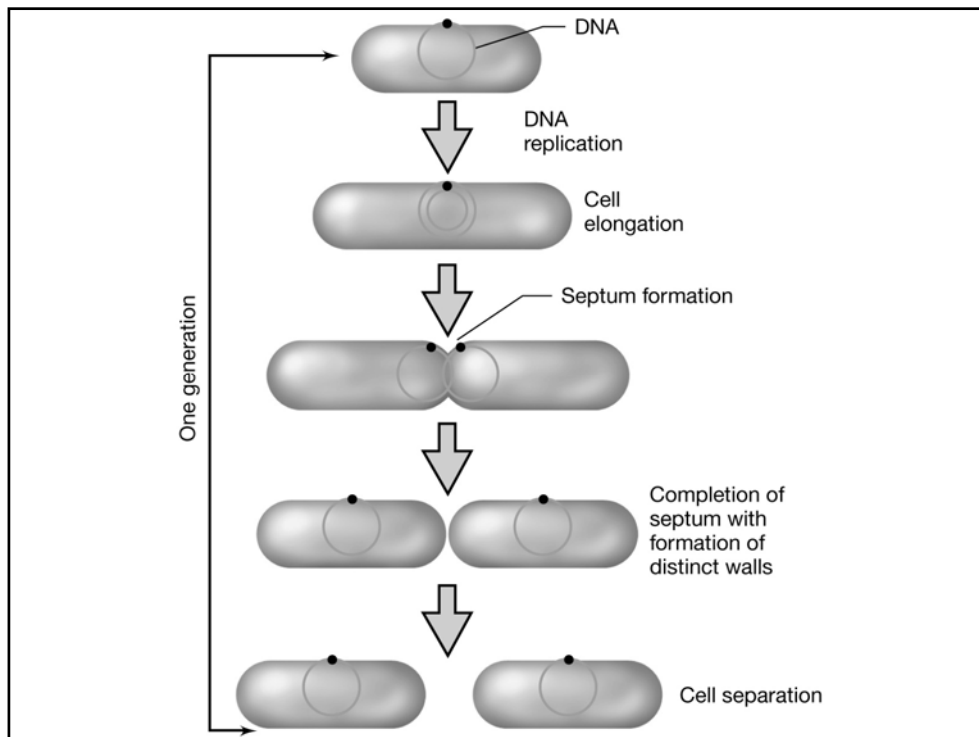


The Process of Growth

- Metabolism required for growth, both anabolic and catabolic; ~2000 reactions!
- Usual Definition: **Increase in cell numbers**
Other definitions possible – spores, UMC's, respiration, viable but nonculturable, morphology changes (life cycle)
- Divide via Binary Fission: 3 mechanisms involved!
Cell Elongation – cell wall
DNA Replication – rate limiting step
Cell Division – septum formation



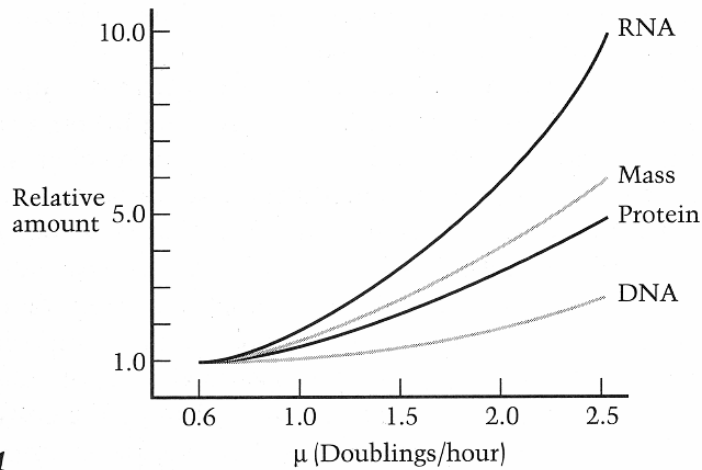


Figure 1

Effect of nutrition-imposed growth rate on the composition of *Escherichia coli* B/r. All values are expressed in amounts per cell normalized to values at $\mu = 0.6$ (mass = 1.48×10^{-13} g; protein = 1.00×10^{-13} g; RNA = 2.0×10^{-14} g; DNA = 6.3×10^{-15} g). (Plotted from data in Bremer and Dennis, 1987.)

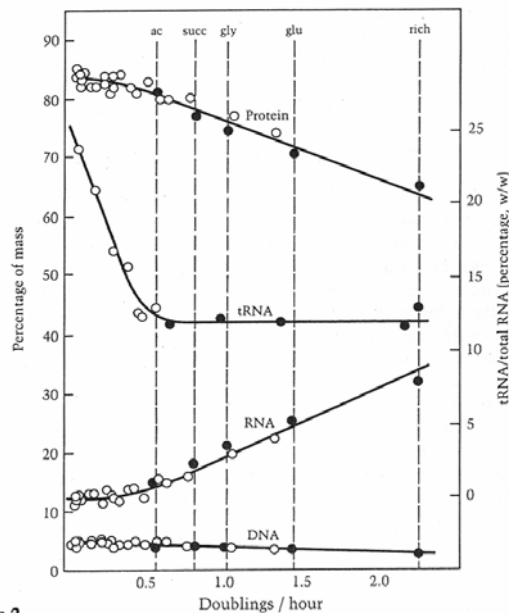
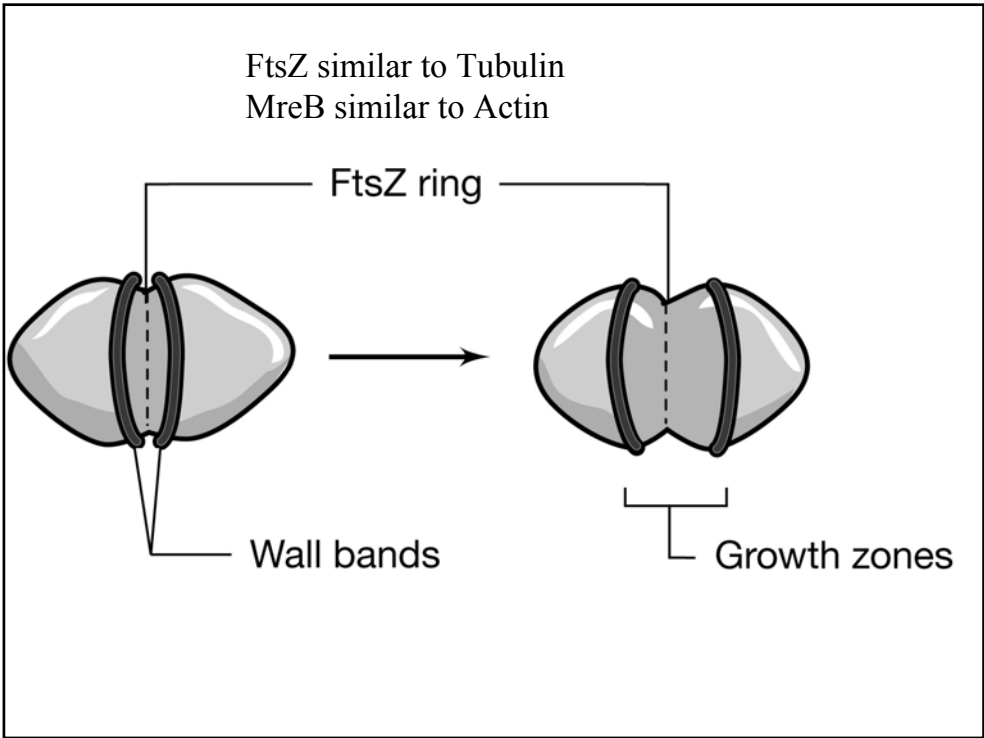
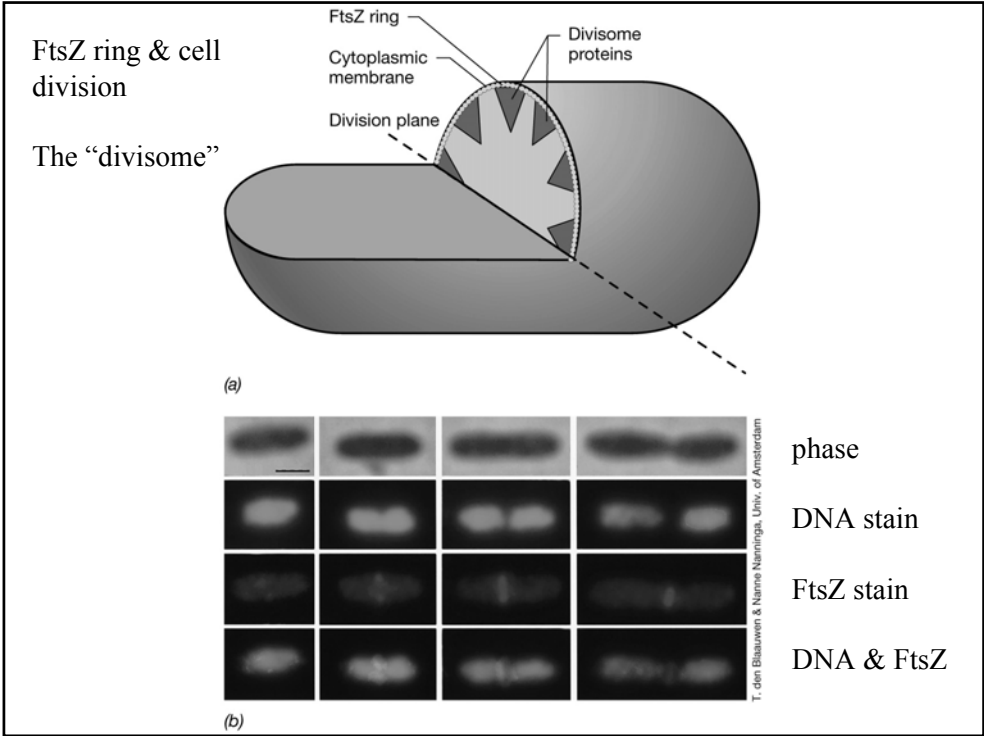


Figure 2

Effect of growth rate on the cellular proportions of protein, RNA, and DNA. Filled circles refer to results from cultures undergoing balanced growth in batch culture in various media; open circles are from cultures growing in a glucose-limited chemostat. (From Jacobsen, 1974.)



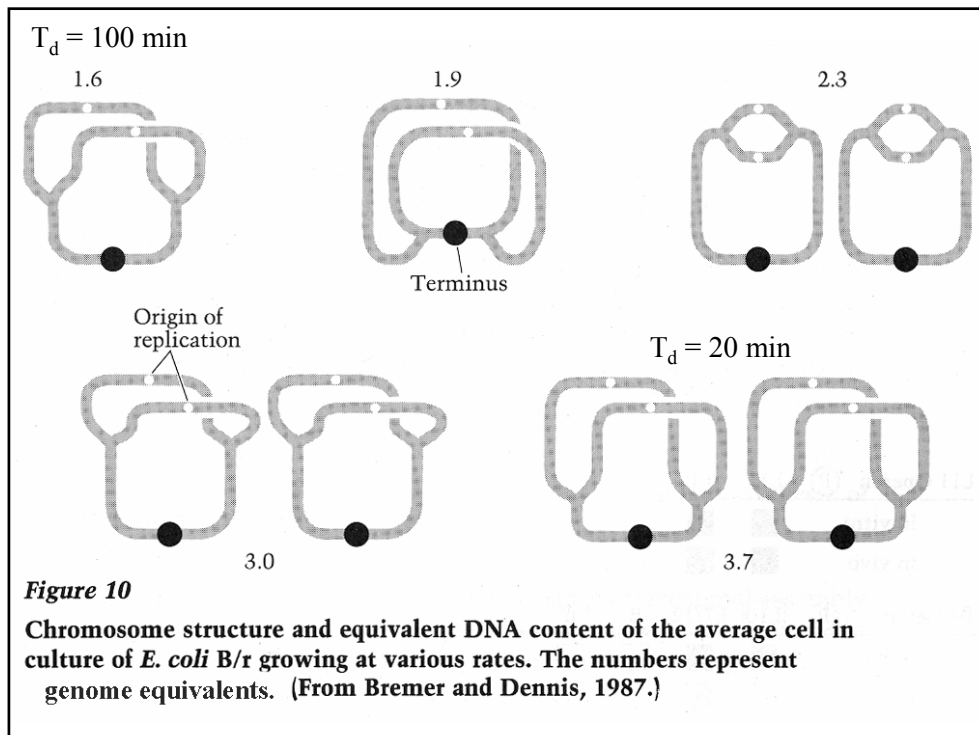
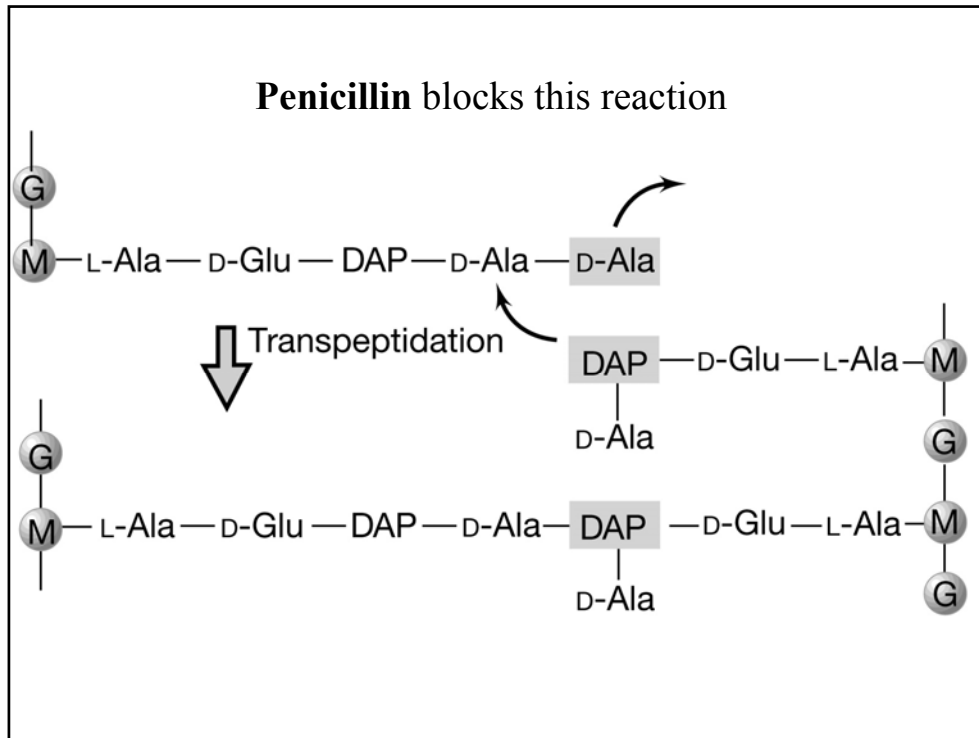




Figure 14

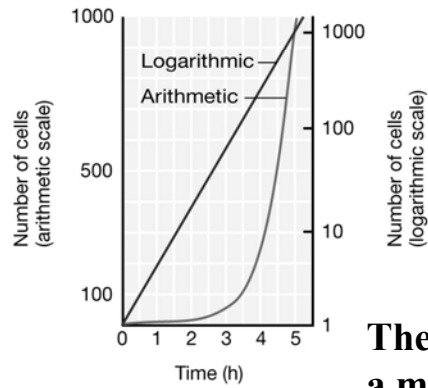
Electron micrograph of a mixture of cells of *E. coli* B/r grown at different rates. The large cells grew with a doubling time of 22 minutes, the small ones with a doubling time of 72 minutes. (From Nanninga and Woldringh, 1985.)

The Process of Growth

- Growth Rate: Time it takes to reproduce
 $t_{1/2} = \ln 2 / \mu = 0.693 / \mu = g$
- Phases of Growth in Batch culture
Lag, Log, Stationary, Death
- Measurement of Growth
Total cell counts
Viable cell counts
Turbidity

Time (h)	Total number of cells	Time (h)	Total number of cells
0	1	4	256
0.5	2	4.5	512
1	4	5	1,024
1.5	8	5.5	2,048
2	16	6	4,096
2.5	32	.	.
3	64	.	.
3.5	128	10	1,048,576

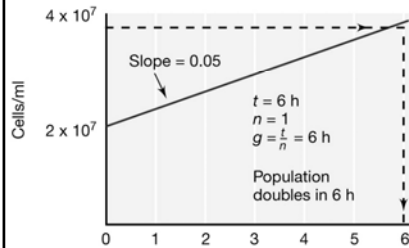
(a)



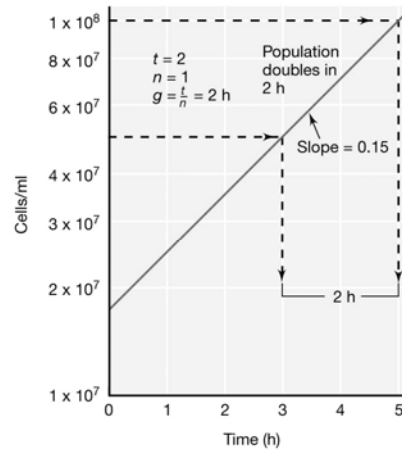
(b)

The growth rate of a microbial culture

How to estimate the generation times of an exponential microbial culture using semi-log plots.



(a)

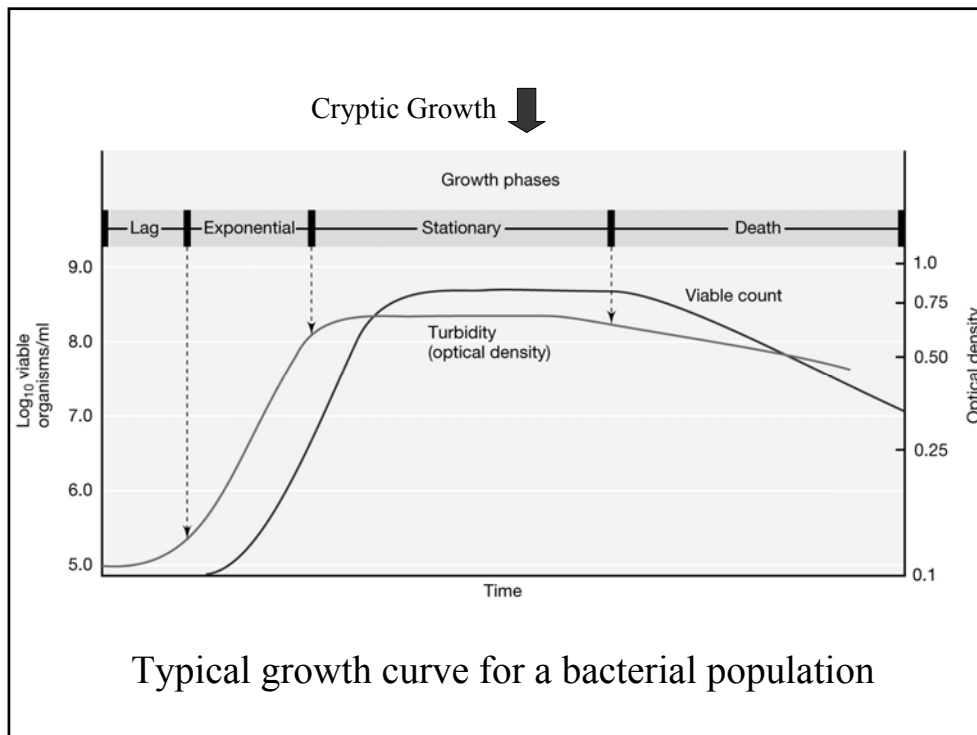


(b)

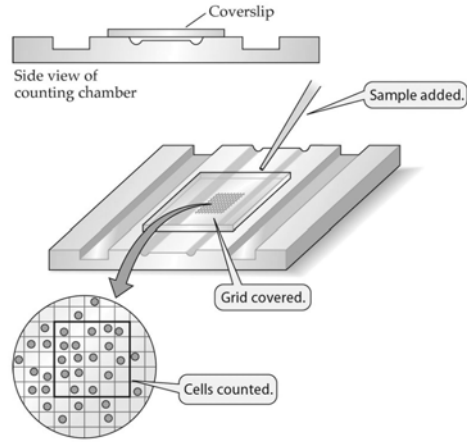
t = time
 n = generations
 g = generation time

Table 6.1 Approximate generation times for several organisms growing in media optimal for growth

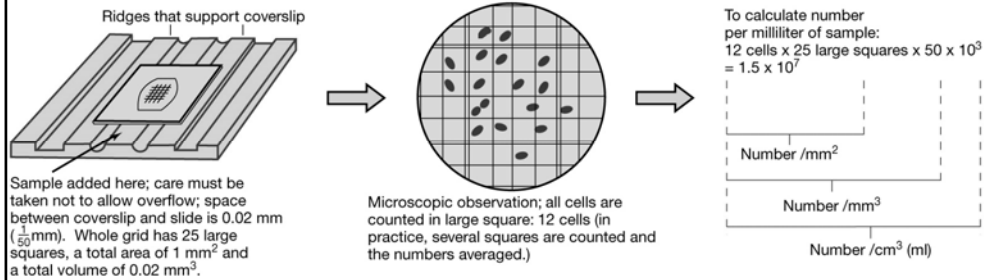
Species	Generation Time
<i>Escherichia coli</i>	20 min
<i>Bacillus subtilis</i>	28 min
<i>Staphylococcus aureus</i>	30 min
<i>Pseudomonas aeruginosa</i>	35 min
<i>Thermus aquaticus</i>	50 min
<i>Thermoproteus tenax</i>	1 hr 40 min
<i>Rhodobacter sphaeroides</i>	2 hr 20 min
<i>Sulfolobus acidocaldarius</i>	4 hr
<i>Thermoleophilum album</i>	6 hr
<i>Thermophilum pendens</i>	10 hr
<i>Mycobacterium tuberculosis</i>	13 hr 20 min



Total Cell counts using the Petroff-Hausser Counter



Total Cell counts using the Petroff-Hausser Counter



Viable cell count methods

30-300 on standard
Petri Dish

Spread-plate method

Sample is pipetted onto surface
of agar plate (0.1 ml or less)

Sample is spread evenly over
surface of agar using sterile
glass spreader

Incubation

Surface colonies
Typical spread-plate results

Pour-plate method

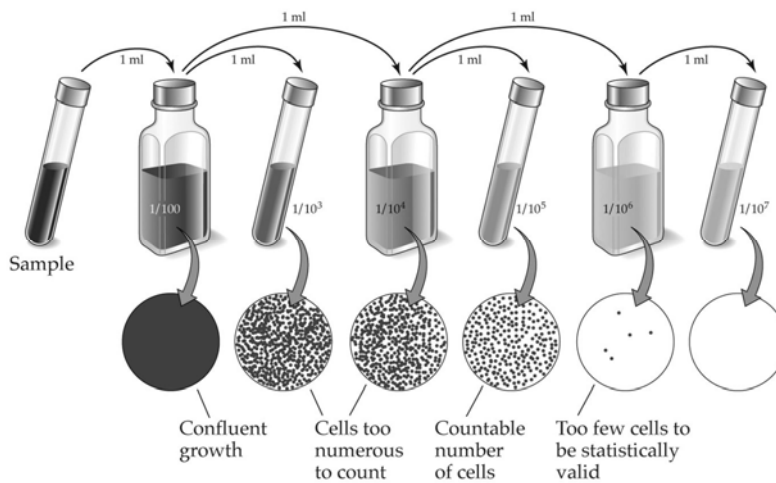
Sample is pipetted into
sterile plate

Sterile medium is added and
mixed well with inoculum

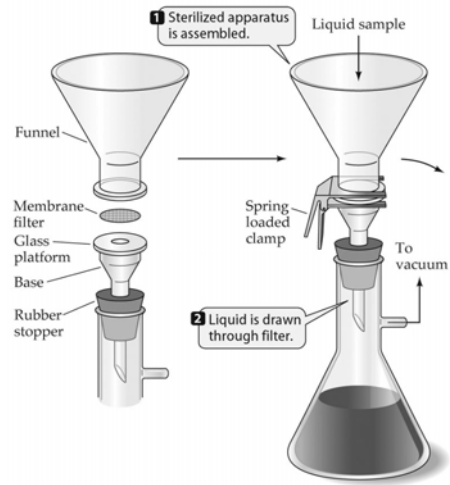
Incubation

Surface colonies
Subsurface colonies
Typical pour-plate results

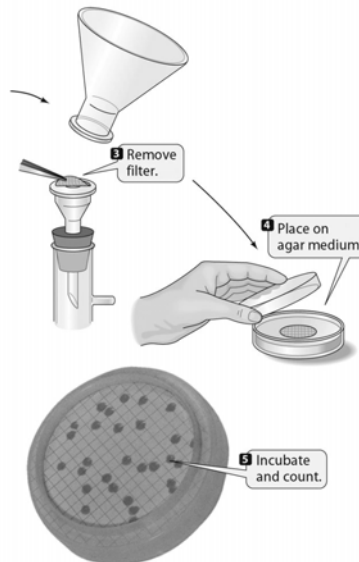
Counting the number of viable cells by serial dilution and plate count



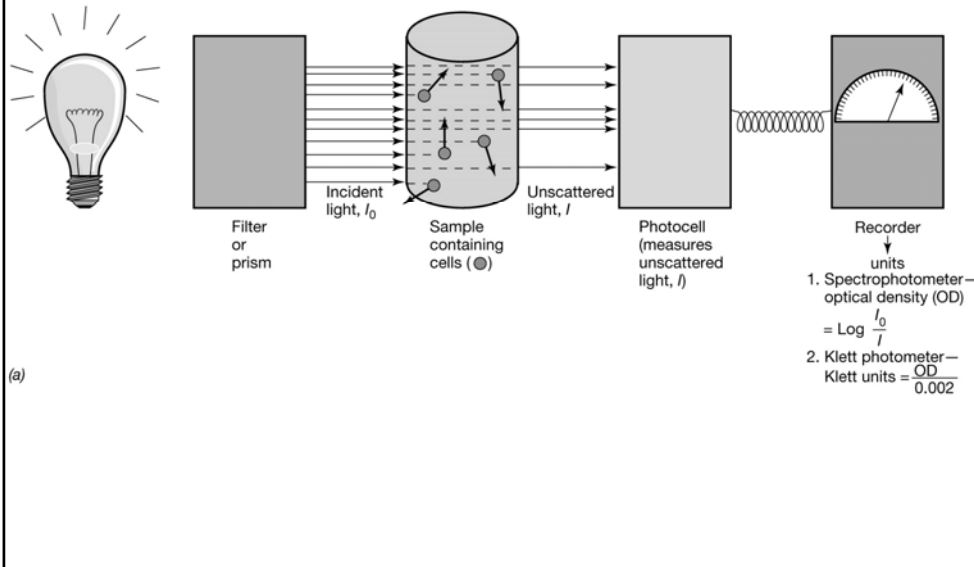
(Part 1) Concentration of cells by membrane filtration



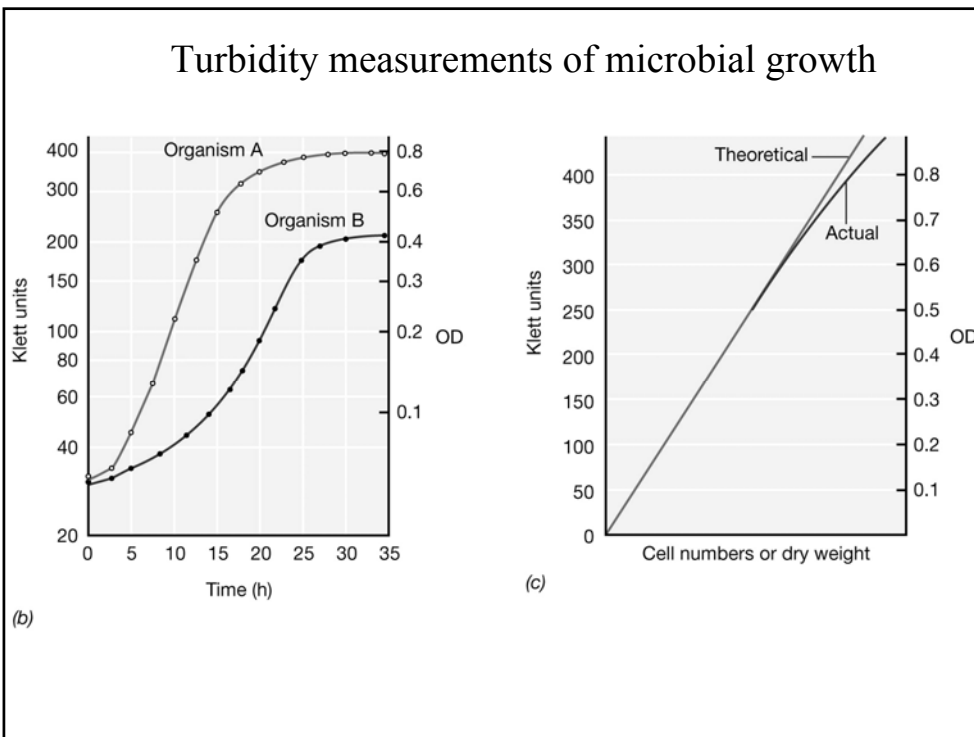
(Part 2) Concentration of cells by membrane filtration



Turbidity measurements of microbial growth



Turbidity measurements of microbial growth

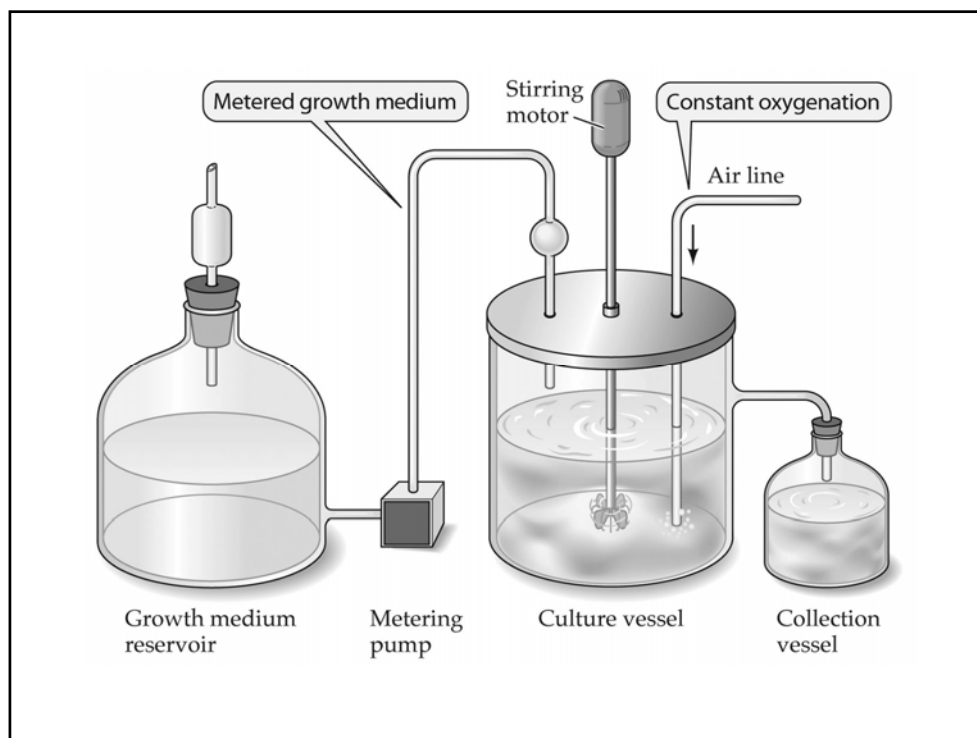


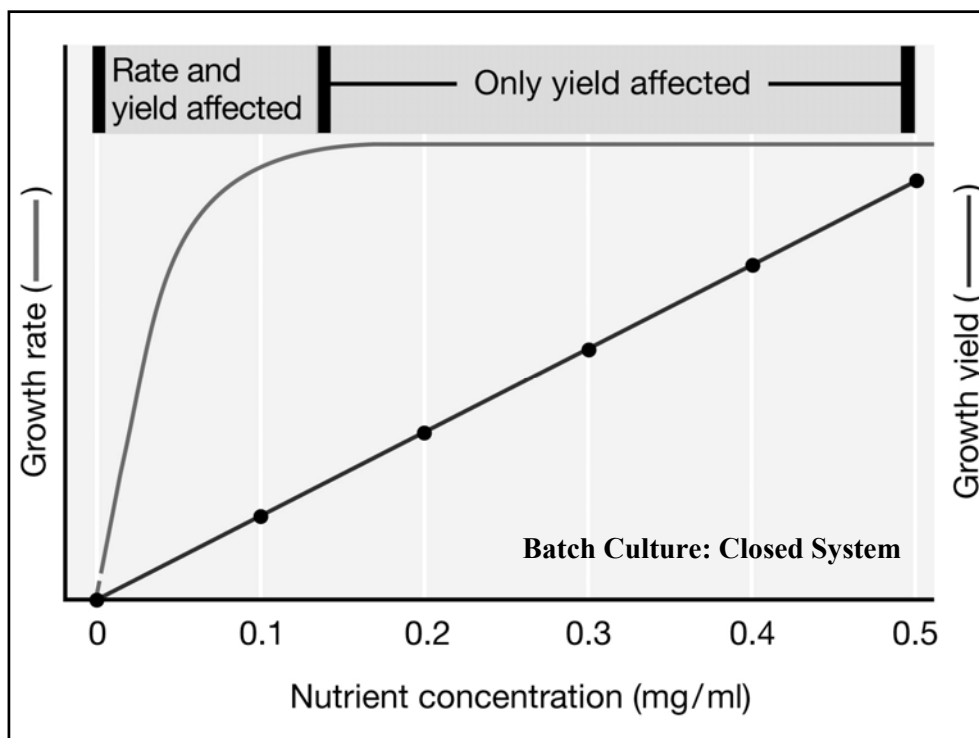
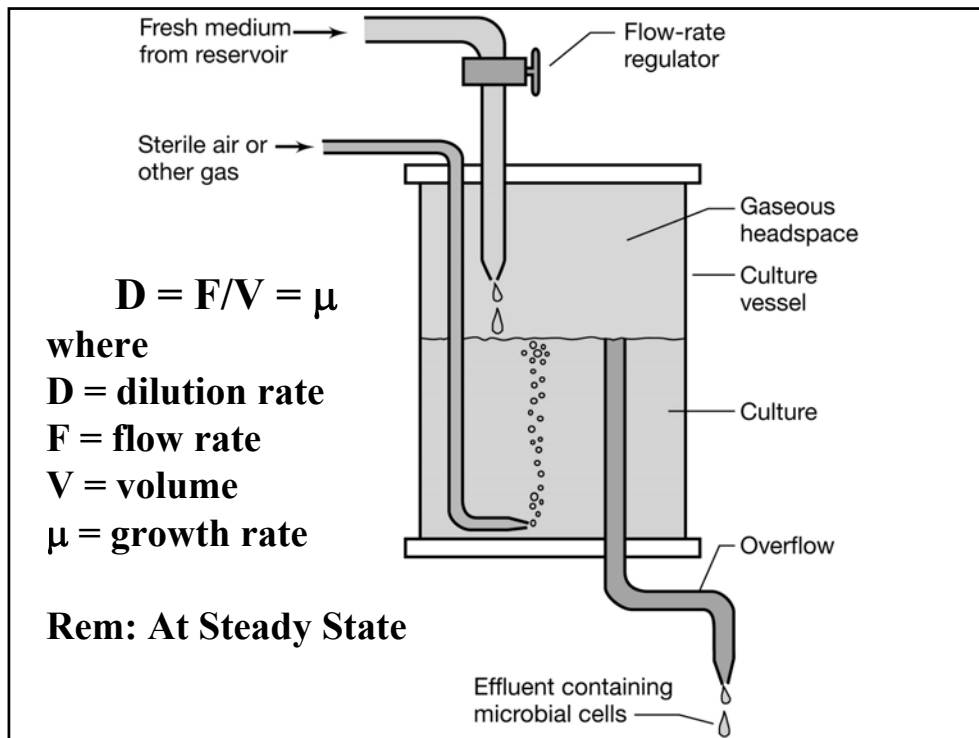
The Process of Growth

- Continuous Culture: The wonders of the Chemostat
Steady State
Reproducible Physiology
Fine control

Key parameters – K_s , μ_{max} , Yield

Closed systems vs. Open systems vs. Nature!
(Batch) (Chemostat)





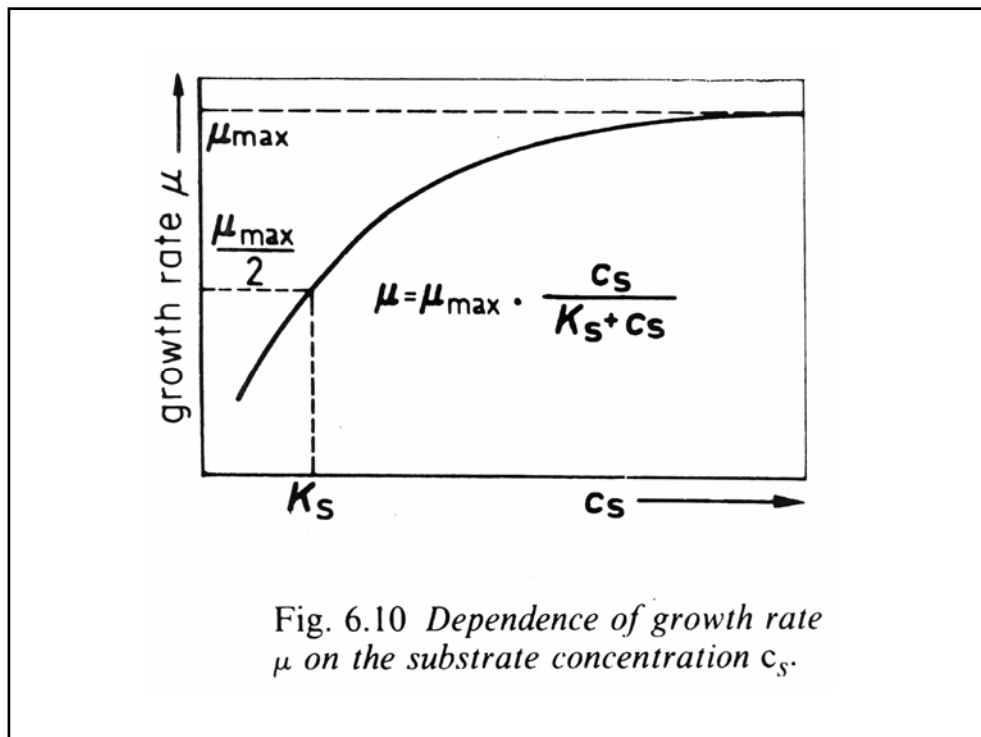
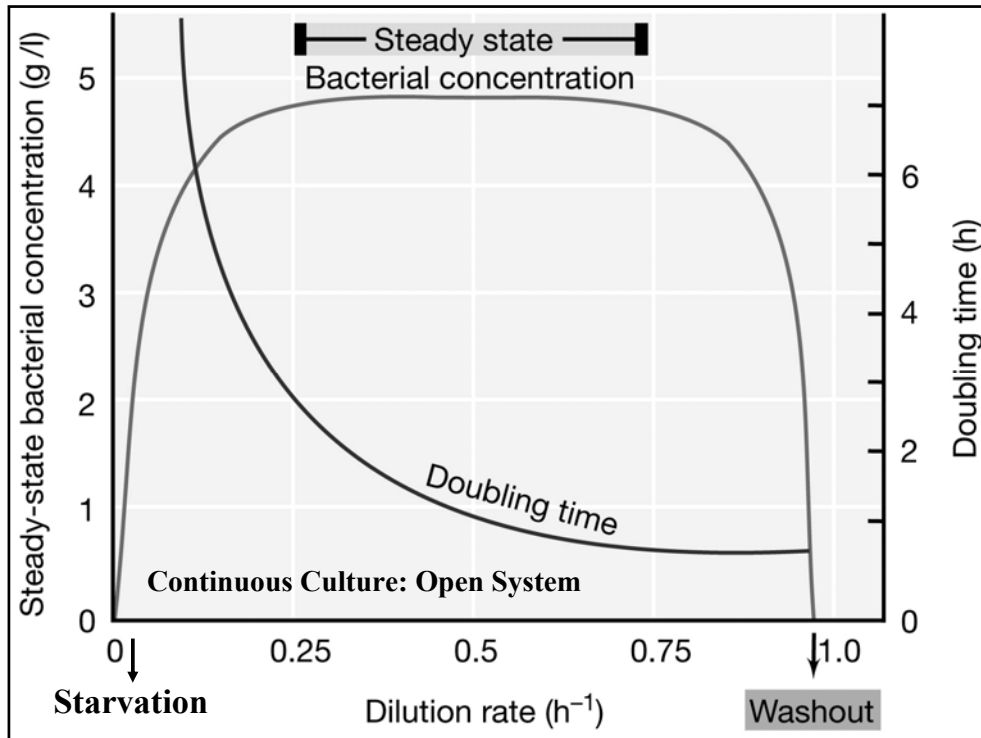


Fig. 6.10 Dependence of growth rate μ on the substrate concentration c_s .

Steady-state relationship between substrate concentration and output of bacterial mass

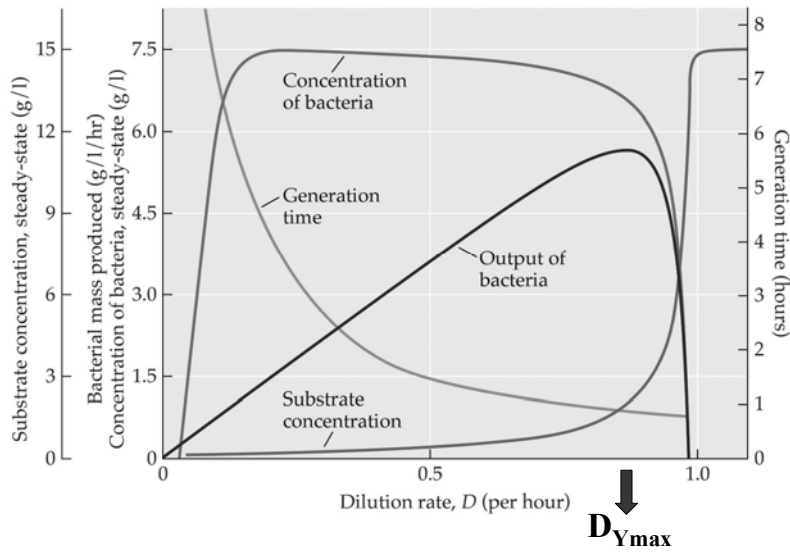


Table 6.2 Growth yields of anaerobic bacteria utilizing glucose as the energy source

	Mol ATP/Mol Glucose	y_{max} (g of cell/mol Glucose)	y_{ATP} (g of cell/mol ATP)
<i>Lactobacillus delbrueckii</i> ^a	2	21	10.5
<i>Enterococcus faecalis</i> ^a	2	20	10
<i>Zymomonas mobilis</i> ^b	1	9	9

^aHomolactic fermentation, Embden–Meyerhof pathway (see Chapter 10).

^bAlcoholic fermentation, Entner–Doudoroff pathway (see Chapter 10).