initiated.

→ Simple kinetic approach

Extended Herbert-Pirt
+ Substrate inhibition

Mixed substrates / diauxy



time period needed to induce ethanol processing enzymes

Substrate-uptake rates (Substrate C_{S1} is preferred and inhibits Substrate C_{S2} consumption)

$$q_{S1} = q_{S1}^{\max} \cdot \frac{C_{S1}}{K_{S1} + C_{S1}}$$

$$q_{S2} = q_{S2}^{\max} \cdot \frac{C_{S2}}{K_{S2} + C_{S2}} \cdot \frac{1}{1 + C_{S1}/K_I}$$

C_{S1} “Product” inhibition
or
Inversed Switching function

Growth rate comes from Extended Herbert-Pirt Eq:

$$\mu = Y_{S1X}^{\max} \cdot q_{S1} + Y_{S2X}^{\max} \cdot q_{S2} - K_d$$

$$-q_s = \frac{1}{Y_{SX}^{\max}} \mu + m_s; \mu = (-q_s - m_s) Y_{SX}^{\max}; k_d = Y_{SX}^{\max} m_s$$

Mixed substrates

Supposing 2 substrates C_{S1} and C_{S2} on which grows biomass

$$q_{S1} = q_{S1}^{\max} \cdot \frac{C_{S1}}{K_{S1} + C_{S1}} \quad q_{S2} = q_{S2}^{\max} \cdot \frac{C_{S2}}{K_{S2} + C_{S2}}$$

Growth rate comes from Extended Herbert-Pirt Eq:

$$\mu = Y_{S1X}^{\max} \cdot q_{S1} + Y_{S2X}^{\max} \cdot q_{S2} - K_d \quad -q_s = \frac{1}{Y_{SX}^{\max}} \mu + m_s; \mu = (-q_s - m_s) Y_{SX}^{\max}; k_d = Y_{SX}^{\max} m_s$$



Ex: $q_{S1}^{\max} = q_{S2}^{\max} = 0.5$; $K_d \approx 0$; $K_{S1} = K_{S2} = 20 \text{ mg/L}$
 $Y_{S1X} = 0.5 \text{ gX/gS1}$; $Y_{S2X} = 0.2 \text{ gX/gS2}$

$$\rightarrow \mu = 0.5 q_{S1} + 0.2 q_{S2} + 0$$

$$C_{S1} = 13.3 \text{ mg/l} \quad C_{S2} = 0 \text{ mg/l} \quad \rightarrow \mu = 0.1$$

$$q_{S1} = 0.2 \quad q_{S2} = 0$$

$$C_{S1} = 5 \text{ mg/l} \quad C_{S2} = 20 \text{ mg/l} \quad \rightarrow \mu = 0.1$$

$$q_{S1} = 0.1 \quad q_{S2} = 0.25$$

$$C_{S1} = 2.2 \text{ mg/l} \quad C_{S2} = 60 \text{ mg/l} \quad \rightarrow \mu = 0.1$$

$$q_{S1} = 0.05 \quad q_{S2} = 0.375$$

Mixed substrates / Conclusion

If:

$$C_{S1} = 13.3 \text{ mg/l} \quad C_{S2} = 0 \text{ mg/l} \quad \rightarrow \mu = 0.1$$

$$q_{S1} = 0.2 \quad q_{S2} = 0$$

$$C_{S1} = 5 \text{ mg/l} \quad C_{S2} = 20 \text{ mg/l} \quad \rightarrow \mu = 0.1$$

$$q_{S1} = 0.1 \quad q_{S2} = 0.25$$

$$C_{S1} = 2.2 \text{ mg/l} \quad C_{S2} = 60 \text{ mg/l} \quad \rightarrow \mu = 0.1$$

$$q_{S1} = 0.05 \quad q_{S2} = 0.375$$

- The same high μ is maintained at lower C_{S1} , due to substrate C_{S2} availability
- Lower q_{S1} is compensated by q_{S2}
- Simultaneous uptake of many substrates allows high μ at very low C_{Si} individual substrate concentrations

Thus, growth under mixed substrates:

1. Gives competitive advantage because **high μ is possible even at low substrate concentrations** $\rightarrow ;-) ?$
2. Is an **extra degree of freedom to manipulate** microbial product formation (due to changes in production of ATP, NADPH or key carbon compounds!)