

Modeling Cell Growth

How much time passes between cellular divisions?

Doubling time,  $t_d$

- $G = 2^{t/t_d}$ 
  - G = number of generations
  - T = time that has passed
  - $t_d$  = doubling time
- Assumption: all cells are the same and have the same doubling time and grow at the same rate etc.
  - This assumption is not always true!!!

Modeling Binary Division

- $X = X_0 2^{t/t_d}$ 
  - X = number of cells
  - $X_0$  = cells at initial conditions
- $\mu_d = \frac{1}{t_d}$
- $X = X_0 2^{\mu_d t}$
- $\mu = \ln 2 \mu_d = \frac{\ln 2}{t_d}$
- $X = X_0 e^{\mu t}$
- $\ln X = \ln X_0 + \mu t$
- $\ln X = \mu t + \ln X_0$
- $\frac{1}{X} \frac{dX}{dt} = \mu$
- $\frac{1}{X} \frac{dX}{dt} = \mu$
- $\mu$  = specific growth rate

Coupling Cell Growth to Substrate Use

Yield coefficient:  $Y_{X/S} = \frac{dX}{dS}$  (g cells/g substrate) [dimensionless]

Assumptions that go into saying that the yield coefficient  $\times dS/dt$  is part of Monod equation (part in parentheses above):

- Accumulation = in - out + generation - consumption
- $\frac{dX}{dt}$  = generation
  - No consumption == no death (not valid... Slow down)
  - for now, only assuming closed system, cells don't die, but this won't always be the case

Y subscript is a hint to which is the numerator vs denominator

- $Y_{X/S} \frac{dS}{dt} = \frac{dX}{dt} \frac{dS}{dX} = \frac{dX}{dt} \frac{dS}{dX}$
- $\frac{dS}{dt} = \frac{1}{Y_{X/S}} \frac{dX}{dt} = Y_{S/X} \frac{dX}{dt}$

Derivation of Growth Expression

- Continuity equation: Accumulation = in - out + generation - consumption
  - Mass is conserved
  - Accumulation =  $\frac{dX}{dt}$
  - In =  $F X_0$
  - Out =  $F X$
  - Generation =  $\mu X$
  - Consumption = 0 (assume no appreciable death)
- $\frac{dX}{dt} = F X_0 - F X + \mu X$ 
  - Divide by  $\frac{V}{dt}$
  - $D = \frac{F}{V}$ 
    - Dilution rate: how quickly are you replacing the volume of the reactor?
  - $D = \frac{1}{t_{residence}}$
  - Assume that feed is sterile—no new cells in
- $\frac{dX}{dt} = (\mu - D)X$ 
  - “Net difference between  $\mu$  and D, times X)
  - Cell growth - cells leaving
  - At steady state,  $\frac{dX}{dt} = 0$ 
    - cell growth = cells leaving

Using Levenspiel Equation

- $\frac{dX}{dt} = \mu_{max} \left[ \frac{S}{S + K_m} \right] \left[ 1 - \frac{P}{P_{max}} \right]^n X$

Maiorella Ethanol Model

- $\mu = \mu_{max} \left[ \frac{S}{S + K_m} \right] \left[ 1 - \frac{P}{P_{max}} \right]^n$ 
  - E = efficiency of cell mass production
  - Specific ethanol production rate
  - Substrate concentration
  - $\mu_{max}$  = maximum specific production rate
  - $K_m$  = Monod constant
  - n = toxic power constant
  - $P_{max}$  = maximum product concentration
- Productivity of product is primary equation, everything else is linked to that

Luedeking-Piret Model

- $\frac{dS}{dt} = \alpha \frac{dX}{dt} + \beta X$ 
  - Substrate consumption = growth associated + non-growth associated
- $\frac{dX}{dt} = \mu X = \frac{\mu_{max} S}{K_S + S} X$ , or other function!
  - $\alpha = Y_{S/X}$  = Yield coefficient
  - $\beta = m_e$  = maintenance coefficient

Monod Equation

$r_X = \frac{dX}{dt} = \mu X$

- Where X = cell concentration (g/L)
- $\mu$  = specific growth rate

$X_t = X_0 e^{\mu t}$

$$\frac{dX}{dt} = \mu X = \frac{\mu_{max} S}{K_S + S} X$$

- X = cell concentration [g/L]
- $\mu_{max}$  = maximum specific growth rate
- S = concentration of limiting nutrient
- $K_S$  = Monod coefficient

Homogeneous

Exponential growth

- Cells grow as quickly as possible

Balanced growth

- All cells are equal metabolically/physiologically
- All cells grow at same rate: “synchronization of cell division” – Campbell 1957
- Because all cells are the same, we can assume that the mass of each cell is the same
  - X = # cells/L becomes X = g cells/L

Substrate controlled

- Some nutrient is the primary driver of how fast cells can grow
- Assume that it is the carbon source, but could also be oxygen

Saturated vs. limited growth

- Fundamental discovery that Monod pointed out in dissertation work was that cells, like enzymes, have a saturation
  - MM enzyme kinetics: 1917, a couple decades before Monod
  - There is some maximum growth rate for cell division; give more food and don't grow any faster
  - Below that threshold, they will grow, but more slowly because there aren't enough resources for them to grow!

Fermentation types

- substrate consumption/product formation proportional to rate of growth – product is part of central metabolism
- substrate and growth proportional; after some time, product formation proportional but not equal – product secondary or intermediate
- no relationship between product formation and cell growth – things just needed to maintain what you have to survive (antibiotics)

Maximum Cell Concentration

- $X_m = Y_{X/S} [S_0 + K_S - (K_S (S_0 + K_S))^{1/2}]$
- $\bar{x}_m = Y_{X/S} S_0$  if  $S_0 \gg K_S$ 
  - Maximum cells = yield coefficient \* substrate available
- Cell concentrations rapidly plummet to zero if dilution rate fluctuates over critical
  - Usually want to run just below in case of fluctuations

Derivation of Substrate-Growth Coupling Expression

$V dS = F S_0 dt - F S dt + 0 - Y_{S/X} \mu X dt$

- Assume that generation = 0
  - Not generating substrate, but not always the case
    - Could feed the reactor a different substrate to generate the substrate to make the final product
  - Think about which terms are useful for each situation
    - Consumption: rate of cells growing multiplied by yield coefficient
  - Don't include dilution because it doesn't affect substrate
  - Divide by volume, dt; Dilution factor stays the same as growth expression
    - $\frac{dS}{dt} = \frac{D(S_0 - S) - Y_{S/X} \mu X}{1}$
    - Assume steady state
    - Net flux of substrate
    - $0 = D(S_0 - S) - Y_{S/X} \mu X$
    - Use what was found above; know that  $\mu = D$
    - $\mu = D = \frac{\mu_{max} S}{K_S + S}$
    - Ideally get equations into this form:  $y = mx + b$ 
      - Y = dependent variable – things we don't control, result of choices we make (i.e. S)
      - X = independent variable – things we have control over (i.e.  $S_0$ )
    - $S = \frac{K_m D}{\mu_{max} - D}$
    - Independent variable = S
    - Dependent variable = D
    - Not exactly  $y = mx + b$ , but can segregate knowns from parameters to discover
  - $\frac{dS}{dt} = 0 = D(S_0 - S) - Y_{S/X} \mu X$ 
    - $\mu(S_0 - S) = Y_{S/X} \mu X$
    - $\mu(S_0 - S) = Y_{S/X} \mu X$
    - $X = \frac{S_0 - S}{Y_{S/X}}$
    - Substitute in S from above
      - $Y_{X/S} = 1/Y_{S/X}$
      - $X = Y_{X/S} (S_0 - S) = \frac{S_0 - S}{Y_{S/X}}$
      - Fix one of D, vary other =  $S_0$  to look at response
        - $X = Y_{X/S} S_0 - \frac{Y_{S/X} D}{\mu_{max} - D}$
        - $Y = mx + b$  format!
        - Slope =  $Y_{X/S}$
        - Independent variable =  $S_0$
        - B = last term
    - Cells consume substrate because they are growing and because they are alive – apparent yield coefficient in data, not actual yield coefficient!
      - In a batch reactor:  $\frac{dS}{dt} = \left( \frac{m_s}{\mu} + Y_{S/X} \right) \mu X = Y_{S/X,app} \mu X$ 
        - $Y_{S/X,app} = \left( \frac{m_s}{\mu} + Y_{S/X} \right)$
        - $Y_{S/X,app}$  = observed disappearance of S per appearance of X
        - In a CSTBR at steady state  $\mu = D$ 
          - $Y_{S/X,app} = \left( \frac{m_s}{D} + Y_{S/X} \right)$
          - $Y_{S/X,app} = m_s \frac{1}{D} + Y_{S/X}$
          - Y =  $m_s \frac{1}{D} + b$  format!
          - Slope is maintenance coefficient ( $m_e$ ), intercept is true yield coefficient

Steady State to Find Critical Condition

- $X = 0 = Y_{X/S} \left( S_0 - \frac{K_S D}{(\mu_{max} - D)} \right)$ 
  - $S_0 = \frac{K_S D_C}{(\mu_{max} - D_C)}$
  - $D_C = \frac{\mu_{max} S_0}{(K_S - S_0)}$  (critical dilution rate)
  - When  $S_0 \gg K_S$ ,  $\mu = D_C = \mu_{max}$
- Graphing biomass output (concentration of cells per hour) vs. D
  - Biomass output = DX
  - Constant  $S_0$
  - Critical point is peak, just before washout when productivity goes to zero
- Zooming in on the roll off point – critical point doesn't quite meet the maximum productivity

Maximum productivity

- Rate of cell output = R = DX
- $R = D Y_{X/S} \left[ S_0 - \frac{K_S D}{(\mu_{max} - D)} \right]$
- $\frac{dR}{dD} = 0 = \frac{d}{dD} \left[ D Y_{X/S} \left( S_0 - \frac{K_S D}{(\mu_{max} - D)} \right) \right]$

2,3-Butanediol

- Plot xylose, 2,3-butanediol concentration, ln(cell weight) vs hours.
  - Ln of cell dry weight because cells grow exponentially, but ln makes it appear linear!
  - When no 2,3-butane diol, just enough oxygen to respire
  - When cell dry weight goes off of the linear, oxygen deprived so 2,3-butanediol was formed
  - Rate of aeration constant throughout experiment
  - When rate of oxygen consumption exceeds rate of oxygen supply → fermentation occurs

Modeling Approach

- Cellular function is ATP constrained
  - “Gibbs Free Energy is electron-rich”
- ATP use is prioritized: cell maintenance then cell growth
  - Cells try to stay alive first, then divide
- Metabolism is regulated to maximize ATP production
  - Cells make as much ATP as they can, given environmental constraints
- O<sub>2</sub> is required for maximum ATP generation
  - Respiration is preferred until O<sub>2</sub> deficiency
- When O<sub>2</sub> is limiting, anaerobic metabolism is induced to make up the ATP deficiency
  - Cell growth is scaled to make up for deficiency

Phase I – Aerobic Growth

- $\frac{dX}{dt} = \mu_{max} X$
- $\frac{dS}{dt} = -(Q_{SA} + Q_{SR} + Q_{SF}) MW_{xylose}$
- $Q_{Si} = \left[ \frac{mol}{L \cdot hr} \right]$
- $Q_{SA} = \frac{1 \text{ mol } Xylose}{120 \text{ g Cells}} \frac{dx}{dt} = \frac{1}{120} \frac{dx}{dt}$
- $Q_{SR} = \frac{3}{70} \left( \frac{dx}{dt} \frac{1}{Y_{ATP,1}} + m_{e,1} \right) X$
- $Q_{SA} = \frac{1}{120} \frac{dx}{dt} \left[ \frac{mol}{L \cdot hr} \right]$
- $MW_{xylose} = 150 \frac{g}{mol}$
- $Q_{SA} 150 \frac{g}{mol} = Y_{\frac{X}{S}} \frac{dX}{dt} = \frac{1}{120} \frac{dX}{dt}$
- $Y_{\frac{X}{S}} = \frac{150}{120} = 1.25$

Phase II – When is O<sub>2</sub> limiting?

- Xylose + 5 O<sub>2</sub> → 5 CO<sub>2</sub> + 70/3 ATP
- Limiting when OUT > Oxygen Transfer Rate
- 5 Q<sub>SR</sub> >  $K_L a C^*$
- $N_A = H k_L a (p - p^*) = k_L a (C - C^*)$ 
  - H = Henry's law constant
  - $K_L$  = lumped transfer coefficient
  - A = surface area of bubble
  - P = partial pressure
  - C = concentration of O<sub>2</sub> in the water/bubble
  - High temperatures = low solubility of a gas
  - Concentration decreases as distance from bubble increases
  - At the limit, no oxygen in solution, rate of mass transfer = C\*
    - Once molecules go into solution, bacteria uptake them
- Metabolism is balanced so that oxidative phosphorylation (electron transport system, ETS) is saturated
  - NADH + ½ O<sub>2</sub> + H<sup>+</sup> → ATP + NAD<sup>+</sup>
    - $\frac{dNADH}{dt} = 0$ 
      - generation = consumption
      - consumption = Q<sub>ETS</sub> = 2  $k_L a C^*$
      - generation = 10 Q<sub>SR</sub> + 5/6 Q<sub>SF</sub>
      - Assumption: cell will self-optimize to make the most ATP possible
        - Only do fermentation out of desperation
    - Q<sub>ETS</sub> =  $\frac{dNADH}{dt}$
    - Q<sub>ETS</sub> = -2  $k_L a C^*$  + 10 Q<sub>SR</sub> + 5/6 Q<sub>SF</sub>
  - Maximize ATP generation!
    - Q<sub>SR</sub> = 1/10 (Q<sub>ETS</sub> - 5/6 Q<sub>SF</sub>)
    - Consumption =  $\mu_{max} X/Y_{ATP} + m_e X$
    - dATP/dt = 0 → generation = consumption
    - generation = 10/3 Q<sub>SR</sub> + 5/3 Q<sub>SF</sub> + 2 Q<sub>ETS</sub>
    - Q<sub>SF</sub> = 18/25 ( $\mu_{max} X/Y_{ATP} + m_e X$  - 7/3 Q<sub>ETS</sub>)

Phase III - Fermentation

- Q<sub>ETS</sub> = 2  $k_L a C^*$
- Q<sub>SF</sub> = 18/25 ( $\mu_{max} X/Y_{ATP} + m_e X$  - 14/3 2  $k_L a C^*$ )
- Q<sub>SR</sub> = 1/10 (2  $k_L a C^*$  - 5/6 Q<sub>SF</sub>)
- Q<sub>SA</sub> 1/120  $\frac{dX}{dt}$
- $\frac{dX}{dt} = \mu_{max} X - k_d P X$
- $\frac{dP}{dt} = 5/6 Q_{SF} MW_{2,3butanediol}$
- MW<sub>2,3butanediol</sub> = 90 g/mol
- $\frac{dS}{dt} = -\frac{1}{120} Q_{SR} + Q_{SF} + Q_{SA} MW_{xylose}$

Effect of Dissolved Oxygen on Cell Growth

- $\mu = \frac{\mu_{max} DO}{K_{DO} + DO}$ 
  - $K_{DO} \ll DO$
  - $\mu \approx \mu_{max}$