

# Population balance modeling -an application in particle technology

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Link to Matlab:

[www.glue.umd.edu/~sehrman/popbal.htm](http://www.glue.umd.edu/~sehrman/popbal.htm)

# Outline

- Aerosol reactors in industry
- Design problem, sneak peak
- Particle collection using cyclones
- Aerosol topics
  - size distributions
  - aggregates and fractals
  - coagulation/breakup
- Population balance equations
- Discrete and sectional approach
- Design problem, revisited

# Acknowledgments

- Population balance lectures and design problems developed in collaboration with Dr. R. Bertrum Diemer, Principal Consultant, Chemical Reaction Engineering, DuPont, Wilmington DE, USA
- Matlab code: Brendan Hoffman, Kelly Tipton, Yechun Wang, Matt McHale, Spring 2003 students
- Support from US National Science Foundation & DuPont
- Purpose
  - increase interest in field of particle technology
  - show practical application of population balance modeling approach

# Important aerosol products

- Silica
- Titania
- Carbon black
- Specialty materials
  - Nano zinc oxide used today in sunscreen
  - High surface area catalyst supports (alumina, zirconia, etc..)
  - Chemical mechanical polishing agents (ceria, silica, etc... )

# General Aerosol Process Schematic

Feed #1  
Preparation

Feed #2  
Preparation

.

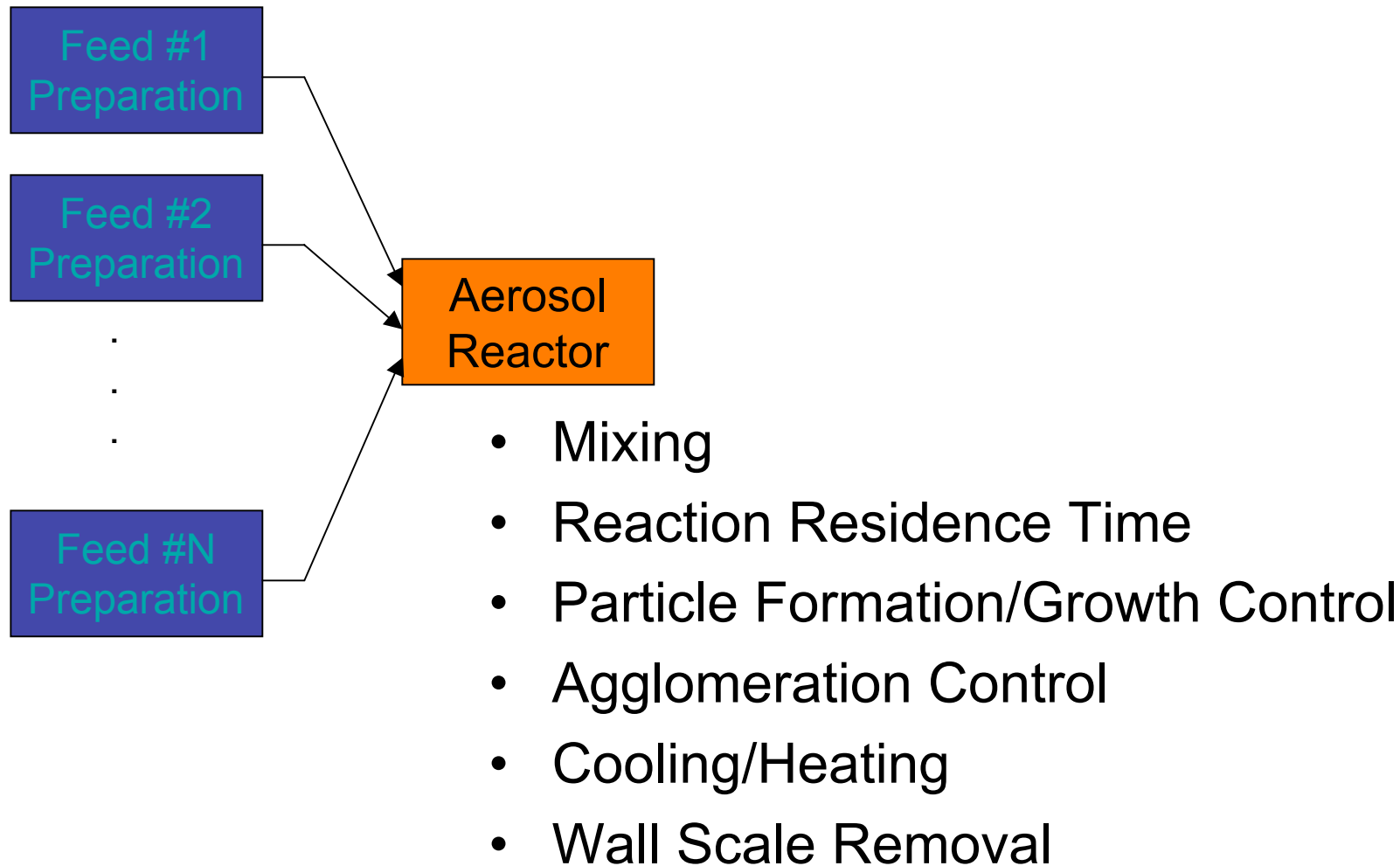
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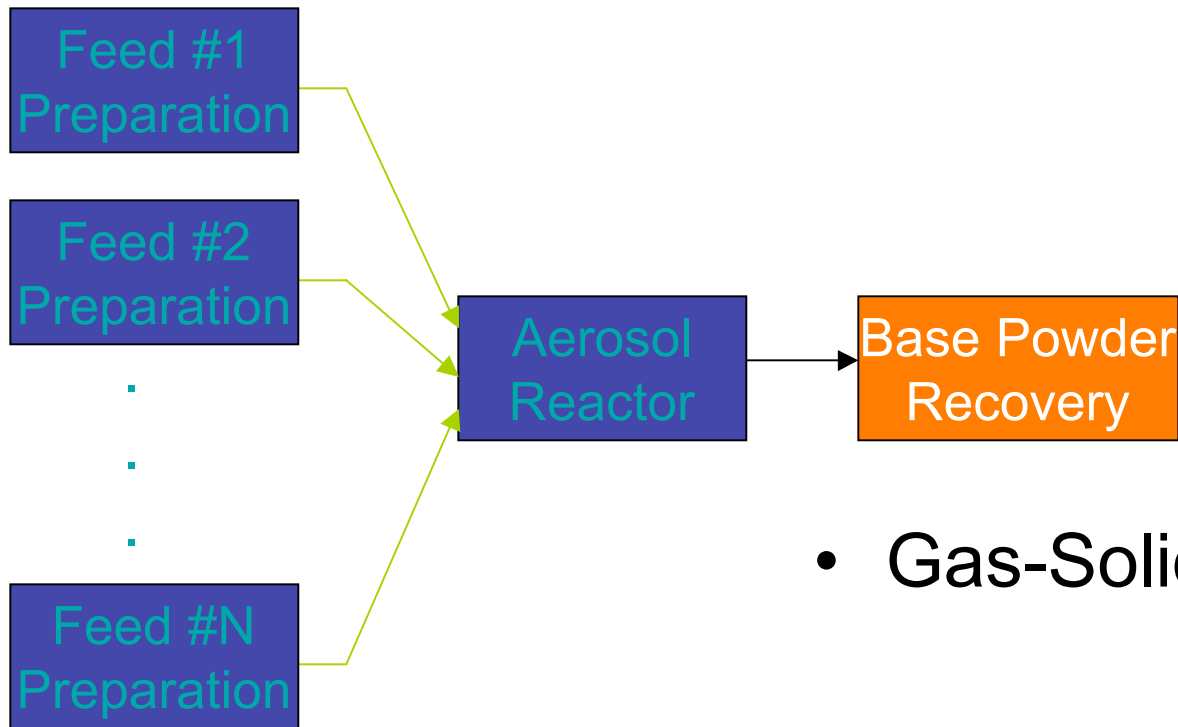
Feed #N  
Preparation

- Vaporization
- Pumping/Compression
- Addition of additives
- Preheating

# General Aerosol Process Schematic

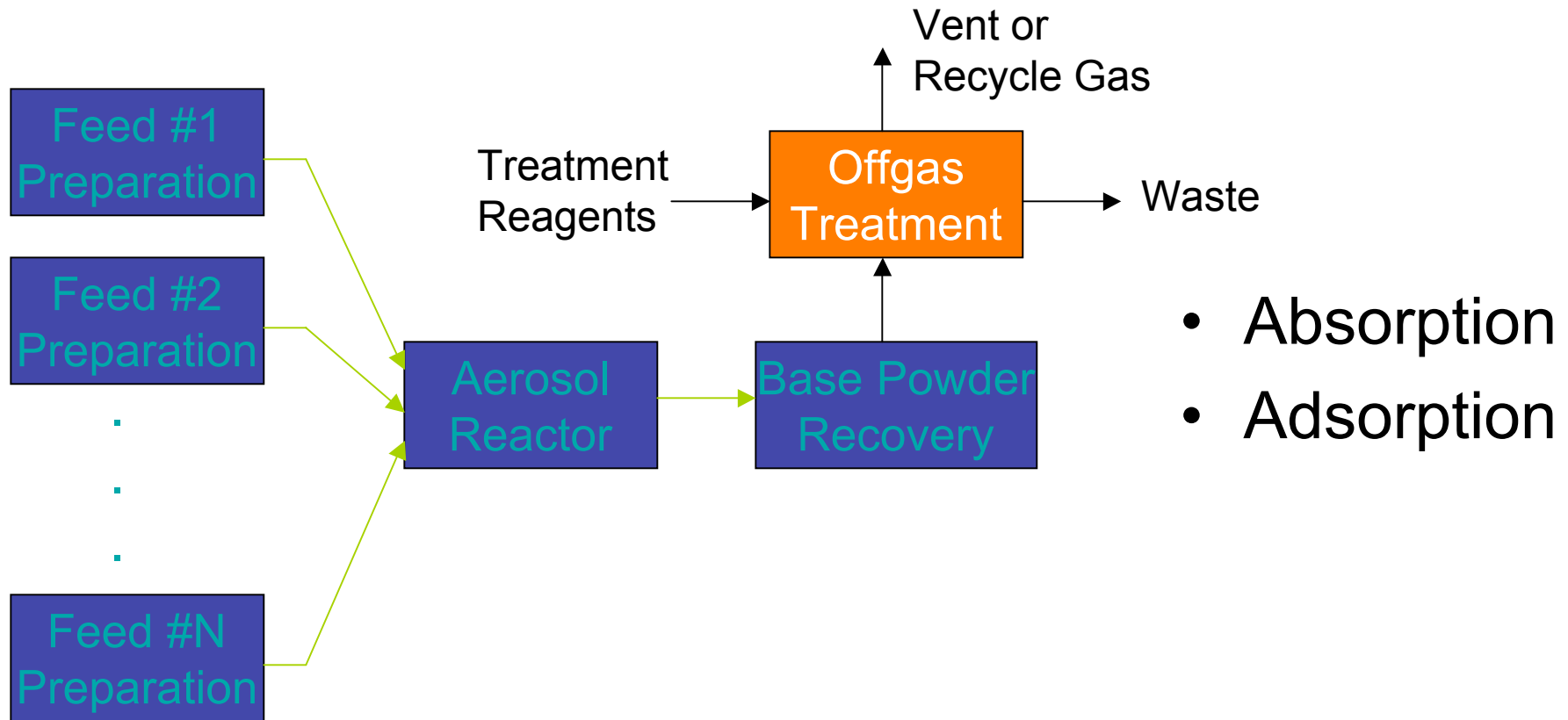


# General Aerosol Process Schematic



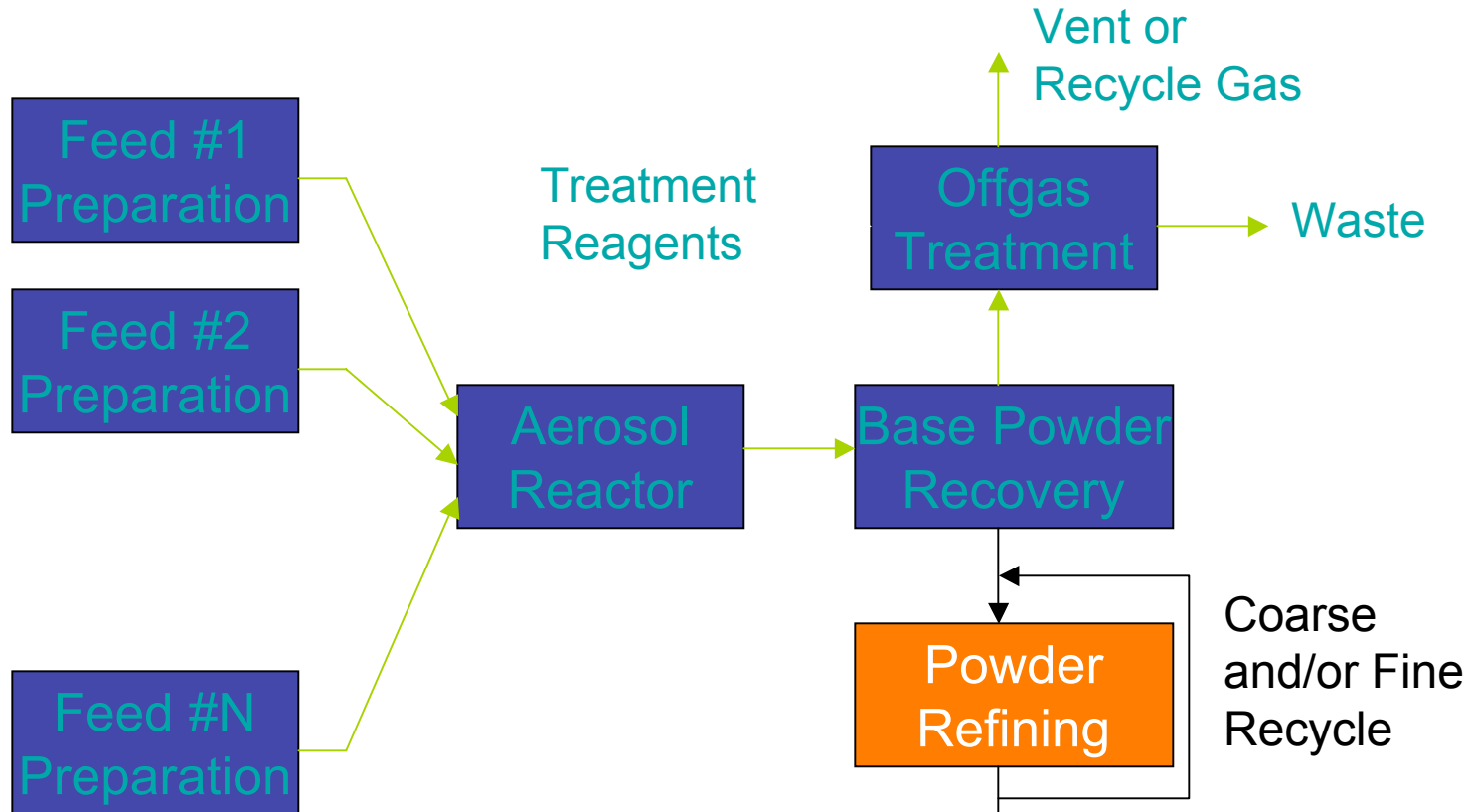
- Gas-Solid Separation

# General Aerosol Process Schematic





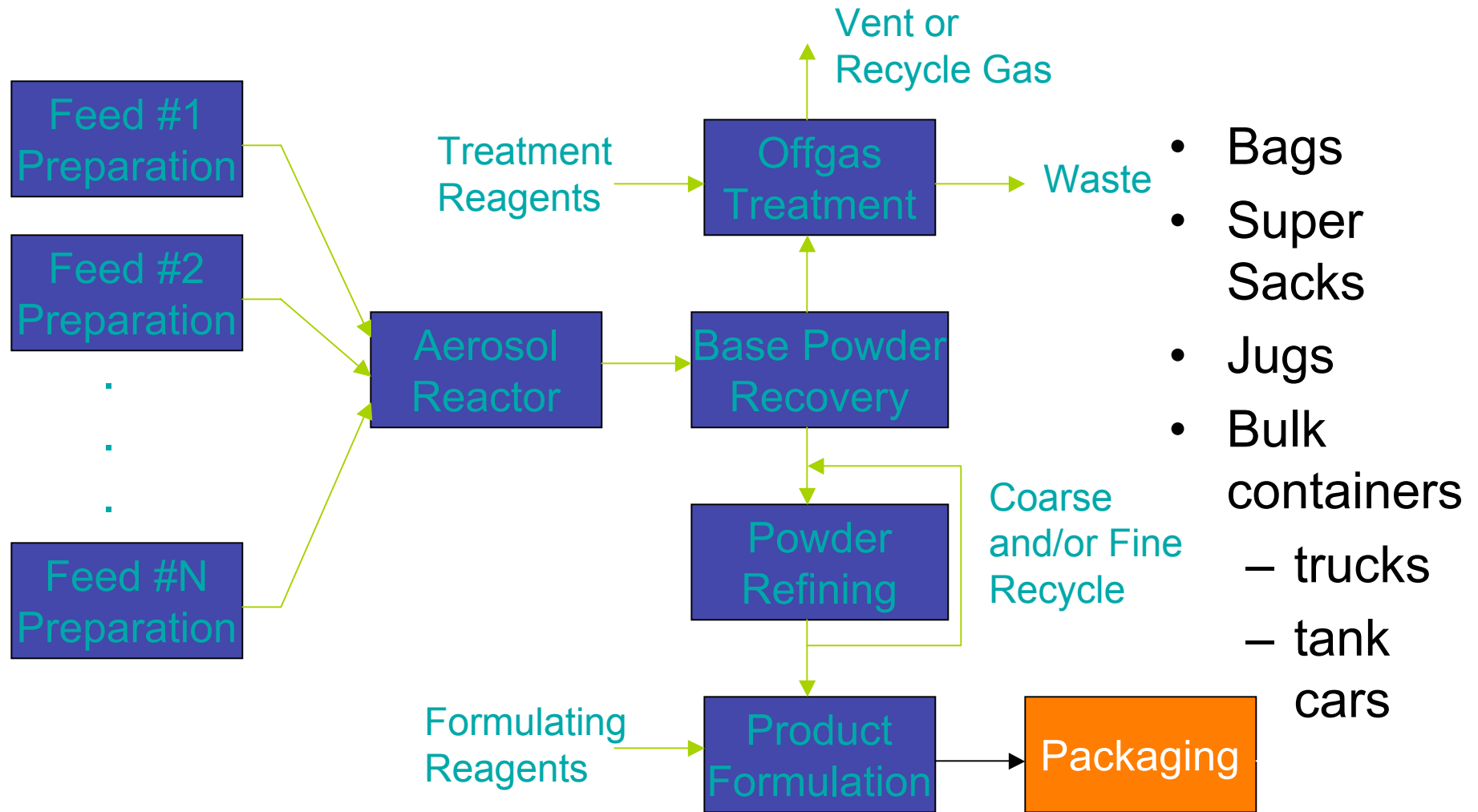
# General Aerosol Process Schematic



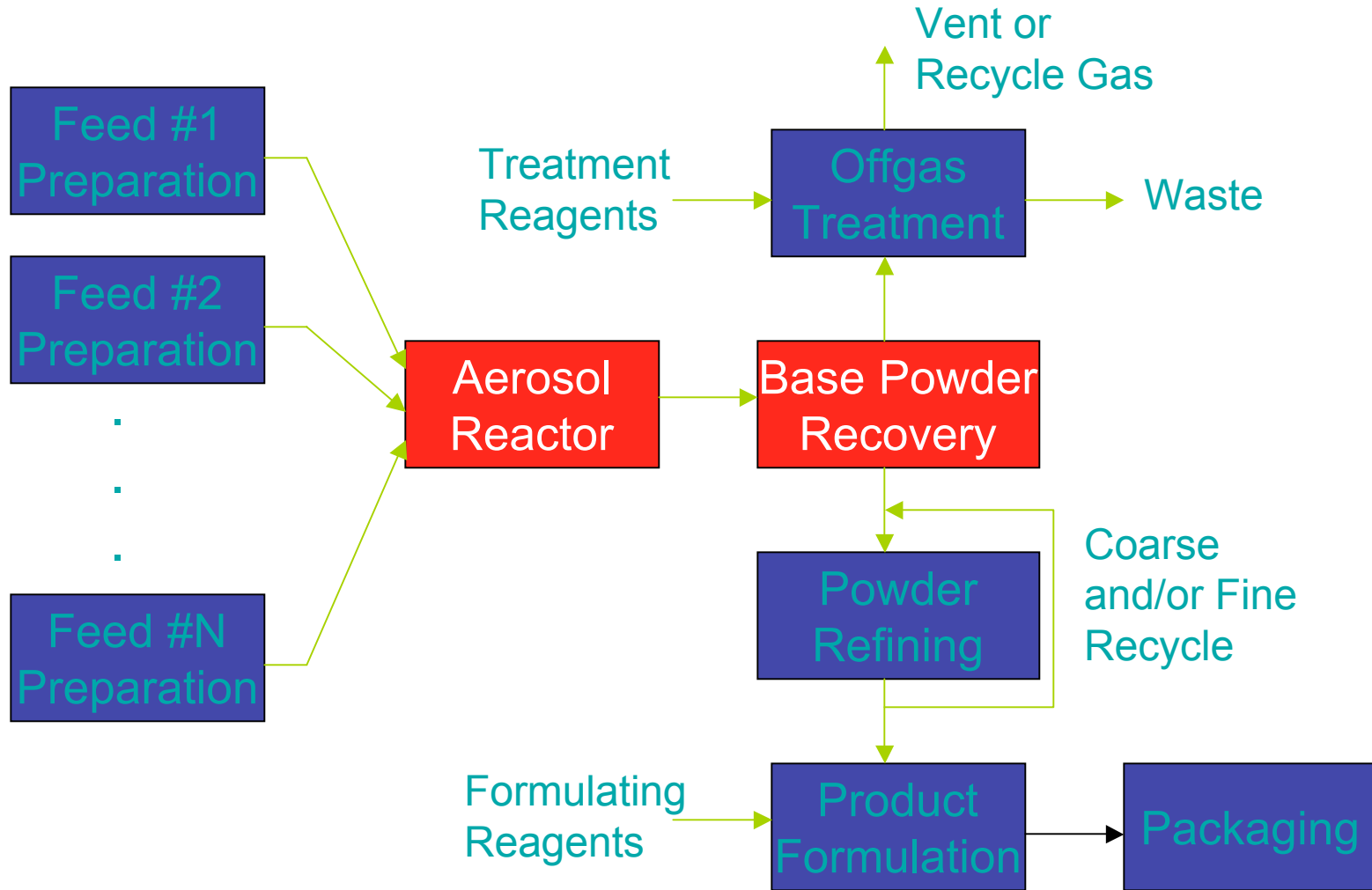
- Degassing
- Desorption
- Conveying
- Size Modification
- Solid Separations



# General Aerosol Process Schematic

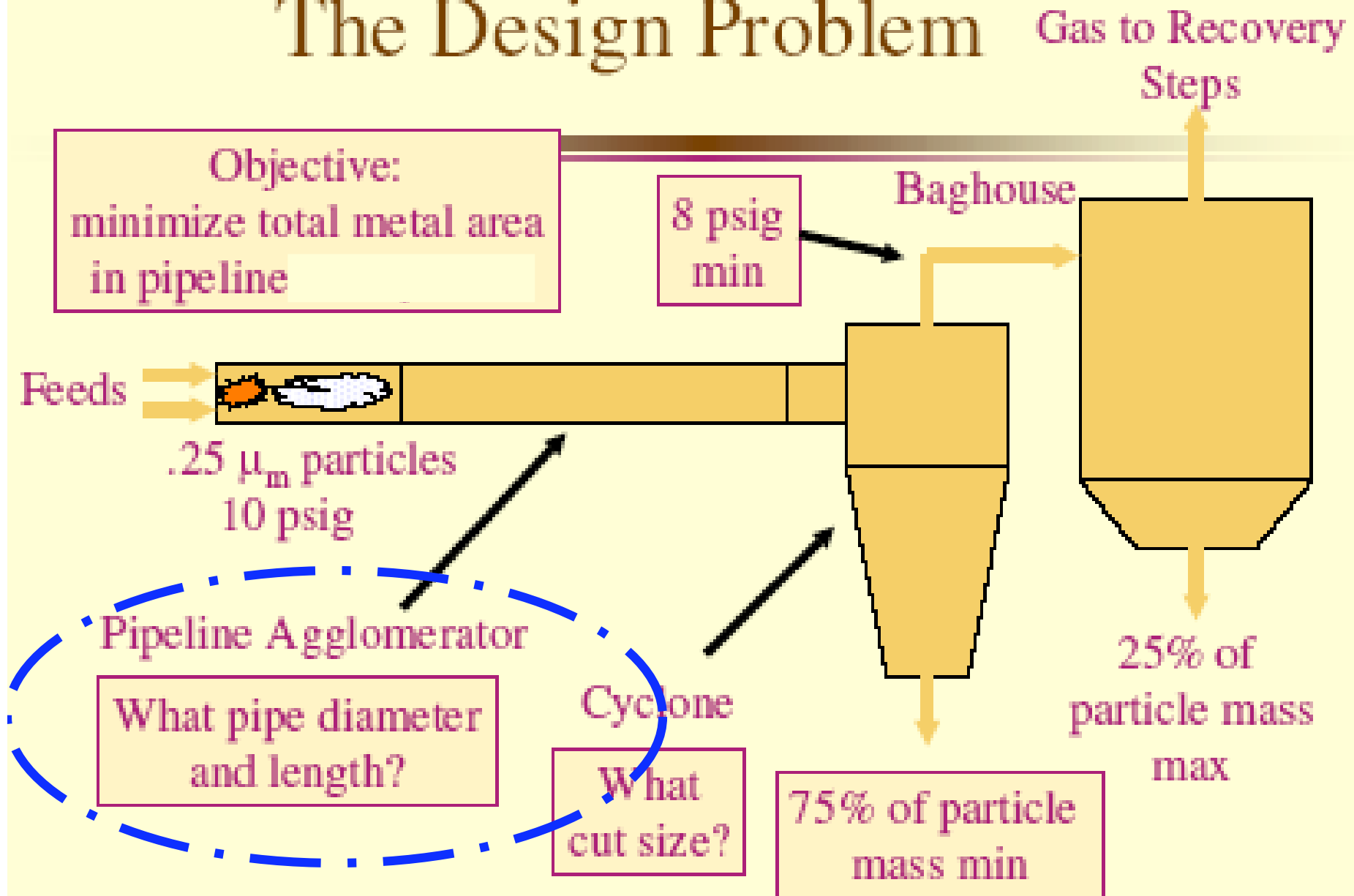


# General Aerosol Process Schematic

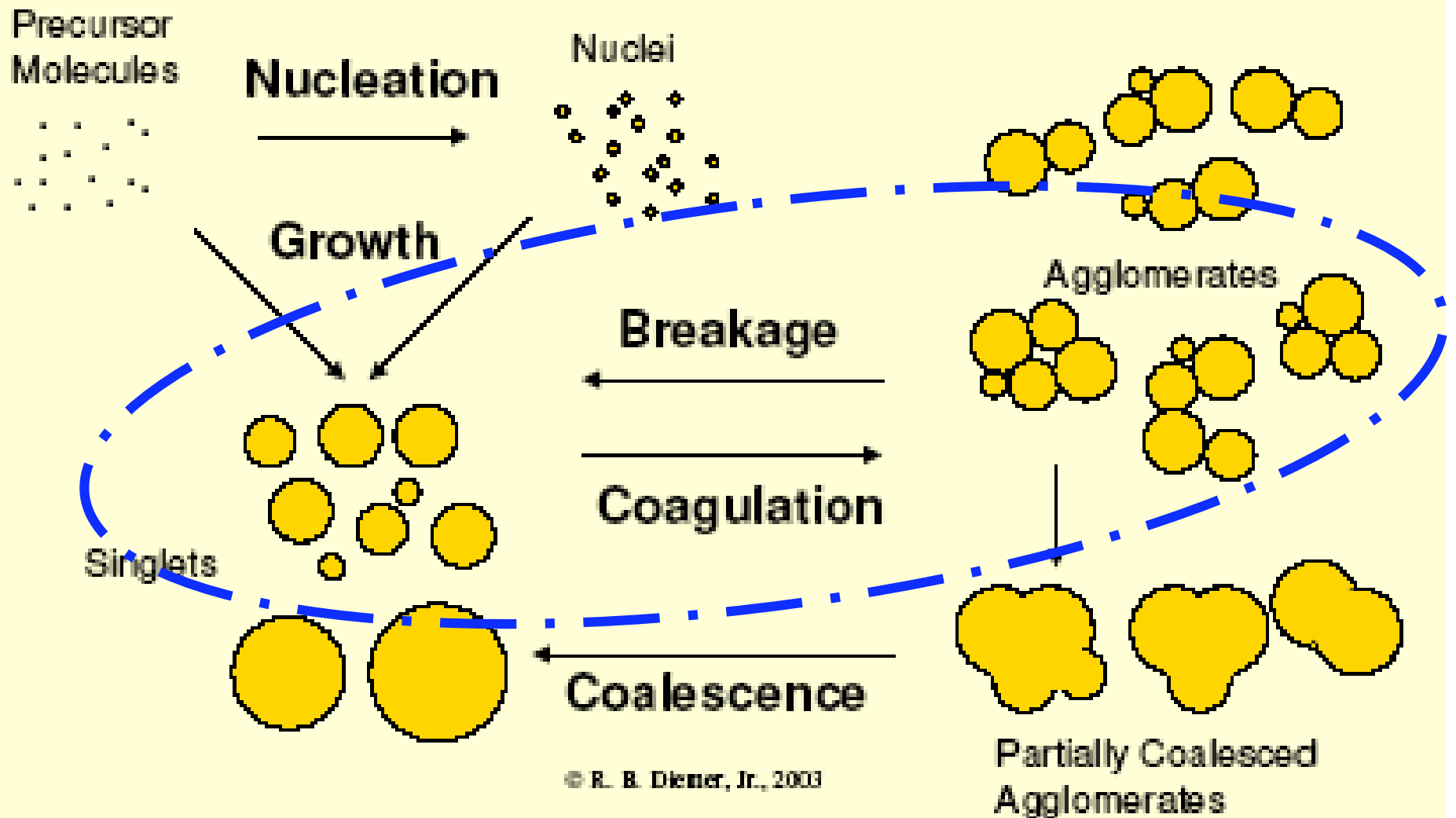


**Our focus, reactor and gas solid separation**

# The Design Problem



# Particle Formation, Growth & Transformation



# Thermal Carbon Black Process

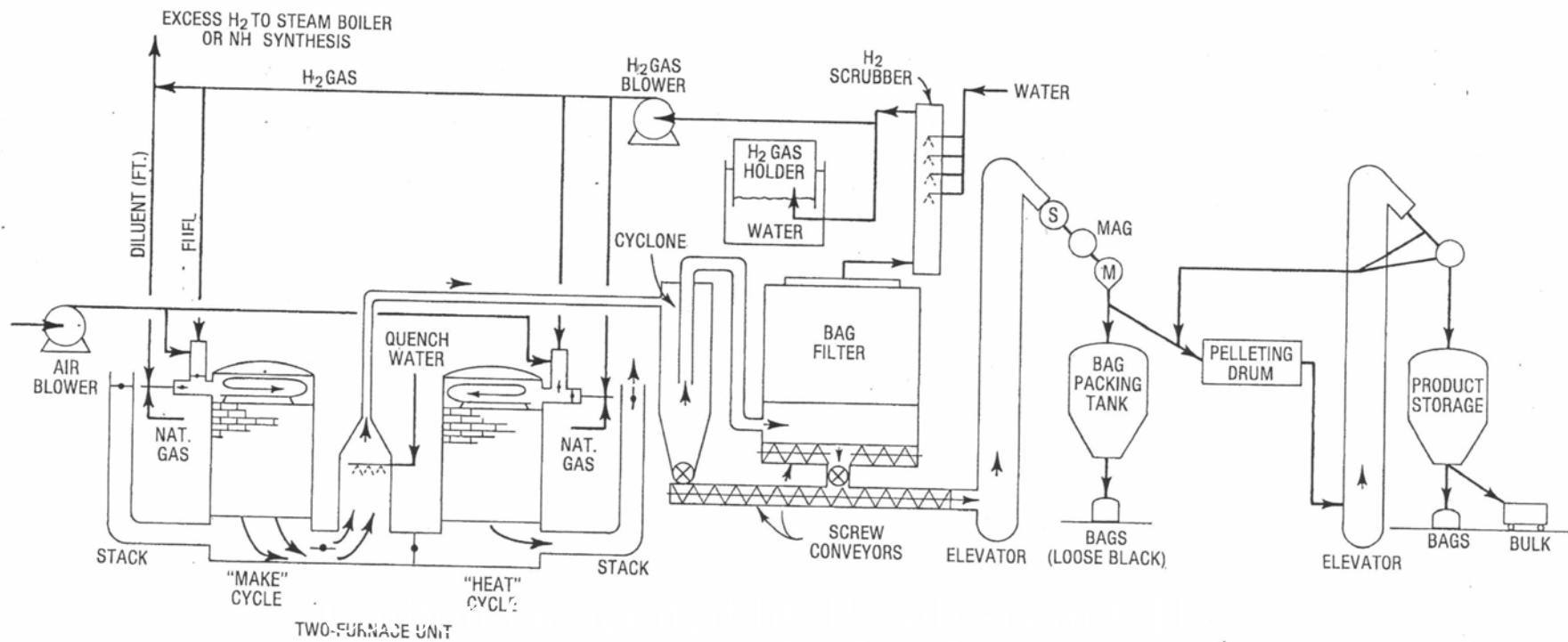


FIG. 26. Thermal process (natural gas feedstock).

Johnson, P. H., and Eberline, C. R., "Carbon Black, Furnace Black", *Encyclopedia of Chemical Processing and Design*, J. J. McKetta, ed., Vol. 6, Marcel Dekker, 1978, pp. 187-257.

# Furnace Carbon Black Process

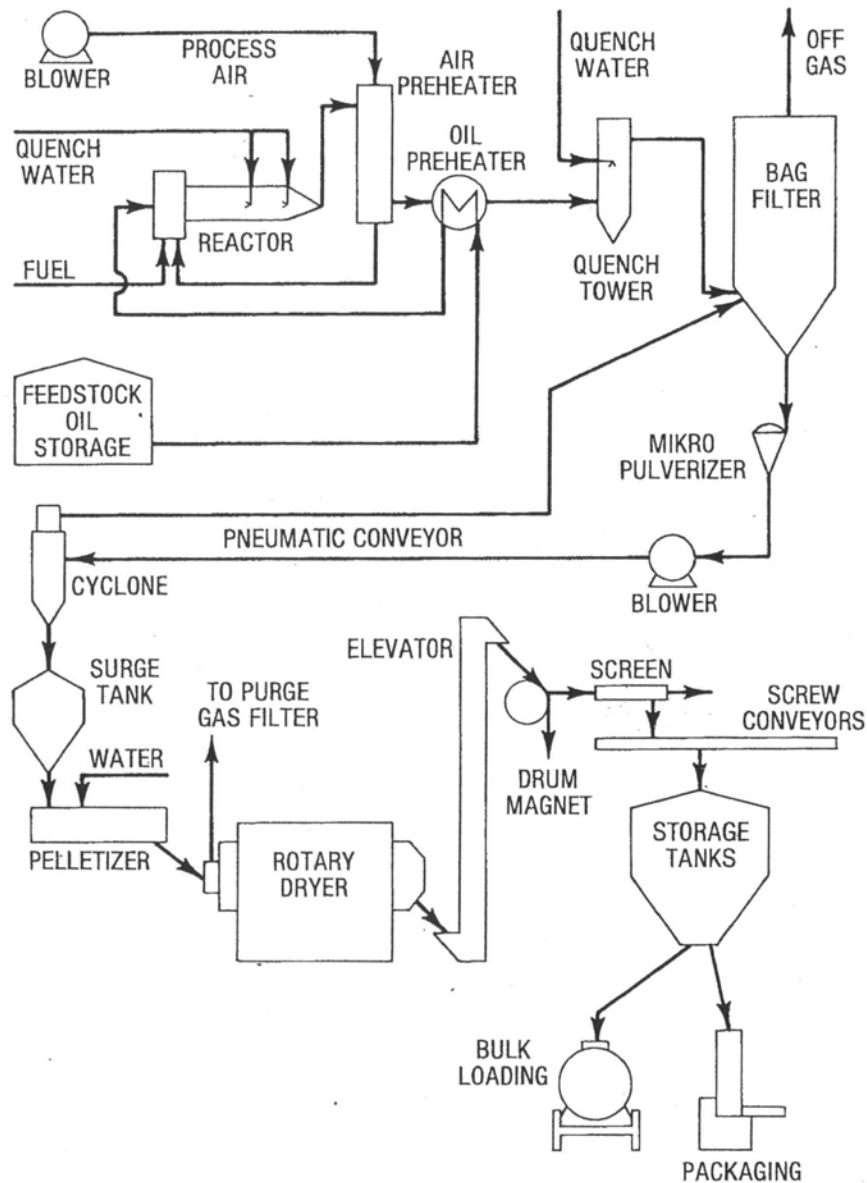


FIG. 18. Oil furnace carbon black process.

Johnson, P. H., and Eberline, C. R.,  
“Carbon Black, Furnace Black”,  
*Encyclopedia of Chemical  
Processing and Design*, J. J.  
McKetta, ed., Vol. 6, Marcel Dekker,  
1978, pp. 187-257.



# Thermal Carbon Black

- ~200 nm Primary Particles
- 4-6 Primaries/  
Agglomerate

Johnson, P. H., and Eberline, C. R., "Carbon Black, Furnace Black", *Encyclopedia of Chemical Processing and Design*, J. J. McKetta, ed., Vol. 6, Marcel Dekker, 1978, pp. 187-257.

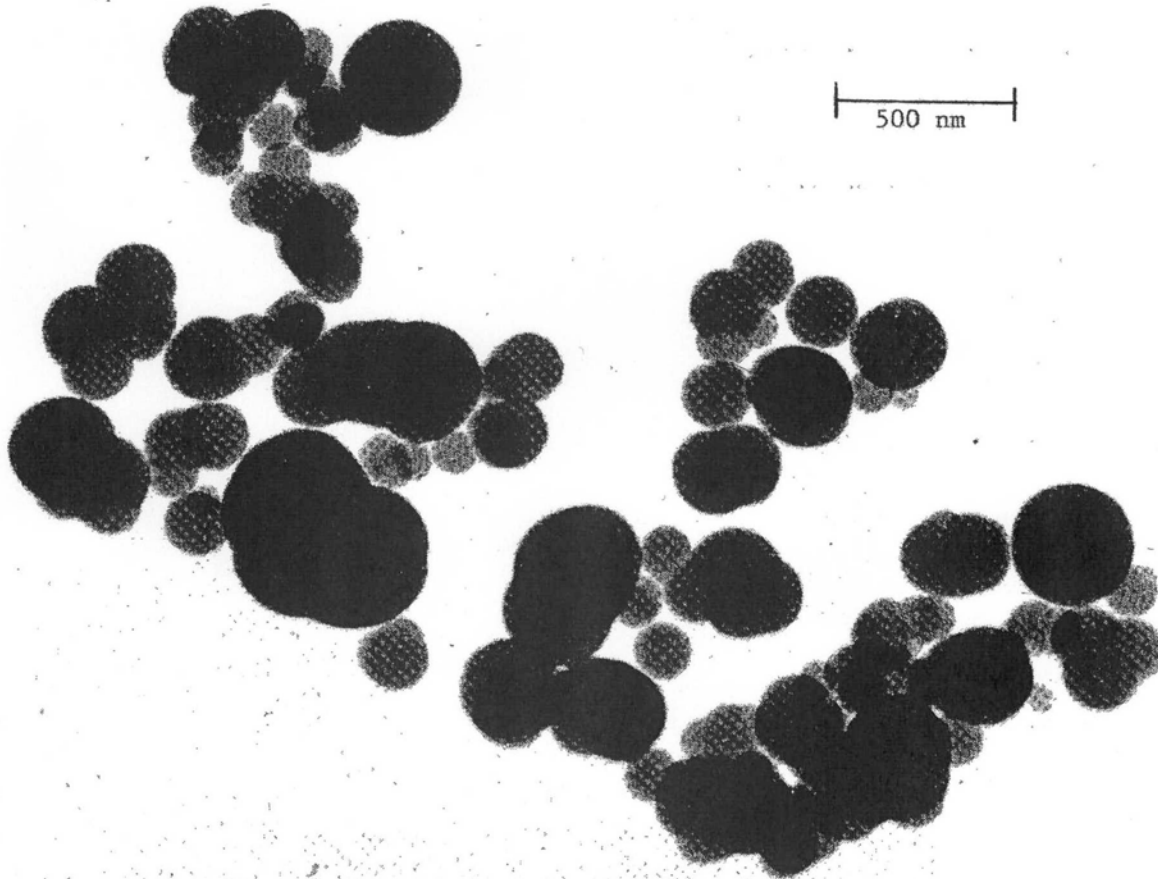


FIG. 7. Thermal black.

# Oil-Furnace Carbon Black

- ~20 nm Primary Particles
- 10-50 Primaries/  
Agglomerate

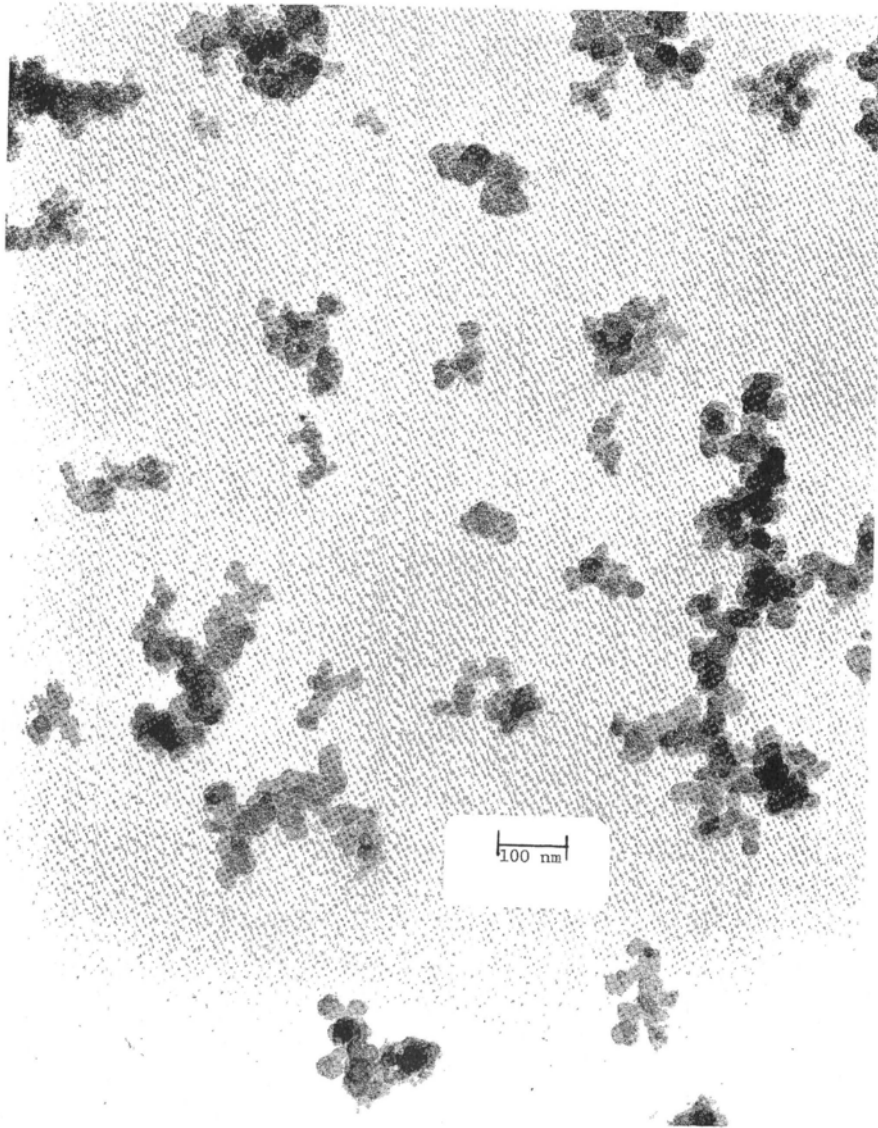


FIG. 8. A structured oil-furnace black.

Johnson, P. H., and Eberline, C. R., "Carbon Black, Furnace Black", *Encyclopedia of Chemical Processing and Design*, J. J. McKetta, ed., Vol. 6, Marcel Dekker, 1978, pp. 187-257.

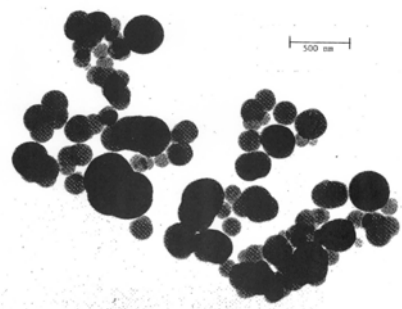


FIG. 7. Thermal black.

## What do you see?

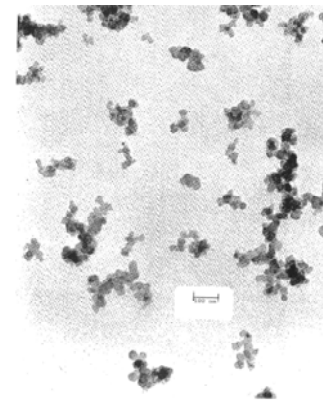


FIG. 8. A different carbon black.

- What do you notice about each sample?
- How do the two images differ?

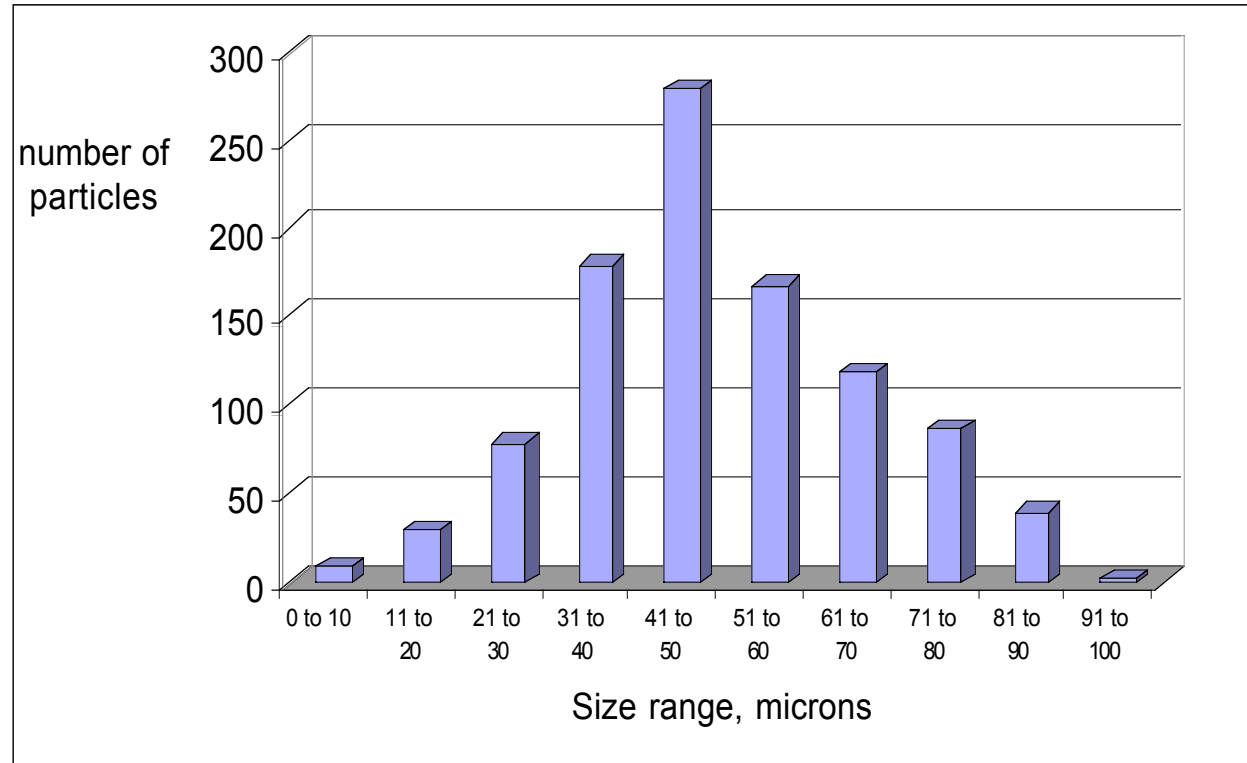
# Particles not all same...

- Diameter
- Volume
- Surface area
- Structure

# How to represent size distributions, histogram example:

Size range, microns	number of particles
0 to 10	10
11 to 20	30
21 to 30	80
31 to 40	180
41 to 50	280
51 to 60	169
61 to 70	120
71 to 80	88
81 to 90	40
91 to 100	3

## Number of particles vs particle diameter



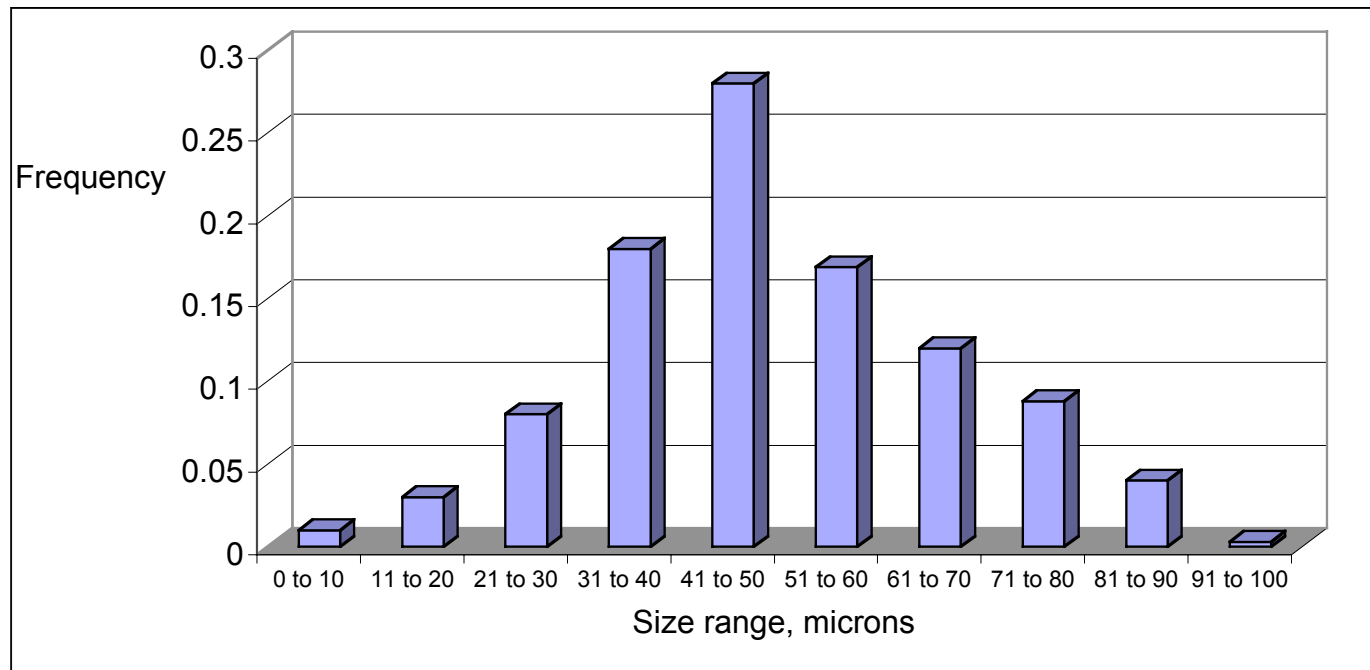
Can create histogram from raw particle size data using Analysis tool pack add-in, with Excel.. After add-in, go to 'tools', then 'data analysis', then 'histogram'.

# How to represent size distributions, histogram example:

Size range, microns	number of pa	Frequency
0 to 10	10	0.01
11 to 20	30	0.03
21 to 30	80	0.08
31 to 40	180	0.18
41 to 50	280	0.28
51 to 60	169	0.169
61 to 70	120	0.12
71 to 80	88	0.088
81 to 90	40	0.04
91 to 100	3	0.003

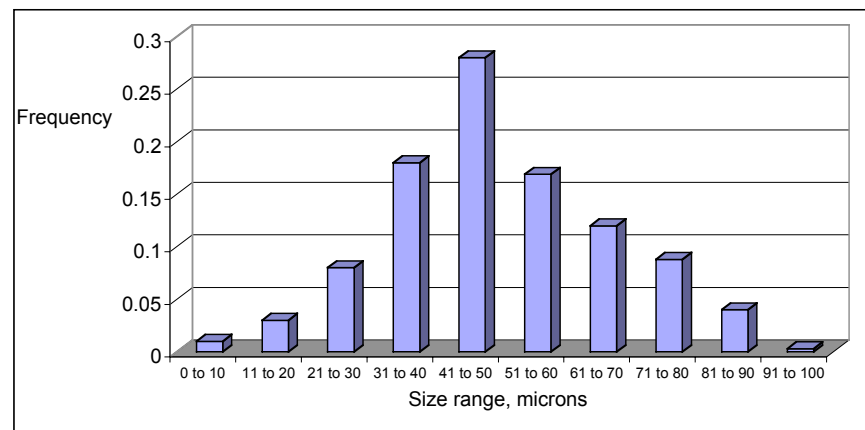
total number 1000

## Number frequency vs particle diameter



# Test yourself...

- If you change the frequency distribution so that is based on the MASS of particles in each size range versus the NUMBER, will the shape of the frequency distribution change?

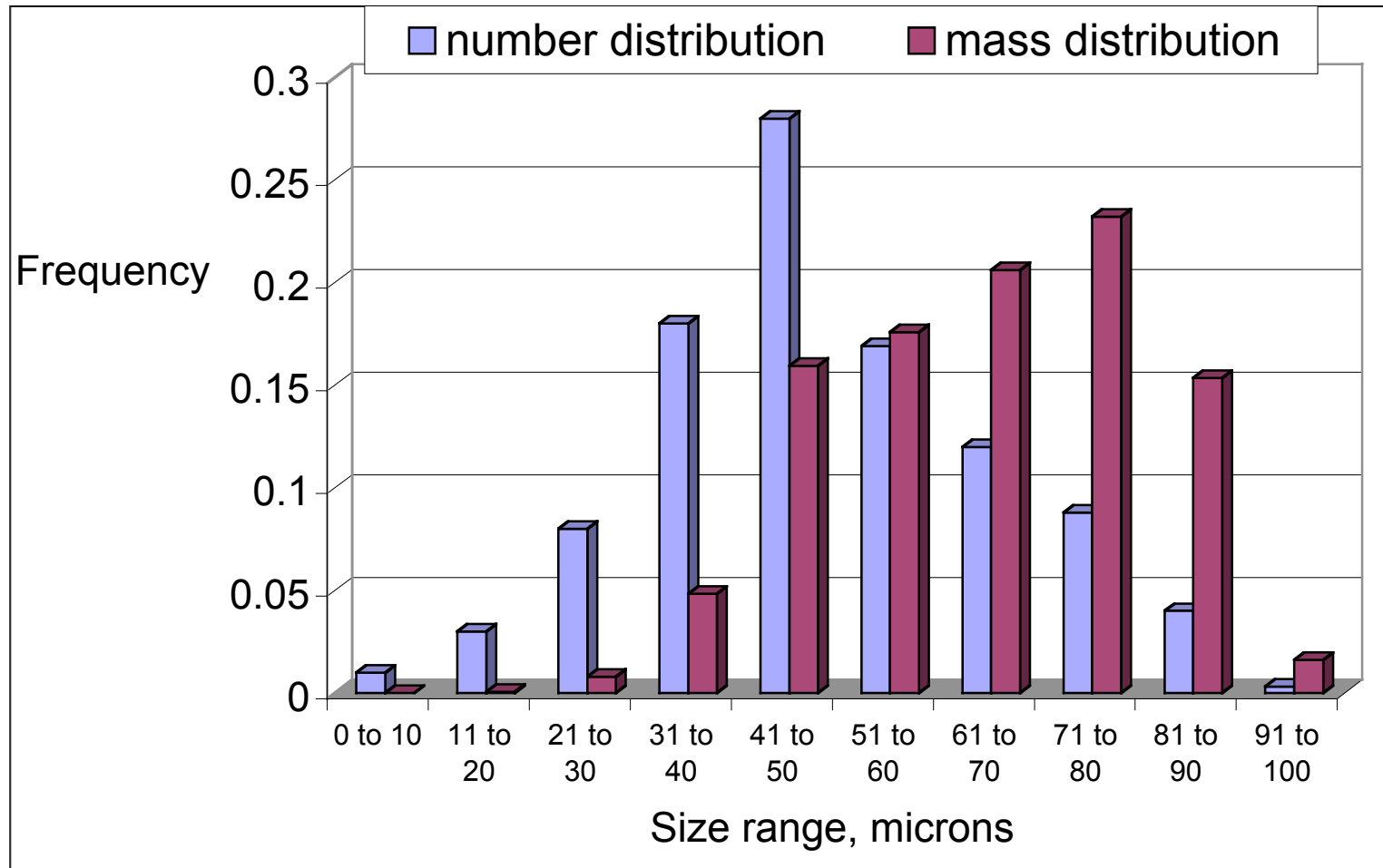


# More spreadsheet manipulations

Size range, microns	number of particles	Frequency	assume diameter	mass of particles in each bin, g	Mass frequency
0 to 10	10	0.01	5	6.54E-10	0.00
11 to 20	30	0.03	15	5.30E-08	0.00
21 to 30	80	0.08	25	6.54E-07	0.01
31 to 40	180	0.18	35	4.04E-06	0.05
41 to 50	280	0.28	45	1.34E-05	0.16
51 to 60	169	0.169	55	1.47E-05	0.18
61 to 70	120	0.12	65	1.72E-05	0.21
71 to 80	88	0.088	75	1.94E-05	0.23
81 to 90	40	0.04	85	1.29E-05	0.15
91 to 100	3	0.003	95	1.35E-06	0.02
total number	1000				
total mass assuming 1g/cc density				8.37E-05	



# Number vs mass distribution



# Continuous distributions

Also useful: continuous distributions, where some function,  $n_d$ , describes the number of particles of some diameter  $d_p$  at a given point, at a given time.

In terms of number concentration:

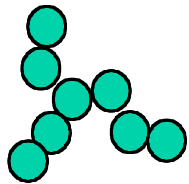
Let  $dN$  = number of particles per unit volume of gas at a given position in space (represented by position vector  $\mathbf{r}$ ), at a given time ( $t$ ), in the particle range  $d_p$  to  $d_p + d(d_p)$ .  $N$  = total number of particles per unit volume of gas at a given position in space at a given time. Size distribution function is defined as:

$$n_d(d_p, \mathbf{r}, t) = \frac{dN}{d(d_p)}$$

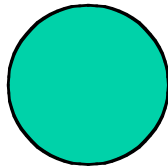
Can also have size distribution function,  $n$ , with particle volume  $v$  as size parameter:  $n(v, \mathbf{r}, t) = \frac{dN}{dv}$

# Aggregates of hard spheres

- When primary particles collide and stick, but do not coalesce, irregular structures are formed



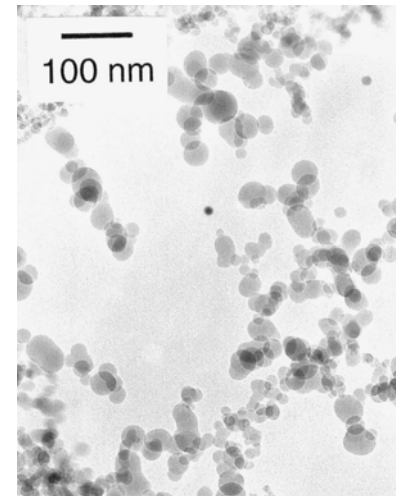
agglomerate



spherical equivalent

how should  
these structures  
be characterized?

- Radius gives space taken up, but no information about mass/actual volume. Using only actual volume doesn't indicate how much space it takes up.
- Real flame generated aerosol:



# Concept of fractal dimension

- Aerosol particles which consist of agglomerates of 'primary particles', (often, combustion generated) may be described using the concept of fractals.
- Fractals - The relationship between diameter of aerosol agglomerates, and the volume of primary particles in the agglomerate can be written:

$$\frac{v}{v_o} = \left( \frac{d}{d_o} \right)^{D_f} \quad \text{where } v_o = \frac{\pi}{6} d_o^3$$

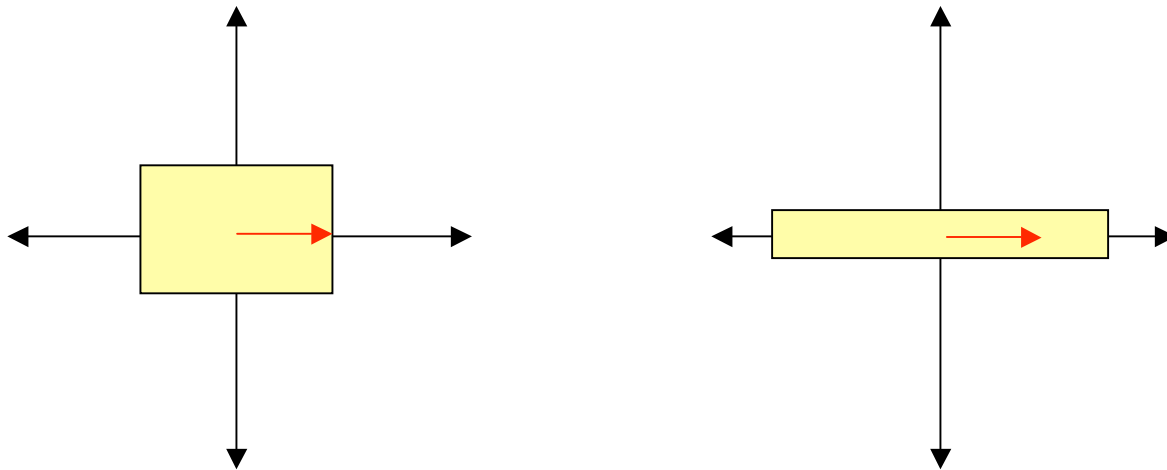
is the volume of the primary particle of diameter  $d_o$

- $d$  typically based on  $2 * \text{radius of gyration}$

# Radius of Gyration

“The Radius of Gyration of an Area about a given axis is a distance  $k$  from the axis. At this distance  $k$  an equivalent area is thought of as a **Line Area** parallel to the original axis. The moment of inertia of this **Line Area** about the original axis is unchanged.”

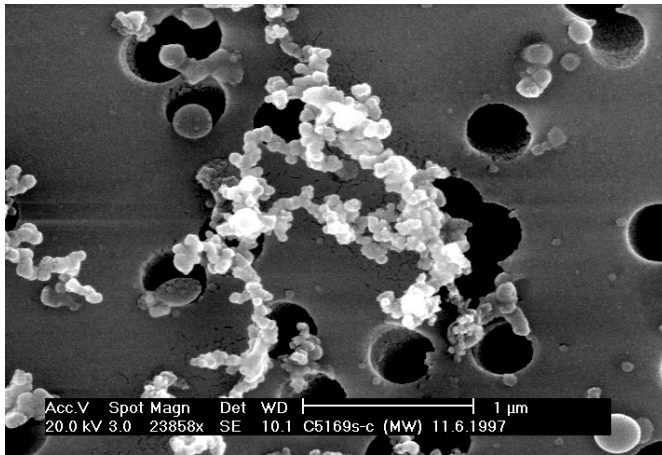
<http://www.efunda.com/math/areas/RadiusOfGyrationDef.cfm>



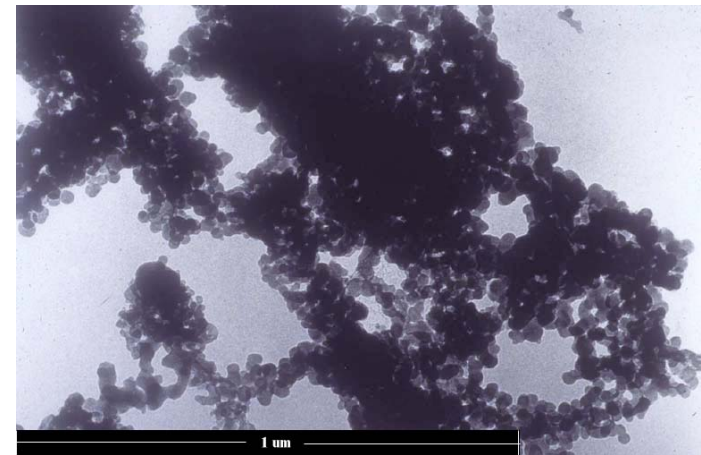
# Fractal dimension

- Fractals -  $D_f = 2$  = uniform density in a plane,  $D_f$  of 3 = uniform density in three dimensions
- Typical values for agglomerates ranges from 1.5 to near 3 depending on mechanism of agglomeration, possible rearrangement, and external field effects
- Small agglomerates (few particles) not really fractal but “fractal-like”
- For hundreds of particles, relationship holds well

# Test yourself...



<http://www.mpch-mainz.mpg.de/~gth/soot1.jpg>



[http://faculty.engineering.ucdavis.edu/jenkins/previous/August2002/16\\_23.jpg](http://faculty.engineering.ucdavis.edu/jenkins/previous/August2002/16_23.jpg)

Which picture shows particle agglomerates with a lower fractal dimension?

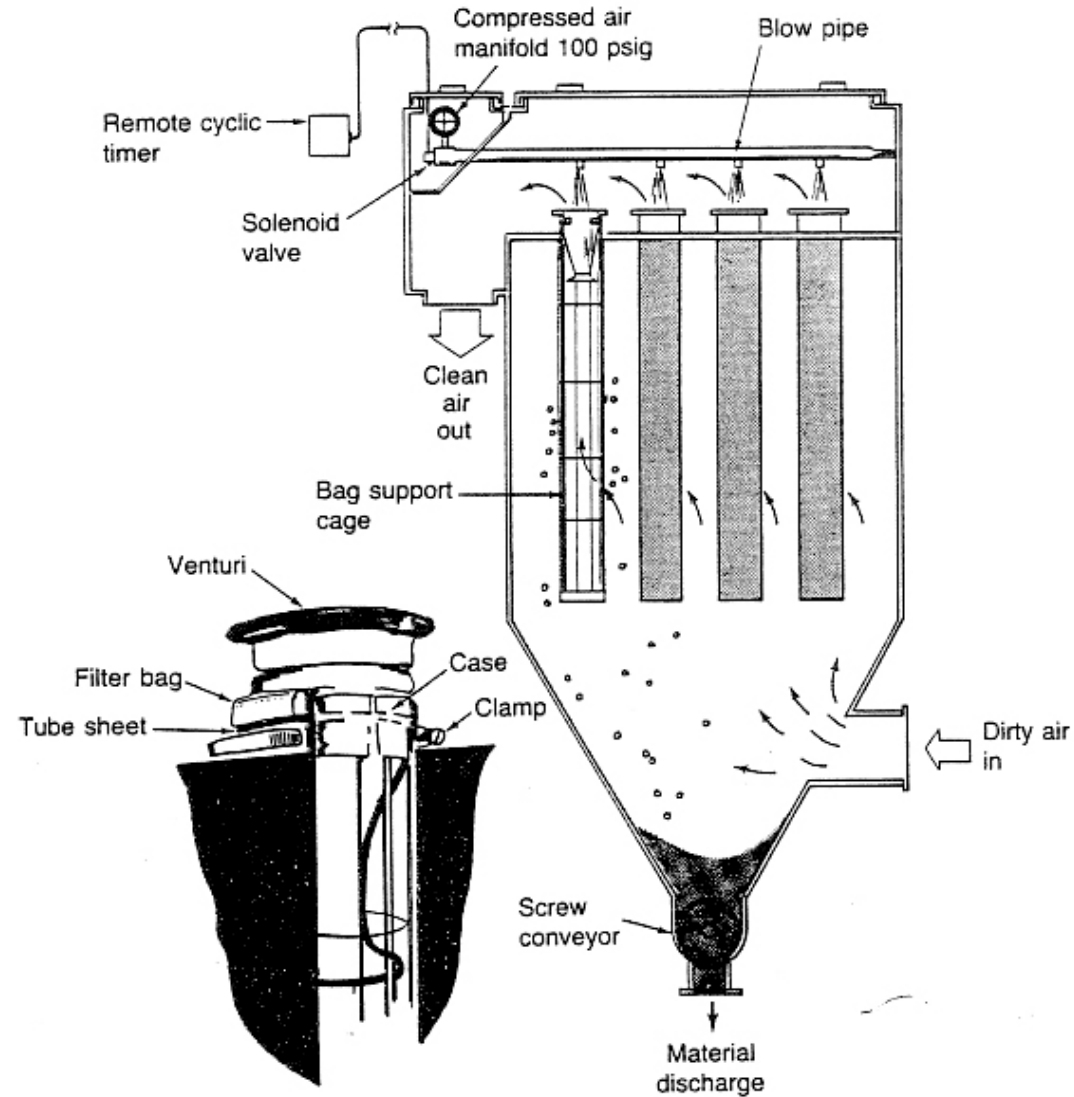
# Working backwards in the problem

- Particle gas separators
- Use different phenomena
  - Gravitational settlers
  - Filters (baghouse)
  - Scrubbers
  - Inertial separators (cyclone)
  - Electrostatic precipitators



# Pulse Jet Baghouse

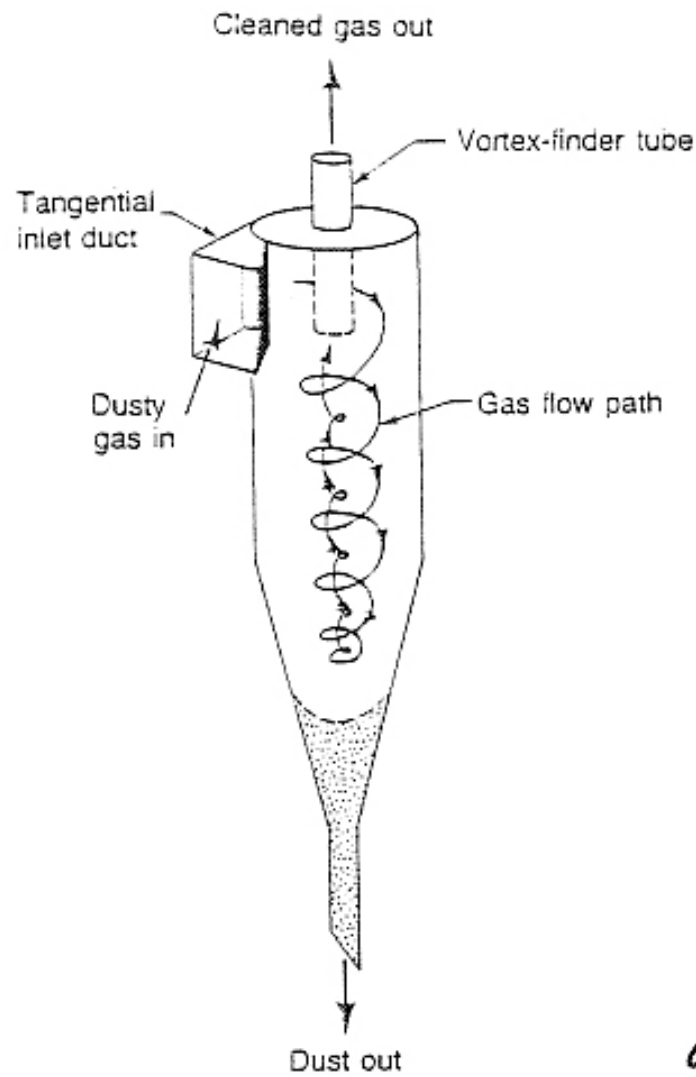
Figure 6.6 Schematic diagram of a pulse-jet baghouse.



Air Pollution Control: A Design Approach  
Cooper & Alley, PWS Eng. Boston 1986

# Standard Cyclone

Figure 4.1 Schematic flow diagram of a standard cyclone.



Air Pollution  
Control  
Cooper & Alley (1986)

Force balance on particle in cyclone

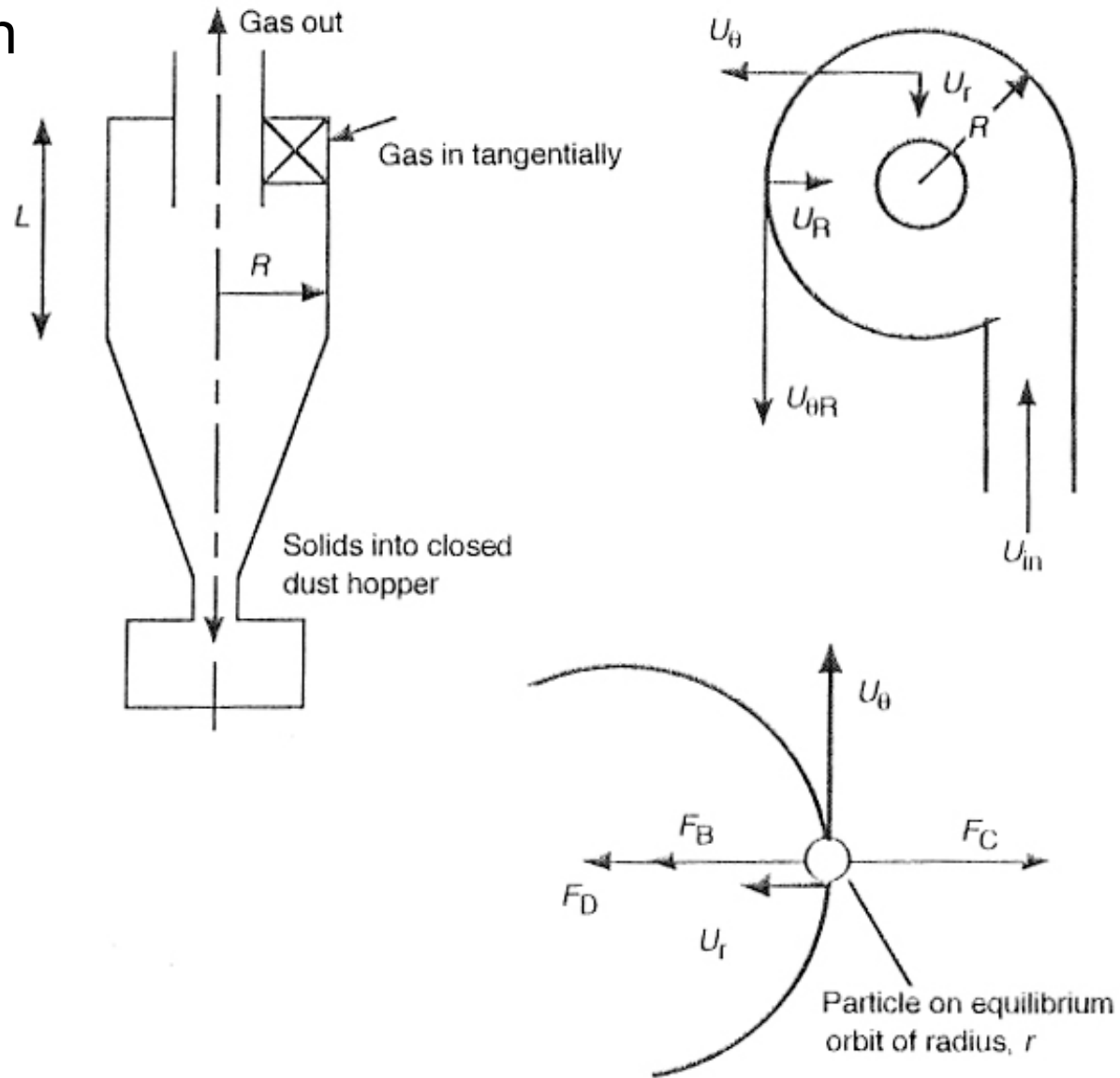
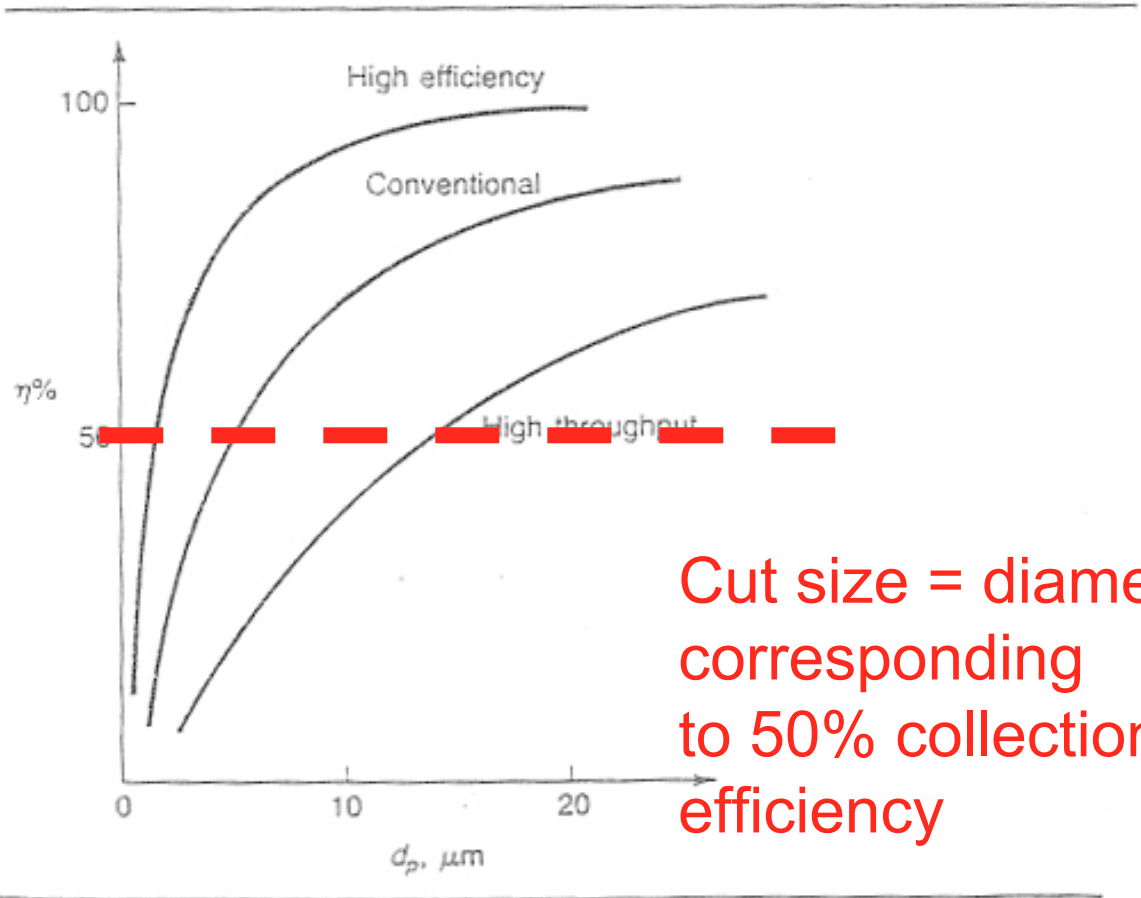


Figure 7.3 Reverse flow cyclone – a simple theory for separation efficiency  
Rhodes, Introduction to Particle Technology, Wiley, 1998

# Cyclone Efficiency as Function of Particle Diameter

Which type has lowest pressure drop?

Figure 4.3 General relationship of collection efficiency versus particle size for cyclones.



Cut size = diameter corresponding to 50% collection efficiency

NOTE: Efficiency versus size curves represent broad generalizations, not exact relationships.

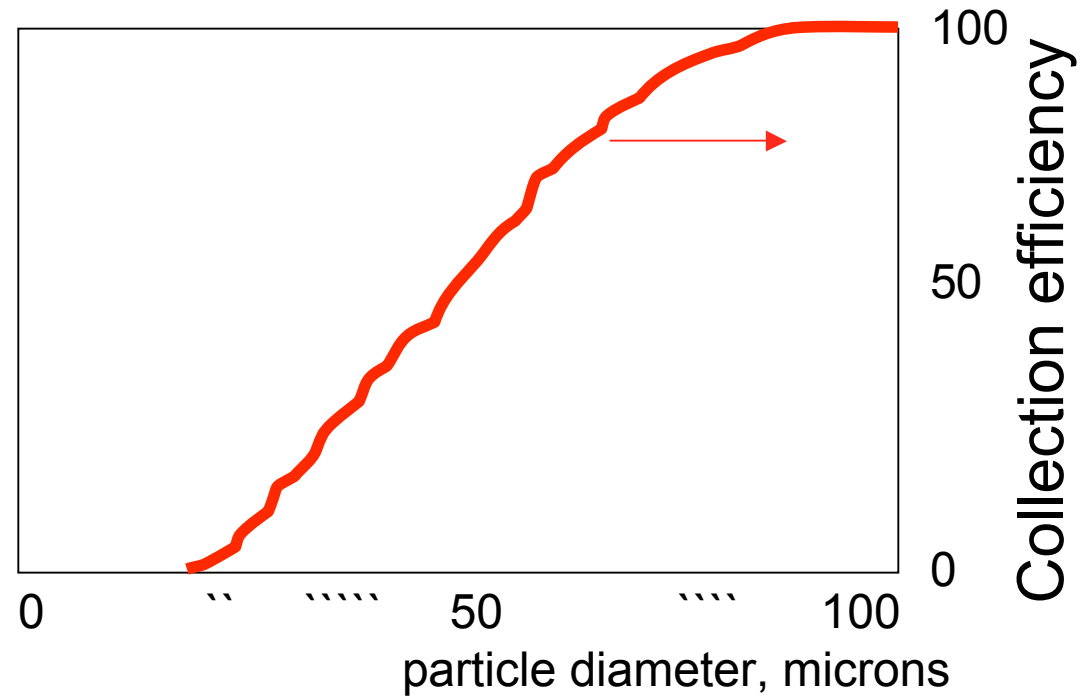
*Air Pollution Control: A design approach  
Cooper & Alley PWS Engineering, Boston—1986*

# Cyclones

- Advantages
  - Low pressure drop
  - Cheap to build /maintain (no moving parts)
- Disadvantages
  - Poor efficiency for smaller particles (less than 10 microns)
  - Not suitable for abrasive particles
- Hence, grow particles with agglomerator to increase efficiency

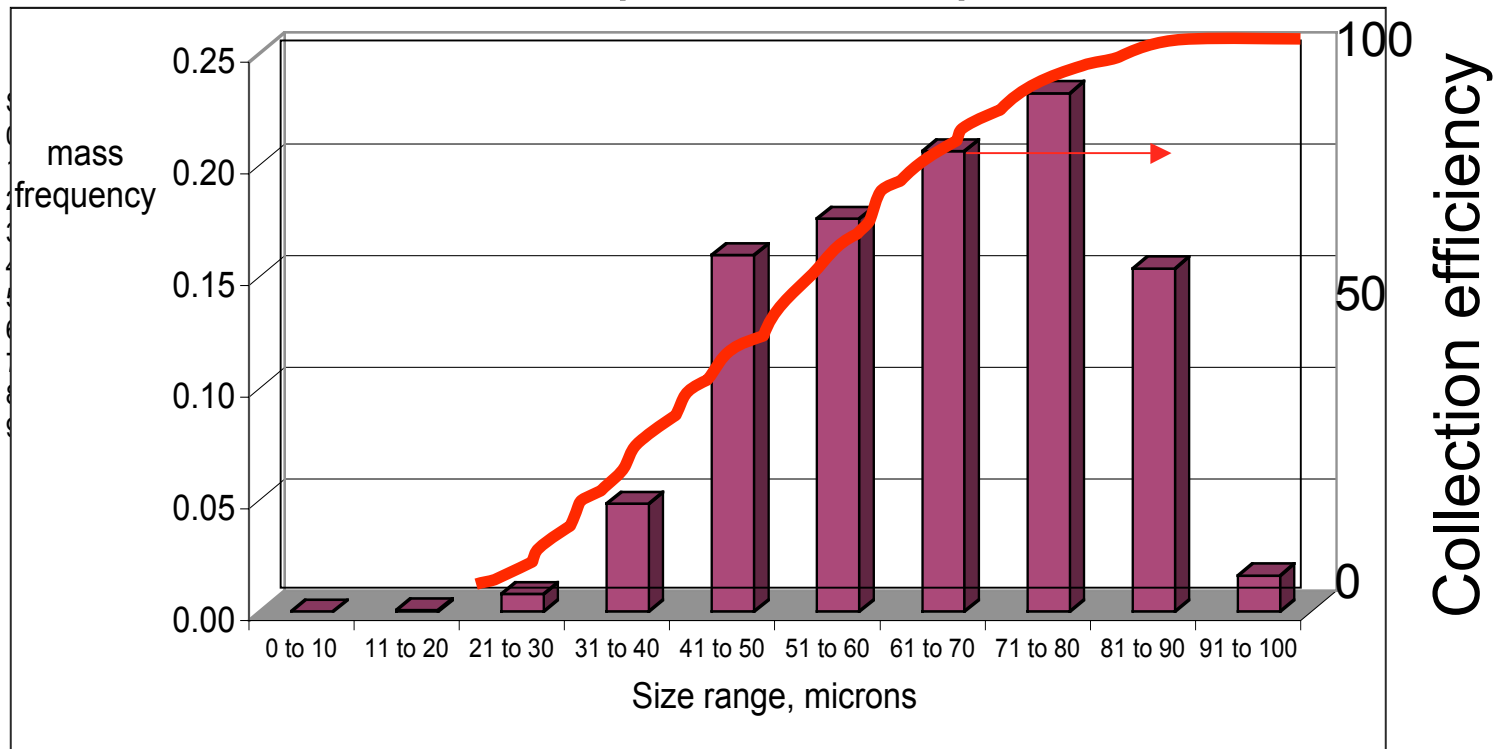
# Collection efficiency

- Example curve

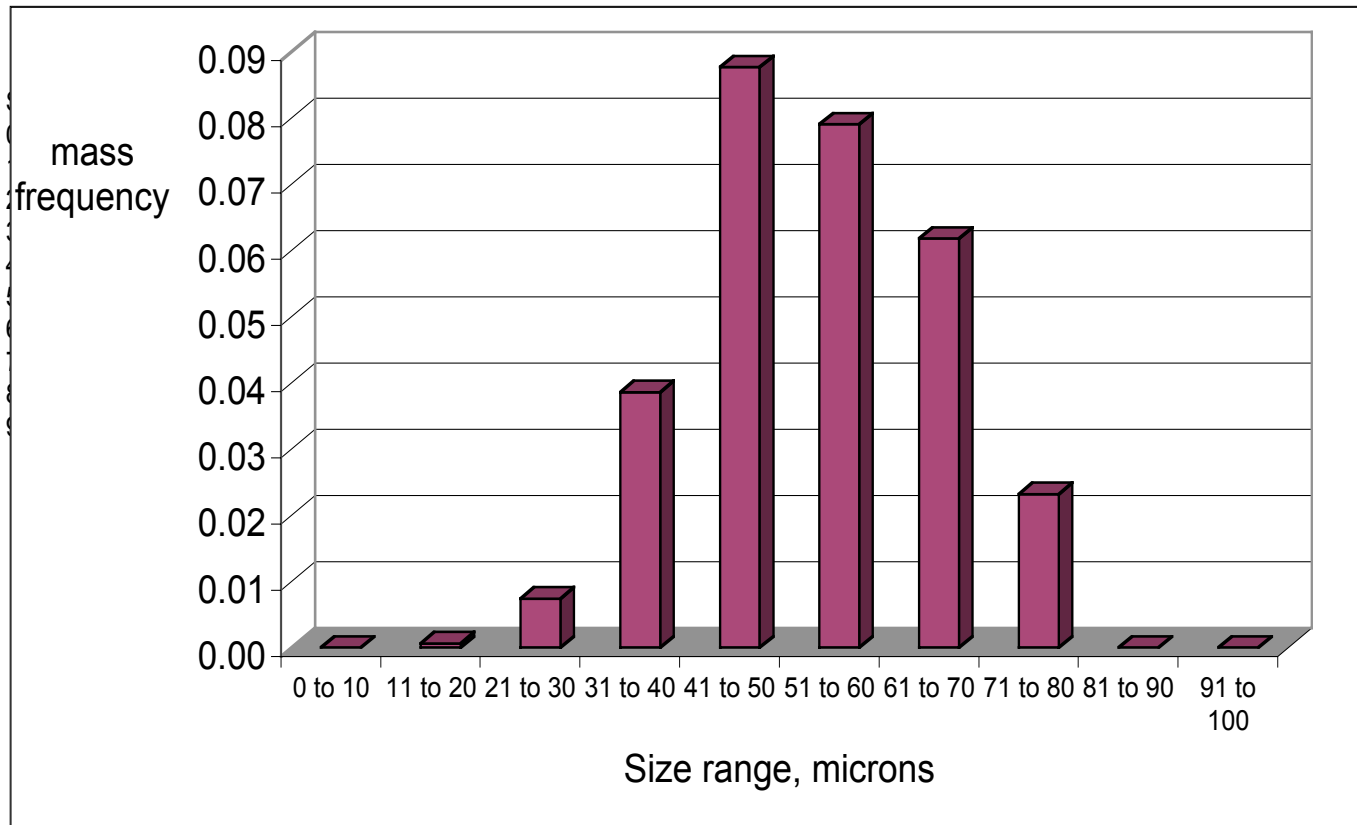


# Histogram before cyclone

Number of particles vs particle diameter



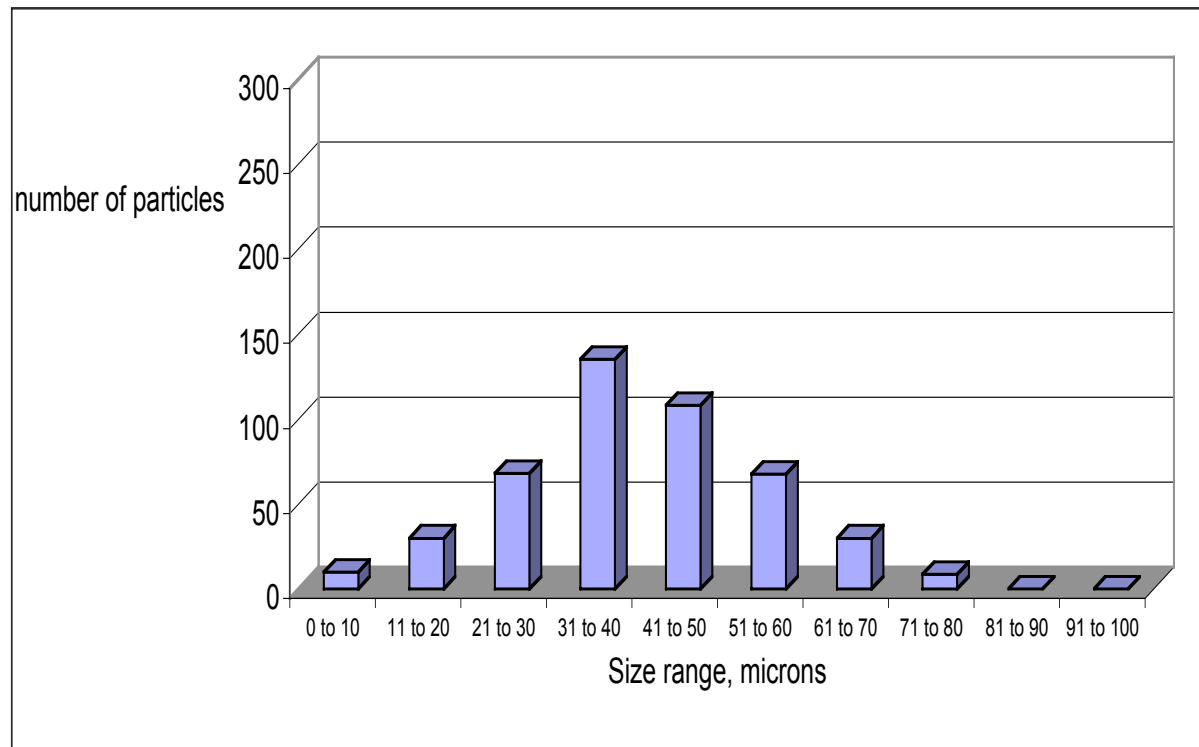
# After cyclone





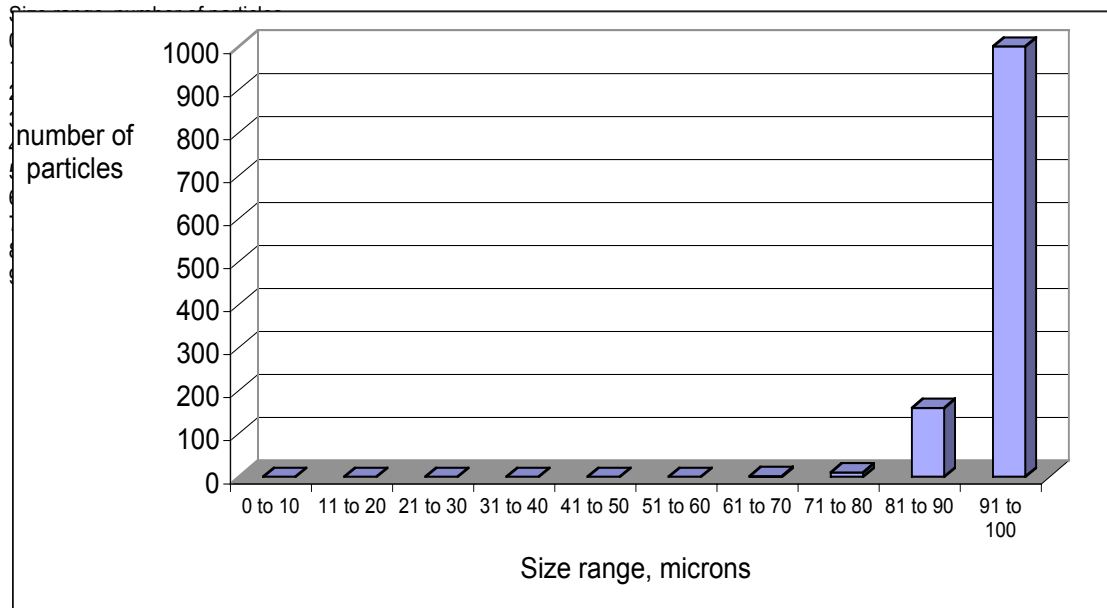
# Histogram after cyclone

Number of particles vs particle diameter



# Quick solution?

- Why not make the tube very very long, and just allow a very very long time for coagulation so that the particles become very big?

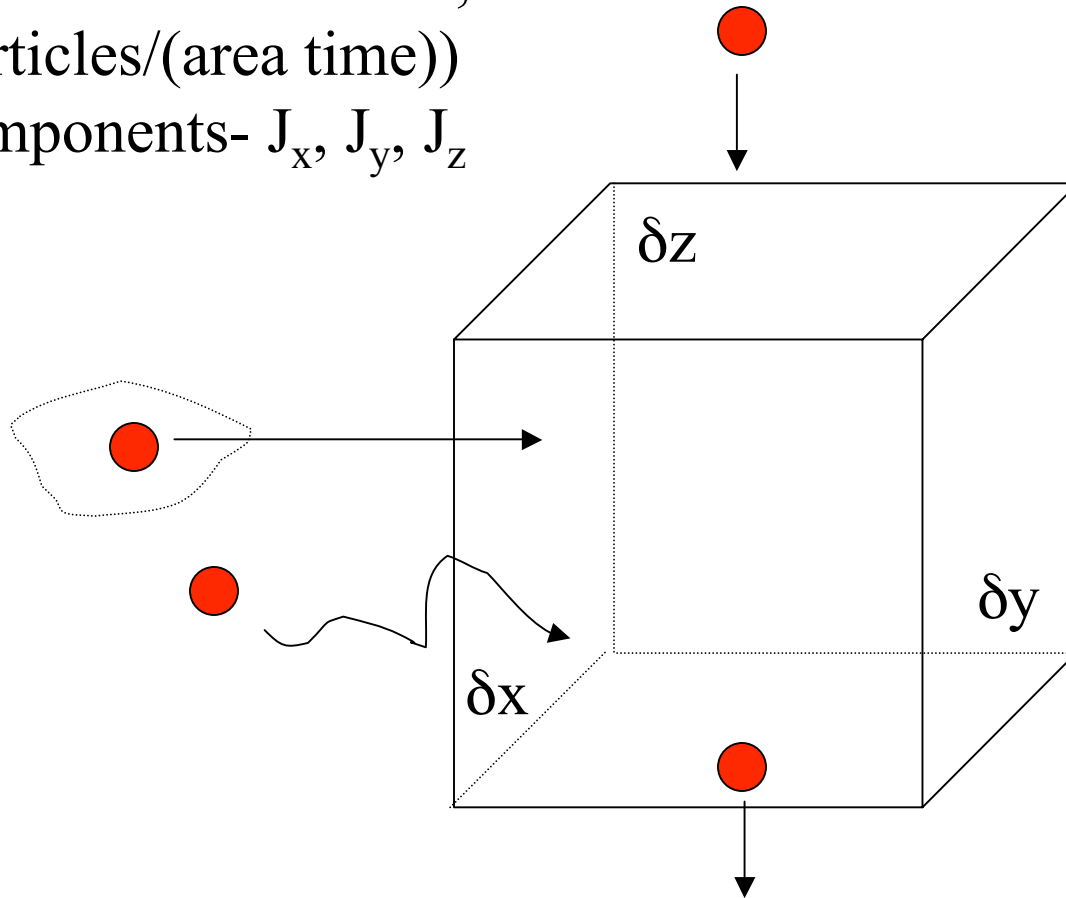


# Dynamics of size distributions

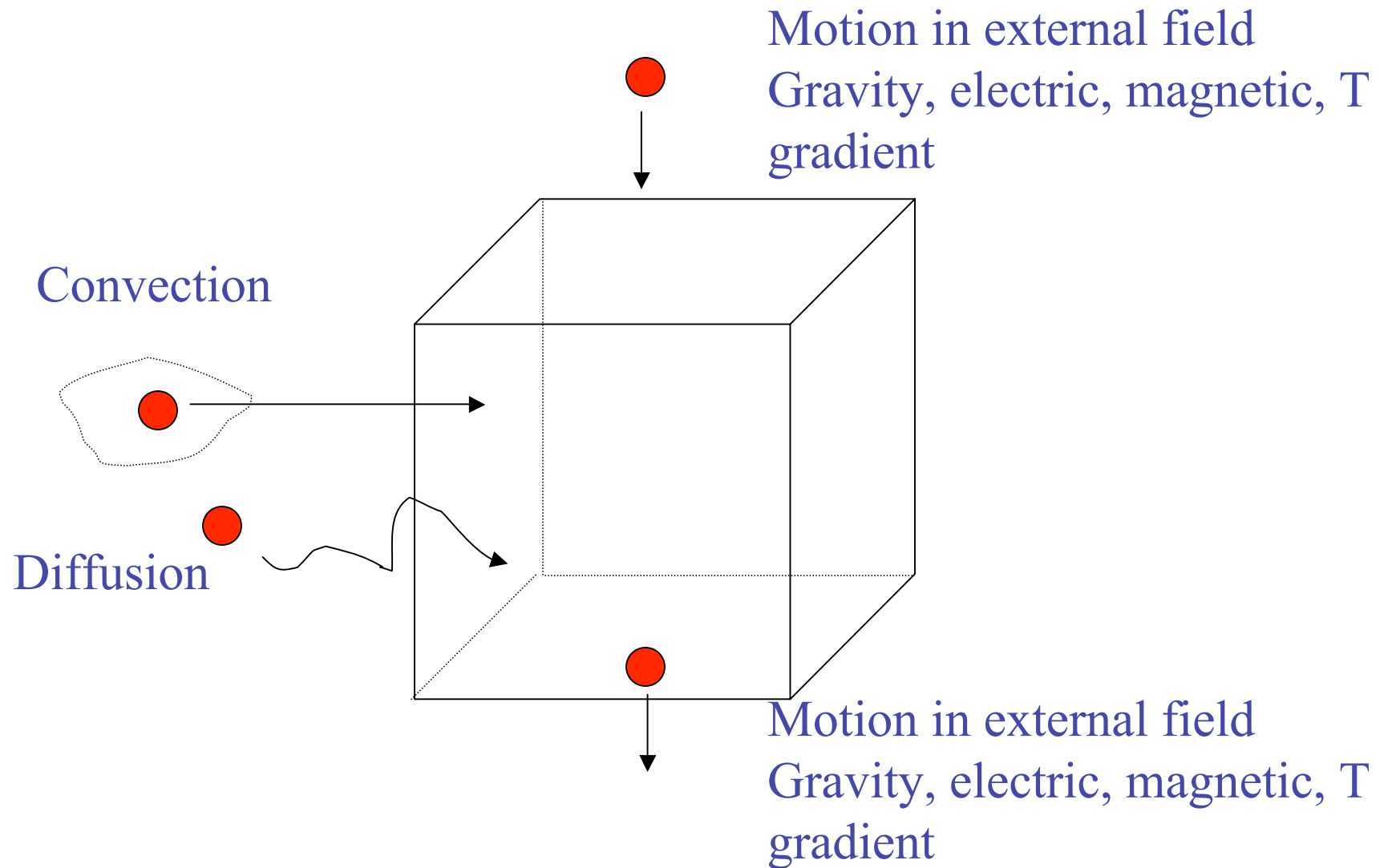
- Control volume approach
- Population balance
- Physics of coagulation/breakup

# Take a control volume....

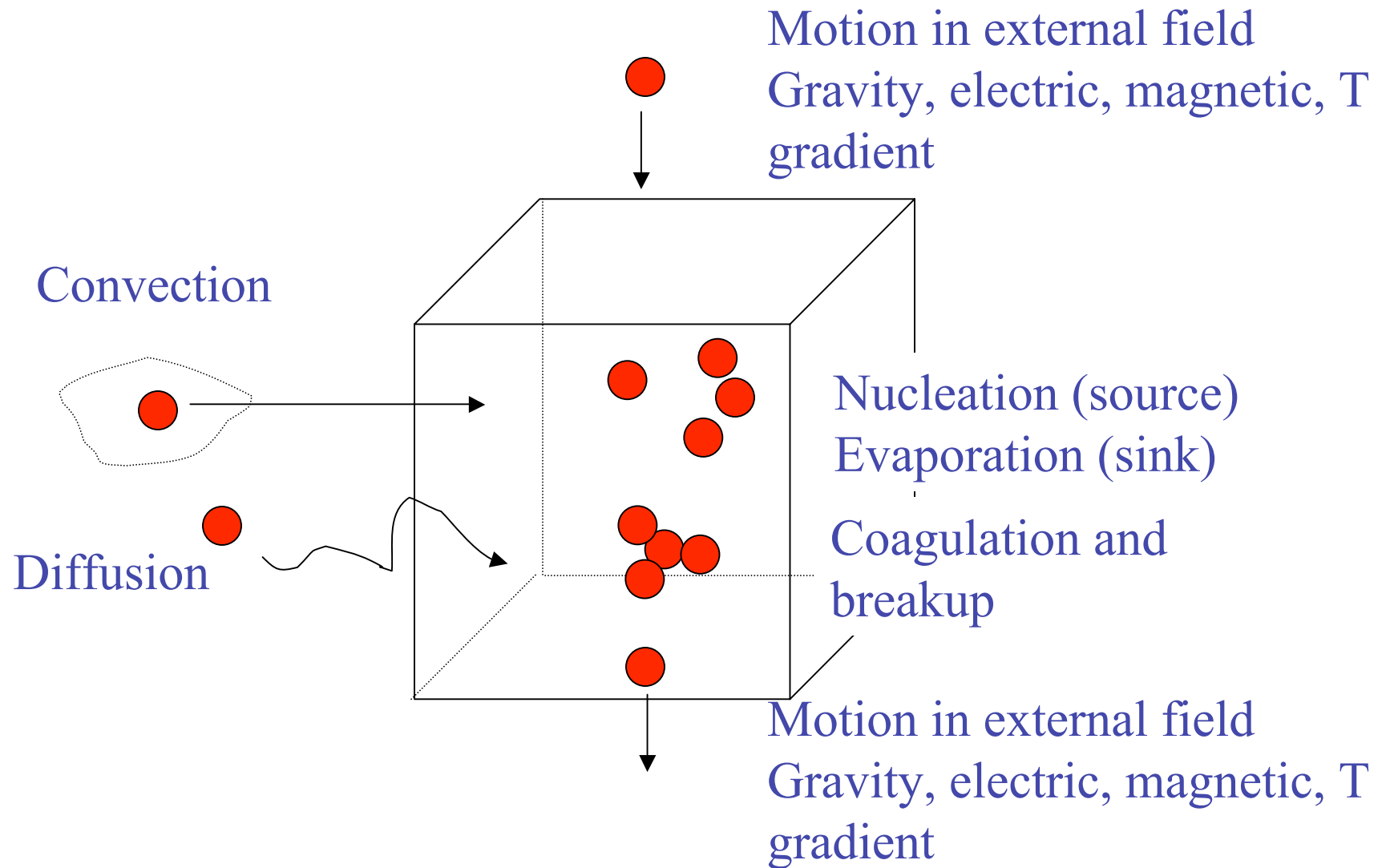
$\mathbf{J}$ , particle flux vector,  
(particles/(area time))  
Components-  $J_x, J_y, J_z$



# Take a control volume....

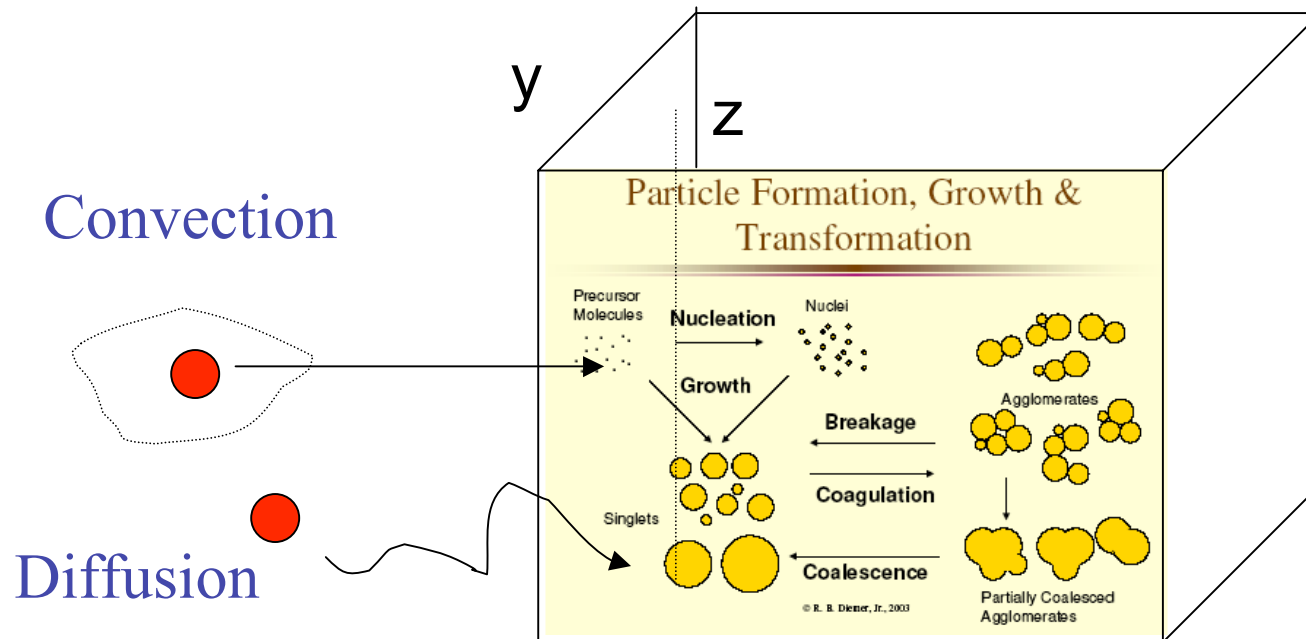


# Take a control volume....



# Take a control volume....

x



In our problem, no external fields

External coordinates,  $x, y, z, t$

Internal coordinate - particle volume  $V$  for 1D population balance

# Population balance for dynamics of distributions

- One dimension
  - **Particles, variable diameter**
  - Polymers, variable molecular weight
- 2 dimensions
  - Particles, variable diameter, surface area
  - Polymers, variable molecular weight, number of branch points



# Simplifying assumptions, initial conditions

- Assumptions:
  - Steady state, axisymmetric flow, incompressible fluid
  - Plug flow, no slip condition at wall
  - Diffusive transport of particles negligible with respect to convective transport
  - Dynamics of size distribution only varying with axial distance in agglomerator
  - No sources or sinks
- Initial conditions:
  - Particles all same size to start
  - Well mixed

# General Differential Form, 1-D Population

$$\begin{aligned}
 & -\nabla \cdot [\mathbf{u}_p(V, \mathbf{x}, t) n(V, \mathbf{x}, t)] && \text{convection} \\
 & +\nabla \cdot [D_p(V, \mathbf{x}, t) \nabla n(V, \mathbf{x}, t)] && \text{diffusion} \\
 & -\frac{\partial}{\partial V} [G(V, \mathbf{x}, t) n(V, \mathbf{x}, t)] && \text{growth ("In - Out" in internal coordinate)} \\
 & +S(V, \mathbf{x}, t) && \text{sources \& sinks (Net Generation)} \\
 & && = \\
 & \frac{\partial n(V, \mathbf{x}, t)}{\partial t} && \text{accumulation}
 \end{aligned}$$

} "In - Out" in external coordinates

*Note: object's velocity may differ from fluid's velocity owing to either slip or action of external forces*

# Full 1-D Population Balance (a partial integrodifferential equation)

$$\begin{aligned}
 \frac{\partial n}{\partial t} + \nabla \cdot (\mathbf{u}_p n) - \nabla \cdot (D_p \nabla n) = & \text{ nucleation term} \\
 N \delta(V - v_0) - \frac{\partial}{\partial V} (G n) & \text{ growth term} \\
 + \frac{1}{2} \int_0^V \beta(v, V-v) n(v) n(V-v) dv - n(V) \int_0^\infty \beta(v, V) n(v) dv & \text{ coagulation terms} \\
 + \int_V^\infty \Gamma(\Phi) b(V; \Phi) n(\Phi) d\Phi - \Gamma(V) n(V) & \text{ breakage terms}
 \end{aligned}$$

N = nucleation rate  
 G = accretion rate  
 $\beta$  = coagulation rate  
 $\Gamma$  = breakage rate  
 b = daughter distribution  
 $v_0$  = nuclei size

# Problem Setup

- Steady-state, incompressible, axisymmetric flow
- Plug flow, no slip
- Neglect diffusion
- Population Balance Model:

$$u_z \frac{\partial n}{\partial z} = \int_V^\infty \Gamma(\Phi) b(V; \Phi) n(\Phi) d\Phi - \Gamma(V) n(V) + \frac{1}{2} \int_0^V \beta(v, V-v) n(v) n(V-v) dv - n(V) \int_0^\infty \beta(v, V) n(v) dv$$

# Partial List of Techniques

- Discrete Methods

- Sectional Methods



Will discuss

---

- Similarity Solutions }

- LaPlace Transforms

- Orthogonal Polynomial Methods

- Spectral Methods

- Moment Methods

- Monte Carlo Methods



Will not  
discuss

# But first, some particle physics

New dimensionless numbers

Knudsen number =  $2\lambda/d_p$

Particle Reynolds number =  $\rho_f d_p U/\mu$

**Free molecular regime:**

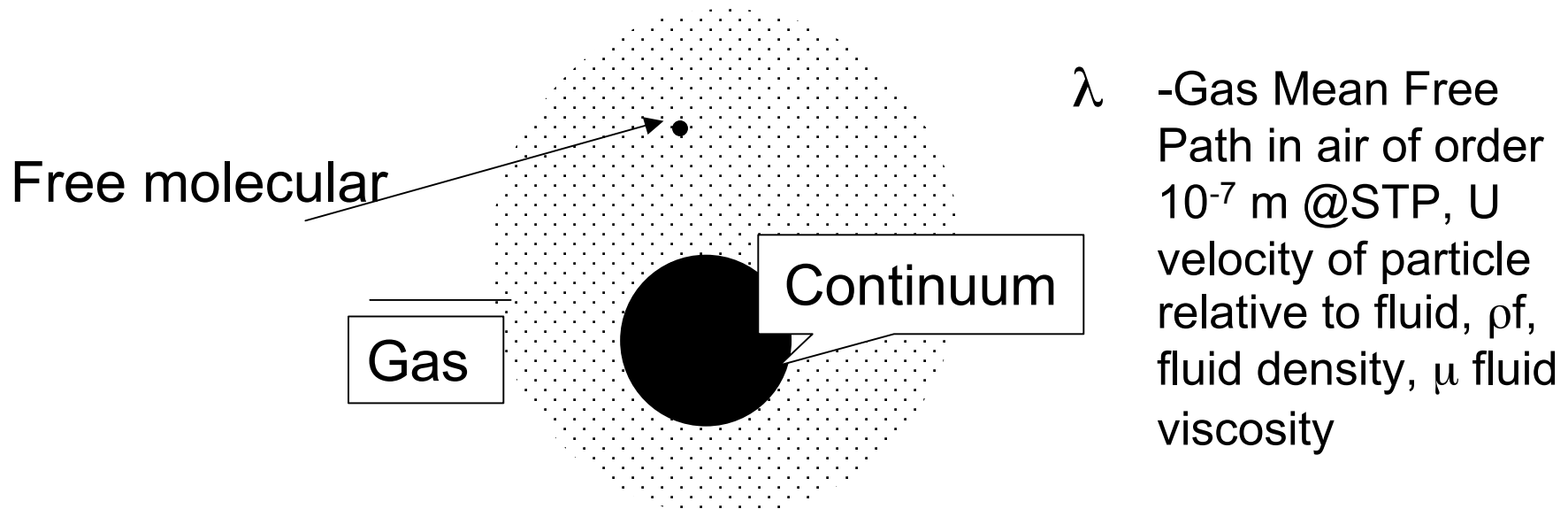
$Kn \gg 1$

**Continuum regime:**

$Kn \ll 1$

**Transition regime:**

in between



# Particles in fluid

- So for small particles, collisions with individual gas molecules affect particle motion, growth dynamics
- If size scale of particles --> size scale of eddies in very turbulent flow, turbulence in fluid may affect motion and growth dynamics
  - Levich (1962) agglomeration rate  $\propto v_{\text{basic}}^{9/4}$
  - Important for particles greater than 1 micron and up

# Coagulation - definitions

- Coagulation of solid particles = agglomeration
- When relative motion of particles is Brownian, process = thermal coagulation
- Relative motion from particle- fluid interactions = turbulent coagulation
- Saffman Turner (1956) divided into 2 processes
  - Turbulent shear agglomeration: particles on different streamlines are traveling at different speeds, enhances collisions
  - Turbulent inertial agglomeration: particle trajectories depart from flow streamlines, and lead to collisions



# Collision frequency function

Consider the change in number concentration of particles of size  $i$ , where  $v_i = v_j + v_{i-j}$

collision frequency - # collisions/time between particles of size  $j$  and size  $i-j$  =

$v_j, v_{i-j}$  are volumes of particles of size  $j$  and  $i-j$

$$N_{j,i-j} = \beta(v_j, v_{i-j}) n_j n_{i-j}$$

$\beta$ , also known as a collision kernel, depends on the size of the colliding particles, and properties of system such as temperature

For our problem, we assume all collisions 'stick'

# Test yourself....

- For small particles (free molecular and continuum) will the collision kernel increase or decrease as the system temperature increases?
- For particles in the continuum regime, will the collision kernel increase or decrease as the viscosity of the surrounding gas increases?

# Coagulation - discrete distribution bookkeeping

For a discrete size distribution, the rate of formation of particles of size  $i$  by collision of particles of size  $j$  and  $i-j$ , is given by:  $\frac{1}{2} \sum_{j=1}^{i-1} N_{j, i-j}$  where the factor  $1/2$  is introduced because each collision is counted twice in summation

Rate of loss of particles of size  $i$  by collision with all other particles is given by:  $\sum_{j=1}^{\infty} N_{i, j}$

Change in number concentration of particles of size  $i$  given by:

$$\frac{dn_i}{dt} = \frac{1}{2} \sum_{j=1}^{i-1} N_{j, i-j} - \sum_{j=1}^{\infty} N_{i, j} = \frac{1}{2} \sum_{j=1}^{i-1} \beta_{j, i-j}(v_j, v_{i-j}) n_j n_{i-j} - n_i \sum_{j=1}^{\infty} \beta_{ij}(v_j, v_i) n_j$$

theory of coagulation for discrete spectrum developed by Smoluchowski (1917) change = formation - loss

# Notation change:

Impt for problem statement: Alternate, more general, way to represent two colliding particles,  $v_j v_{i-j} \rightarrow$

$\phi, V-\phi$  where  $V$  is final size of pair after collisions

# Coagulation, continuous distributions

continuous nomenclature

$n(v)$  = number of particles per unit volume of size  $v$ , a continuous distribution

collision rate:

$$N_{\phi, v-\phi} = \beta(\phi, v - \phi)n(\phi)n(v - \phi)d\phi d(v - \phi)$$

where  $\beta$  is the collision frequency function described earlier

The rate of formation of particles of size  $v$  by collision of smaller particles of size  $\phi$  and  $v-\phi$  is given by:

$$\text{formation in range } dv = \frac{1}{2} \left[ \int_0^v \beta(\phi, v - \phi)n(\phi)n(v - \phi)d(v - \phi) \right] dv$$

$$\text{loss in range } dv = \left[ \int_0^\infty \beta(\phi, v)n(\phi)n(v)d\phi \right] dv$$

Here, loss is from collisions with all other particles, so must integrate over entire size range

# Collision frequency functions

for particles in  
continuum regime:  
(Stokes-Einstein  
relationship valid)

$$\beta(\phi, V - \phi) = \frac{2kT}{3\mu} \left[ 2 + \left( \frac{\phi}{V - \phi} \right)^{1/3} + \left( \frac{V - \phi}{\phi} \right)^{1/3} \right]$$

for particles in  
free molecular regime: (derived from kinetic theory of collisions  
between hard spheres)

$$\beta_{ij} = \left( \frac{3}{4\pi} \right)^{1/6} \left( \frac{6kT}{\rho_p} \right)^{1/2} \left( \frac{1}{\phi} + \frac{1}{v - \phi} \right)^{1/2} \left( \phi^{1/3} + (v - \phi)^{1/3} \right)^2$$

interpolation formulas between regimes given by  
Fuchs (1964) *The Mechanics of Aerosols*

# Collision frequency values:

	$10^{10}\beta$ $\text{cm}^3/\text{sec}$		
$d_1/d_2$	0.01	0.1	1.0
0.01	18		
0.1	240	14.4	
1.0	3200	48	6.8

where particle diameters,  $d_1$  and  $d_2$  are in microns

# Test yourself...

- Why are the collision kernel values greater for collisions between smaller and larger particles compared to particles of the same size?



# Turbulent coagulation

$$\beta(\phi, V - \phi) = 0.31 \sqrt{\frac{\varepsilon}{\nu}} \left[ V + 3\phi^{1/3} (V - \phi)^{2/3} + 3\phi^{2/3} (V - \phi)^{1/3} \right]$$

- $\nu$  = kinematic viscosity
- $\varepsilon$  = energy dissipation rate (rate of conversion of turbulence into heat by molecular viscosity,  $\text{m}^2/\text{s}^3$ )

# Comparison... brownian vs. turbulent

- For collisions between 1 micron diameter particles, Brownian kernel  $\sim 100$  x turbulent kernel
- But for collisions between 500 micron diameter particles, turbulent kernel  $\sim 10$  x Brownian kernel

# Breakup

- Definitions:
  - Parent: starting agglomerate, size  $\Phi$
  - Daughter: resulting fragments, size  $V$
- Different causes of breakup
  - Thermal/Brownian
  - Flow induced
- Simplifying assumption for our problem
- Breakup into equal sized daughter fragments (size  $V$ ), rate given by:

$$\Gamma(V) = \Gamma_o \left( \frac{\varepsilon}{\nu} \right)^{3/2} V^{1/3}$$
$$b(V; \Phi) = 2\delta \left( V - \frac{\Phi}{2} \right)$$

# Discrete Methods

- Size is integer multiple of fundamental size
- Write balance equations for every size
- Gives distribution directly
- Huge number of equations to solve
- Have to decide what the largest size is
- Example for coagulation and breakage:

$$V_i = iV_0; \quad V_0 = \frac{\pi d_0^3}{6}$$

$$u_z \frac{dn_i}{dz} = \frac{1}{2} \sum_{j=1}^{i-1} \beta_{j,i-j} n_j n_{i-j} - n_i \sum_{j=1}^{\infty} \beta_{i,j} n_j + \sum_{j=i+1}^{\infty} \Gamma_j b(i; j) n_j - \Gamma_i n_i$$

# Discrete Example Problem Setup

$$\beta_{c,i,j} = \frac{2kT}{3\mu} \left[ 2 + \left(\frac{i}{j}\right)^{2/3} + \left(\frac{j}{i}\right)^{1/3} \right]; \quad \beta_{t,i,j} = .31 \sqrt{\frac{\epsilon}{\nu}} V_0 (i + 3i^{2/3} j^{2/3} + 3i^{1/3} j^{2/3} + j)$$

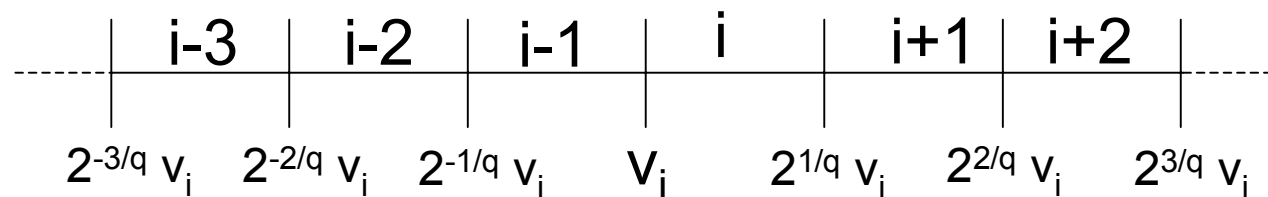
$$\Gamma_i = \begin{cases} 1 \times 10^{-8} \text{ s}^2/\text{cm} \left(\frac{\epsilon}{\nu}\right)^{3/2} V_0^{2/3} i^{4/3}, & i > 1 \\ 0, & i = 1 \end{cases}; \quad b(i,j) = \begin{cases} 2, & j = 2i \\ 1, & j = 2i+1, 2i-1 \\ 0, & j < 2i-1, j > 2i+1 \end{cases}$$

$$u_z \frac{dn_i}{dz} = \frac{1}{2} \sum_{j=1}^{i-1} (\beta_{c,j,i-j} + \beta_{t,j,i-j}) n_j n_{i-j} - n_i \sum_{j=1}^{\infty} (\beta_{c,i,j} + \beta_{t,i,j}) n_j \\ + \Gamma_{2i+1} n_{2i+1} + 2\Gamma_{2i} n_{2i} + \Gamma_{2i-1} n_{2i-1} - \Gamma_i n_i$$

Need slightly more than  $2 \times 10^6$  cells to cover entire mass distribution range!

# Sectional Method

- Best rendering due to Litster, Smit and Hounslow
- Collect particles in bins or size classes, with upper/lower size= $2^{1/q}$ , “q” optimized for physics



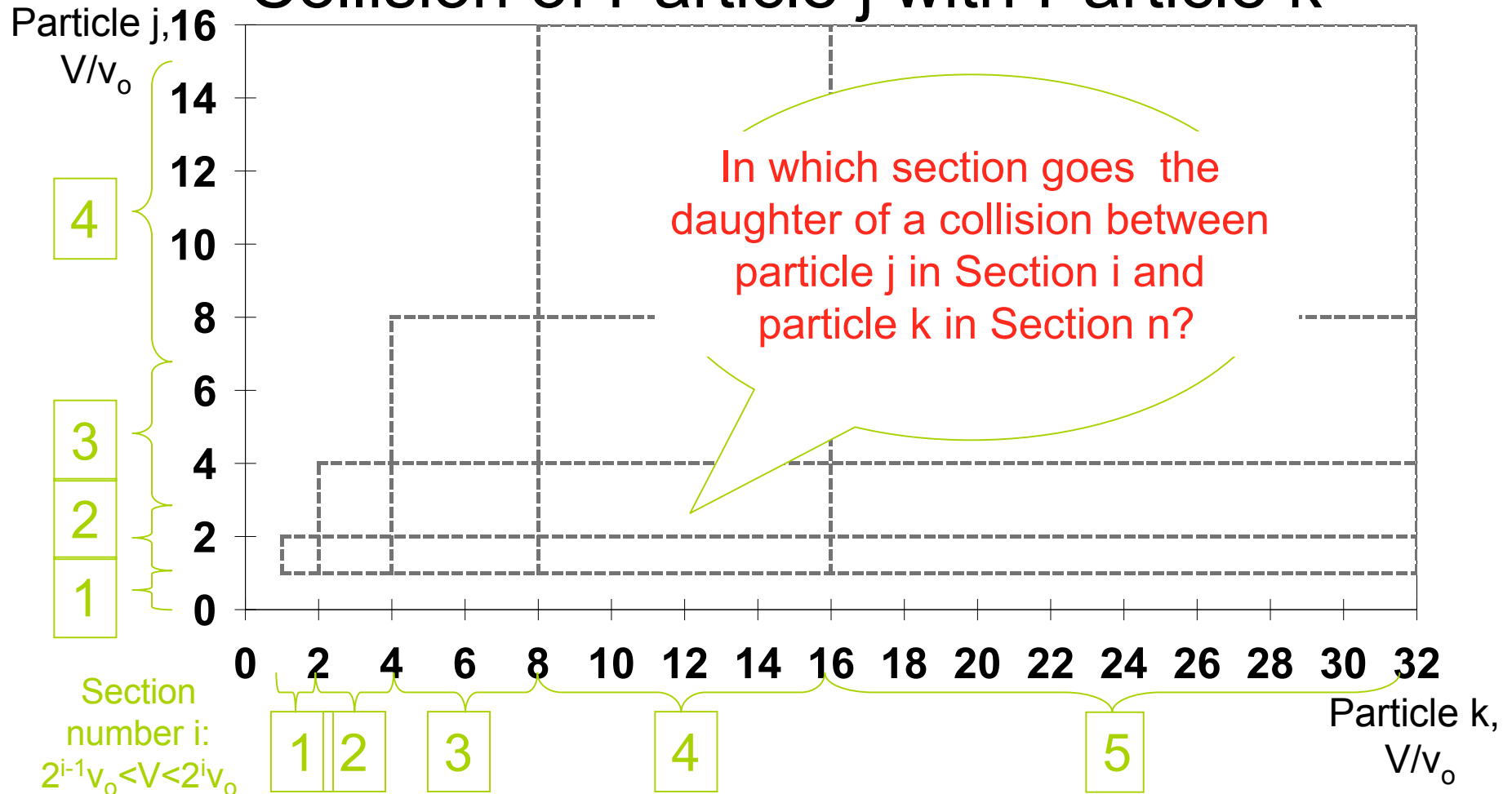
- Balances are written for each size class reducing the number of equations, but too few bins loses resolution
- And... now the equations get more complicated to get the balances right
- Still have problem of growing too large for top class
- Directly computes distribution

# Sectional Interaction Types

- Type 1:
  - some particles land in the  $i^{\text{th}}$  interval and some in a smaller interval
- Type 2:
  - all particles land in the  $i^{\text{th}}$  interval
- Type 3:
  - some particles land in the  $i^{\text{th}}$  interval and some in a larger interval
- Type 4:
  - some particles are removed from the  $i^{\text{th}}$  interval and some from other intervals
- Type 5:
  - particles are removed only from  $i^{\text{th}}$  interval

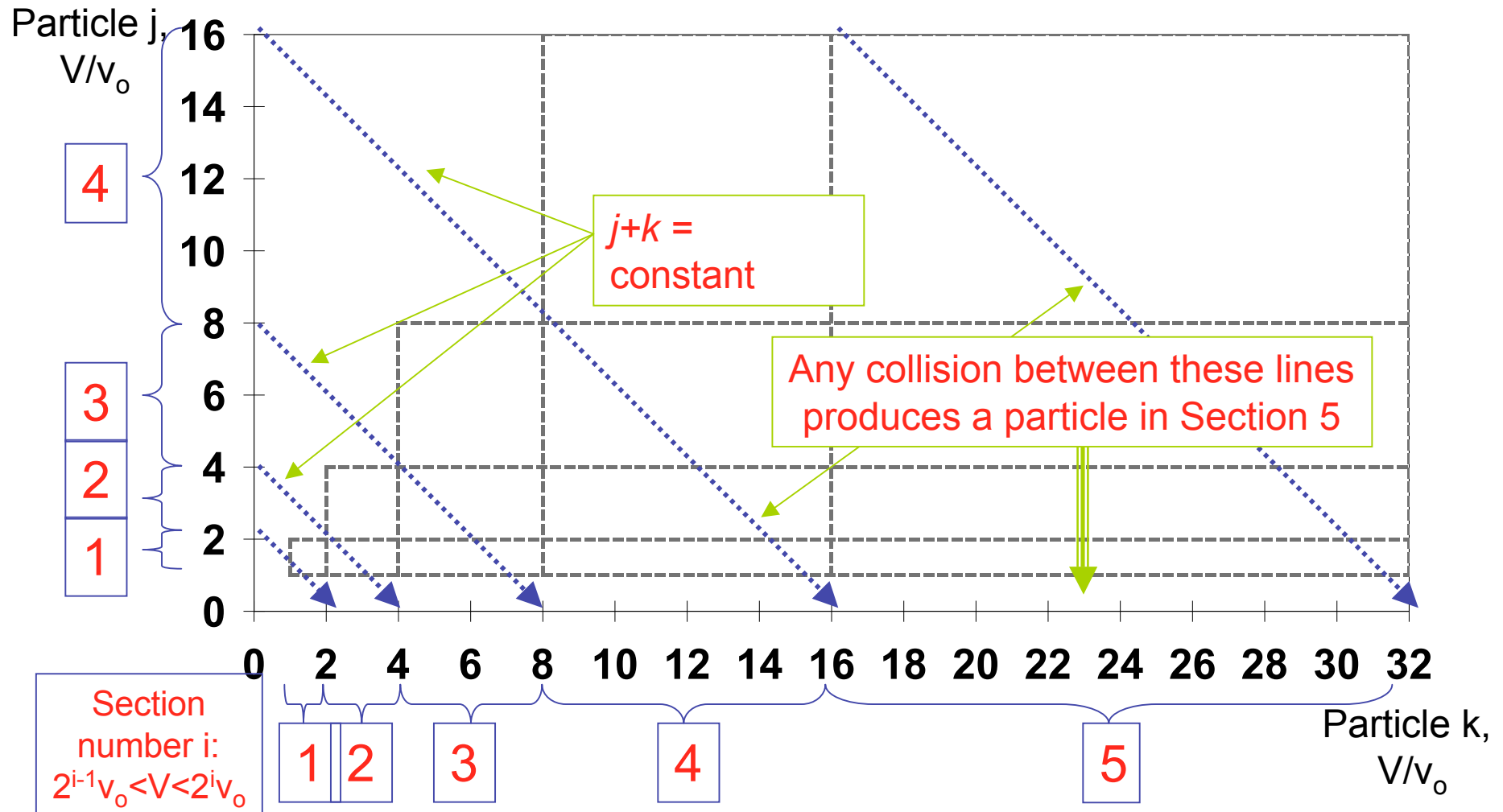
# Sectionalization Example: $q=1$

## Collision of Particle j with Particle k



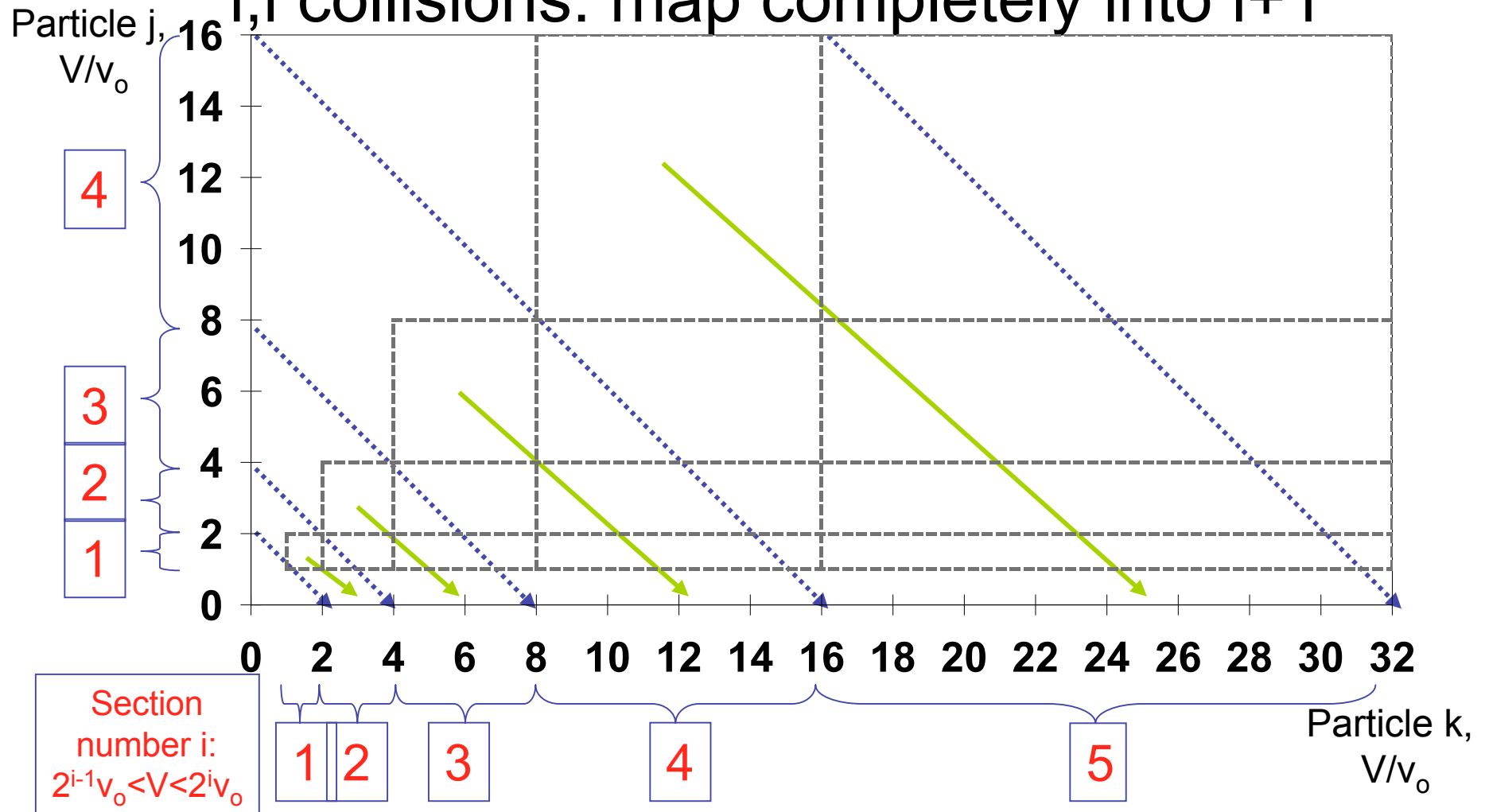


# Sectionalization Example: $q=1$



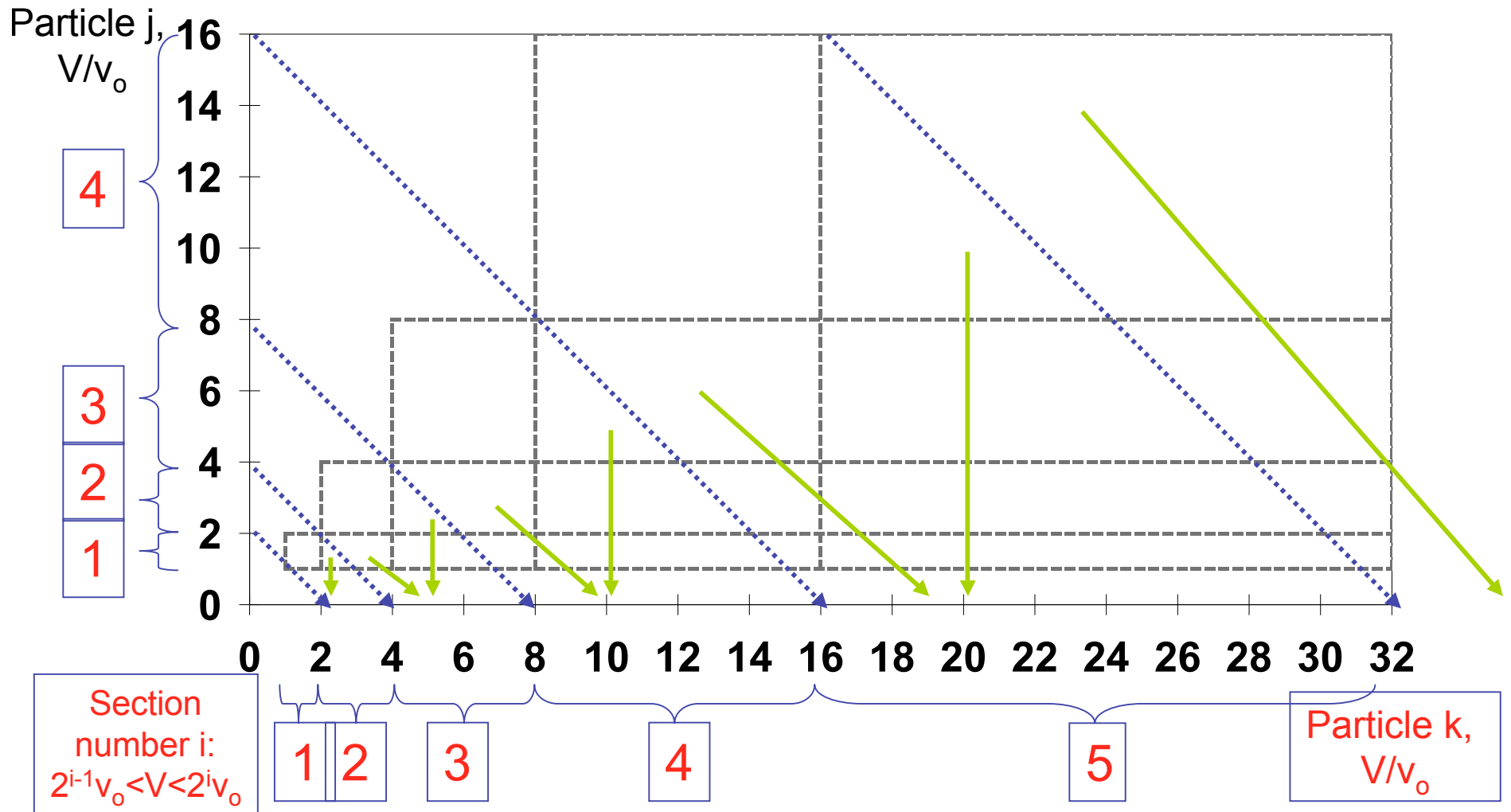
# Sectionalization Example: $q=1$

$i, j$  collisions: map completely into  $i+1$



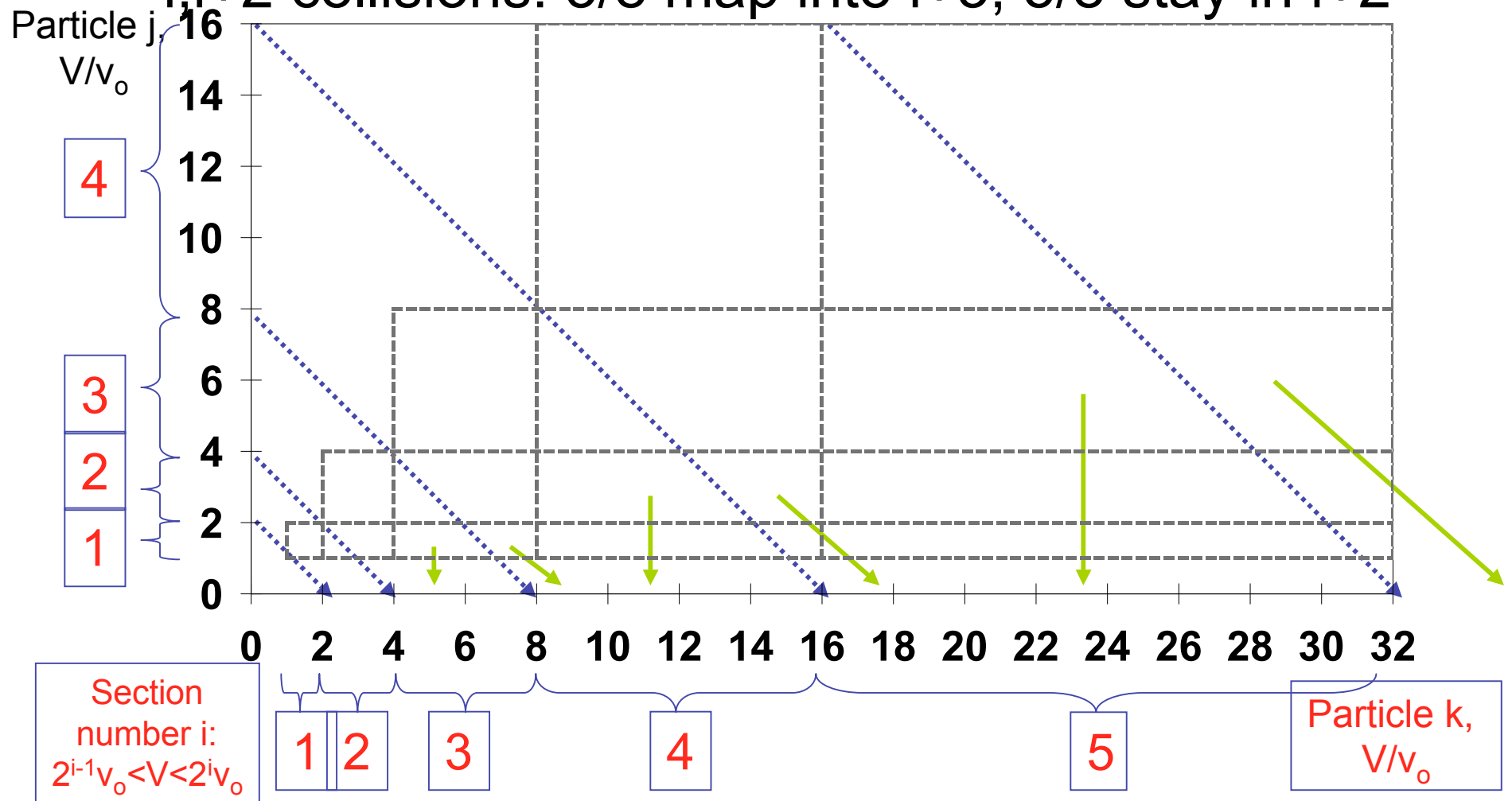
# Sectionalization Example: $q=1$

$i, i+1$  collisions:  $3/4$  map into  $i+2$ ,  $1/4$  stay in  $i+1$

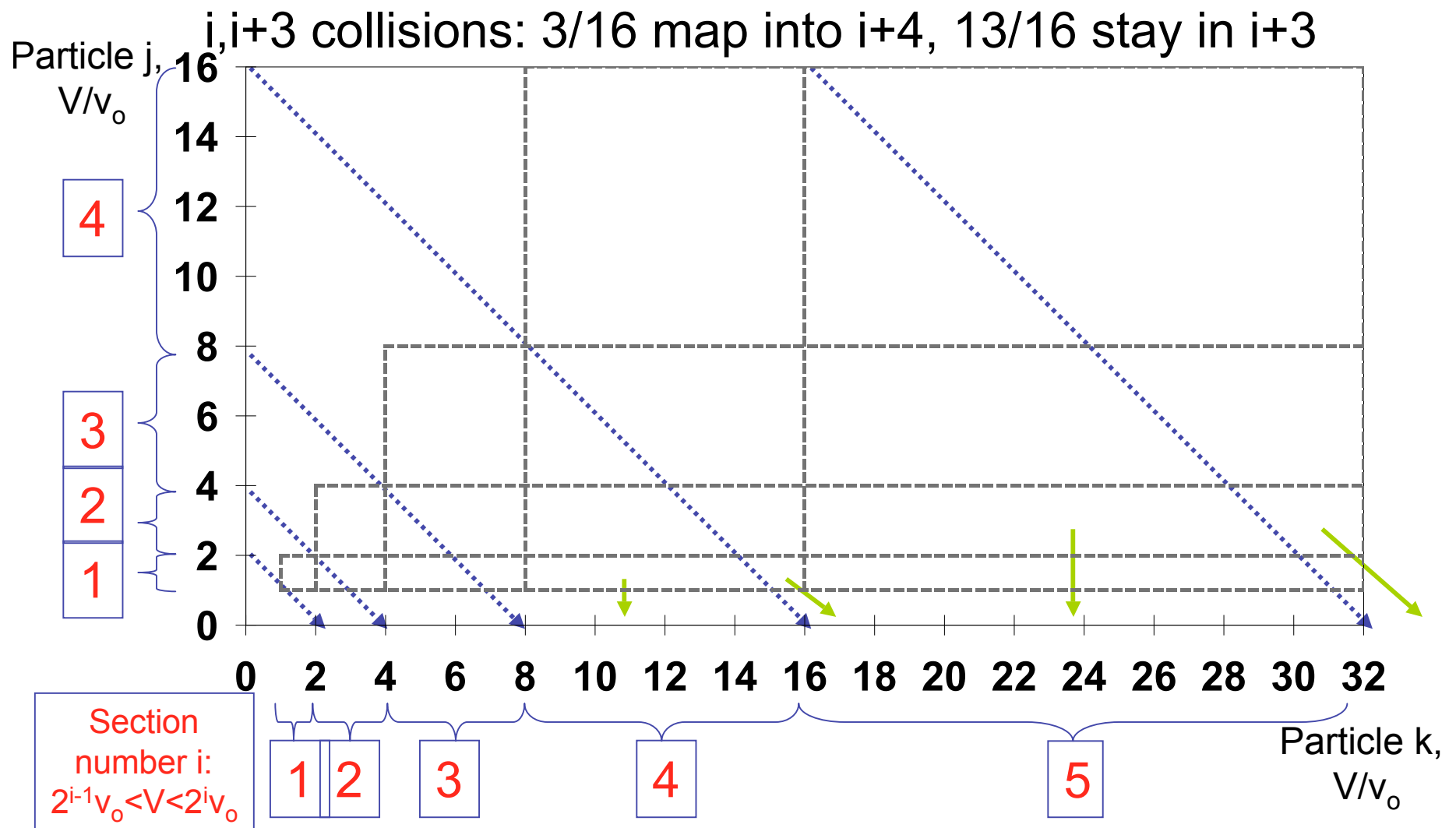


# Sectionalization Example: $q=1$

$i, i+2$  collisions:  $3/8$  map into  $i+3$ ,  $5/8$  stay in  $i+2$

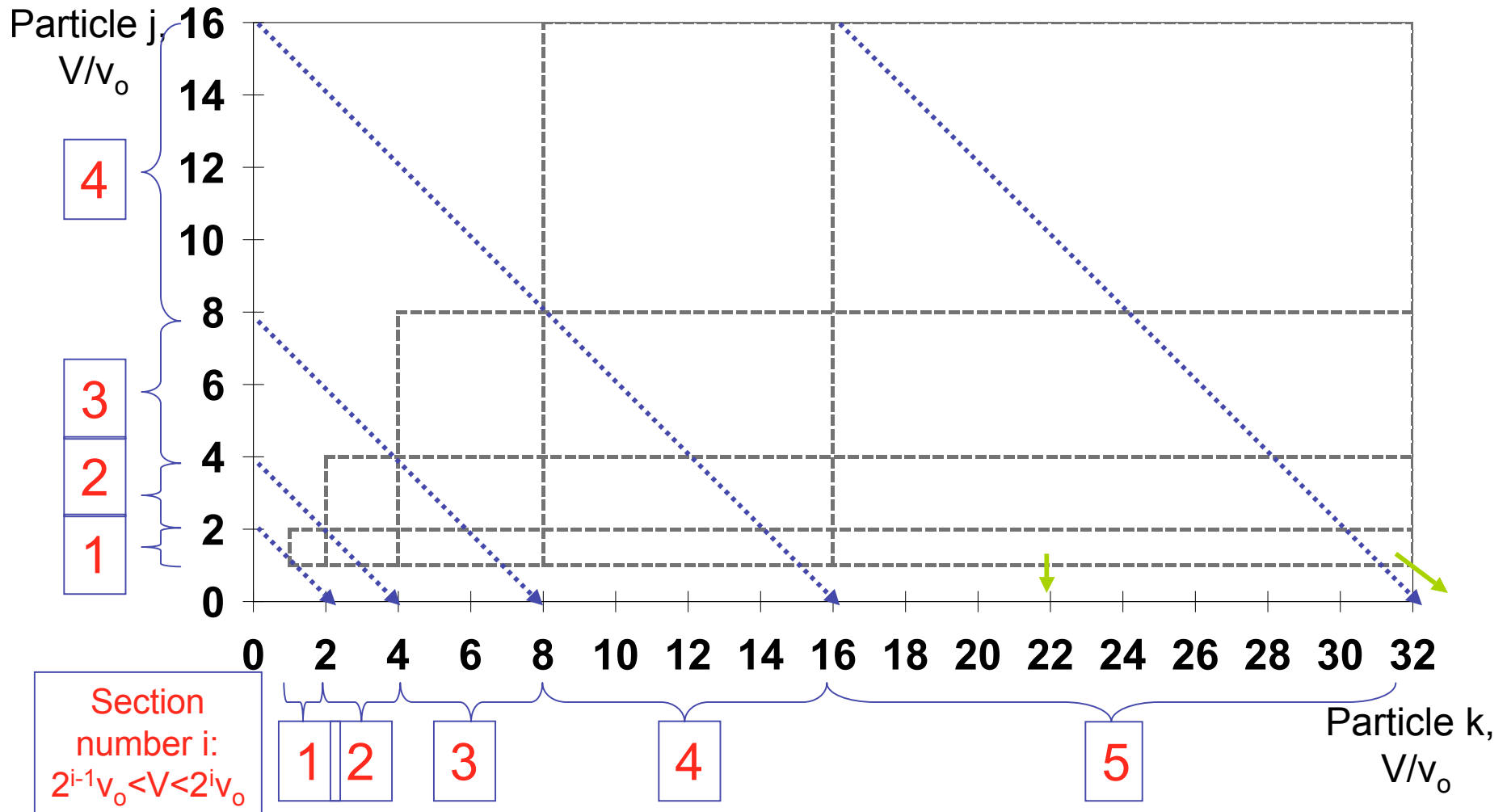


# Sectionalization Example: $q=1$



# Sectionalization Example: $q=1$

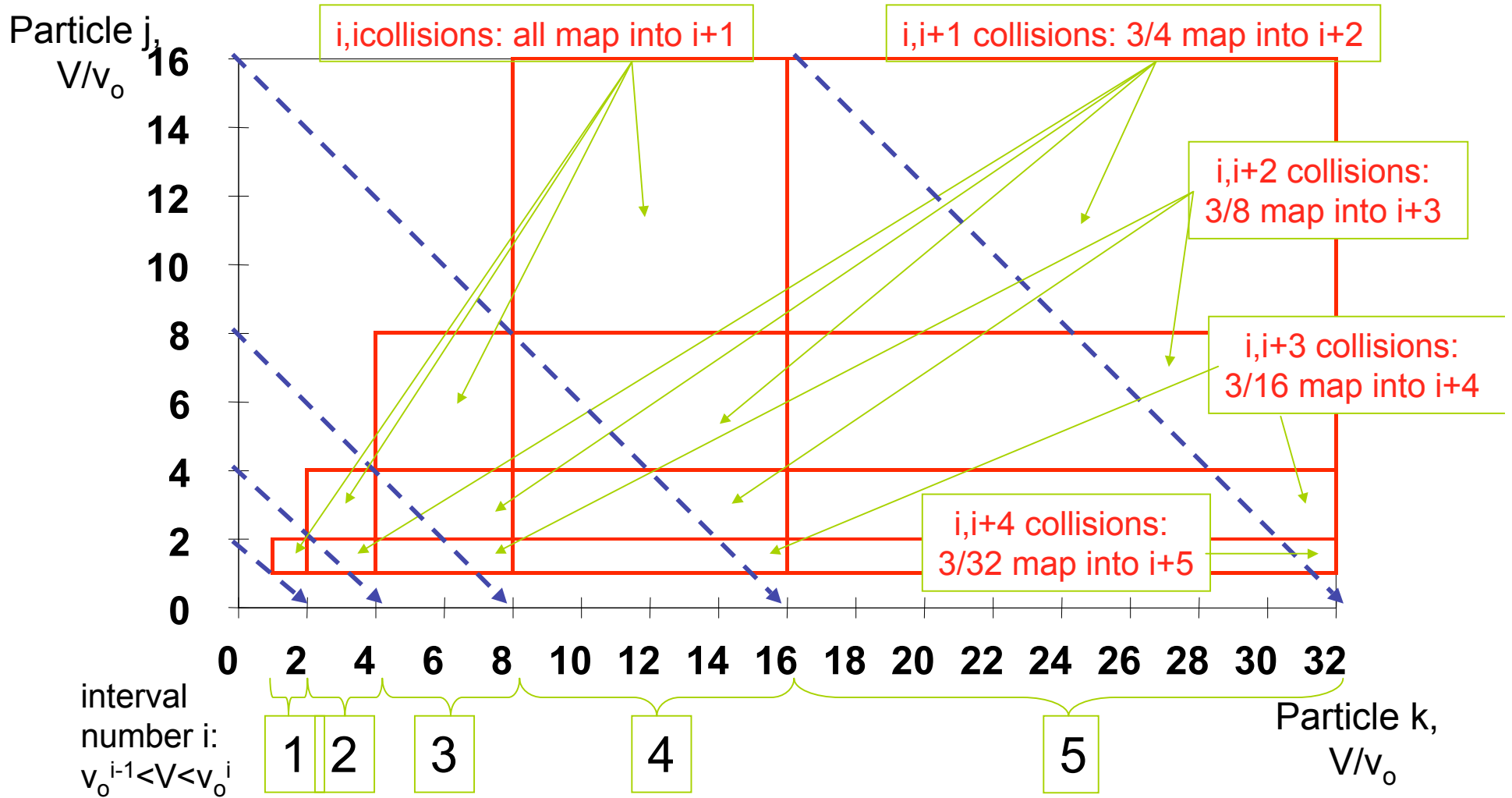
$i, i+4$  collisions: 3/32 map into  $i+5$ , 29/32 stay in  $i+4$



# Sectionalization Example: $q=1$

$i, n$  collisions:  $3/2^{n-i+1}$  map into  $n+1$ ,  $n > i > 0$

$i, i$  collisions: all map into  $i+1$



# Sectional Coagulation Model, $q=1$

- Model Equation:

$$u_z \frac{dN_i}{dz} = N_{i-1} \sum_{j=1}^{i-2} \beta_{i-1,j} \theta_{i-1,j} N_j + \frac{1}{2} \beta_{i-1,i-1} N_{i-1}^2 - N_i \left[ \sum_{j=1}^{i-1} \beta_{i,j} \theta_{i,j} N_j + \sum_{j=i}^{\infty} \beta_{i,j} N_j \right]$$

- Tentatively:

$$\theta_{i,j} = C \frac{3}{2} 2^{j-i}$$

- Can show (via 0<sup>th</sup> and 1<sup>st</sup> moments) that:

- number balance gives correct general form for arbitrary  $\theta_{ij}$
- mass balance only closes for  $C=2/3$  when  $V_i/V_j=2^{i-j}$
- final expression:  $\theta_{i,j} = 2^{j-i}$
- kernels evaluated via:

$$\bar{V}_i = 2^{i-1} \bar{V}_1 \quad \text{with} \quad \bar{V}_1 = \frac{V_0 + 2V_0}{2} = \frac{3V_0}{2} \quad (\text{recovers } 3/2 \text{ factor})$$



# Gaa!

- At least breakup is easier to visualize
- True equi-sized daughter distributions
- All particles from bin  $i$  that breakup will map to lower bin  $i-1$

## Sectional Example Problem Setup (for q=1)

$$\beta_{i,j} = \frac{2kT}{3\mu} \left[ 2 + 2^{(i-j)/3} + 2^{(j-i)/3} \right] + .31 \sqrt{\frac{\epsilon}{v}} \left( \frac{3V_0}{2} \right) \left[ 2^{i-1} + 3 \left( 2^{(2i+j)/3-1} + 2^{(i+2j)/3-1} \right) + 2^{j-1} \right]$$

$$\Gamma_i = \begin{cases} 1 \times 10^{-8} \text{ s}^2/\text{cm} \left( \frac{\epsilon}{v} \right)^{3/2} \left( \frac{3V_0}{2} \right)^{1/3} 2^{(i-1)/3}, & i > 1; \\ 0, & i = 1 \end{cases}; \quad b(i,j) = \begin{cases} 2, & j = i + 1 \\ 0, & j \neq i + 1 \end{cases}$$

$$u_z \frac{dN_i}{dz} = N_{i-1} \sum_{j=1}^{i-2} \frac{\beta_{i-1,j}}{2^{i-j-1}} N_j + \frac{1}{2} \beta_{i-1,i-1} N_{i-1}^2 - N_i \left[ \sum_{j=1}^{i-1} \frac{\beta_{i,j}}{2^{i-j}} N_j + \sum_{j=1}^{\infty} \beta_{i,j} N_j \right] + 2\Gamma_{i+1} N_{i+1} - \Gamma_i N_i$$

Need about 22 sections to cover entire mass distribution range, suggest using 25-30

# General $2^{1/q}$ Sectional Coagulation Model

$$\frac{dN_i}{dt} = N_{i-1} \sum_{j=1}^{i-S(q)-1} \beta_{i-1,j} \theta_{i-1,j} N_j + \frac{1}{2} \beta_{i-q,i-q} N_{i-q}^2 - N_i \left[ \sum_{j=1}^{i-S(q)} \beta_{i,j} \theta_{i,j} N_j + \sum_{j=i-S(q)+1}^{\infty} \beta_{i,j} N_j \right]$$

$$+ \sum_{k=2}^q \sum_{j=i-S(q-k+2)-k+1}^{i-S(q-k+1)-k} \beta_{i-k,j} (\theta_{i-1,j} + \psi_k) N_{i-k} N_j$$

$$- \sum_{k=2}^q \sum_{j=i-S(q-k+2)-k+2}^{i-S(q-k+1)-k+1} \beta_{i-k+1,j} (\theta_{i,j} + 2^{1/q} \psi_{k+1}) N_{i-k+1} N_j$$

} 2 new terms

$$S(q) = \sum_{m=1}^q m; \quad \theta_{i,j} = \frac{2^{(j-0)/q}}{2^{1/q} - 1}; \quad \psi_k = \frac{2^{(1-k)/q} - 1}{2^{1/q} - 1}$$

# Sectional Example Problem Setup (for $q=1$ )

## Nondimensionalization

$$\beta_{i,j} = \beta_c^\circ \Psi_{c,i,j} + \beta_i^\circ \left( \frac{3V_0}{2} \right) \Psi_{t,i,j}; \quad \Gamma_i = \Gamma^\circ \left( \frac{3V_0}{2} \right)^{1/3} 2^{(i-0)/3}$$

$$\beta_c^\circ = \frac{2kT}{3\mu}; \quad \Psi_{c,i,j} = 2 + 2^{(i-j)/3} + 2^{(j-i)/3}; \quad \Gamma^\circ = 1 \times 10^{-8} \text{ s}^2/\text{cm} \left( \frac{\varepsilon}{V} \right)^{3/2}$$

$$\beta_i^\circ = .31 \sqrt{\frac{\varepsilon}{V}}; \quad \Psi_{t,i,j} = 2^{i-1} + 3 \left( 2^{(2i+j)/3-1} + 2^{(i+2j)/3-1} \right) + 2^{j-1}$$

$$\Theta = \frac{\tau}{\tau_c} = \frac{\beta_c^\circ M_0^\circ z}{u_s}; \quad \Theta_i = \frac{\tau_i}{\tau_c} = \frac{2\beta_c^\circ}{3\beta_i^\circ V_0}; \quad \Theta_b = \frac{\tau_b}{\tau_c} = \frac{\beta_c^\circ M_0^\circ}{\Gamma^\circ} \left( \frac{2}{3V_0} \right)^{1/3}$$

$$\Psi_{i,j} = \Psi_{c,i,j} + \frac{\Psi_{t,i,j}}{\Theta_i}; \quad n_i = \frac{N_i}{M_0^\circ}; \quad M_0^\circ = \frac{2M_1}{3V_0} = \frac{4M_1}{\pi d_0^3}$$

$$\frac{dn_1}{d\Theta} = n_{1-1} \sum_{j=1}^{1-2} \frac{\Psi_{1-1,j}}{2^{1-j-1}} n_j + \frac{1}{2} \Psi_{1-1,1-1} n_{1-1}^2 - n_1 \left[ \sum_{j=1}^{1-1} \frac{\Psi_{1,j}}{2^{1-j}} n_j + \sum_{j=1}^{\infty} \Psi_{1,j} n_j \right] + \frac{2^{1-1/3}}{\Theta_b} (2^{4/3} n_{1+1} - n_1)$$

# Solution tools:

- YOU need to figure out, and modify
  - Change # sections
  - change pipe dimensions
- popbal.m
  - Main, runs functions, checks efficiency, adjusts pipe length till 75% efficiency in cyclone met, checks mass closure
- numdist.m
  - Function, solves ODEs
- gamma.m
  - Function, calculates breakup kernels
- beta.m
  - Function, calculates coagulation kernels

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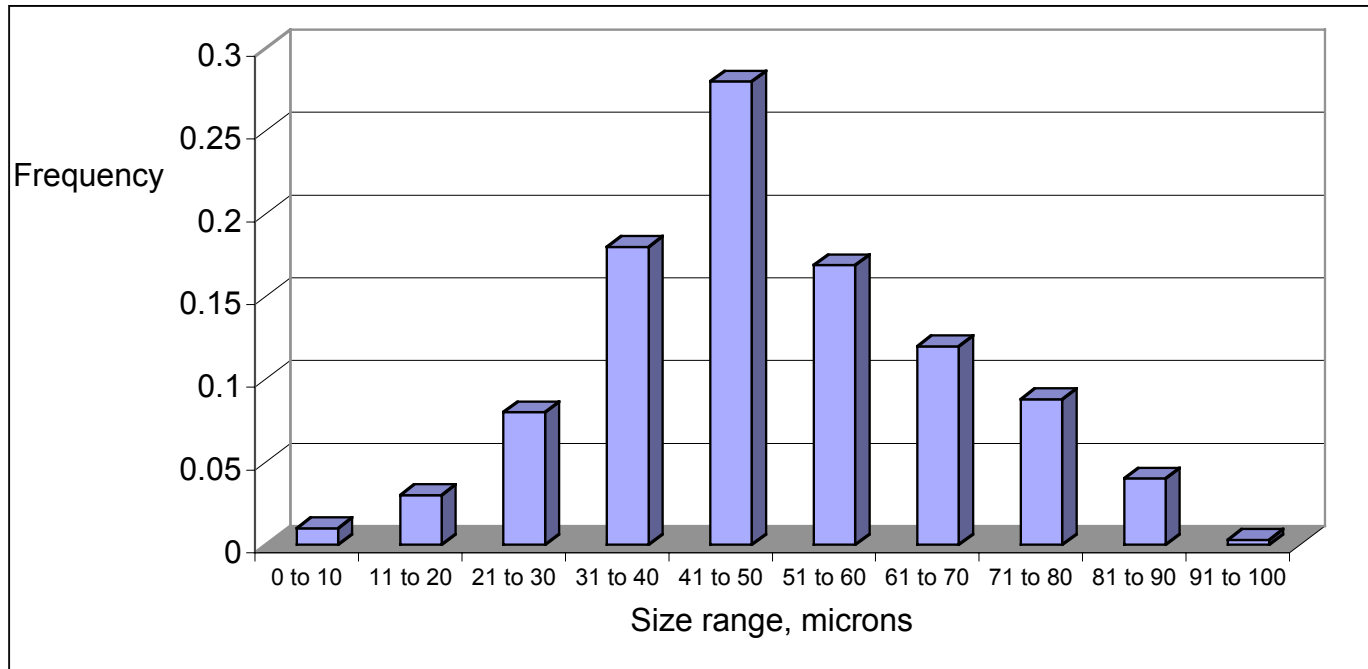
# Embedded spreadsheet for histogram

Size range, microns	number of particles	Frequency	assume diameter	mass of particles in each bin, g	Mass frequency
0 to 10	10	0.01	5	6.54E-10	0.00
11 to 20	30	0.03	15	5.30E-08	0.00
21 to 30	80	0.08	25	6.54E-07	0.01
31 to 40	180	0.18	35	4.04E-06	0.05
41 to 50	280	0.28	45	1.34E-05	0.16
51 to 60	169	0.169	55	1.47E-05	0.18
61 to 70	120	0.12	65	1.72E-05	0.21
71 to 80	88	0.088	75	1.94E-05	0.23
81 to 90	40	0.04	85	1.29E-05	0.15
91 to 100	3	0.003	95	1.35E-06	0.02

total number 1000

total mass assuming 1g/cc density

## Number frequency vs particle diameter



# Population balance equation

**Population balance equations (PBEs)** have been introduced in several branches of modern science, mainly in Chemical Engineering, to describe the evolution of a population of particles. This includes topics like crystallization, liquid-liquid extraction, gas-liquid dispersions, liquid-liquid reactions, comminution, aerosol engineering, biology (where the separate entities are cells), polymerization, etc. Population balance equations can be said to be derived as an extension of the Smoluchowski coagulation equation which describes only the coalescence of particles. PBEs, more generally, define how populations of separate entities develop in specific properties over time. They are a set of Integro-partial differential equations which gives the mean-field behavior of a population of particles from the analysis of behavior of single particle in local conditions.<sup>[1]</sup> Particulate systems are characterized by the birth and death of particles. For example, consider precipitation process (formation of solid from liquid solution) which has the subprocesses nucleation, agglomeration, breakage, etc., that result in the increase or decrease of the number of particles of a particular radius (assuming formation of spherical particles). Population balance is nothing but a *balance on the number of particles of a particular state* (in this example, *size*).

## Formulation of PBE

Consider the average number of particles with particle properties denoted by a particle state vector  $(\mathbf{x}, \mathbf{r})$  (where  $\mathbf{x}$  corresponds to particle properties like size, density, etc. also known as internal coordinates and,  $\mathbf{r}$  corresponds to spatial position or external coordinates) dispersed in a continuous phase defined by a phase vector  $\mathbf{Y}(\mathbf{r}, t)$  (which again is a function of all such vectors which denote the phase properties at various locations) is denoted by  $f(\mathbf{x}, \mathbf{r}, t)$ . Hence it gives the particle characteristics in property and space domains. Let  $h(\mathbf{x}, \mathbf{r}, \mathbf{Y}, t)$  denote the birth rate of particles per unit volume of particle state space, so the number conservation can be written as<sup>[1]</sup>

$$\frac{d}{dt} \int_{\Omega_{\mathbf{x}}(t)} dV_{\mathbf{x}} \int_{\Omega_{\mathbf{r}}(t)} dV_{\mathbf{r}} f(\mathbf{x}, \mathbf{r}, t) = \int_{\Omega_{\mathbf{x}}(t)} dV_{\mathbf{x}} \int_{\Omega_{\mathbf{r}}(t)} dV_{\mathbf{r}} h(\mathbf{x}, \mathbf{r}, \mathbf{Y}, t)$$

This is a generalized form of PBE.<sup>[1]</sup>

## Solution to PBE

Monte Carlo methods <sup>[2]</sup>, <sup>[3]</sup> discretization methods and moment methods <sup>[2]</sup><sup>[3]</sup><sup>[4]</sup><sup>[5]</sup><sup>[6]</sup> are mainly used to solve these equations. The choice depends on the application and computing infrastructure.

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# 12

## Population Balance Models

**Keywords: Models, Population Balance**

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All the previous models discussed in last lectures have been based on the basic mass, momentum and energy balances. The equations that involve dispersion are most representative of a process when the mixing is on a relatively small scale. For mixing in stirred tanks (in turbulence), dispersion models are not effective. Certain categories of models which cannot successfully be treated within the framework of the models based on transport phenomena can be treated by the population balance concept.

The application of population -balance principles to the modelling of flow and mixing characteristics in vessels was formally organized by Danckwert's. He defined certain distribution functions for the residence time of the fluid elements in a process vessel. RTD functions give information about the fraction of the fluid that spends a certain time in a process vessel. RTD models do not give point- to – point variation of the dependent variables. Danckwert's description was based on macroscopic lumped-population-balance. Population balance models represents macro-mixing that are sufficient to give adequate estimates of the behavior of the process.

### Description of Flow Pattern's in Process Vessel's

There are two types of ideal flow patterns in our process vessels;

1. Plug flow pattern – Piston Flow
2. Bochmix Flow Pattern – Perfectly mixed

Plug flow occurs when the fluid velocity is uniform over the entire cross-section of the vessel.

Perfect mixing assumes that the vessel contents are completely homogenous down to a molecular scale.

Between these two extreme's lie flow patterns in actual process.

In channeling (known as by-passing) some elements of fluid slip on pass through the vessel considerably faster than others do. Channeling may be found in flow through purely packed

vessels, through vessels of small length-to-diameter ratios or through heat exchanger's with proper baffling.

Stagnant packets (dead space) may occur in header's, at the base of pressure gauges, or in the odd-shaped corner's, represents regions with extremely pure contacting.

## AGE DISTRIBUTION FUNCTIONS

The residence time of a fluid element is the time that elapses from the time the element enters the vessel to the time it leaves it. The age of a fluid element at a given instant of time is the time that elapses between the element's entrance into the vessel and the given instant, and is of course, less than or equal to the residence time. The age is equal to the residence time for those molecules that are just leaving the vessel.

### INTERNAL AGE DISTRIBUTION OF A FLUID IN A CLOSED VESSEL

The functional notation  $I(t)$  will be used for the internal age distribution frequency of fluid elements in a vessel.  $I(t)$  have the unit- fraction of ages per unit time.

$$\int_0^{\infty} I(t) dt = 1 \quad \dots 12.1$$

The time  $t=0$  refers to an arbitrary initial time and not the start of the flow of fluid into the vessel. In physical terms, eq. (12.1) states that all fluid has an age between 0 and  $\infty$ .

As a consequence of the above, the fraction of vessel contents younger than a specified age  $t$  is,

$$\int_0^t I(t') dt' = 1 \quad \dots 12.2$$

While the fraction older than  $t$  is  $\int_t^{\infty} I(t') dt'$ ,

$$\int_t^{\infty} I(t') dt' = 1 - \int_0^t I(t') dt' \quad \dots 12.3$$

### Age Distribution of the Exit Stream; The Residence Distribution of fluid in a closed vessel, $E(t)$

The function  $E(t)$  is the age distribution frequency of the fluid elements leaving the vessel and has the units of fraction of ages per unit time. The fraction of exit ages itself is  $E(t) t$ . The function is normalized to;

$$\int_0^{\infty} E(t) dt = 1 \quad \dots 12.4$$

The fraction of fluid in the exit stream younger than age  $t$  is

$$\int_0^t E(t') dt' = 1 \quad \dots 12.5$$

while the fraction of material older than  $t$  is

$$\int_t^{\infty} E(t') dt' = 1 - \int_0^t E(t') dt' = 1 \quad \dots 12.6$$

The mean RTD is found from the first moment

$$\bar{t} = \int_0^{\infty} tE(t) dt = \frac{V}{Q} \quad \dots 12.7$$

$$\frac{V}{Q} = \bar{t} = \tau$$

In a similar fashion the mean age of fluid elements inside the vessel is,

$$\bar{t}_1 = \int_0^{\infty} tI(t) dt \quad \dots 12.8$$

### **Intensity Function, $\Lambda(t)$**

It is defined as fraction of fluid in the vessel of age  $t$  that will leave at time between  $t$  and  $t + \Delta t$ . The intensity function is useful in detecting the existence of dead space and by passing.

## Relations between Age Distribution Functions:

The three age distribution frequency functions described above are related through the unsteady state macroscopic age population balance. The balance is made in the units of time. The input age distribution is zero. The general macroscopic population balances can be used to relate  $E(t)$  and  $I(t)$ , but a simpler alternate method is used here.

Consider a constant volume vessel,  $V$ , with constant flow rate  $Q$  and call all fluid entering the vessel at  $t > 0$  new fluid. The existing contents at  $t = 0$  are the old fluid. At some time  $t$ , eq. 12.2 gives the function of new fluid.

$$\text{Amount of new fluid in vessel} = V \int_0^t I(t') dt'$$

Eq. 12.6 gives the fraction of out flowing fluid at any instant of time, that has an age greater than  $t$ ; the amount of old fluid that has left the vessel during all times from 0 to  $t$  is

$$\text{Amount of old fluid gone from the vessel} = \int_0^t Q dt' \int_{t'}^{\infty} E(t'') dt''$$

Then by a simple balance, the old fluid left must have been replaced by the new fluid,

$$V \int_0^t I(t') dt' = \int_0^t Q dt' \int_{t'}^{\infty} E(t'') dt'' \quad \dots 12.9$$

Differentiation of both sides of eq. 12.9 with respect to time with the introduction of eq. 12.7 gives,

$$\bar{t}I(t) = \int_t^{\infty} E(t') dt' = 1 - \int_0^t E(t') dt' \quad \dots 12.10$$

Differentiating once more,

$$E(t) = -\bar{t} \frac{dI(t)}{dt} \quad \dots 12.11$$

The intensity function can also be related to the  $E(t)$  and  $I(t)$  functions from

$$\left\{ \begin{array}{l} \text{Amount of fluid} \\ \text{leaving between} \\ t \text{ and } t+\Delta t \end{array} \right\} = \left\{ \begin{array}{l} \text{Amount not leaving} \\ \text{before time } t \end{array} \right\} \left[ \begin{array}{l} \text{Fraction of age } t \\ \text{that will leave} \\ \text{between times } t, t+ \Delta t \end{array} \right]$$

$$QE(t)dt = [VI(t)][\Lambda(t)dt] \quad \dots 12.12$$

$$\Lambda(t) = \frac{1}{\bar{t}} \frac{E(t)}{I(t)} = - \frac{d[\ln \bar{t} I(t)]}{dt} \quad \dots 12.13$$

Each of the age distribution functions can be expressed in dimensionless form.

$$\theta = \frac{t}{\bar{t}} \quad \dots 12.14$$

Thus,  $E(\theta)d\theta = E(t)dt, I(\theta)d\theta = I(t)dt$

$$E(\theta) = \bar{t}E(t) \quad \dots 12.15$$

$$I(\theta) = \bar{t}I(t) \quad \dots 12.16$$

$$\Lambda(\theta) = \bar{t}\Lambda(t) \quad \dots 12.17$$

$$E(\theta) = - \frac{d I(\theta)}{d\theta} I(\theta) \quad \dots 12.18$$

$$\Lambda(\theta) = \frac{E(\theta)}{I(\theta)} = - \frac{\ln(\theta)}{\theta} \quad \dots 12.19$$

# Population Balances in Engineering

- Crystallization, Precipitation
- Coagulation, Agglomeration and Flocculation
- Particle Breakage, Particle Attrition
- Dissolution/Leaching
- Cell Growth
  - Fermentation
  - Tissue Engineering

# Constant Stirred Tank Crystallizer (MSMPR)

- On-line Analysis
  - Feed Flow Rate Controlled
  - Product Flow Rate Controlled
  - Liquid Level Controlled
  - Reactor
    - pH -  $\checkmark$
    - Ionic Strength -  $\checkmark$
    - Temperature -  $\checkmark$
    - Stirrer RPM -  $\checkmark$
    - Stirrer Torque -  $\checkmark$
    - Heat Balance -  $\checkmark$
  - Particle Size
    - Movie/Stills of Particles
- Off-line Analysis
  - Beckman-Coulter LS230
  - AA + ICP of feed and output
  - Yield





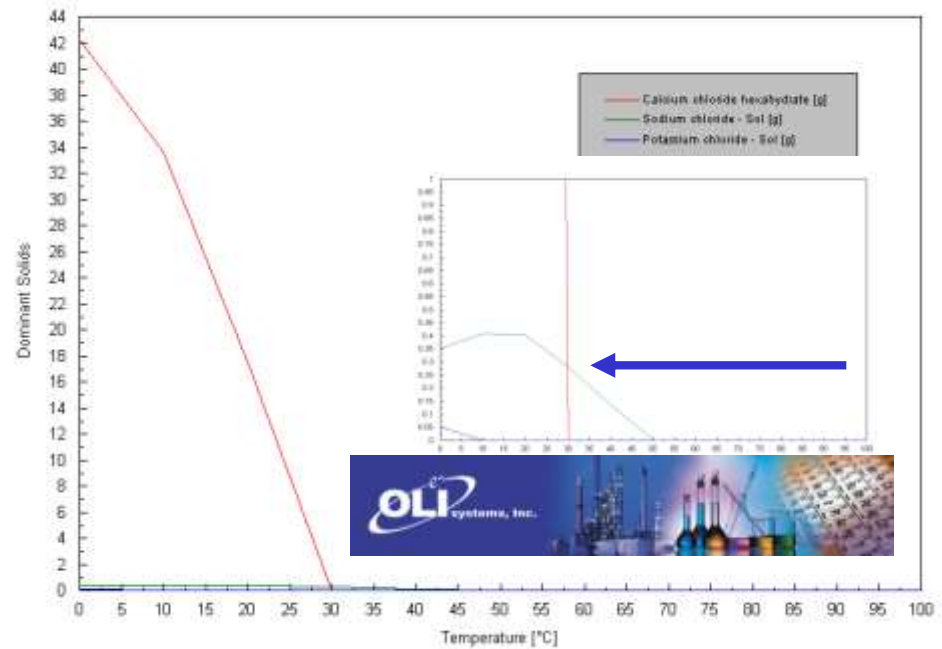
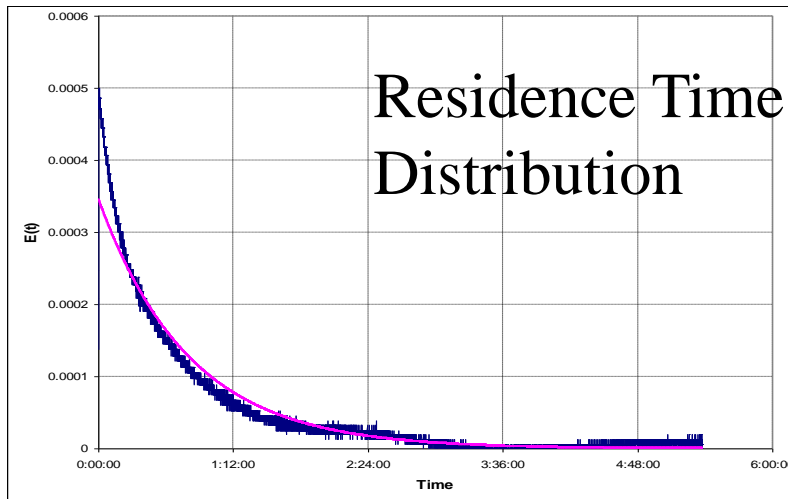
# CaCl<sub>2</sub>-NaCl-KCl

- 45% CaCl<sub>2</sub>
  - NaCl+KCl

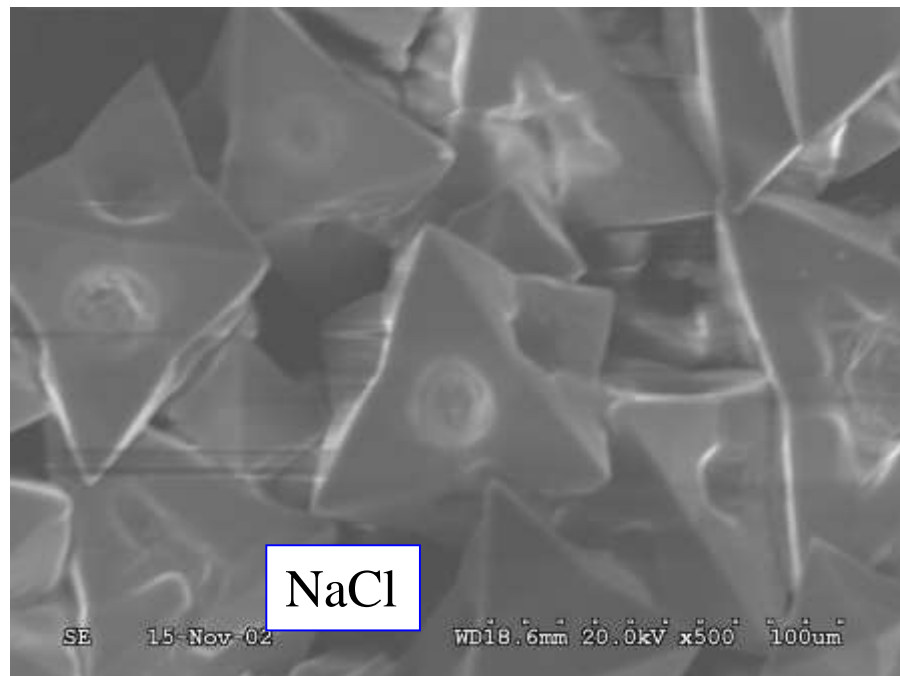
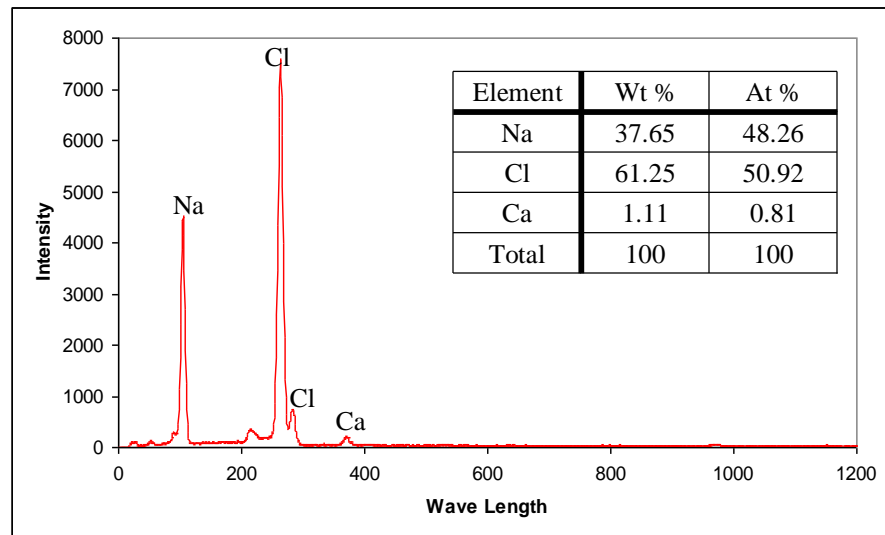
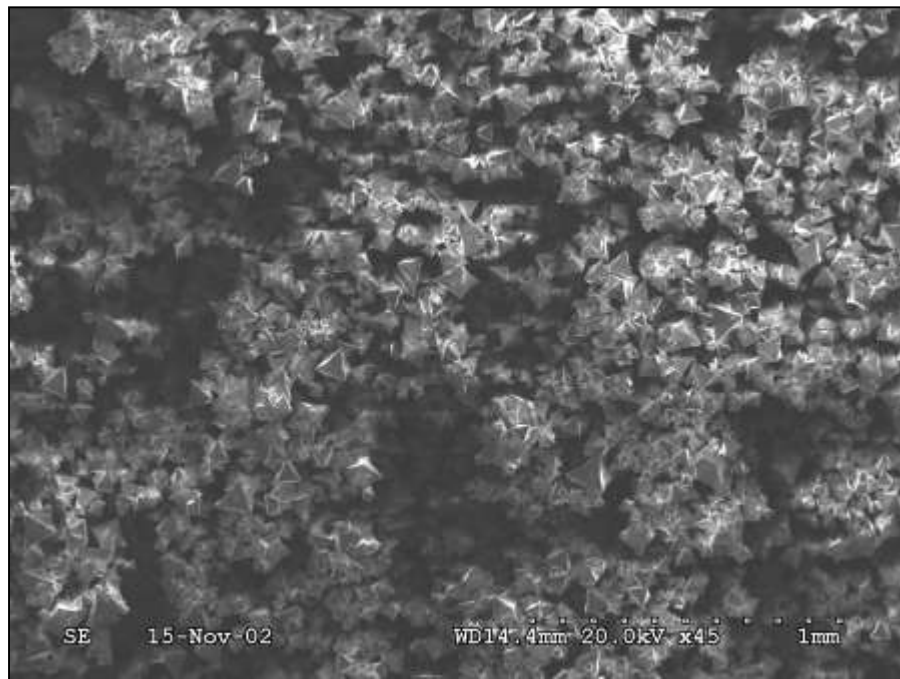


Living.  
Improved daily.

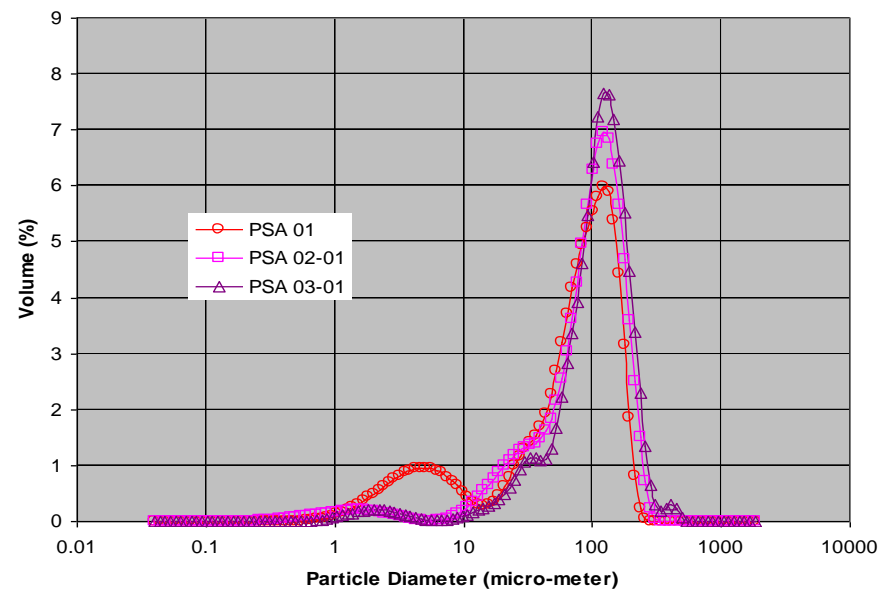
- Baffled Stirred Tank
  - Cooling
  - Impurity Removal



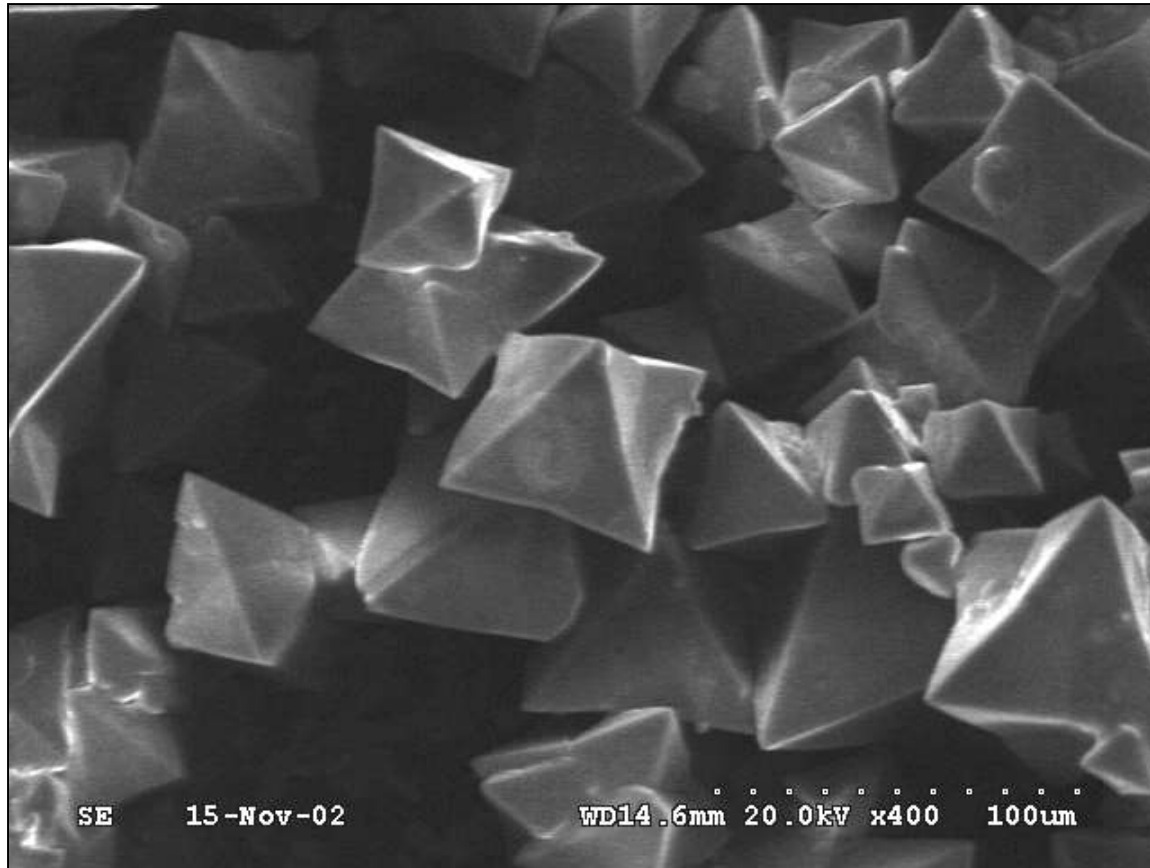
# CaCl<sub>2</sub> Results - Sample #10-01

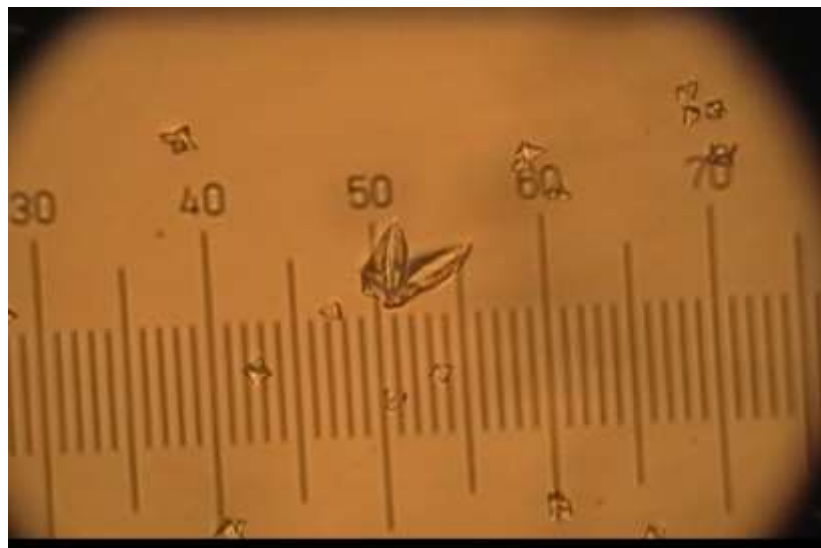


## Particle Size Analysis with Beckman Coulter LS230

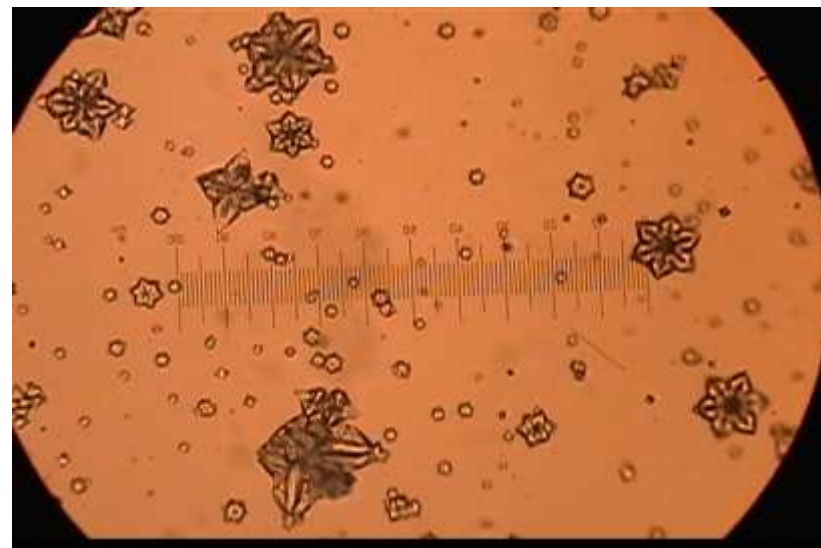


# Sample #11-02

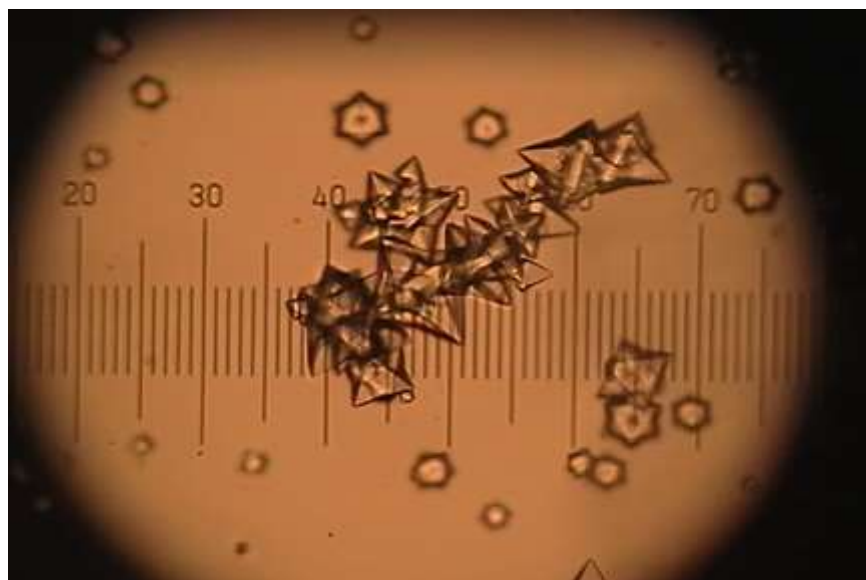




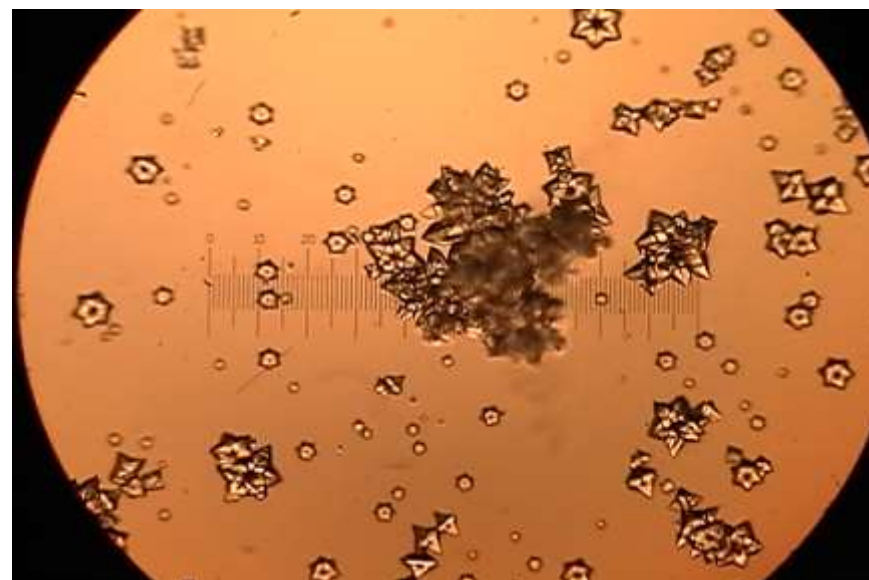
30 min.



90 min.

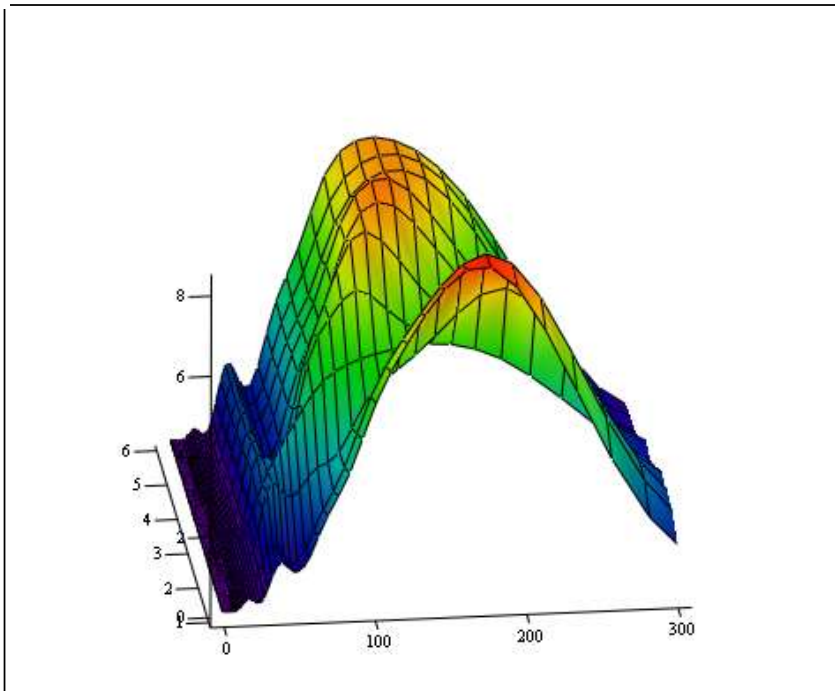


6 hr.

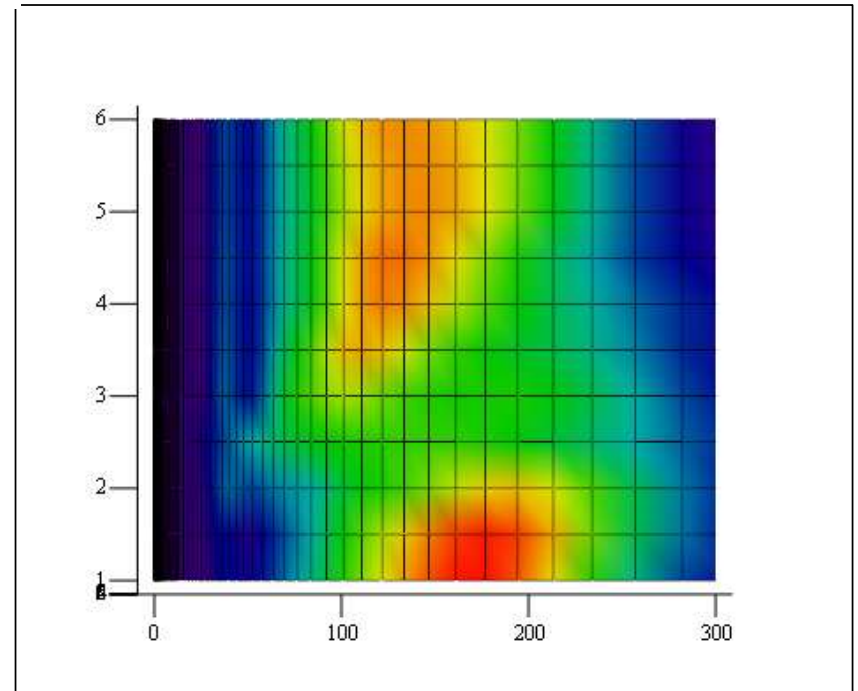


6 hr.

# Particle Size Distribution with Time



(x, y, z)



(x, y, z)

07/25/03

Reaction Temp: 30.1°C, Flow Rate: 45 ml/min, Mixing Power: 287.7 rpm, Reactor Volume: 1400 ml

# Particulate Mechanisms

- Nucleation
  - Heterogeneous
  - Secondary
    - (crystal impact with impeller creates nuclei)
- Particle Growth
  - Diffusion Limit
  - Other mechanisms
- Particle Breakage/Attrition
- Particle Aggregation

## Traditional Population Balance – differential number

$$\begin{aligned} \frac{\partial f_N(x,t)}{\partial t} + \frac{\partial [G(x,t) f_N(x,t)]}{\partial x} &= \text{Birth} - \text{Death} = \\ + \int_x^\infty \zeta(x') b(x') p(x|x') f_N(x',t) dx' - b(x) f_N(x,t) &\quad \text{Breakage} \\ + \frac{1}{2} \int_0^x a(x-x') f_N(x-x',t) f_N(x',t) dx' - f_N(x,t) \int_0^x a(x,x') f_N(x',t) dx' &\quad \text{Aggregation} \end{aligned}$$

with boundary conditions

$$f_N(x,t) = 0 \quad \text{as} \quad x \Rightarrow \infty$$

$$\left. \frac{\partial [G(x,t) f_N(x,t)]}{\partial x} \right|_{x=x_0} = \frac{\partial f_N(x=x_0,t)}{\partial t} \quad \underline{\text{Nucleation Rate}} \quad \text{at size } x_0$$

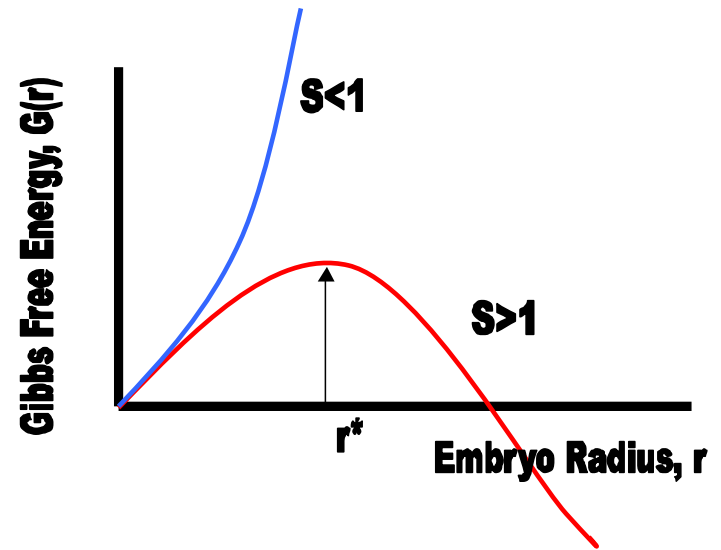
and initial condition

$$f_N(x,t) = f_{N,o}(x) \quad \text{at} \quad t = 0.$$



# Nucleation

- primary homogeneous
  - Classical Nucleation
    - $J(\#/m^3\text{sec})=J_{\text{max}} \exp(-G(r^*)/RT)$
- primary heterogeneous
  - Classical Nucleation + new surface energy
    - $J(\#/m^3\text{sec})=J_{\text{max}} \exp(-G_{\text{het}}(r^*)/RT)$
- secondary
  - $J(\#/m^3\text{sec}) = f(\text{rpm}, S, \text{Solids Mass Fraction})$





# Crystal Growth Rate Mechanisms

$$dR/dt = K * f(S) * g(R)$$

Growth Mechanism	C	f(S)	g(R)	Ref
Diffusion Bulk	$\hat{V}DC_{eq}$	S-1	1/R	1
Mono Surface Nucleation	$\beta_A D d^{-3}$	$\exp[\Delta G_s^*/k_B T]^{\S}$	R <sup>2</sup>	2
Poly Surface Nucleation	$\frac{(Dd/3)}{(N_A C_{eq})^{2/3}}$	$(S-1)^{2/3} \exp[\Delta G_s^*/k_B T]^{\S}$	1	3
Screw Dislocation	$D_s n_{se} \beta / (y_o^2 \rho)$	$S^2/S_1 \tanh(S_1/S)^{\S\S}$	1	4
Heat Conduction	$\hat{V}k_H R T^2/\Delta H_f$	$\ln S^{\S\S\S}$	1/R	-
Chemical Reaction	$\eta \hat{V}DC_{eq}$	S-1	1/R	-

$$\S \Delta G_s^* = \beta L^2 \gamma_e^2 d^2 / (1 \beta_A k_B T \ln S)$$

$$\S\S S_1 = (y_o/y_s) S$$

$$\S\S\S \ln S = \int_{T_0}^T \Delta H_f / (RT^2) dT$$

# Particle Collision Rate due to Turbulence

Regime	size $a_i + a_j$	Collision Velocity $\Delta u(a_i, a_j)$	Collision Frequency Factor $\beta(a_i, a_j, r, t)$
<b>Brownian Diffusion</b>	$0 > a_i + a_j > \eta$		$\frac{2}{3}(a_i + a_j) \left( \frac{k_B T}{\mu} \right) \left( \frac{1}{a_i} + \frac{1}{a_j} \right)$
<b>Turbulent subrange</b>			
<b>Viscous</b>	$\eta > a_i + a_j > 6\eta$	$0.257 \left( \frac{\varepsilon}{\nu} \right)^{1/2} (a_i + a_j)$	<b>1.29</b> $(\langle \varepsilon^{1/2} \rangle / \nu^{1/2}) (a_i + a_j)^3$ Equivalent to <b>(4/3)</b> $\gamma (a_i + a_j)^3$
<b>Transition</b>	$6\eta \leq a_i + a_j < 25\eta$	$0.471 \frac{\varepsilon^{5/12}}{\nu^{1/4}} (a_i + a_j)^{2/3}$	<b>2.36</b> $\langle \varepsilon^{5/12} \rangle \nu^{-1/4} (a_i + a_j)^{8/3}$
<b>Inertial</b>	$25\eta \leq a_i + a_j < L/2$	$1.37 (\varepsilon)^{1/3} (a_i + a_j)^{1/3}$	<b>6.87</b> $\langle \varepsilon^{1/3} \rangle (a_i + a_j)^{7/3}$
<b>Macro</b>	$L/2 > a_i + a_j \sim L$	$\sqrt{2} (\varepsilon L)^{1/3}$	<b>7.09</b> $\langle \varepsilon^{1/3} \rangle L^{1/3} (a_i + a_j)^2$

# Breakage Kinetics

- Breakage Rate =

Collision Frequency \* Target Efficiency \* Breakage Probability

Collisions

Collision Efficiency

Particle-wall

Particle-impeller

Particle-baffle

Particle-Particle due to either diffusion or turbulence

# Crystallizers are Complicated

- Size Increases by
  - Aggregation
  - Growth
- Size Decreases by
  - Breakage
  - Dissolution
- Number of Particles Increases by
  - Nucleation
  - Breakage
  - Less Aggregation

HOW TO DECOUPLE?

Do not try to solve all at once!

# Decomposition of the Problem

- Using standard operating conditions for tank,
  - Study aggregation separately
  - Study breakage separately
- Focus on aggregation and breakage under batch conditions after crystallization has been run to steady state

## Traditional Population Balance – differential number

$$\begin{aligned} \frac{\partial f_N(x,t)}{\partial t} + \frac{\partial [G(x,t)f_N(x,t)]}{\partial x} &= \text{Birth} - \text{Death} = \\ &+ \int_x^\infty \zeta(x')b(x')p(x|x')f_N(x',t)dx' - b(x)f_N(x,t) \quad \text{Breakage} \\ &+ \frac{1}{2} \int_0^x a(x-x')f_N(x-x',t)f_N(x',t)dx' - f_N(x,t) \int_0^\infty a(x,x')f_N(x',t)dx' \quad \text{Aggregation} \end{aligned}$$

with boundary conditions

$$f_N(x,t) = 0 \quad \text{as} \quad x \Rightarrow \infty$$

$$\left. \frac{\partial [G(x,t)f_N(x,t)]}{\partial x} \right|_{x=x_o} = \frac{\partial f_N(x=x_o,t)}{\partial t} \quad \text{Nucleation Rate at size } x_o$$

and initial condition

$$f_N(x,t) = f_{N,o}(x) \quad \text{at} \quad t = 0.$$

# Stieltjes Formulation<sup>1</sup> – Cumulative Mass

Ramkrishna, D., “Population Balances,”

Academic Press, 2000, p.56 and private communications.

$$\frac{\partial F_m(x, t)}{\partial t} + \int_0^x x' \frac{\partial}{\partial x'} \left[ \frac{g_M(x', t)}{x'} \frac{\partial F_m(x, t)}{\partial x'} \right] dx' = \hat{B} - \hat{D} =$$

$$\frac{\partial F_m(x, t)}{\partial t} + g_M(x', t) \frac{\partial F_m(x, t)}{\partial x'} - \int_0^x \frac{g_M(x', t)}{x'} \partial_x F_m(x', t) = \hat{B} - \hat{D} =$$

$$\int_x^\infty b(x') P_m(x|x') \partial_x F(x', t) \quad \text{Breakage}$$

$$- \mu_1(t) \int_{s=0}^x \int_{u=x-s}^\infty \frac{a(s, u)}{u} \partial_x F(u, t) \partial_x F(s, t) \quad \text{Agglomeration}$$

with boundary conditions

$$\left. \frac{\partial F_m(x, t)}{\partial t} \right|_{x=x_0} \quad \text{Mass Nucleation Rate at size } , x_0$$

$$F_m(x, t) = 0 \quad \text{at} \quad x = 0, \quad F_m(x, t) = 1 \quad \text{at} \quad x = \infty$$

and initial condition

$$F_m(x, t) = F_{m,0}(x) \quad \text{at} \quad t = 0$$

# Sinc Approximation

$$F_m(x, t) \approx \sum_{j=-N}^N \sum_{k=-M}^M \operatorname{sinc}\left(\frac{\varphi(x)}{h_x} - j\right) \operatorname{sinc}\left(\frac{\phi(t)}{h_t} - k\right) F_m(x_j, t_k)$$

$$\varphi(x) = \ln(x)$$

$$x_j = \varphi^{-1}(jh_x) = \exp(jh_x)$$

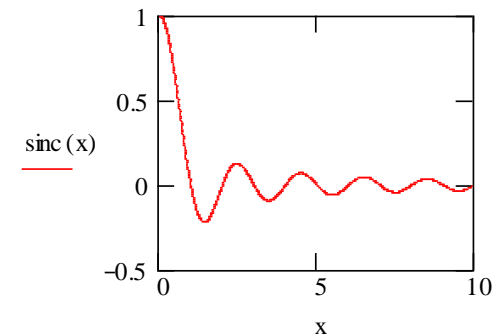
$$h_x = \frac{c_1}{\sqrt{N}}$$

$$\phi(t) = \ln[t/(T-t)]$$

$$t_k = \phi^{-1}(kh_t) = \frac{T \exp(kh_t)}{1 + \exp(kh_t)}$$

$$h_t = \frac{c_2}{\sqrt{M}}$$

$$\operatorname{sinc}(z) = \frac{\sin(\pi z)}{\pi z}$$





# Final Formulation

$$\begin{aligned}
 F_m(x, t) = & F_{m,o}(x, t = 0) + \\
 & - \int_0^t g_M(x', t') \frac{\partial F_m(x', t')}{\partial x'} dt' + \int_0^t \int_0^x \frac{g_M(x', t')}{x'} \partial_x F_m(x', t') dt' \quad \text{growth} \\
 & + \int_0^t \int_x^\infty b(x') P_m(x|x') \partial_x F(x', t') dt' \quad \text{Breakage} \\
 & - \int_0^t \mu_1(t') \int_{s=0}^x \int_{u=x-s}^\infty \frac{a(s, u)}{u} \partial_x F(u, t') \partial_x F(s, t') dt' \quad \text{Agglomeration}
 \end{aligned}$$


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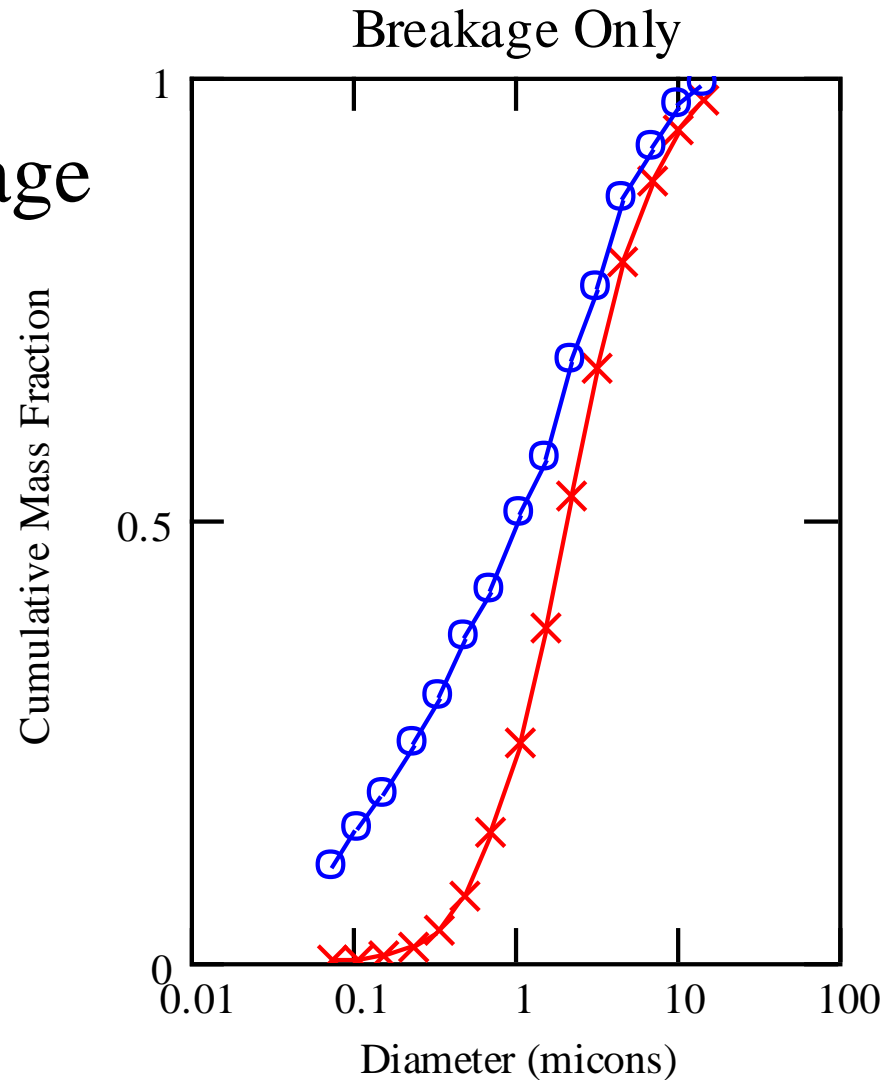
$$\begin{aligned}
 \overline{\overline{F_m}} = & \overline{\overline{F_{m,o}}} + \\
 & - \overline{\overline{At_{I+}}} \bullet \overline{\overline{[g_M]}} \bullet \overline{\overline{[Ax_D \bullet F_m]}} + \overline{\overline{At_{I+}}} \bullet \overline{\overline{Ax_{I+}}} \bullet \overline{\overline{\left[ \frac{g_M}{\bar{x}} \right]}} \bullet \overline{\overline{[Ax_D \bullet F_m]}} \quad \text{Growth} \\
 & + \overline{\overline{At_{I+}}} \bullet \overline{\overline{Ax_{I+}}} \bullet \left\{ \overline{\overline{[bP_m]}} \bullet \overline{\overline{[Ax_D \bullet F_m]}} \right\} \quad \text{Breakage} \\
 & - \overline{\overline{At_{I+}}} \bullet \left[ \overline{\overline{diag[\mu_1(\bar{t})]} \overline{\overline{X}} \overline{\overline{P}} \right] \quad \text{Agglomeration}
 \end{aligned}$$

# Sinc Results – Batch

- Power Law Breakage

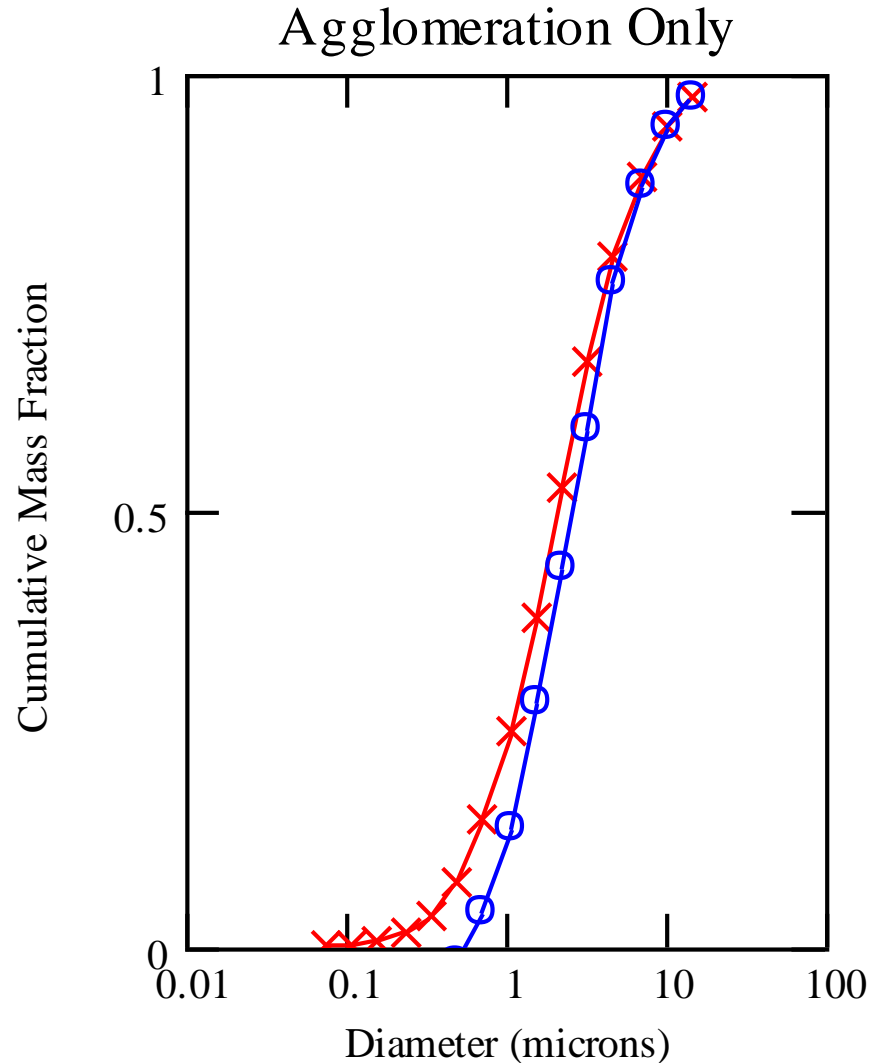
- $S(x)=x^{1.2}$

- $Kt=0.1$



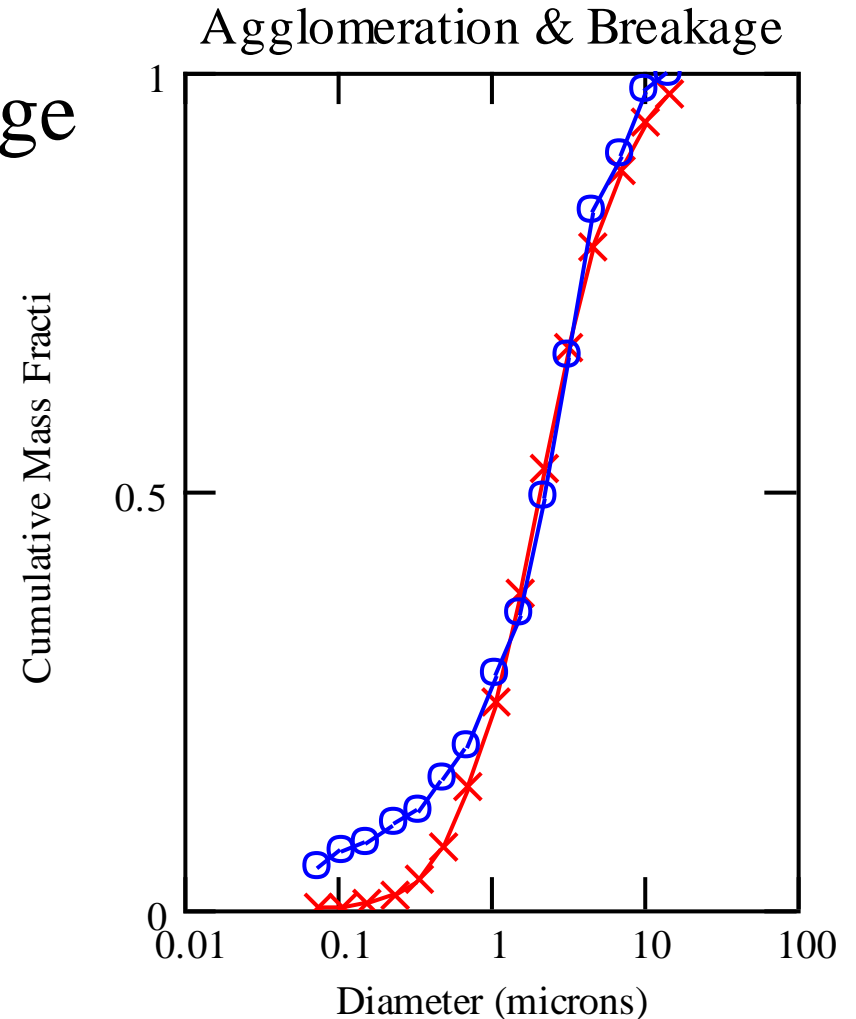
# Sinc Result -Batch

- Aggregation Only
  - Sum Kernal
  - $Kt=0.5$
  - $\mu = 0.1$



# Sinc Result- Batch

- Aggregation & Breakage
  - Compensating Effects



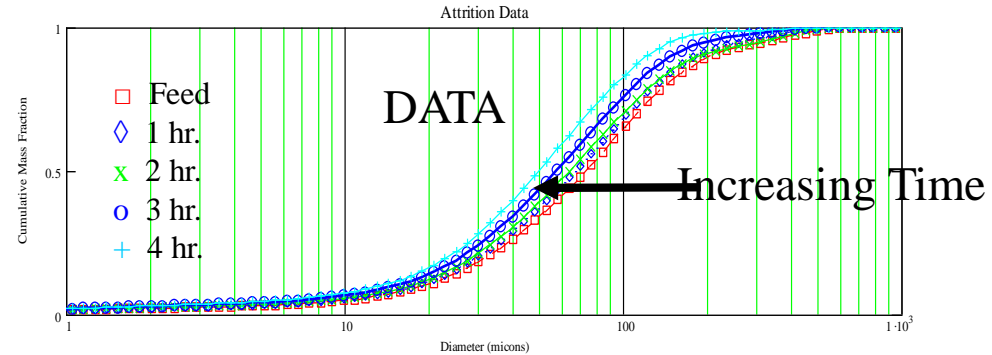
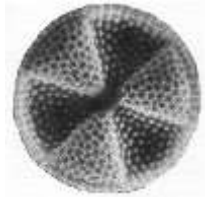
# Need to Know

- Aggregation Rate Constant
  - Breakage Rate Constant
  - Breakage Selectivity
  - Breakage Daughter Distribution
- 
- Where are we going to get these?

# Focused Experimentation

- Using standard operating conditions for tank,
  - Study aggregation separately
  - Study breakage separately
- Forced Fitting of the Data

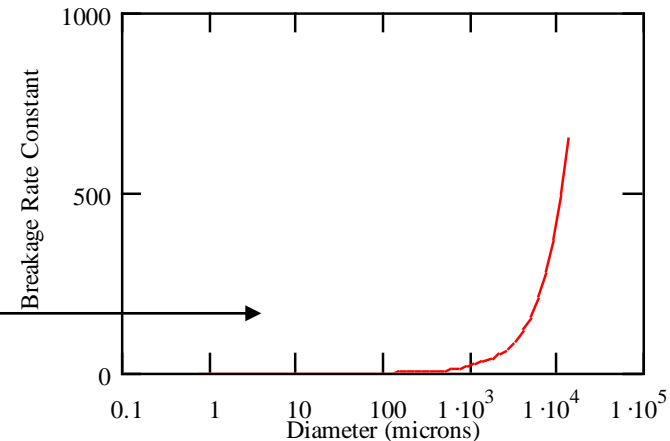
# Diatomite Breakage



- Batch Experiment for Breakage
  - 10% wt. Commercial Diatomite
- 1.4 L baffled batch stirred tank
  - 300 rpm
  - Sampling every hr.
- Progeny Function
- Rate Function

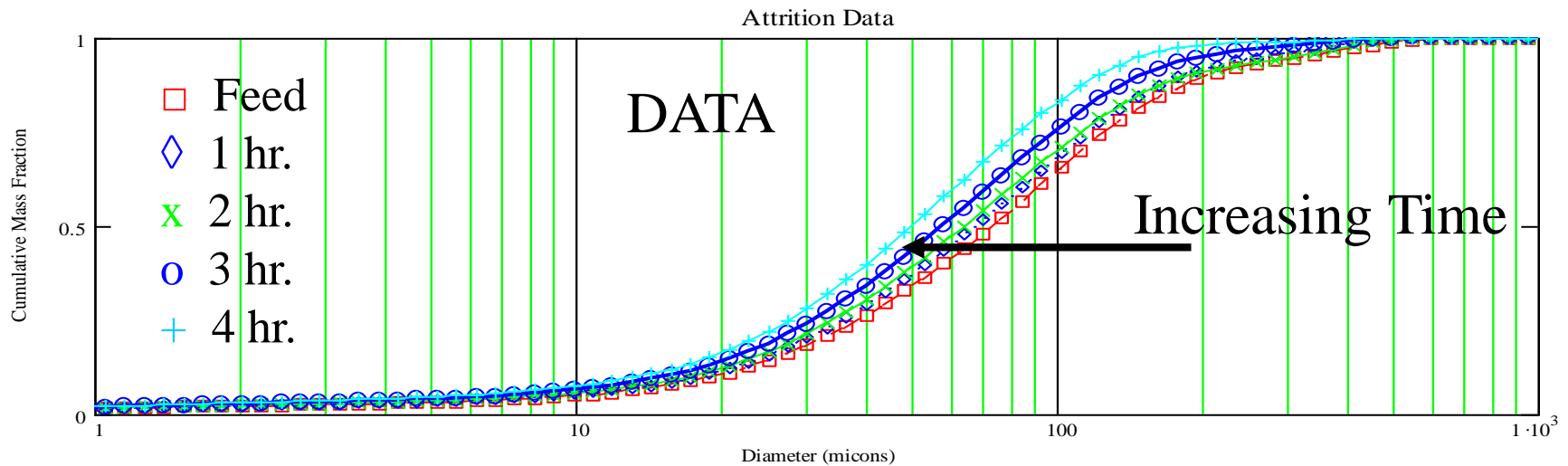
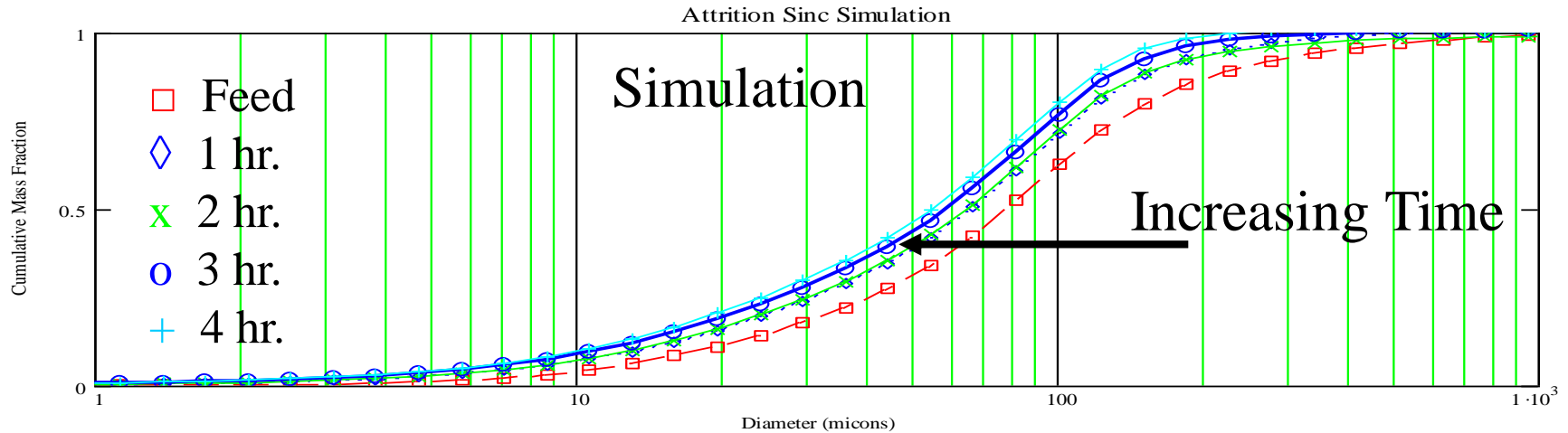
$$P_m(x|x') = \frac{1 - \exp(-x/x')}{1 - \exp(-1)}$$

$$b(x) := \left( \frac{x}{2 \cdot x_0} \right)^{1.4} \cdot \Phi(x - 2 \cdot x_0)$$



# Diatomite Attrition

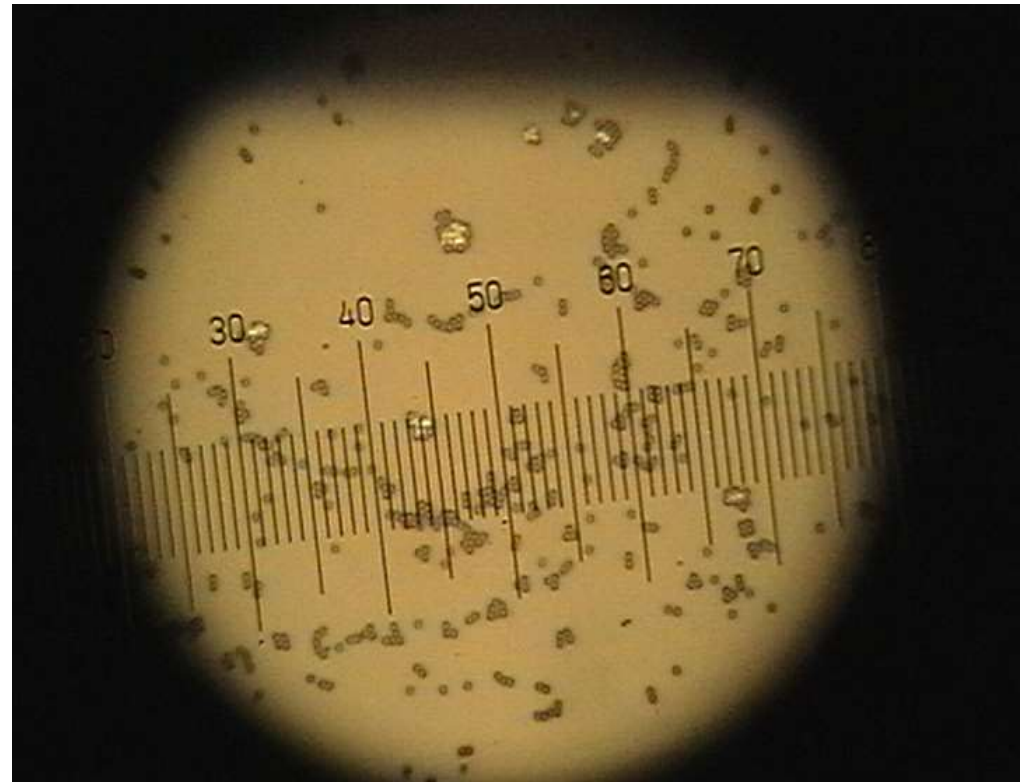
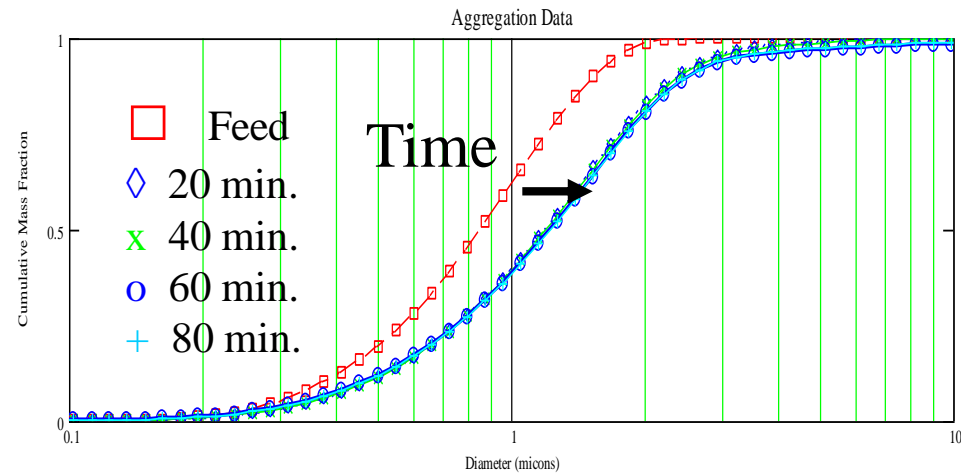
$T \cdot hr = 240 \text{ min}$



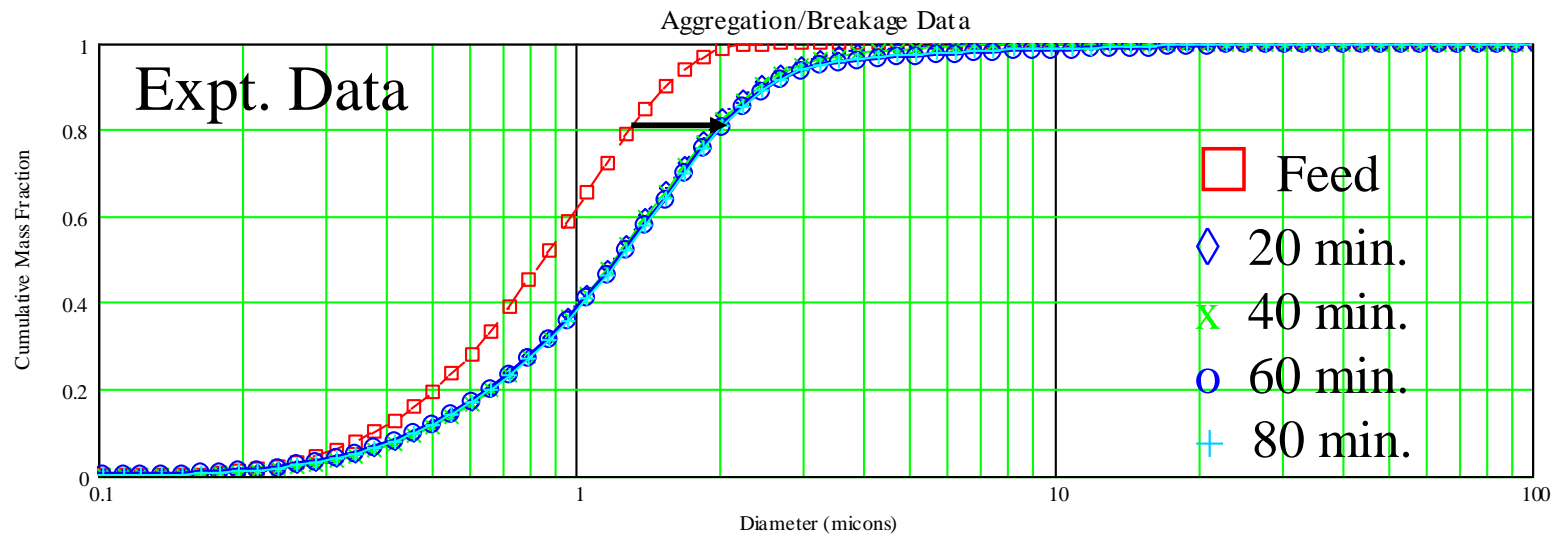
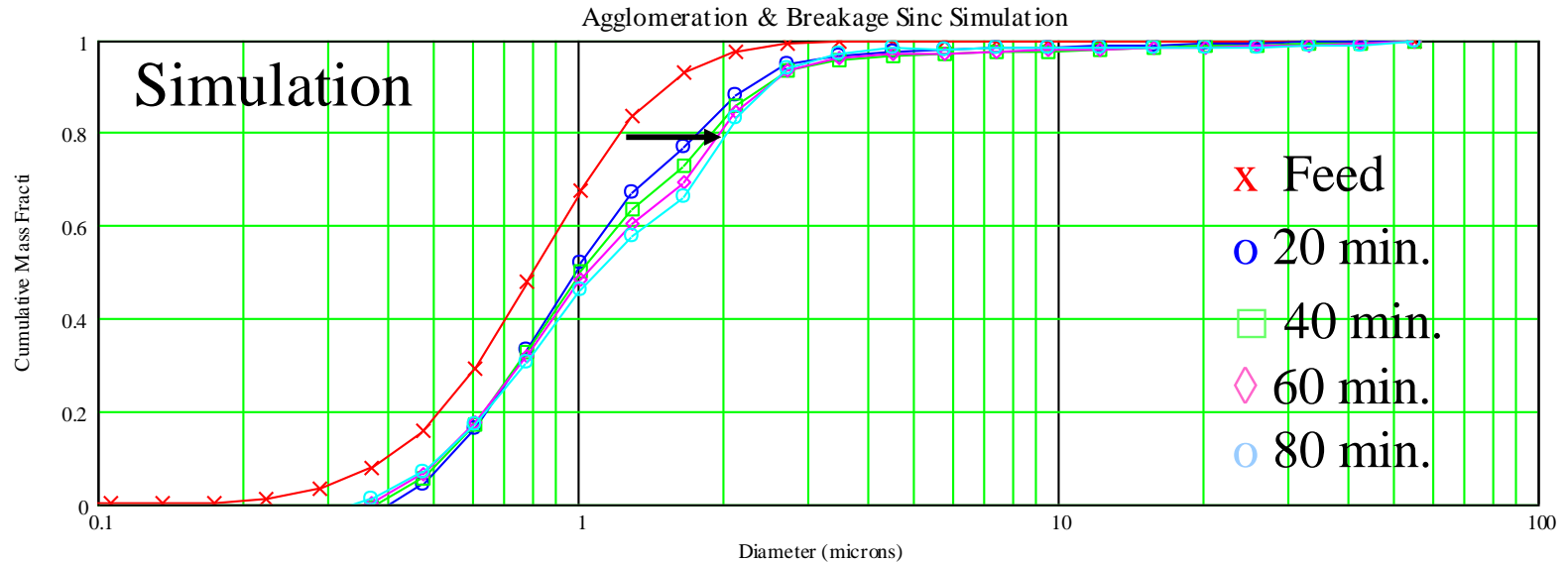


# Polystyrene Latex Aggregation

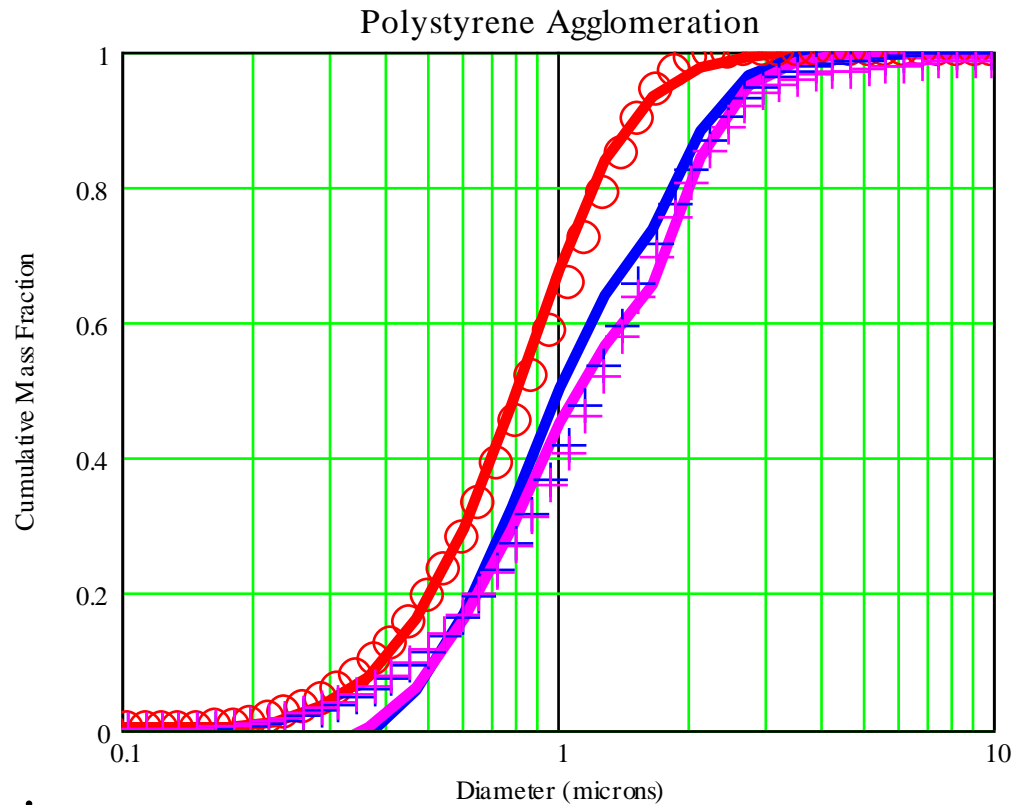
- Batch Experiment for Aggregation
  - 1% wt. Commercial Polystyrene Latex (1 micron) in  $10^{-3}$  M NaCl
- 1.4 L baffled batch stirred tank
  - 100 rpm
  - Sampling every 10 min.
- Dynamic Balance
  - Breakage (diatomite expt.)
  - Aggregation
    - $a(x,y) = x + y$



# Dynamic Balance for Polystyrene Latex



# Polystyrene Latex Aggregation



Data = Points

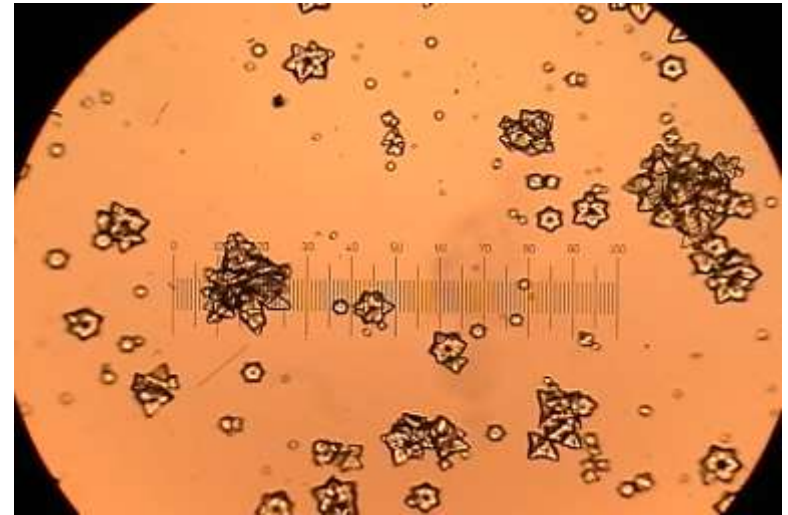
Sinc Simulation = Lines

# Focus on Aggregation and Breakage after Crystallization

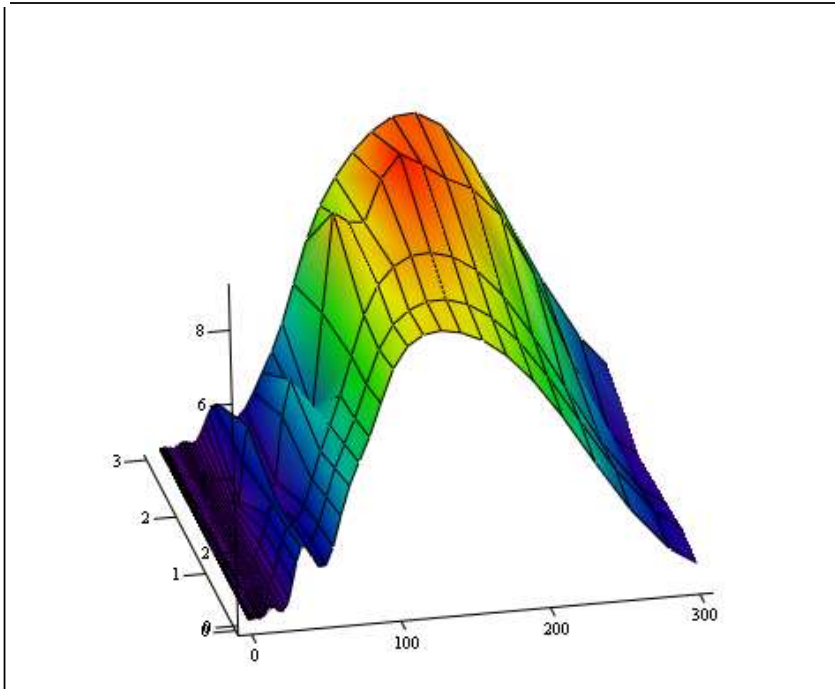
- Stop Feed – Batch Tank
  - Constant RPM, Constant Temperature
    - Supersaturation is Constant
      - No nucleation
      - No particle growth
  - Measure Changes in Particle Size Distribution

# NaCl Aggregation

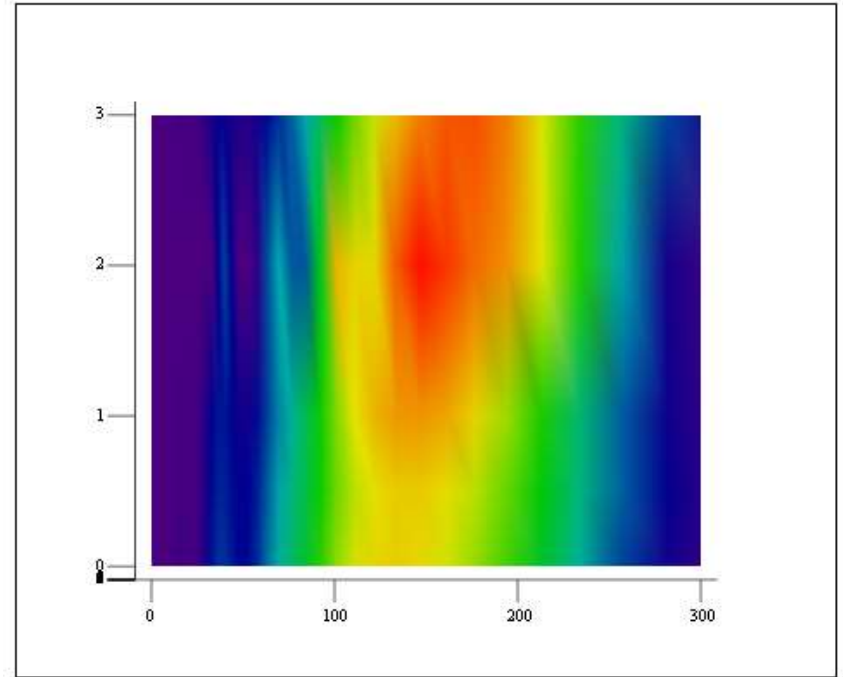
- 1.4 L Batch, Baffled Stirred Tank
  - 287 rpm
- Charge from Crystallization
  - 2% NaCl particles in 47% CaCl<sub>2</sub> solution
  - Aggregation/Attrition over 3 hrs.



# Aggregation/Breakage Data



(x, y, z)

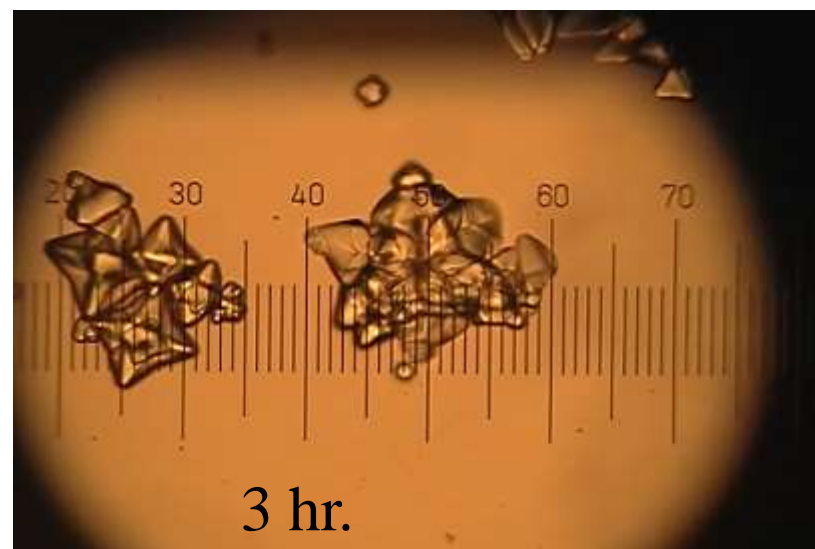
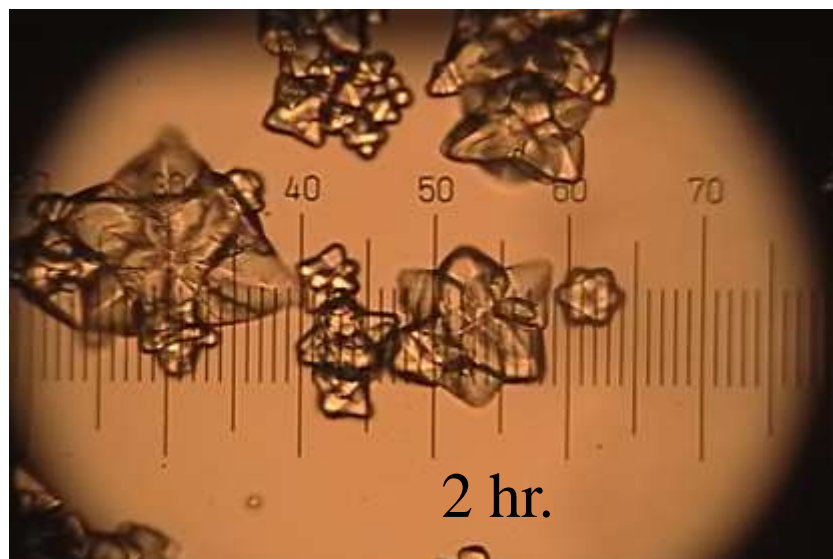
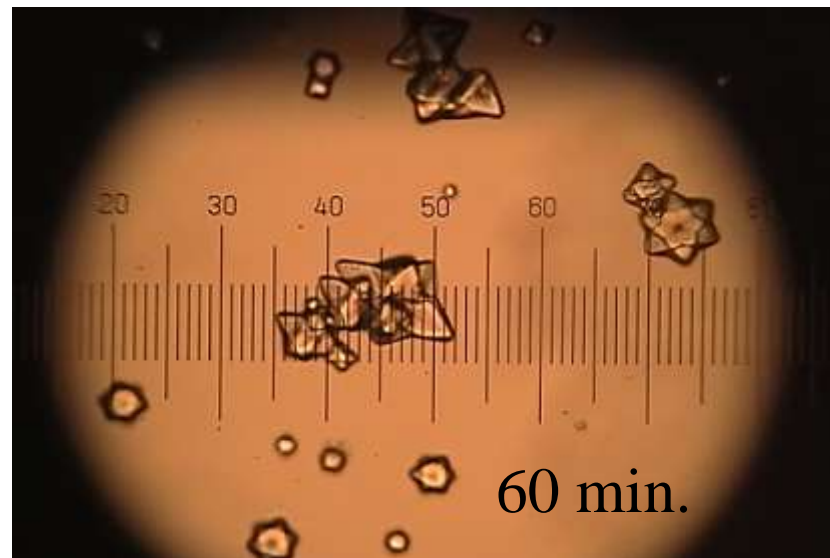
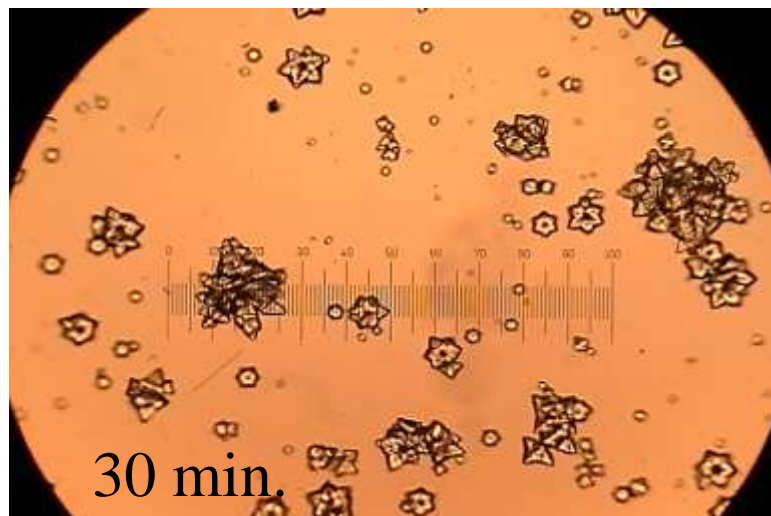


(x, y, z)

07/25/03

Reaction Temp: 30.1°C, Flow Rate: 45 ml/min, Mixing Power: 287.7 rpm, Reactor Volume: 1400 ml

# Aggregation/Breakage



## Traditional Population Balance – differential number

$$\begin{aligned} \frac{\partial f_N(x,t)}{\partial t} + \frac{\partial [g(x,t)f_N(x,t)]}{\partial x} &= \text{Birth} - \text{Death} = \\ &+ \int_x^\infty \zeta(x')b(x')p(x|x')f_N(x',t)dx' - b(x)f_N(x,t) \quad \text{Breakage} \\ &+ \frac{1}{2} \int_0^x a(x-x')f_N(x-x',t)f_N(x',t)dx' - f_N(x,t) \int_0^\infty a(x,x')f_N(x',t)dx' \quad \text{Aggregation} \end{aligned}$$

with boundary conditions

$$f_N(x,t) = 0 \quad \text{as} \quad x \Rightarrow \infty$$

$$\left. \frac{\partial [g(x,t)f_N(x,t)]}{\partial x} \right|_{x=x_o} = \frac{\partial f_N(x=x_o,t)}{\partial t} \quad \text{Nucleation Rate at size } x_o$$

and initial condition

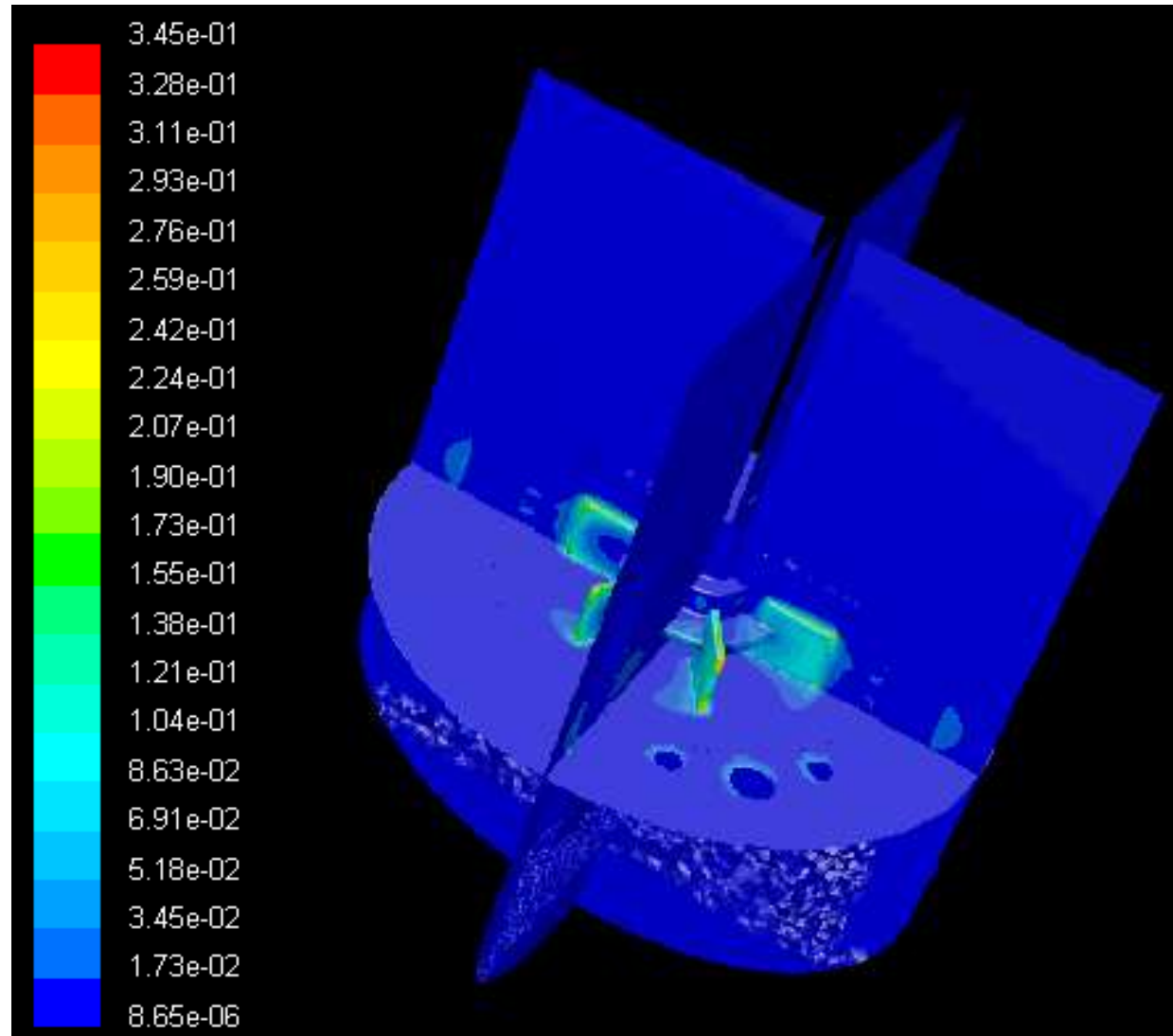
$$f_N(x,t) = f_{N,o}(x) \quad \text{at} \quad t = 0.$$



# Particle Collision Rate due to Turbulence

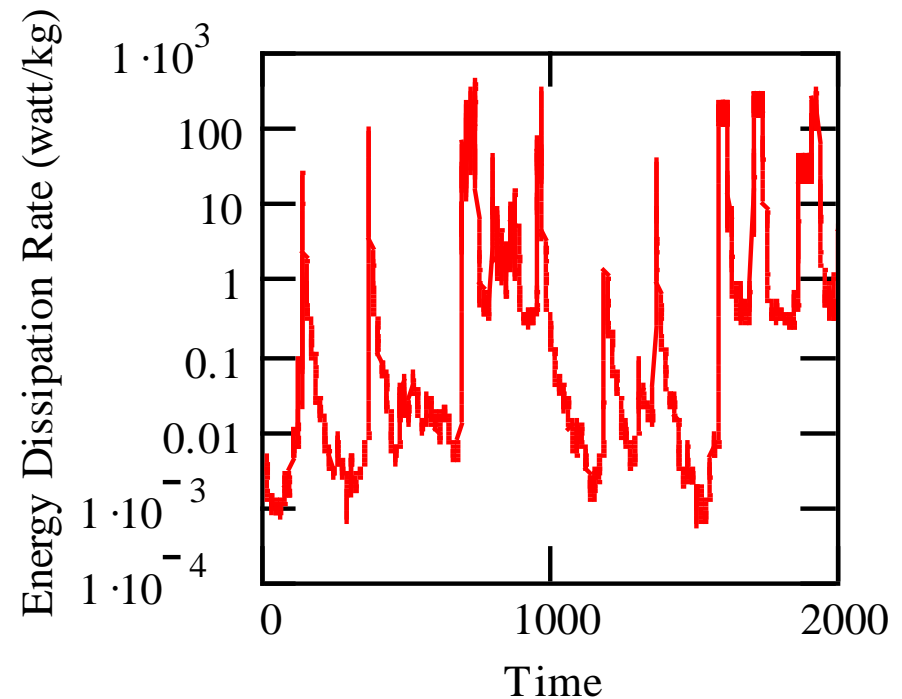
Regime	size $a_i + a_j$	Collision Velocity $\Delta u(a_i, a_j)$	Collision Frequency Factor $\beta(a_i, a_j, r, t)$
<b>Brownian Diffusion</b>	$0 > a_i + a_j > \eta$		$\frac{2}{3}(a_i + a_j) \left( \frac{k_B T}{\mu} \right) \left( \frac{1}{a_i} + \frac{1}{a_j} \right)$
<b>Turbulent subrange</b>			
<b>Viscous</b>	$\eta > a_i + a_j > 6\eta$	$0.257 \left( \frac{\varepsilon}{\nu} \right)^{1/2} (a_i + a_j)$	<b>1.29</b> $(\langle \varepsilon^{1/2} \rangle / \nu^{1/2}) (a_i + a_j)^3$ Equivalent to <b>(4/3) <math>\gamma (a_i + a_j)^3</math></b>
<b>Transition</b>	$6\eta \leq a_i + a_j < 25\eta$	$0.471 \frac{\varepsilon^{5/12}}{\nu^{1/4}} (a_i + a_j)^{2/3}$	<b>2.36</b> $\langle \varepsilon^{5/12} \rangle \nu^{-1/4} (a_i + a_j)^{8/3}$
<b>Inertial</b>	$25\eta \leq a_i + a_j < L/2$	$1.37 (\varepsilon)^{1/3} (a_i + a_j)^{1/3}$	<b>6.87</b> $\langle \varepsilon^{1/3} \rangle (a_i + a_j)^{7/3}$
<b>Macro</b>	$L/2 > a_i + a_j \sim L$	$\sqrt{2} (\varepsilon L)^{1/3}$	<b>7.09</b> $\langle \varepsilon^{1/3} \rangle L^{1/3} (a_i + a_j)^2$

# Energy Dissipation Rate [w/kg] ( $\epsilon$ ) Profile



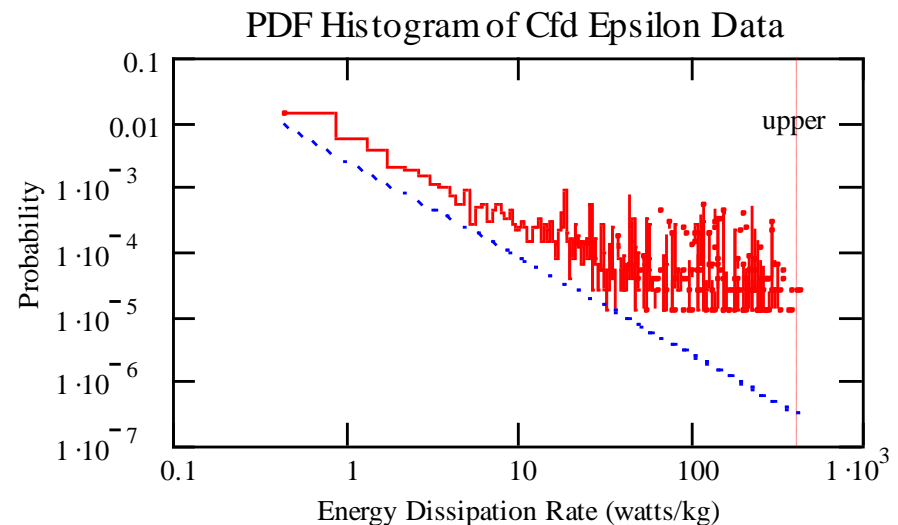
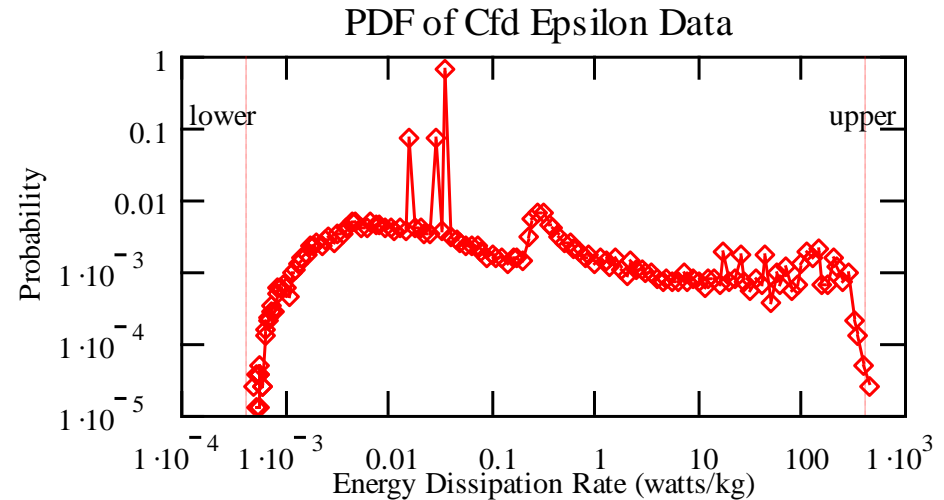
# Energy Dissipation Rate on Individual Particle Track

- High  $\varepsilon$  when passes through impeller zone
- Effectively
  - $\varepsilon = 0$  elsewhere

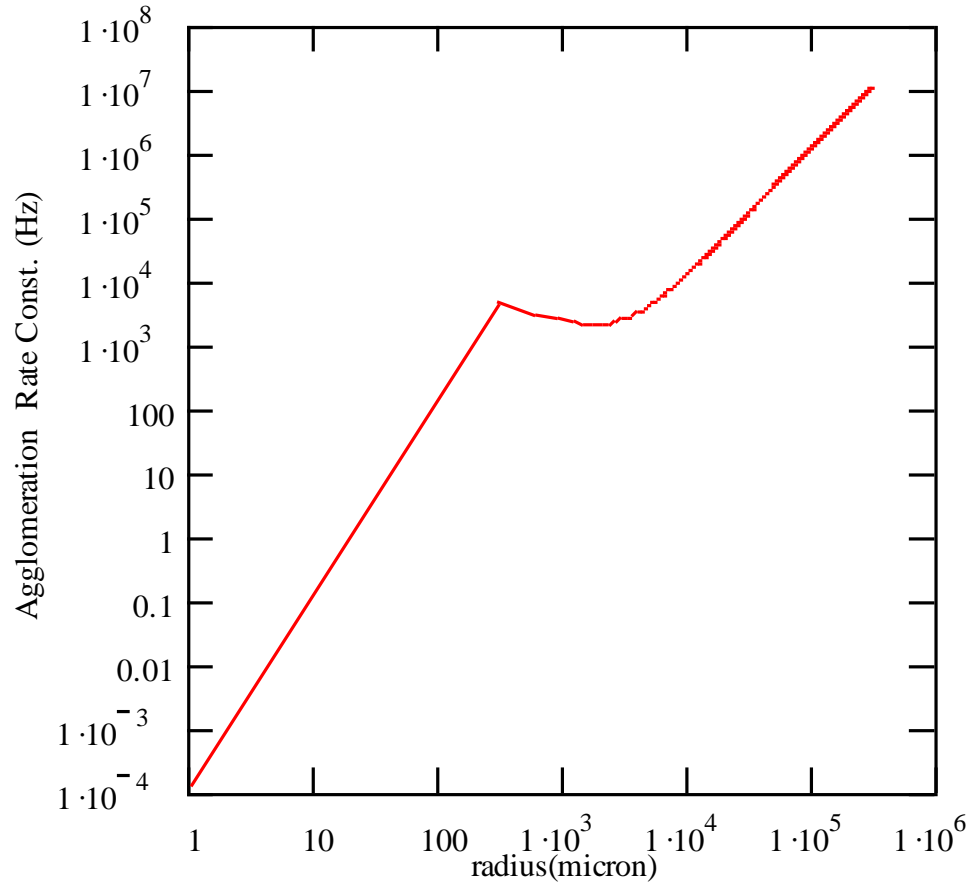


# Energy Dissipation Rate Statistics

- $\epsilon_{\max} = 429 \text{ w/kg}$
- Multimode Distribution
- Probability  $\propto \epsilon^{-3/2}$ 
  - Decay of Isotropic Turbulence
- $V = 1.4 \text{ L}$ ,  $Q = 40 \text{ ml/min.}$ ,  $80 \text{ rpm}$



# Aggregation Rate



07/25/03

Reaction Temp: 30.1°C, Flow Rate: 45 ml/min, Mixing Power: 287.7 rpm, Reactor Volume: 1400 ml

# Breakage Kinetics

- Breakage Rate =

Collision Frequency \* Target Efficiency \* Breakage Probability

Collisions

Collision Efficiency

Particle-wall

Particle-impeller

Particle-baffle

Particle-Particle due to either diffusion or turbulence

# Use Fluid Dynamics with Sinc Methods

- Particle Breakage

- Particle Collisions with

- Particles (same equations as aggregation)
  - depend upon turbulent energy dissipation rate

- Baffles/walls – impaction efficiency

- Impeller – collision frequency

$$\eta_{target} = \left( \frac{N_{St}}{0.32 + N_{St}} \right)^{2.1}$$

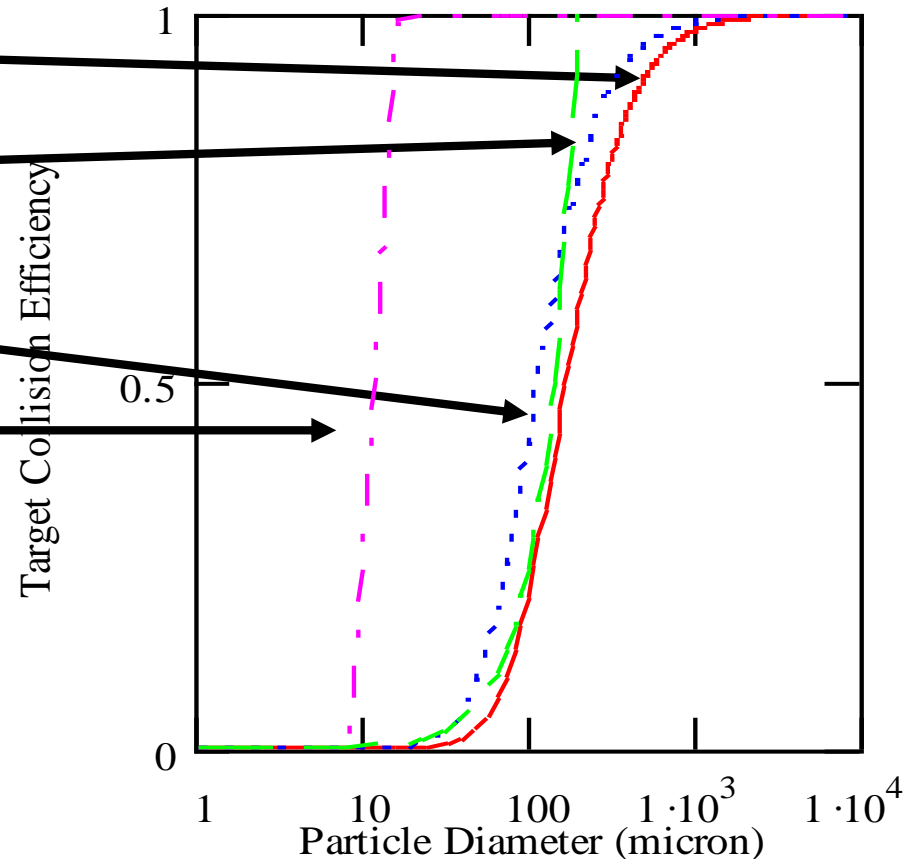
- Sufficient Energy of Collision to Break Particle

$$W_{p,\min} = 64 \frac{G_2^3 \Gamma_2^3}{H_2^5 K_r^3}$$

$$p(x) = \frac{2.25 x^{-3.25}}{x_{\min}^{-2.25} - x_{\max}^{-2.25}}$$

# Particle Breakage Efficiency

- Baffle
- Wall
- Impeller
- Other Particles

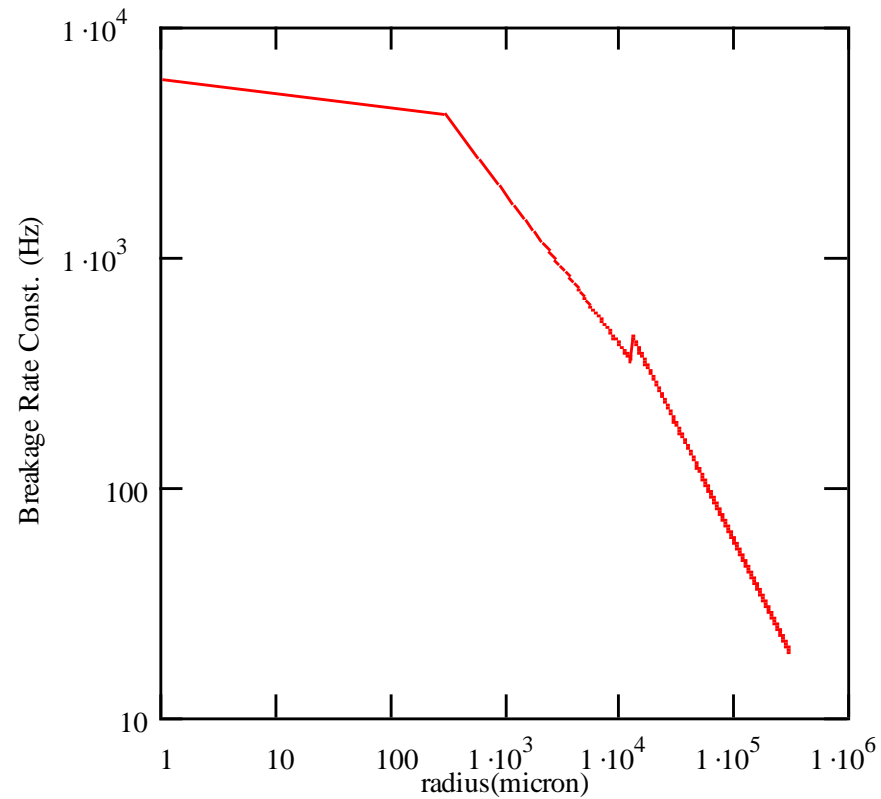


07/25/03

Reaction Temp: 30.1°C, Flow Rate: 45 ml/min, Mixing Power: 287.7 rpm, Reactor Volume: 1400 ml



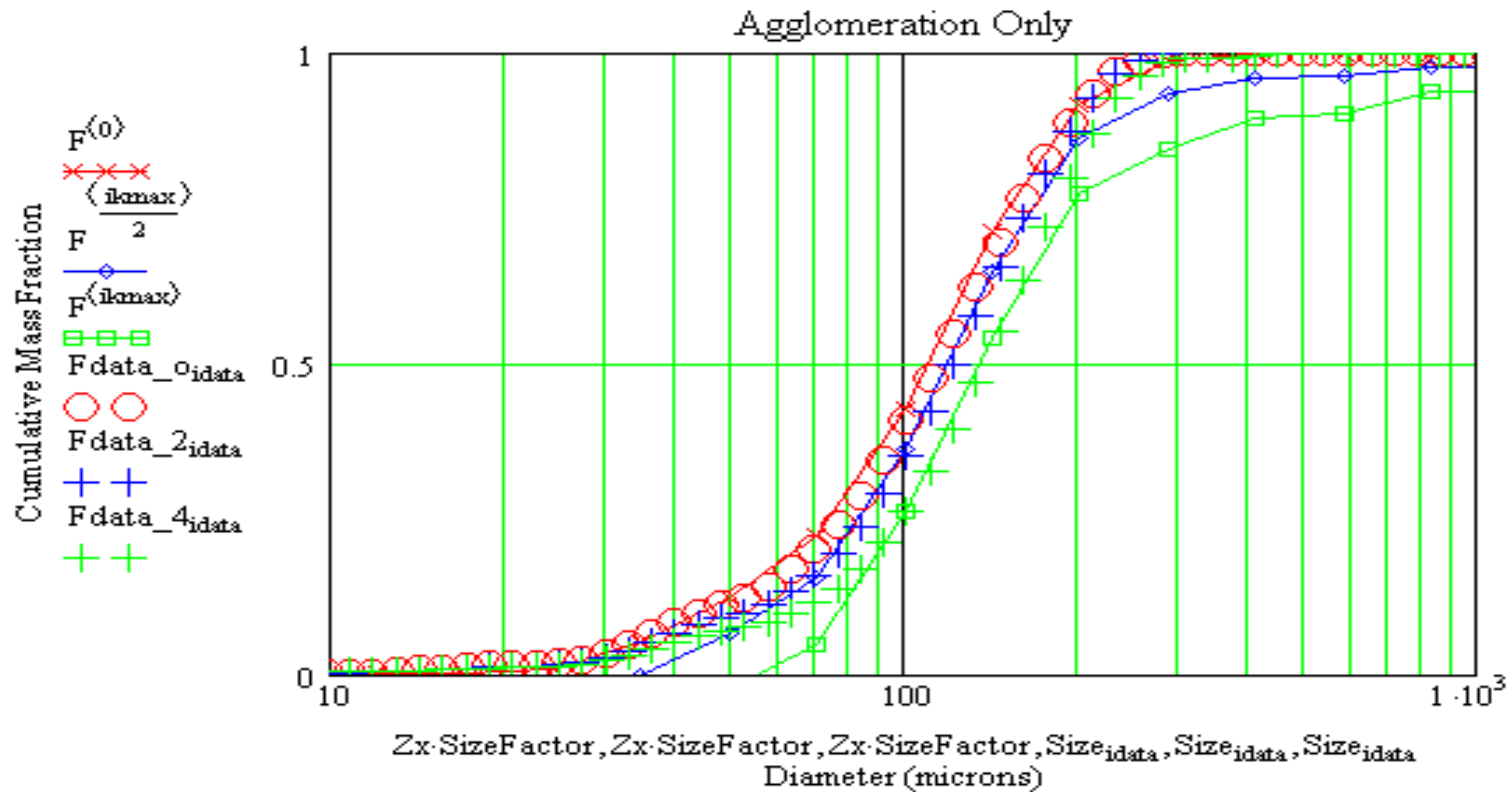
# Breakage Rate



07/25/03

Reaction Temp: 30.1°C, Flow Rate: 45 ml/min, Mixing Power: 287.7 rpm, Reactor Volume: 1400 ml

# Model Results – Aggregation/Breakage



A\_Factor=400, a colloid stability parameter

# Conclusions

- Getting Crystallization Parameters from Experimental Data
  - Requires
    - Decomposition of the multiplicity of phenomenon
    - Knowledge of fundamental particle mechanisms
    - Simple calculation tool for stirred tank

# Introduction to Population Balance Modeling

# Topics

- Modeling Philosophies
  - Where Population Balance Models (PBM) fit in
- Important Characteristics
- Framework
- Uses
  - Cancer Examples
- Parameter Identification
- Potential Applications

# Variations of Scale

- Tumor scale properties
  - Physical characteristics
  - Disease class

- Cellular level
  - Growth and death rates
  - Mutation rates

- Subcellular characteristics
  - Genetic profile
  - Reaction networks (metabolic)
  - CD markers

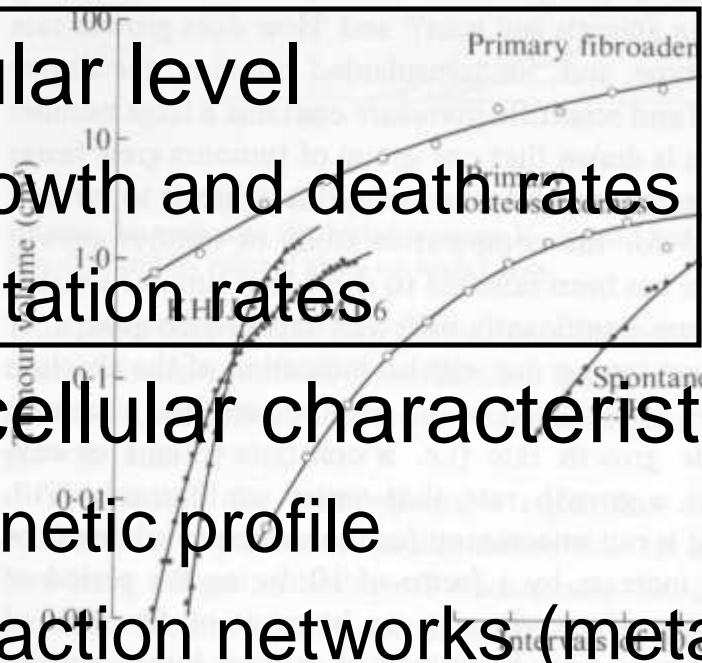


FIG. 1.8. Growth curves for seven experimental tumours whose growth has been studied in detail. In each case the data have been fitted by a Gompertz equation and the corresponding parameters are given in Table 1.1, together with the relevant sources.

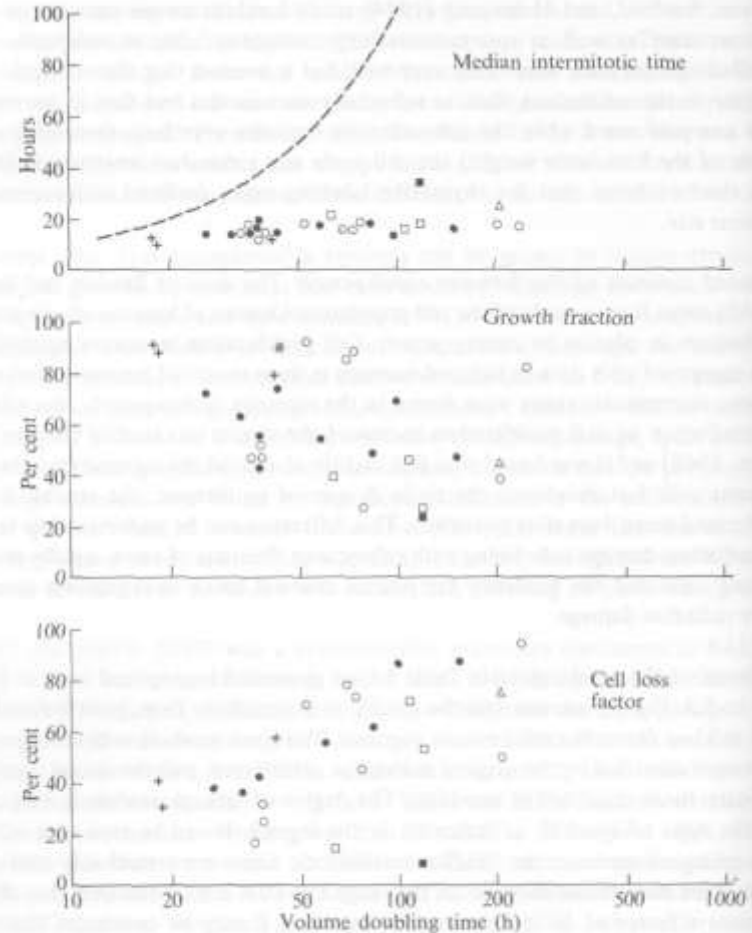
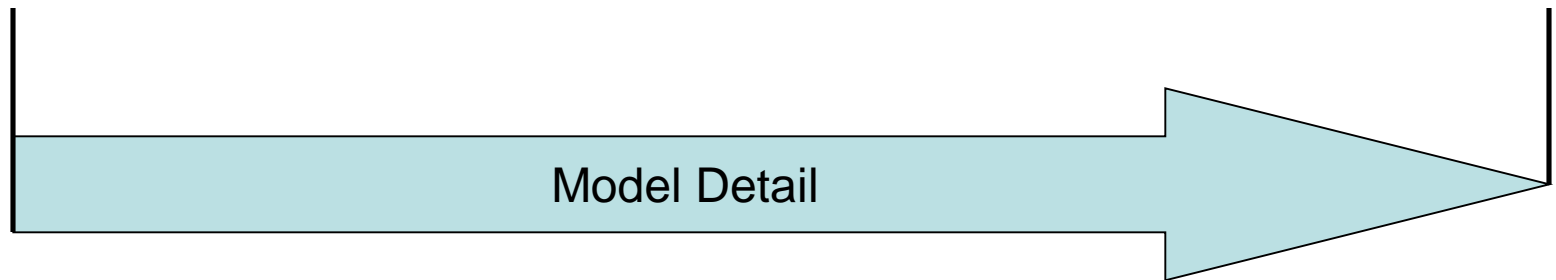


FIG. 5.1. Estimates of the median intermitotic time (the cell cycle time), growth fraction, and cell loss factor for tumours in the mouse:  $\Delta$  autochthonous tumours;  $\square$  early transplants;  $\circ$  frequently passaged transplanted tumours. Open symbols indicate carcinomas, closed symbols are sarcomas, miscellaneous tumours indicated by + (see Table 5.1). The broken line indicates the limit where the mean intermitotic time equals the doubling time.

# Level of Model Detail – Population Growth

- Empirical Models
  - Course structure
  - Averaged cell behavior
  - Gompertz growth
- Mechanistic Models
  - Fine structure
  - Cdc/Cdk interactions



- Population Balance Models
  - “key” parameters (i.e. DNA, volume, age)
  - Details are lumped and averaged
  - Heterogeneous behavior

# State vector

- Completely averaged
  - $X = [ \quad ]$
- Few state variables
  - $X = [\text{size spatial\_location internal\_drug\_level}]$
- More mechanistic
  - $X = [\text{Cdk1 CycA CycE...}]$
- Population Balance
  - Expected distribution among cells



# Population Balance Model Requirements

- Transition rates (e.g. division rate)

$$\Gamma_{i,j}(x_1, x_2, x_3)$$

- Rates of change – growth rates

$$\dot{\mathbf{X}}(x_1, x_2, x_3, \mathbf{y})$$

- Constitutive model or experiments

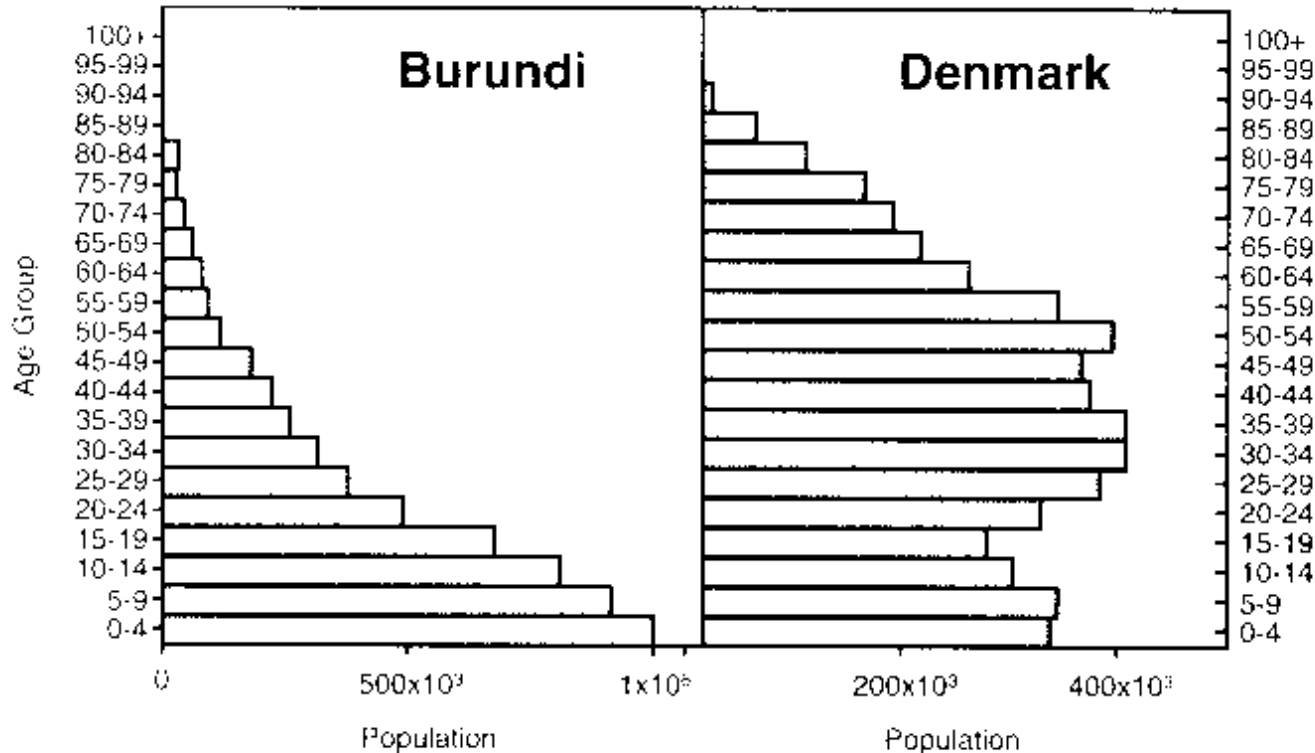
- Advantages

- Simplified description of system
- Flexible framework

- Disadvantages

- Proper identification of system (Burundi not the same as Denmark)
- Identification of rates

# Modeling Philosophies



Hjortsø 2005

**Figure 1.1** Population pyramids for Burundi and Denmark, 2000. (Source: U.S. Census Bureau)

- Birth and death rates

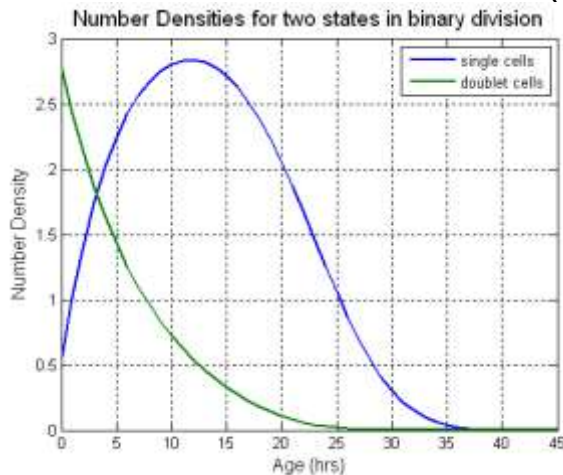
- Empirical
- PBM – vary with age and country
- Mechanistic – need to know the causes behind age-dependence

# Expected Number Density

- Distribution of cell states

2 discrete states (i.e. Denmark)

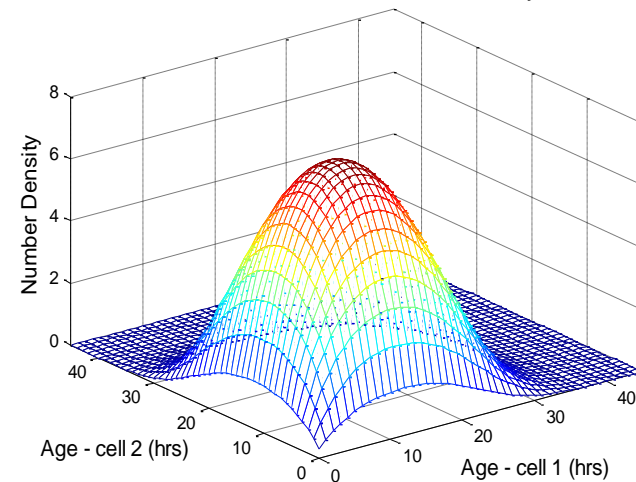
1 continuous variable (i.e. age)



2 continuous variables

(i.e. age and weight)

Multi-dimensional number density



- Total number density

$$E[N(a,b;t)] = N_1(a,b;t) = \int_a^b n_1(x,t) dx$$

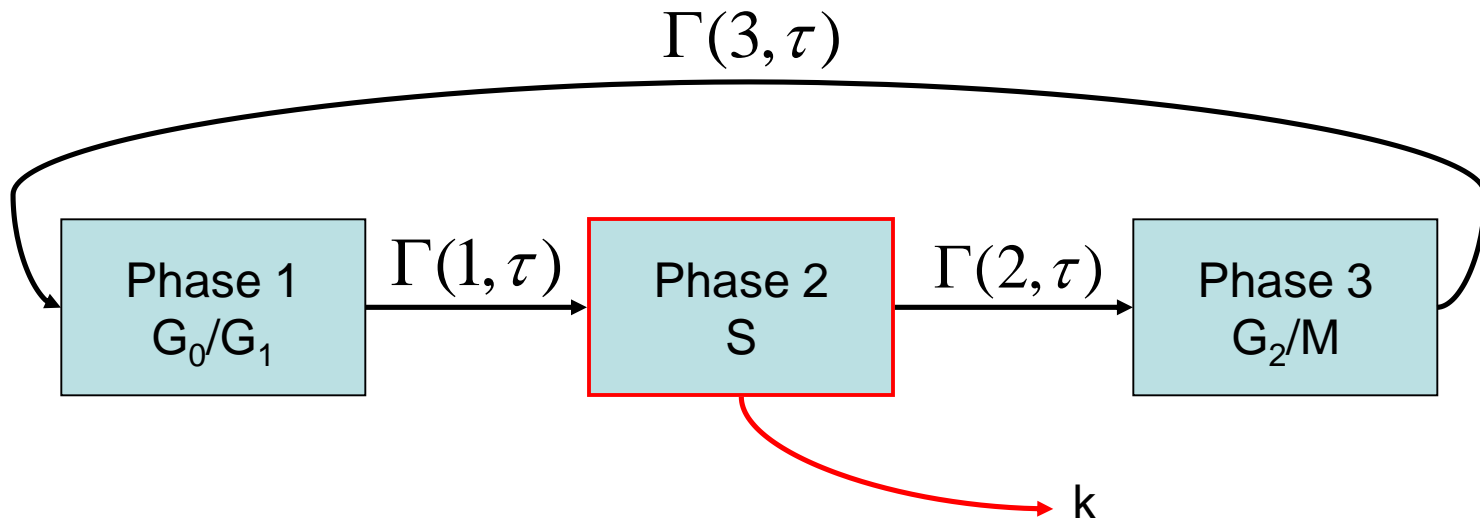
$$E[N(a,b;c,d;t)] = N_1(a,b;c,d;t) = \int_c^d \int_a^b n_1(x,t) dx dx'$$

# Population Description

- Continuous variables
- Time
- Cell
  - Mass
  - Volume
  - Age
  - DNA or RNA
  - Protein
- Patient
  - Age
- Discrete indices
- Cell
  - Cell cycle phase
  - Genetic mutations
  - Differentiation state
- Patient
  - M / F
  - Race / ethnicity

# Example – Cell Cycle Specific Behaviors

- Cell cycle specific drug
  - Discrete – cell cycle phase,  $p$
  - Continuous – age,  $\tau$ , (time since last transition)
- $n_1(p, \tau, t)$



# Cell Cycle Control (Tyson and Novak 2004)

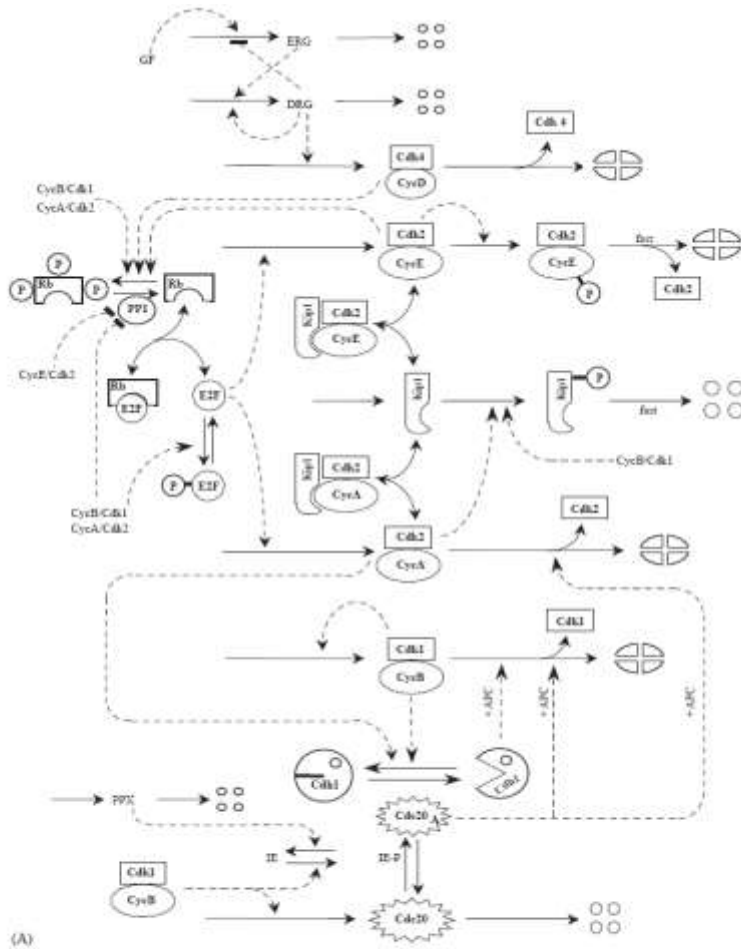


Fig. 1. Molecular network regulating the progression of mammalian cells through the cell cycle. (A) In the center of the diagram we propose a yeast-like cell-cycle "engine" composed of Cdk2/CycE, Cdk2/CycA, Cdk1/CycB, and some ancillary proteins (Kip1, Cdh1 and Cdc20). This part of the diagram should be compared to Chen et al. (2000). To the engine we attach three components characteristic of mammalian cell-cycle controls: (i) retinoblastoma protein, Rb, which binds to and inhibits E2F, a transcription factor for production of CycA and CycE, (ii) the cyclin-dependent kinase, Cdk4/CycD, which phosphorylates and inactivates Rb, and (iii) the signal-transduction pathway, GF-ERG-DRG, which controls C synthesis in response to GF stimulation. Although not indicated on the wiring diagram, the model includes the fact that Kip1 binds to CycD/C but does not inhibit its activity. (B) A more detailed representation of the binding and phosphorylation reactions that govern Rb-E2F interactions. In deriving the model equations (Table 3), we assume that binding and release reactions are fast compared to phosphorylation and dephosphorylation reactions.

Table 3  
Mathematical model of mammalian cell-cycle controls

$$\frac{d[ERG]}{dt} = \frac{k_{15}}{1 + ([DRG]/J_1)^2} - k_{14}[ERG] \quad (1)$$

$$\frac{d[DRG]}{dt} = \tau \left( k'_{11}[ERG] + \frac{k_{11}([DRG]/J_1)^2}{1 + ([DRG]/J_1)^2} \right) - k_{11}[DRG] \quad (2)$$

$$\frac{d[CycD]}{dt} = \tau k_8[DRG] + V_2[CycD : Kip1] + k_{24}[CycD : Kip1] - k_{23}[CycD][Kip1] - k_{14}[CycD] \quad (3)$$

$$\frac{d[CycD : Kip1]}{dt} = k_{23}[CycD][Kip1] - k_{24}[CycD : Kip1] - V_2[CycD : Kip1] - k_{14}[CycD : Kip1] \quad (4)$$

$$\frac{d[CycE]}{dt} = \tau(k'_7 + k_7[E2F]_d) - V_3[CycE] - k_{25}[CycE][Kip1] + k_{25}[CycE : Kip1] + V_4[CycE : Kip1] \quad (5)$$

$$\frac{d[CycE : Kip1]}{dt} = k_{25}[CycE][Kip1] - k_{25}[CycE : Kip1] - V_4[CycE : Kip1] - V_3[CycE : Kip1] \quad (6)$$

$$\frac{d[CycA]}{dt} = \tau k_{26}[E2F]_a[mas] - k_{26}[Cdc20][CycA] - k_{27}[CycA][Kip1] + k_{27}[CycA : Kip1] + V_4[CycA : Kip1] \quad (7)$$

$$\frac{d[CycA : Kip1]}{dt} = k_{27}[CycA][Kip1] - k_{27}[CycA : Kip1] - V_4[CycA : Kip1] - k_{26}[Cdc20][CycA : Kip1] \quad (8)$$

$$\frac{d[Kip1]}{dt} = \tau k_9 - V_4[Kip1] - k_{24}[CycD][Kip1] + k_{24}[CycD : Kip1] + k_{14}[CycD : Kip1] - k_{23}[Kip1][CycE] + [CycA] + k_{25}[CycE : Kip1] + [CycA : Kip1] + V_3[CycE : Kip1] + V_4[CycE : Kip1] + k_{14}[Cdc20][CycA : Kip1] \quad (9)$$

$$\frac{d[E2F]}{dt} = k_{22}[E2F]_T - [E2F] - (k'_{21} + k_{21}[CycA] + [CycB])[E2F] \quad (10)$$

$$\frac{d[CycB]}{dt} = \tau \left( k'_1 + \frac{k_1[CycB]/J_1^2}{1 + ([CycB]/J_1)^2} \right) - V_2[CycB] \quad (11)$$

$$\frac{d[Cdh1]}{dt} = (k'_3 + k_3[Cdc20]) \frac{1 - [Cdh1]}{J_3 + 1 - [Cdh1]} - V_2 \frac{[Cdh1]}{J_3 + [Cdh1]} \quad (12)$$

$$\frac{d[Cdc20_T]}{dt} = \tau k'_{11} + k_{11}[CycB] - k_{12}[Cdc20_T] \quad (13)$$

$$\frac{d[Cdc20]}{dt} = k_{11}[E2F] \frac{[Cdc20_T] - [Cdc20]}{J_{11} + [Cdc20_T] - [Cdc20]} - k_{12} \frac{[Cdc20]}{k_4 + [Cdc20]} - k_{12}[Cdc20] \quad (14)$$

$$\frac{d[PPN]}{dt} = k_{13} - k_{14}[PPN] \quad (15)$$

$$\frac{d[E2F]}{dt} = k_{16}[CycB] \frac{1 - [E2F]}{J_{16} + 1 - [E2F]} - k_{16}[PPN] \frac{[E2F]}{J_{16} + [E2F]} \quad (16)$$

$$\frac{d[GM]}{dt} = k_2[mas]H \left( \frac{[Rb_{tot}]}{[Rb_T]} \right) - k_2[GM] \quad (17)$$

$$\frac{d[mas]}{dt} = \tau_3[GM] \quad (18)$$

Steady-state relations

$$[PPN]_s = \frac{[PPN]_T}{1 + k_{14}(\phi_T([CycE] + [CycA]) + \phi_d[CycB])} \quad (19)$$

$$[Rb_{tot}]_s = \frac{[Rb_T]}{1 + k_{20}/J_{10}([CycD]_T + k_4[CycE] + k_4[CycA] + J_{10}[CycB]) + k_{10}[PPN]_s - [PPN]_s + k_{13}[PPN]_s} \quad (20)$$

$$[E2F]_d = \frac{[E2F]_T - [E2F : Rb][E2F]}{[E2F]_T} \quad (21)$$

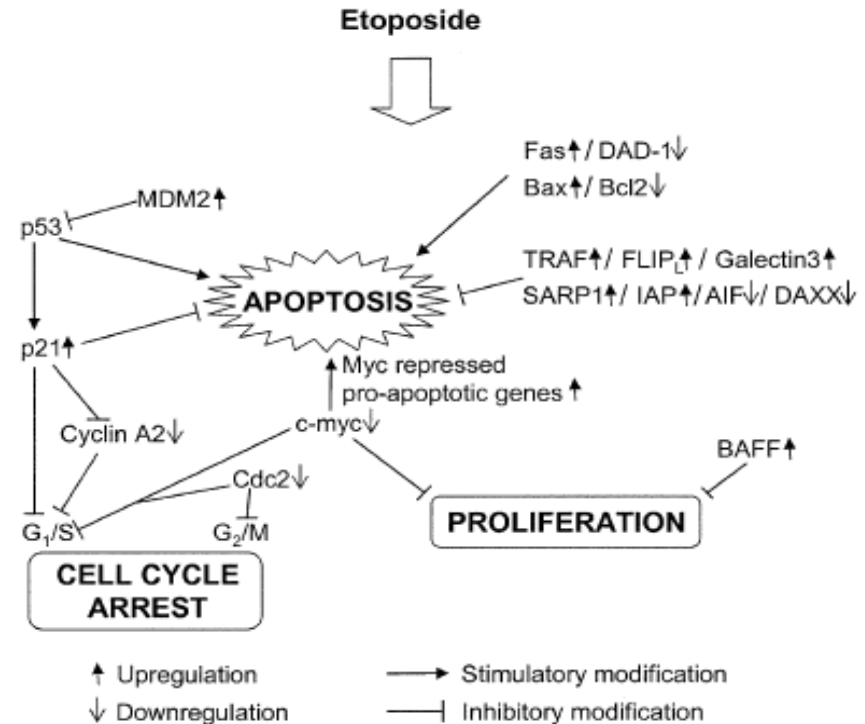
# Cell cycle arrest (Tao et al. 2003)

**Table 1.** Genes involved in cell-cycle regulation and apoptosis whose expression is upregulated or downregulated in etoposide-treated human CD34<sup>+</sup> cord blood cells compared to control as analyzed by cDNA array

Gene Name	Accession No.	Description	Relative* Expression	Fold Change*
<b>Cell Cycle Regulators</b>				
<i>MDM2</i>	NM_002392	p53-associated gene.	0.17	21.3 (up)
<i>P21</i>	U03106	Wild-type p53 activated fragment-1 (WAF1).	0.25	6.6 (up)
<i>Cyclin A2</i>	NM_001237	Cyclin A.	0.78	-2.8 (down)
<i>CDC2</i>	NM_001786	Cell cycle control gene CDC2.	0.20	-1.3 (down)
<i>c-myc</i>	V00568	C-myc oncogene.	1.28	-5.9 (down)
<b>TNF Superfamily</b>				
<i>Fas</i>	M67454	Fas antigen (fas).	0.06	10.0 (up)
<i>TRAIL_R1</i>	NM_003844	Cytotoxic ligand TRAIL receptor.	0.04	8.6 (up)
<i>TRAIL_R2</i>	AF012535	Death receptor 5 (DR5).	0.32	3.2 (up)
<i>TRAIL_R4</i>	NM_003840	Death receptor 2 (DR2).	0.02	13.2 (up)
<i>4-1BBL</i>	U03398	Receptor 4-1BB ligand.	0.91	1.7 (up)
<i>THANK</i>	AF116456	B-cell activating factor (BAFF).	0.12	-4.9 (down)
<i>TNF_R1</i>	M63121	Tumor necrosis factor receptor.	0.29	-1.4 (down)
<b>Apoptosis-Related and <i>bcl-2</i>-Related</b>				
<i>Cox-2</i>	M90100	Cyclooxygenase-2 (Cox-2).	0.11	1.8 (up)
<i>FLIPL</i>	AF015452	Usurpin-γ.	0.31	1.7 (up)
<i>GALECTIN-3</i>	AB006780	Lectin, galactoside-binding, soluble, 3 (galectin-3)	0.10	2.2 (up)
<i>IAP-1</i>	U45878	Inhibitor of apoptosis protein 1.	0.15	1.9 (up)
<i>nNOS</i>	U17327	Neuronal nitric oxide synthase (NOS1).	0.19	2.8 (up)
<i>SARP-1</i>	AF017986	Secreted apoptosis-related protein 1 (SARP1).	0.64	1.9 (up)
<i>AIF</i>	NM_004208	Apoptosis-inducing factor (AIF), nuclear gene encoding mitochondrial protein.	0.66	-1.3 (down)
<i>DAD-1</i>	NM_001344	Defender against cell death 1 (DAD1).	0.74	-1.6 (down)
<i>DAXX</i>	NM_005453	mRNA for death-associated protein, Daxx.	0.54	-1.3 (down)
<i>GAPDH</i>	M33197	Glyceraldehyde-3-phosphate dehydrogenase (GAPDH).	19.52	-1.5 (down)
<i>Bax-α</i>	L22473	Baxα.	0.34	2.4 (up)
<i>bcl-w</i>	NM_004050	<i>bcl-w</i> ( <i>bcl-w</i> ).	0.58	1.6 (up)
<i>Mcl-1</i>	L08246	Myeloid cell differentiation protein (MCL1).	0.63	1.7 (up)
<i>bcl-2</i>	NM_000633	B-cell CLL/lymphoma 2 (BCL-2).	0.24	-1.3 (down)

\*The numbers denote that expression of a gene relative to the expression of gene encoding transferin receptor in control CD34<sup>+</sup> cord blood cells. Relative expression is calculated as a ratio of detected mRNA of a gene and that of the transferin receptor gene.

\*A positive number indicates that expression of a gene is upregulated while a negative value indicates downregulation of a gene in etoposide-treated CD34<sup>+</sup> cord blood cells.



**Figure 6.** Schematic representation of basic molecular components existing in and putative pathways utilized by human CD34<sup>+</sup> cells in their response to etoposide treatment based on cDNA array and quantitative real-time RT-PCR data. Myc and p53 are main pathways elicited by human CD34<sup>+</sup> cells in their response to etoposide treatment. Multiple known pro-survival and pro-apoptotic pathways were simultaneously activated.

# PBMs in biological settings

- Cell cycle-specific chemo
- Budding yeast dynamics
- Rate of monoclonal antibody production in hybridoma cells
- Bioreactor productivity under changing substrate conditions
- Ecological models
  - Predator-prey



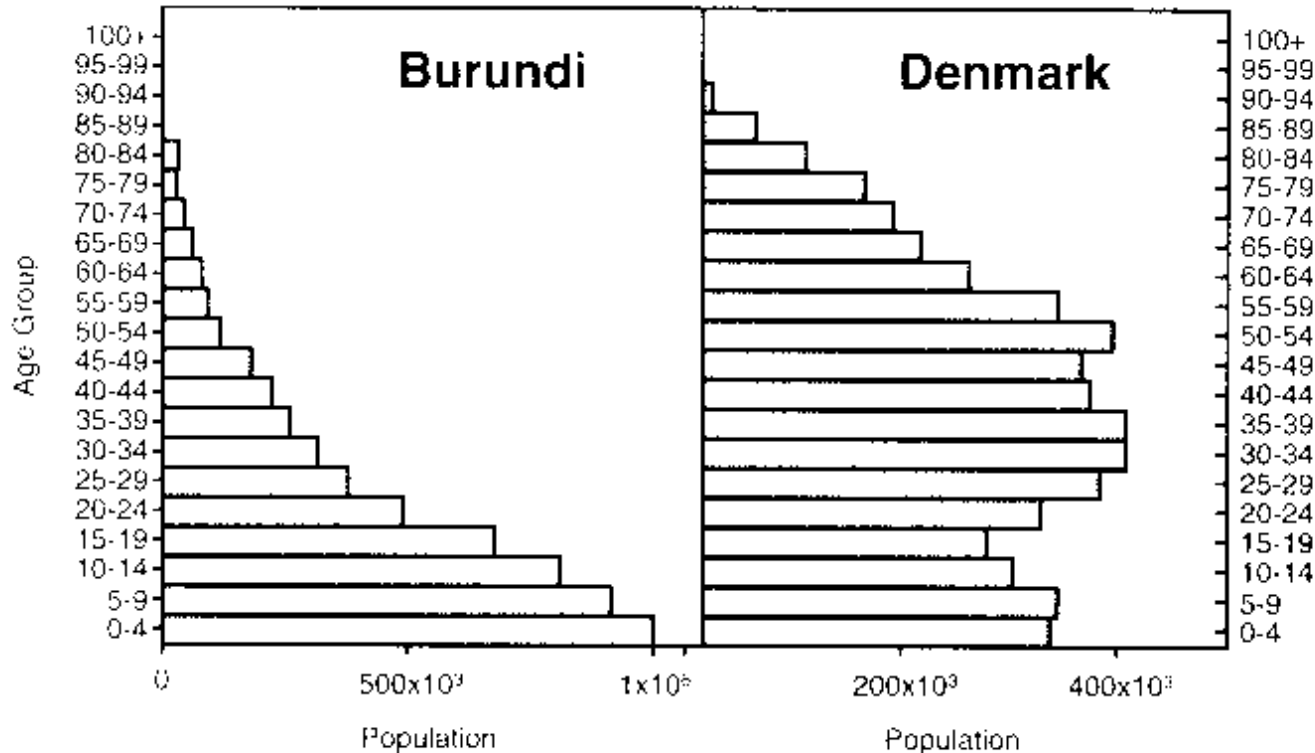
# Topics

- Modeling Philosophies
  - Where Population Balance Models (PBM) fit in
- Important Characteristics
- Framework
- Uses
  - Cancer Examples
- Parameter Identification
- Potential Applications

# Variations of Scale

- Tumor scale properties
  - Physical characteristics
  - Disease class
- Cellular level
  - Growth and death rates
  - Mutation rates
- Subcellular characteristics
  - Genetic profile
  - Reaction networks (metabol- and prote-omics)
  - CD markers

# Modeling Philosophies



Hjortsø 2005

**Figure 1.1** Population pyramids for Burundi and Denmark, 2000. (Source: U.S. Census Bureau)

- Birth and death rates

- Empirical
- PBM – vary with age and country (STATE VECTOR)
- Mechanistic – need to know the causes behind age-dependence

# Population Balance Model Requirements

- Transition rates (e.g. division and rates)

$$\Gamma_{i,j}(x_1, x_2, x_3)$$

- Rates of change – growth rates

$$\dot{\mathbf{X}}(x_1, x_2, x_3, \mathbf{y})$$

- Constitutive model or experiments

- Advantages

- Simplified description of system
- Flexible framework

- Disadvantages

- Proper identification of system (Burundi not the same as Denmark)
- Identification of rates

# Cell Cycle Control (Tyson and Novak 2004)

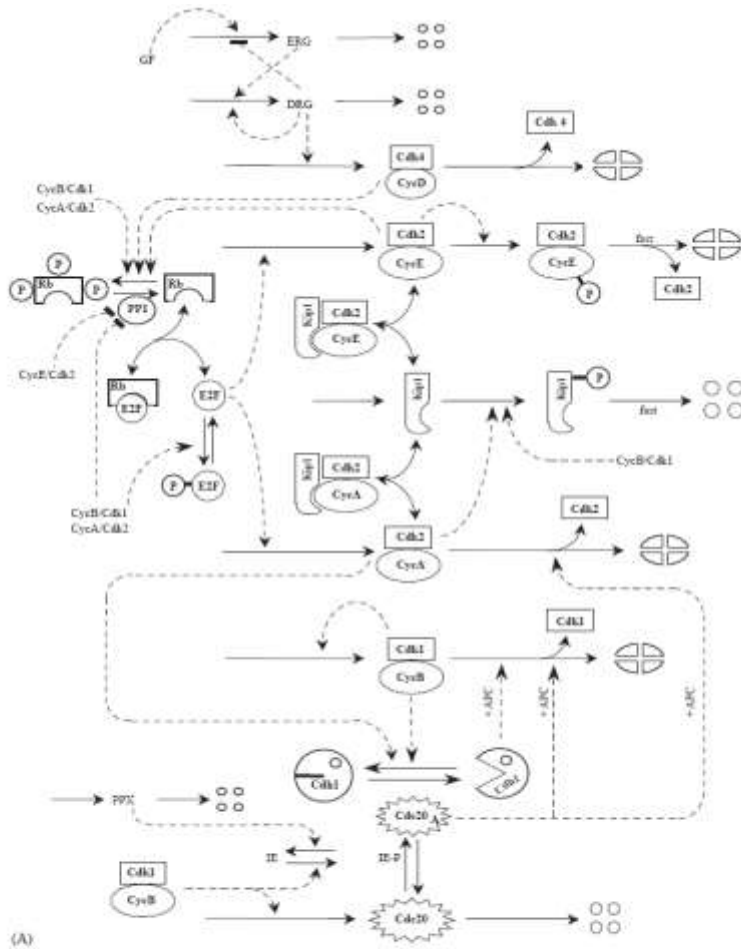


Fig. 1. Molecular network regulating the progression of mammalian cells through the cell cycle. (A) In the center of the diagram we propose a yeast-like cell-cycle "engine" composed of Cdk2/CycE, Cdk2/CycA, Cdk1/CycB, and Cdk1/CycD, and some ancillary proteins (Kip1, Cdh1 and Cdc20). This part of the diagram should be compared to Chen et al. (2000). To the engine we attach three components characteristic of mammalian cell-cycle controls: (i) retinoblastoma protein, Rb, which binds to and inhibits E2F, a transcription factor for production of CycA and CycE, (ii) the cyclin-dependent kinase, Cdk4/CycD, which phosphorylates and inactivates Rb, and (iii) the signal-transduction pathway, GF-ERG-DRG, which controls C synthesis in response to GF stimulation. Although not indicated on the wiring diagram, the model includes the fact that Kip1 binds to CycD/C but does not inhibit its activity. (B) A more detailed representation of the binding and phosphorylation reactions that govern Rb-E2F interactions. In deriving the model equations (Table 3), we assume that binding and release reactions are fast compared to phosphorylation and dephosphorylation reactions.

Table 3  
Mathematical model of mammalian cell-cycle controls

$$\frac{d[ERG]}{dt} = \frac{k_{15}}{1 + ([DRG]/J_1)^2} - k_{14}[ERG] \quad (1)$$

$$\frac{d[DRG]}{dt} = \tau \left( k'_{11}[ERG] + \frac{k_{11}([DRG]/J_1)^2}{1 + ([DRG]/J_1)^2} \right) - k_{11}[DRG] \quad (2)$$

$$\frac{d[CycD]}{dt} = \tau k_8[DRG] + V_2[CycD : Kip1] + k_{24}[CycD : Kip1] - k_{23}[CycD][Kip1] - k_{14}[CycD] \quad (3)$$

$$\frac{d[CycD : Kip1]}{dt} = k_{23}[CycD][Kip1] - k_{24}[CycD : Kip1] - V_2[CycD : Kip1] - k_{14}[CycD : Kip1] \quad (4)$$

$$\frac{d[CycE]}{dt} = \tau(k'_7 + k_7[E2F_d]) - V_4[CycE] - k_{25}[CycE][Kip1] + k_{25}[CycE : Kip1] + V_4[CycE : Kip1] \quad (5)$$

$$\frac{d[CycE : Kip1]}{dt} = k_{25}[CycE][Kip1] - k_{25}[CycE : Kip1] - V_4[CycE : Kip1] - V_4[CycE : Kip1] \quad (6)$$

$$\frac{d[CycA]}{dt} = \tau k_{26}[E2F_A][mass] - k_{26}[Cdc20][CycA] - k_{27}[CycA][Kip1] + k_{27}[CycA : Kip1] + V_4[CycA : Kip1] \quad (7)$$

$$\frac{d[CycA : Kip1]}{dt} = k_{27}[CycA][Kip1] - k_{27}[CycA : Kip1] - V_4[CycA : Kip1] - k_{26}[Cdc20][CycA : Kip1] \quad (8)$$

$$\frac{d[Kip1]}{dt} = \tau k_9 - V_4[Kip1] - k_{24}[CycD][Kip1] + k_{24}[CycD : Kip1] + k_{14}[CycD : Kip1] - k_{23}[Kip1][CycE] + [CycE A] + k_{25}[CycE : Kip1] + [CycA : Kip1] + V_4[CycE : Kip1] + k_{24}[Cdc20][CycA : Kip1] \quad (9)$$

$$\frac{d[E2F]}{dt} = k_{22}[E2F_T] - [E2F] - (k'_{21} + k_{21}[CycA] + [CycB])[E2F] \quad (10)$$

$$\frac{d[CycB]}{dt} = \tau \left( k'_1 + \frac{k_1[CycB]/J_1^2}{1 + ([CycB]/J_1)^2} \right) - V_3[CycB] \quad (11)$$

$$\frac{d[Cdh1]}{dt} = (k'_3 + k_3[Cdc20]) \frac{1 - [Cdh1]}{J_3 + 1 - [Cdh1]} - V_2 \frac{[Cdh1]}{J_4 + [Cdh1]} \quad (12)$$

$$\frac{d[Cdc20_T]}{dt} = \tau k'_{11} + k_{11}[CycB] - k_{12}[Cdc20_T] \quad (13)$$

$$\frac{d[Cdc20]}{dt} = k_{11}[E2F] \frac{[Cdc20_T] - [Cdc20]}{J_{11} + [Cdc20_T] - [Cdc20]} - k_{12} \frac{[Cdc20]}{k_4 + [Cdc20]} - k_{12}[Cdc20] \quad (14)$$

$$\frac{d[PPN]}{dt} = k_{13} - k_{14}[PPN] \quad (15)$$

$$\frac{d[E2F]}{dt} = k_{16}[CycB] \frac{1 - [E2F]}{J_{16} + 1 - [E2F]} - k_{16}[PPN] \frac{[E2F]}{J_{16} + [E2F]} \quad (16)$$

$$\frac{d[GM]}{dt} = k_2[mass]H \left( \frac{[Rb_{tot}]}{[Rb_T]} \right) - k_2[GM] \quad (17)$$

$$\frac{d[mass]}{dt} = \tau_3[GM] \quad (18)$$

Steady-state relations

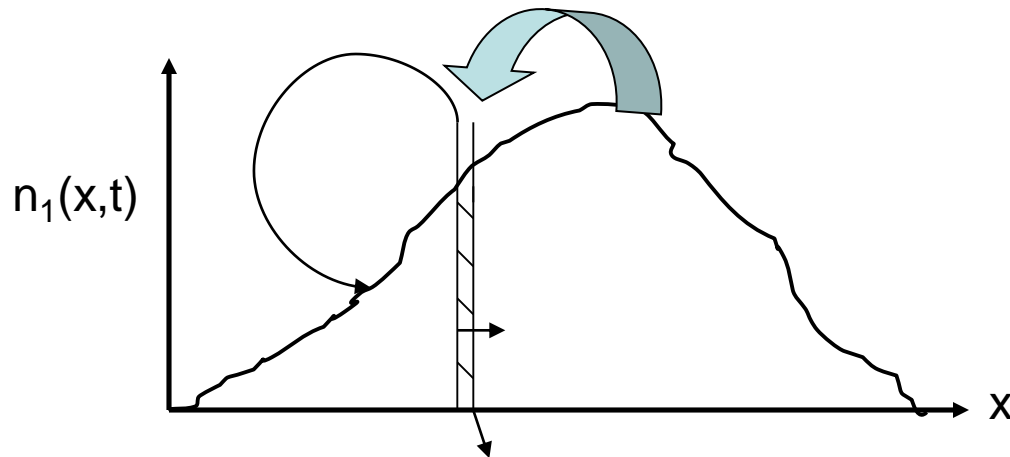
$$[PPN]_s = \frac{[PPN_T]}{1 + k_{14}(\phi_T([CycE] + [CycA]) + \phi_d[CycB])} \quad (19)$$

$$[Rb_{tot}]_s = \frac{[Rb_T]}{1 + k_{20}/J_8([CycD_T] + k_4[CycE] + k_4[CycA] + J_8[CycB])} \quad (20)$$

$$[E2F]_s = \frac{[E2F_T] - [E2F : Rb][E2F]}{[E2F_T]} \quad (21)$$

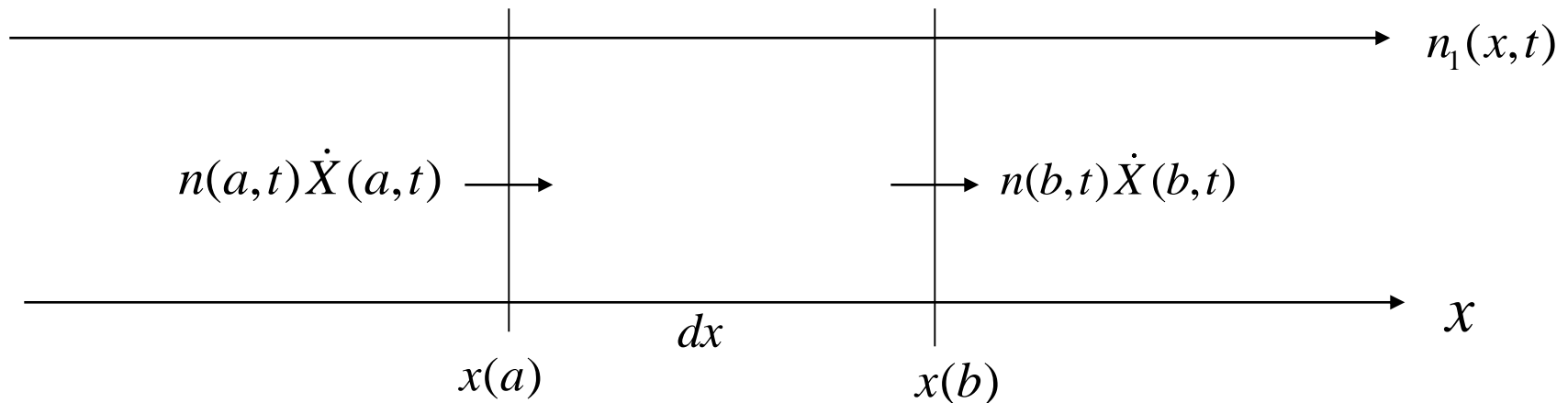
# Changes Number Density

- Changes in cell states



- Cell behavior (growth, division and death, and phase transitions) are functions of a cell's state

# Population Balance – pure growth



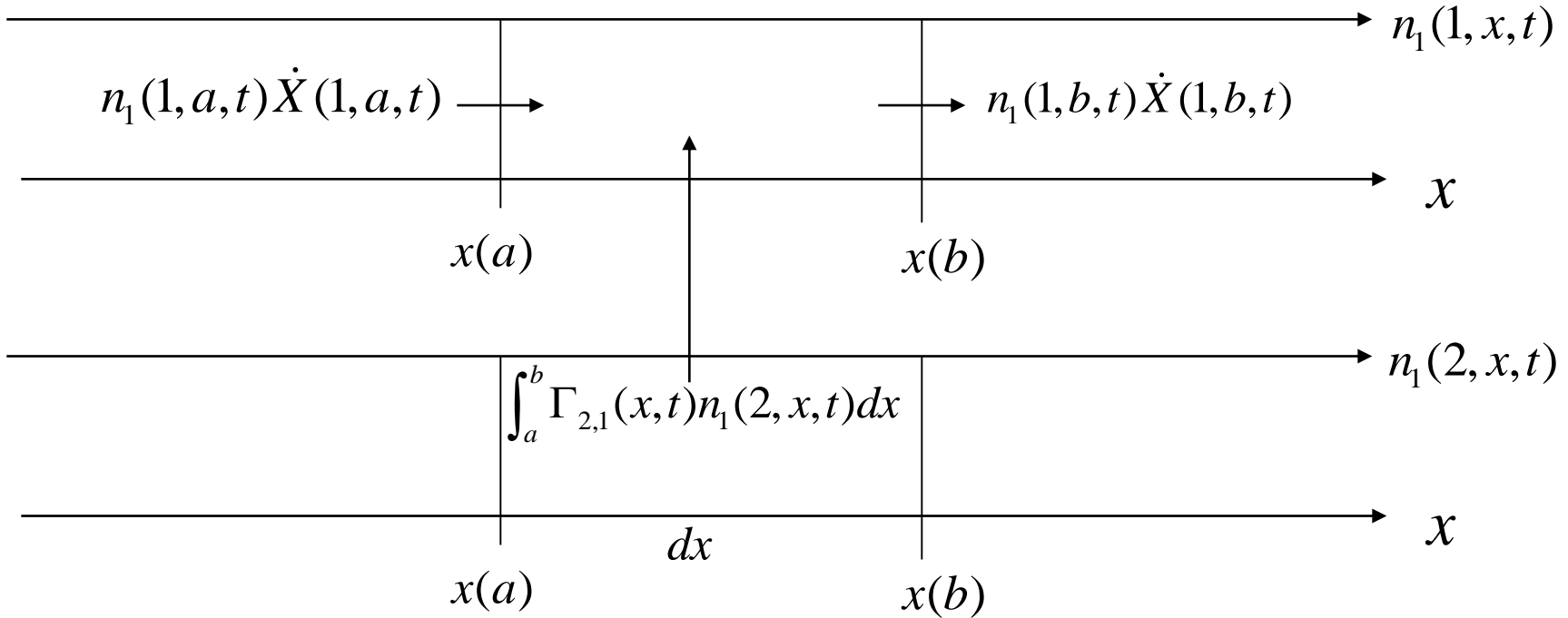
$$\frac{d}{dt} \int_a^b n_1(x, t) dx = n_1(a, t)\dot{X}(a, t) - n_1(b, t)\dot{X}(b, t)$$

$$\int_a^b \frac{\partial}{\partial t} n_1(x, t) dx = - \int_a^b \frac{\partial}{\partial x} (\dot{X}(x, t) n_1(x, t)) dx$$

$$\frac{\partial n_1(x, t)}{\partial t} + \frac{\partial}{\partial x} (\dot{X}(x, t) n_1(x, t)) = 0$$

Growth rate

# Transitions between discrete states



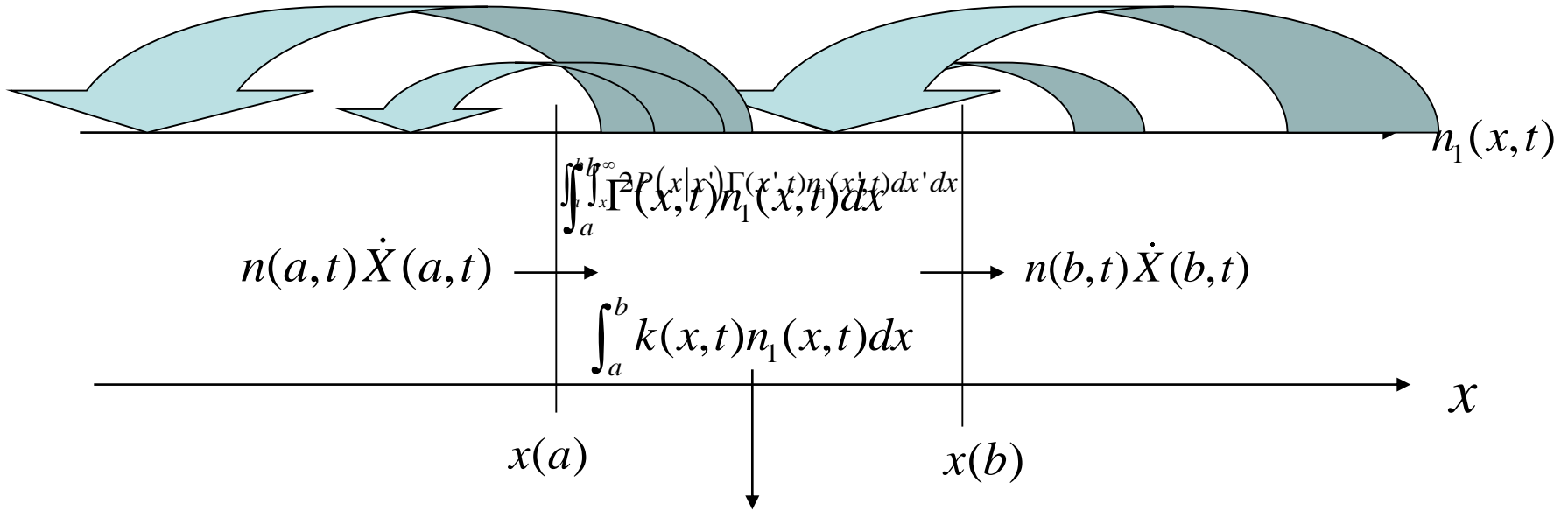
$$\frac{\partial n_1(1, x, t)}{\partial t} + \frac{\partial}{\partial x} \left( \dot{X}(1, x, t) n_1(1, x, t) \right) = \sum_{i,1} \Gamma_{i,1}(i, x, t) n_1(i, x, t) - \sum_{i,1} \Gamma_{1,i}(x, t) n_1(1, x, t)$$

Growth rate

Transition rates



# Cell division and death



$$\frac{\partial n_1(x, t)}{\partial t} + \frac{\partial}{\partial x} \left( \dot{X}(x, t) n_1(x, t) \right) = \int_x^\infty 2P(x|x') \Gamma(x', t) n_1(x', t) dx' - (\Gamma(x, t) + k(x, t)) n_1(x, t)$$

Growth rate

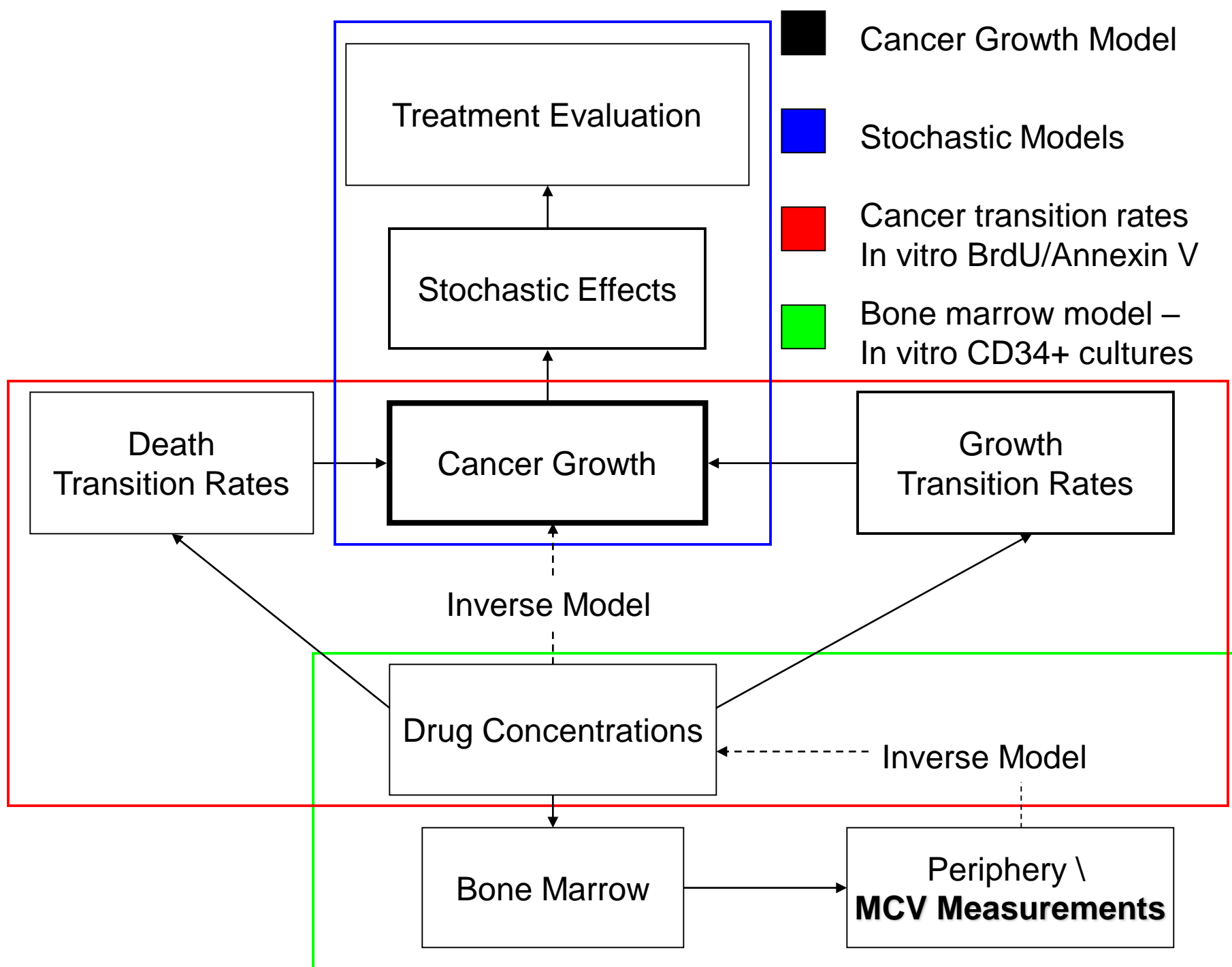
Division rate

Death rate

# Mathematical Model

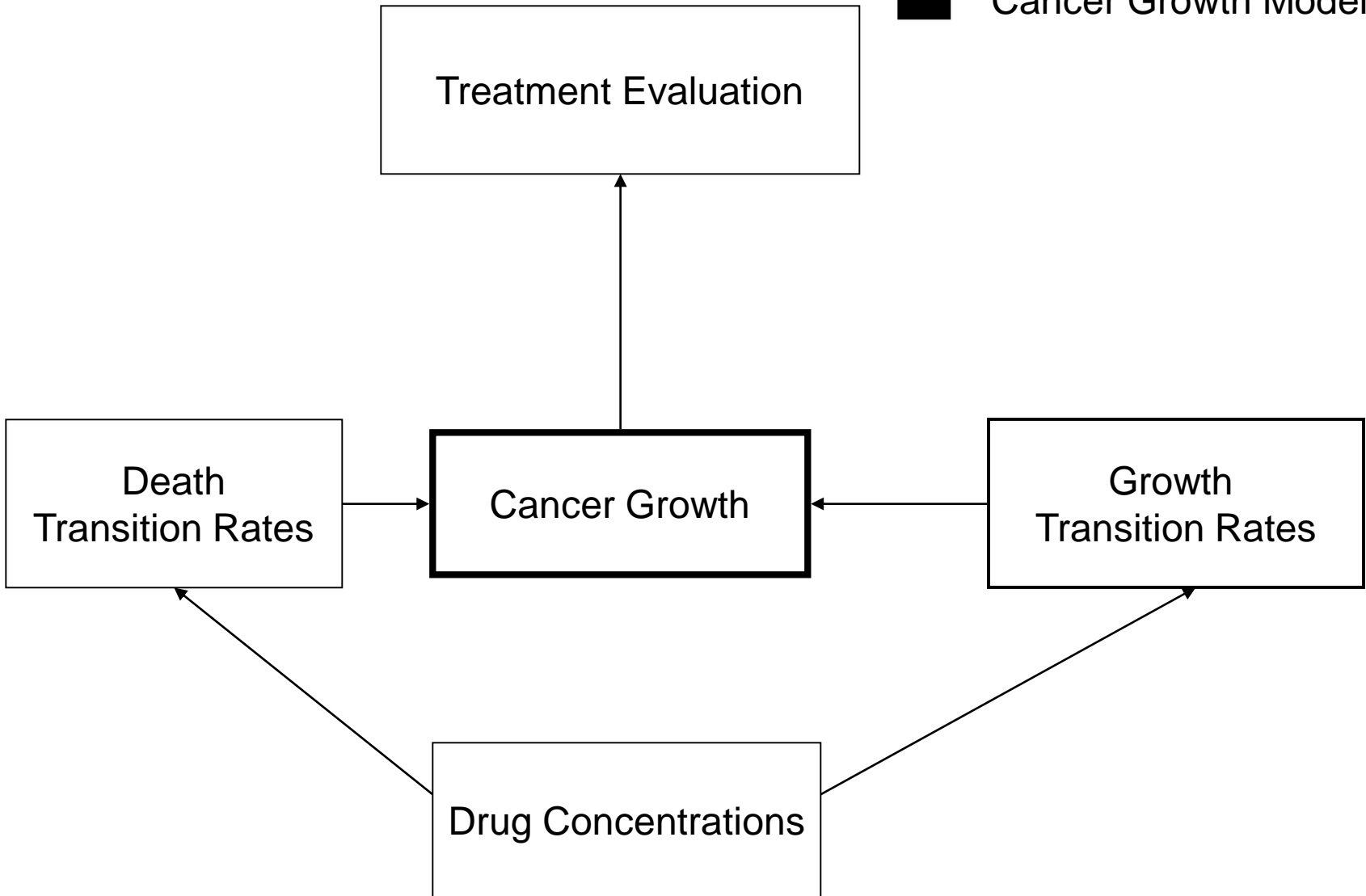
## Cellular -> Macroscopic

- Tumor size and character
  - Simulate cells and observe bulk behavior
- Cell behavior
  - Mutations
  - Spatial effects
  - Cell cycle phase
  - Quiescence
  - Pharmacodynamics
- Mostly theoretical work
  - Occasionally some parameters available
- Optimization of chemo
  - Dosage
  - Frequency
  - Drug combinations
- Solid tumor
  - Size limitations due to nutrients and inhibitors
- Leukemia
  - Quiescent and proliferating populations
  - Cell cycle-specific chemotherapy
- Mutations
  - Branching process



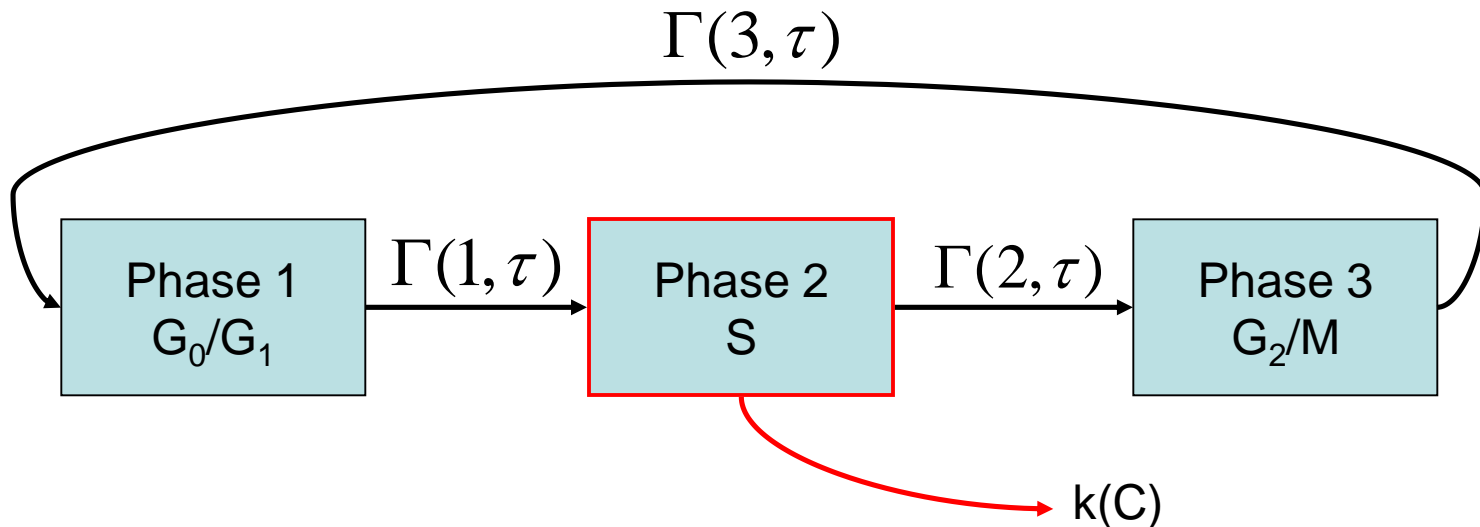


Cancer Growth Model



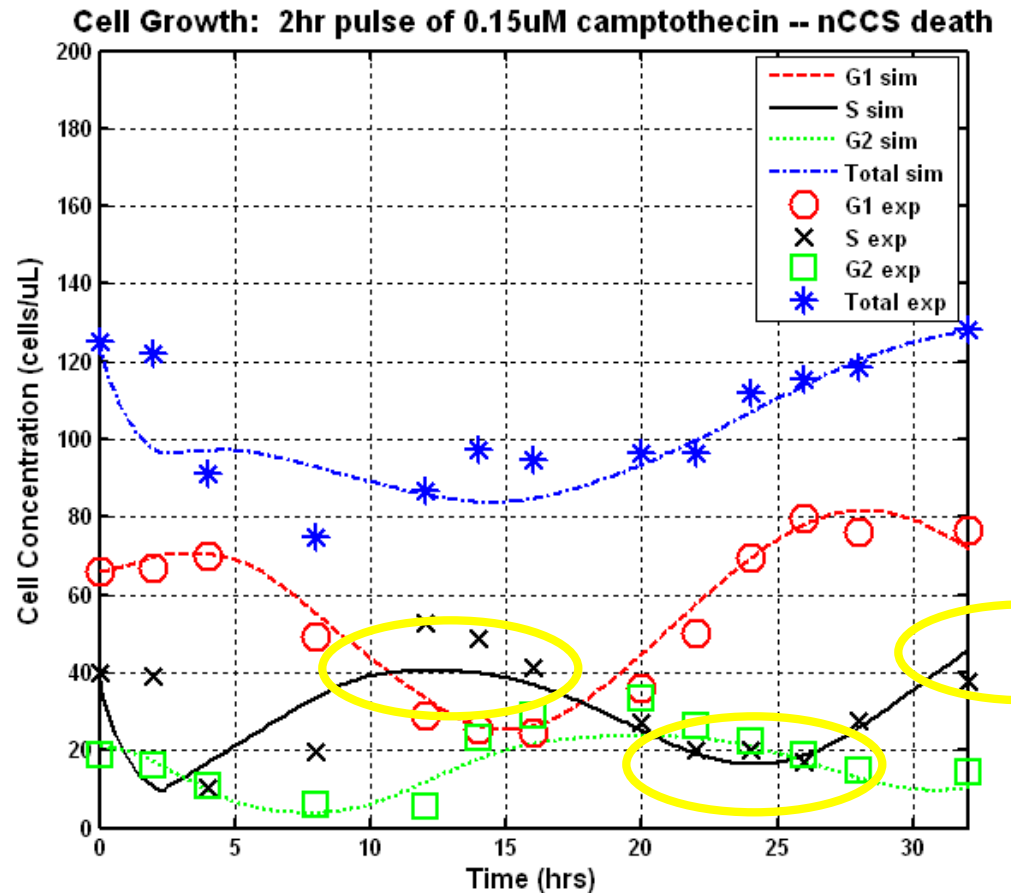
# Example – Cell Cycle Specific Behaviors

- Cell cycle specific drug
  - Discrete – cell cycle phase,  $p$
  - Continuous – age,  $\tau$ , (time since last transition)
- $n_1(p, \tau, t)$



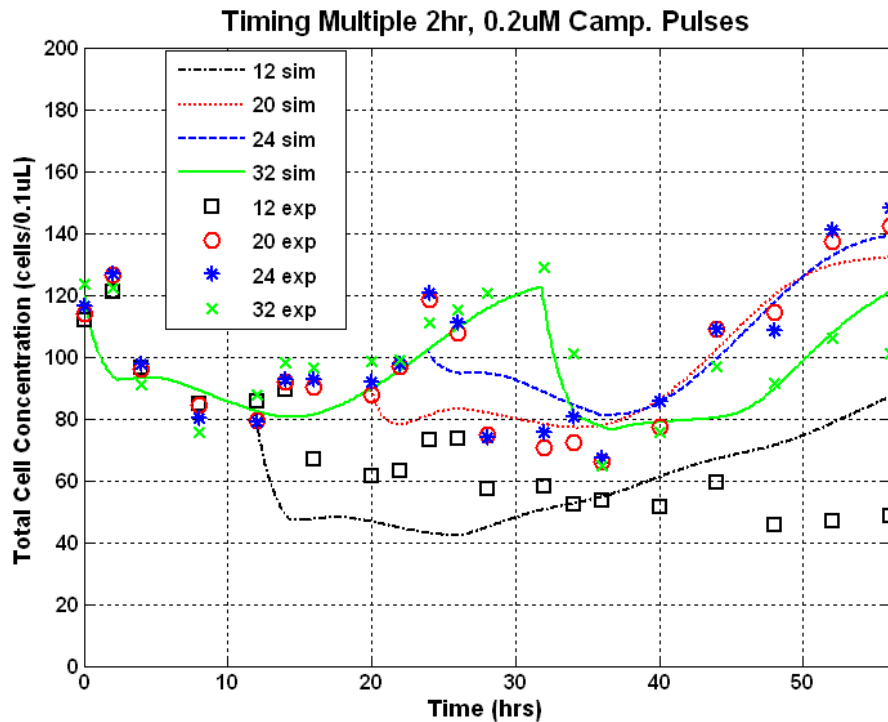
# In vitro verification

- Total population dynamics
- Phase oscillations
  - Period
  - Amplitude
  - Dampening
- Co-culture of Jurkat and HL60 performed for selective treatment



# Use in Treatment Designs

Timing effects



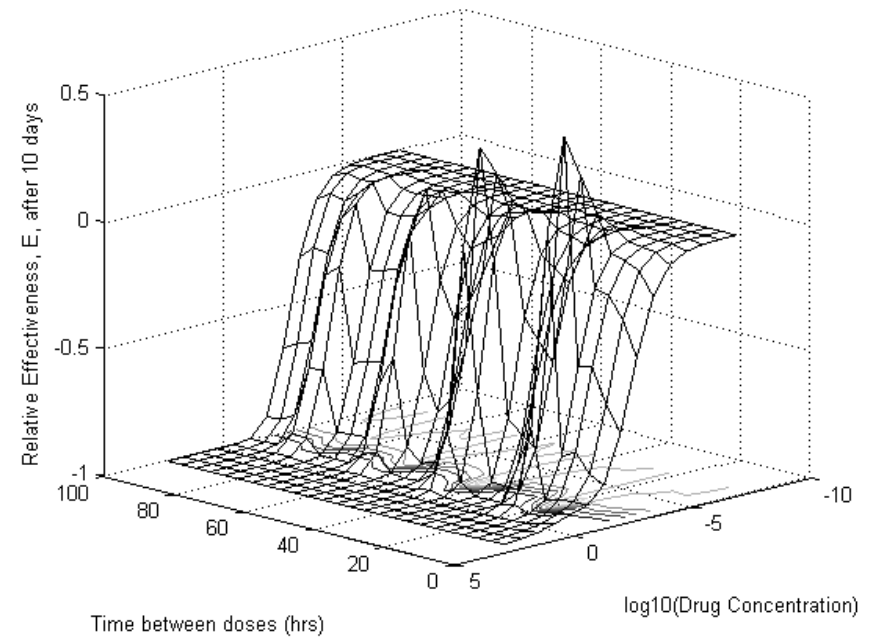
Treatment Optimization

Heathy cells vs. cancerous

Drug Dosage

Administration Timings

Treatment Design with similar S phase percentages

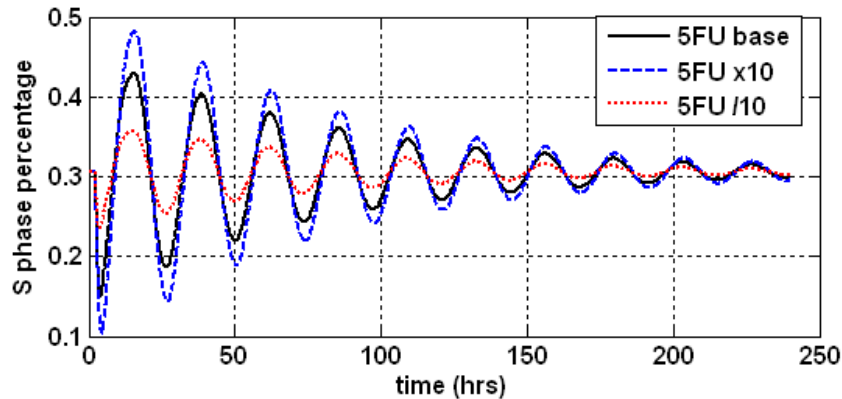


# Model Extrapolation

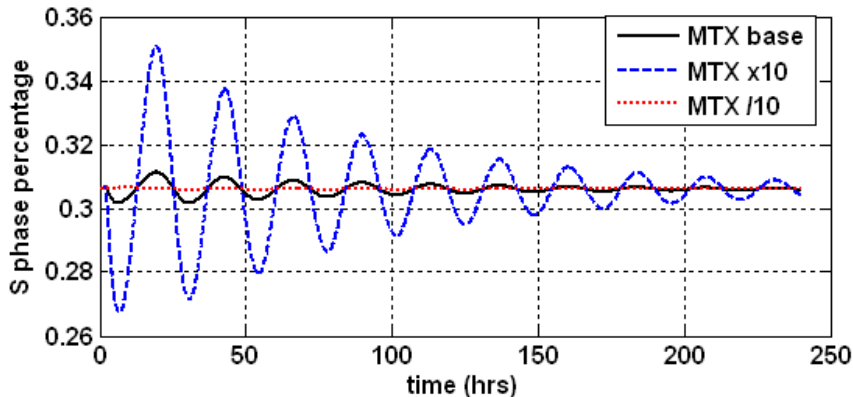
## In vivo factors

Drug half-life

**FU oscillations**

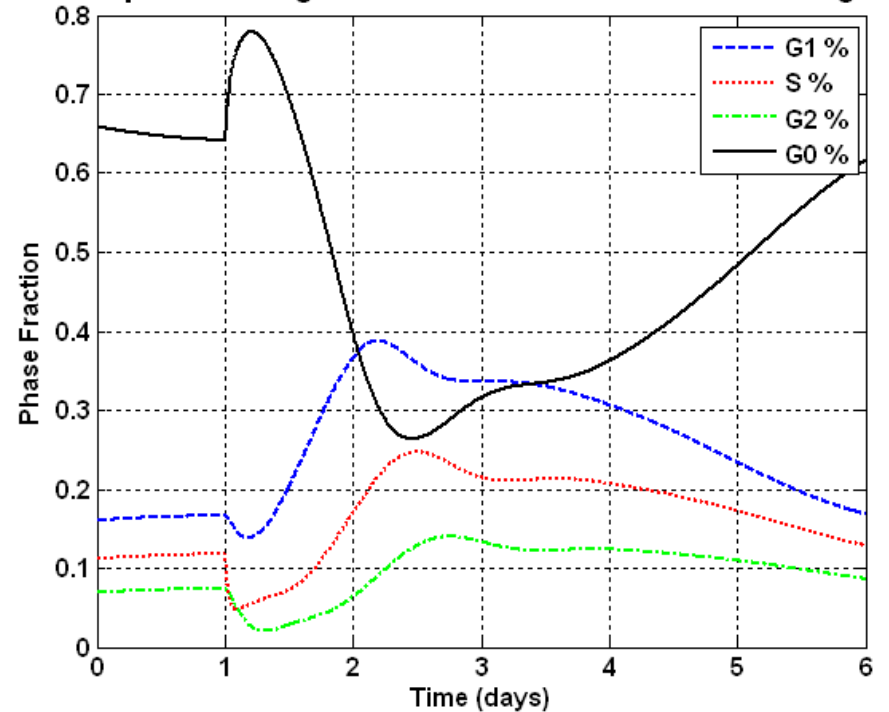


**MTX oscillations**



Activation of quiescent population in bone marrow

**Response to single CAF treatment - All Phase Percentages**





# Age-averaged PBM

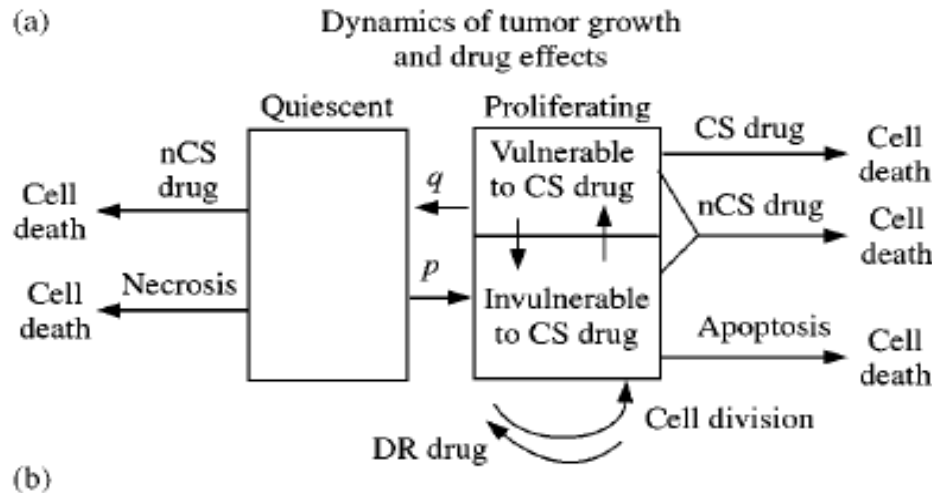
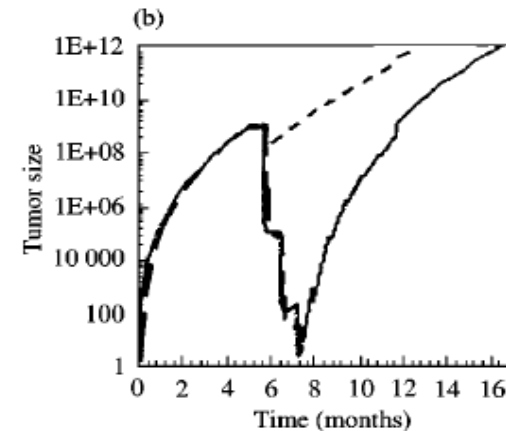
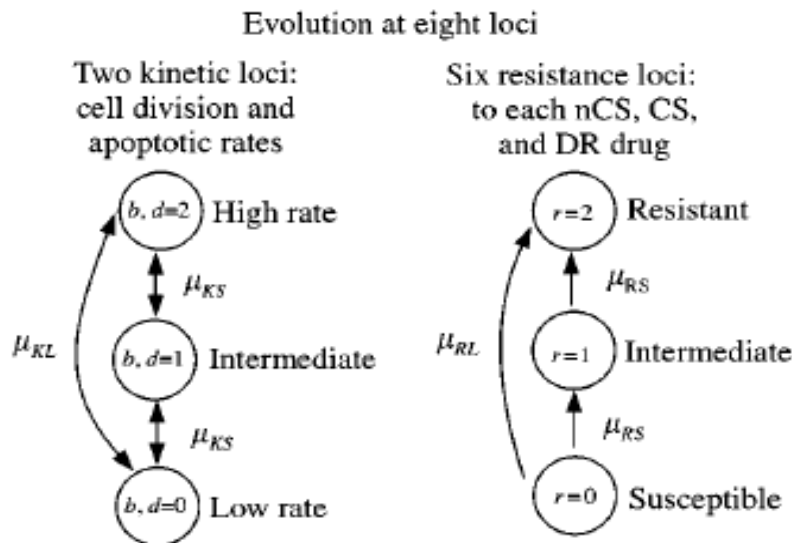


FIG. 1. (a) Diagram of the cellular dynamics of the model. Cells in the proliferating state may divide, die via apoptosis, or enter the quiescent state. Cells in the resting state may die via necrosis or transfer back to the proliferating state. nCS drugs kill cells in both the resting and the proliferating states, while CS drugs kill only a fraction of proliferating cells in a particular phase of the cell cycle. DR drugs slow the progression of cells through cell division. (b) Diagram of mutations in the model that generate intra-tumor heterogeneity. Mutations may occur at loci affecting the cell division and apoptotic rates, as well as at six loci each controlling drug resistance to a particular drug. Mutations of large effect occur with probability  $\mu_{RL} = \mu_{KL} = 10^{-6}$ , and those of small effect with probability  $\mu_{RS} = \mu_{KS} = 10^{-4}$ .



# Prognosis Tree

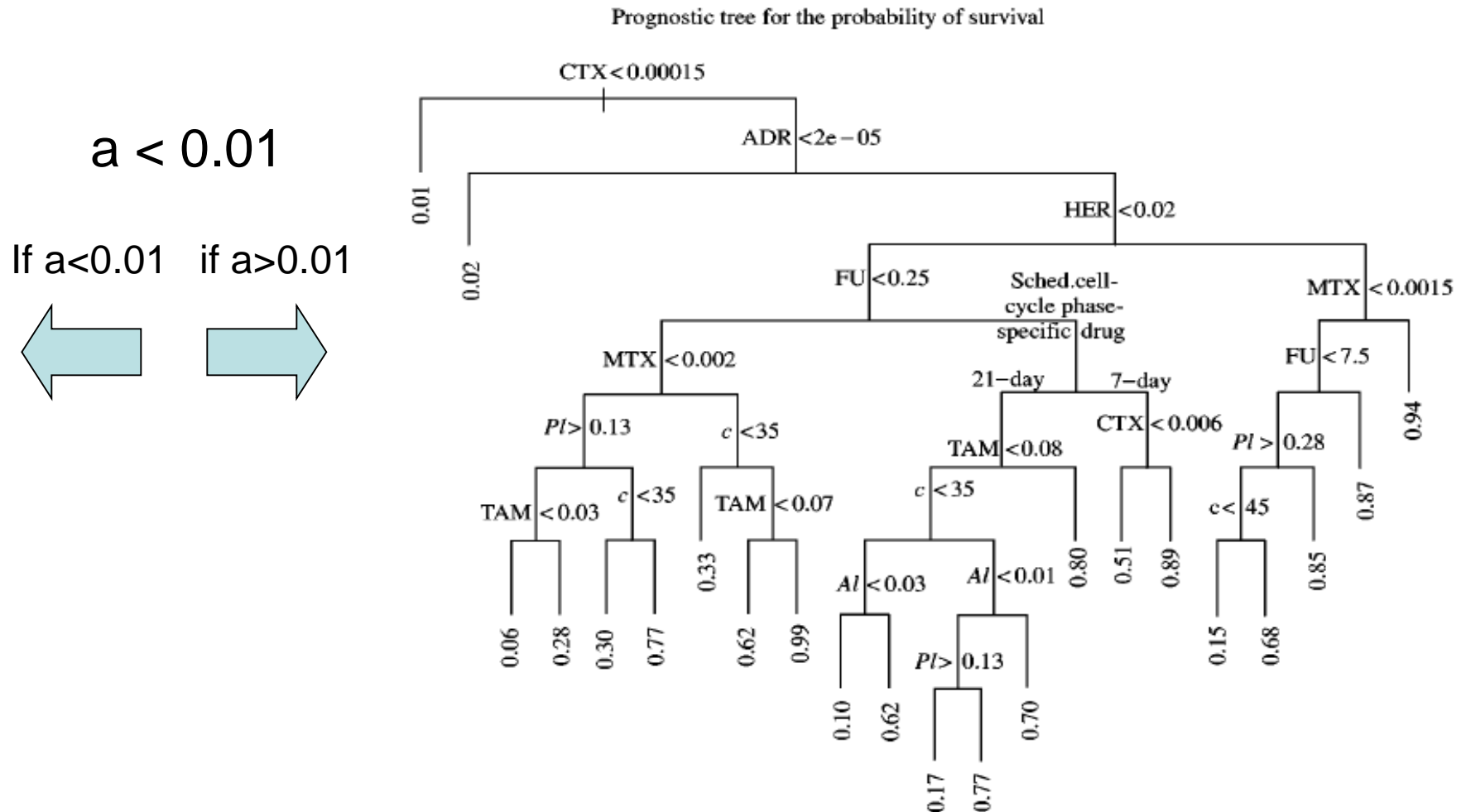
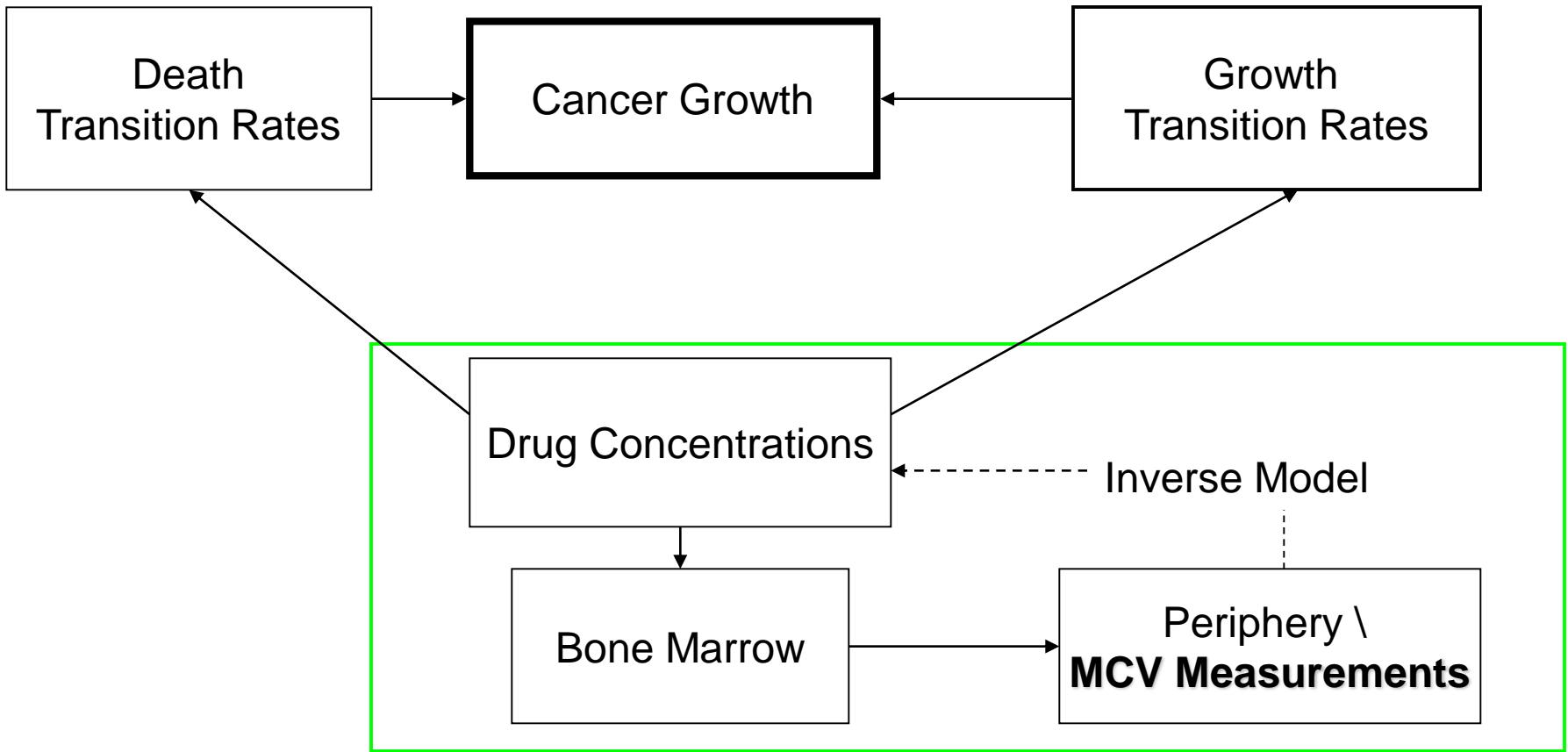


FIG. 6. Decision tree constructed from the simulation of 26 896 tumors. The physician-controlled factors of drug dose and schedule are illustrated in bold, and the tumor-specific factors of *AI*, *PI*, and *c* are in italics. (a) Probability of survival. (b) Duration in months from tumor detection to patient death.

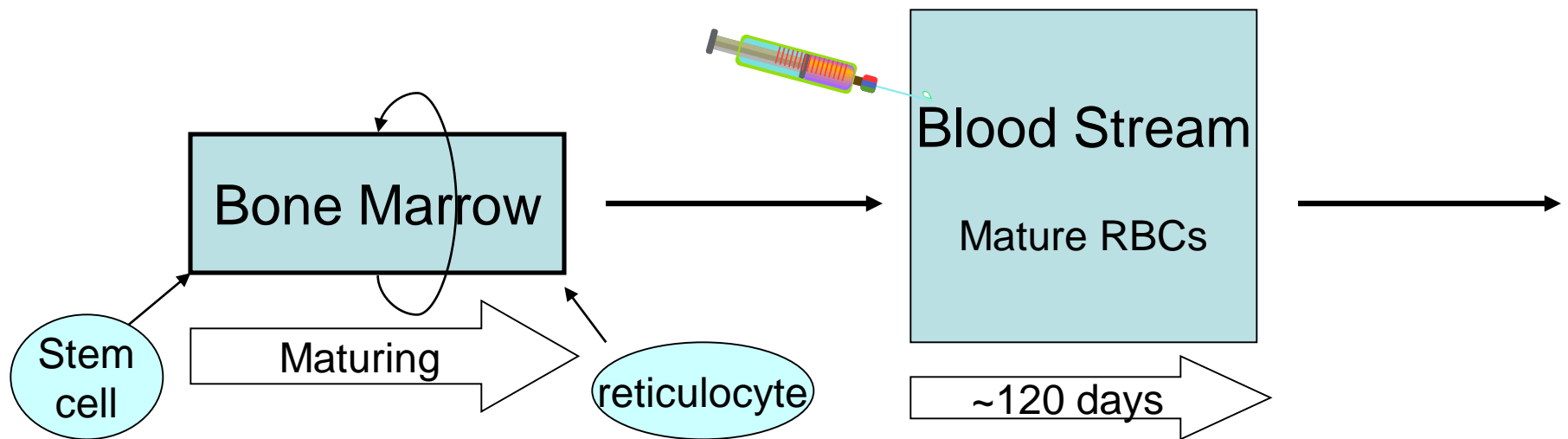
■ Cancer Growth Model

■ Bone marrow model –  
In vitro CD34+ cultures



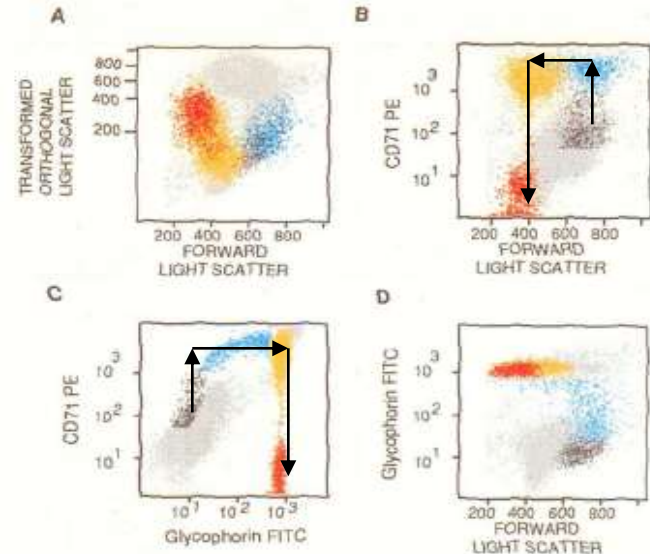
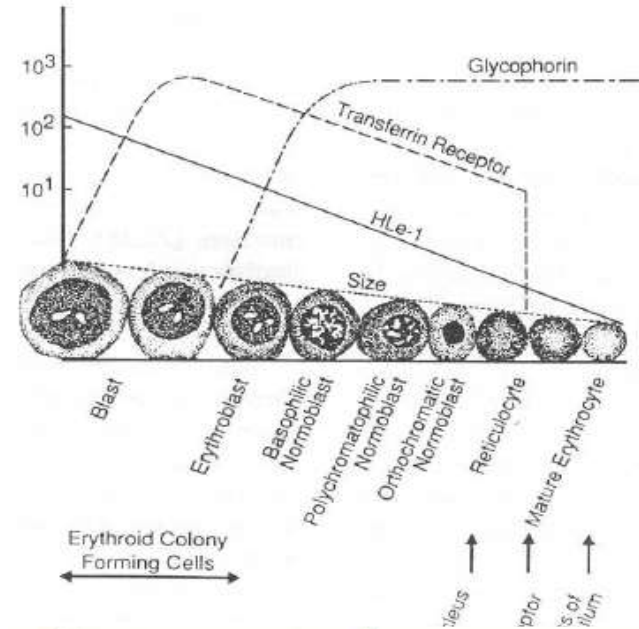
# Modeling a surrogate marker for the drug 6MP

- Bone marrow
  - 6MP inhibits DNA synthesis
- Blood stream
  - Red blood cells (RBCs) become larger
  - RBC size correlates with steady-state 6MP level



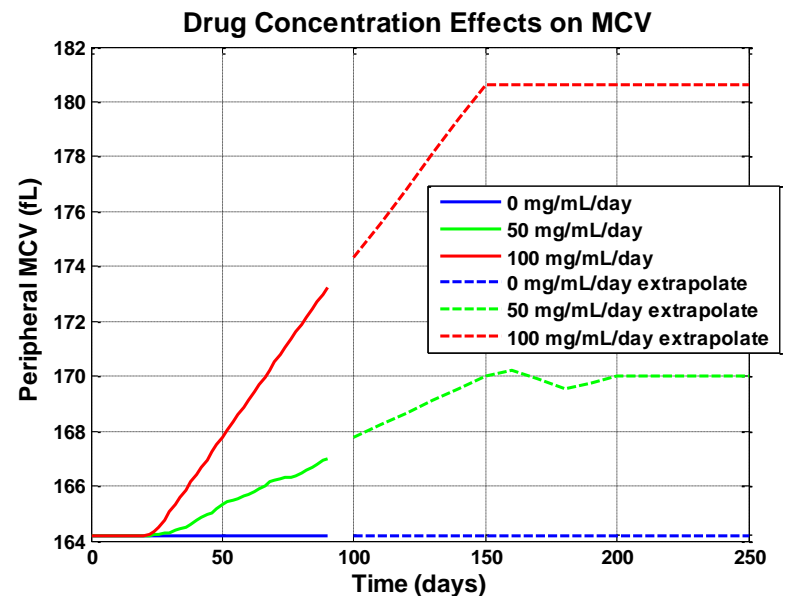
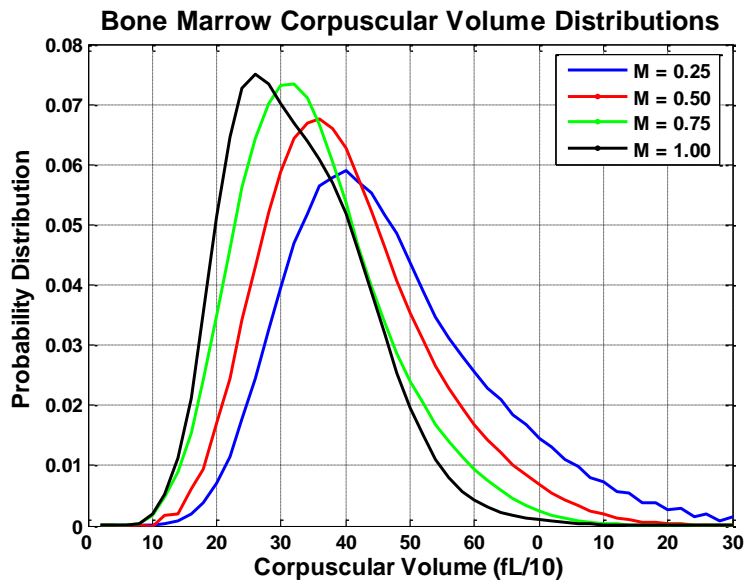
# RBC Maturation

- Cell volume
- DNA inhibition
  - Time since division
  - *DNA synthesis rate*
- Maturation
  - Discrete state
    - Transferrin and glycoprotein A
  - *Continuous*
    - Time spent in state



# Forward Model (hypothetical)

- Days: 6TGN reaches steady-state (SS)
- Weeks: Marrow maturation in new SS
- Months: Periphery in new SS





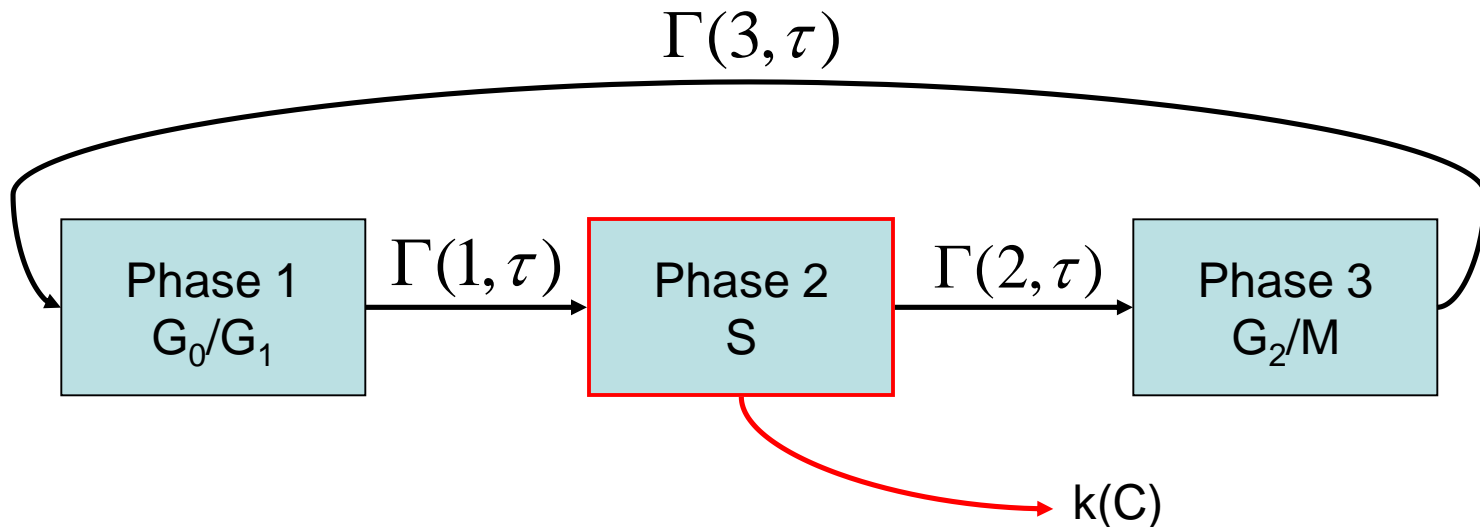
# Topics

- Modeling Philosophies
  - Where Population Balance Models (PBM) fit in
- Important Characteristics
- Framework
- Uses
  - Cancer Examples
- Parameter Identification
- Potential Applications



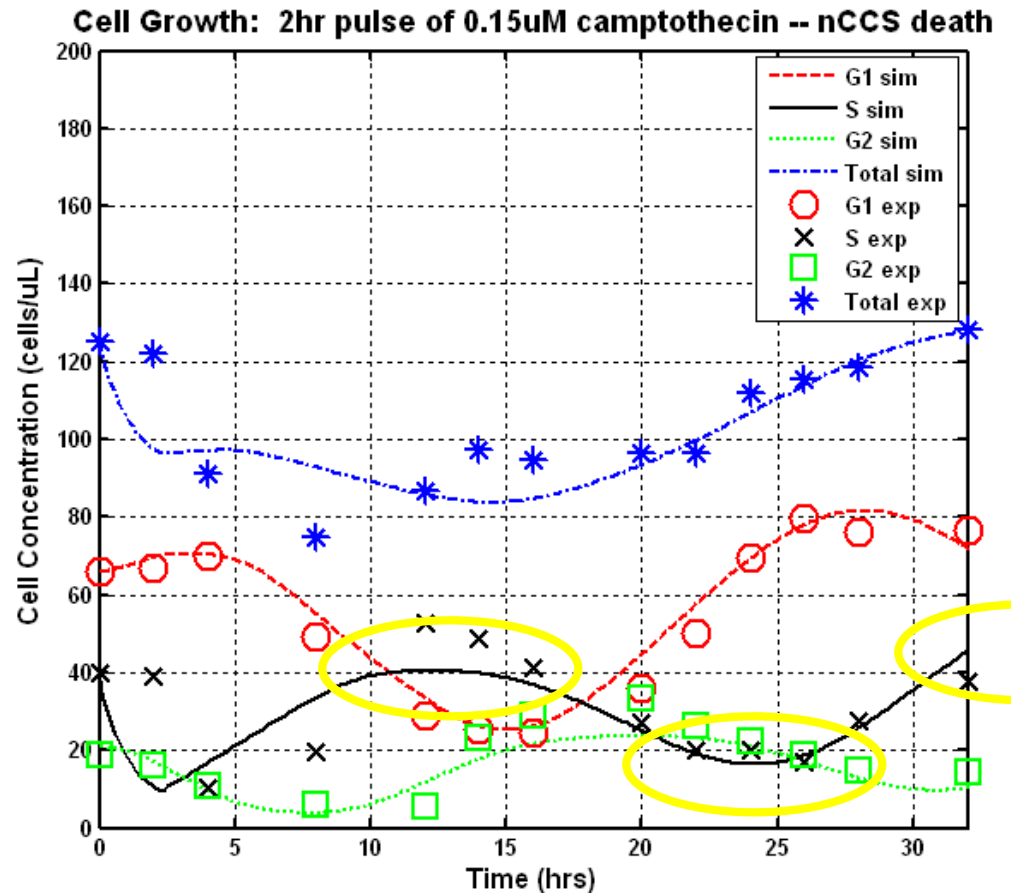
# Example – Cell Cycle Specific Behaviors

- Cell cycle specific drug
  - Discrete – cell cycle phase,  $p$
  - Continuous – age,  $\tau$ , (time since last transition)
- $n_1(p, \tau, t)$



# In vitro verification

- Total population dynamics
- Phase oscillations
  - Period
  - Amplitude
  - Dampening
- Co-culture of Jurkat and HL60 performed for selective treatment



# Age-averaged PBM

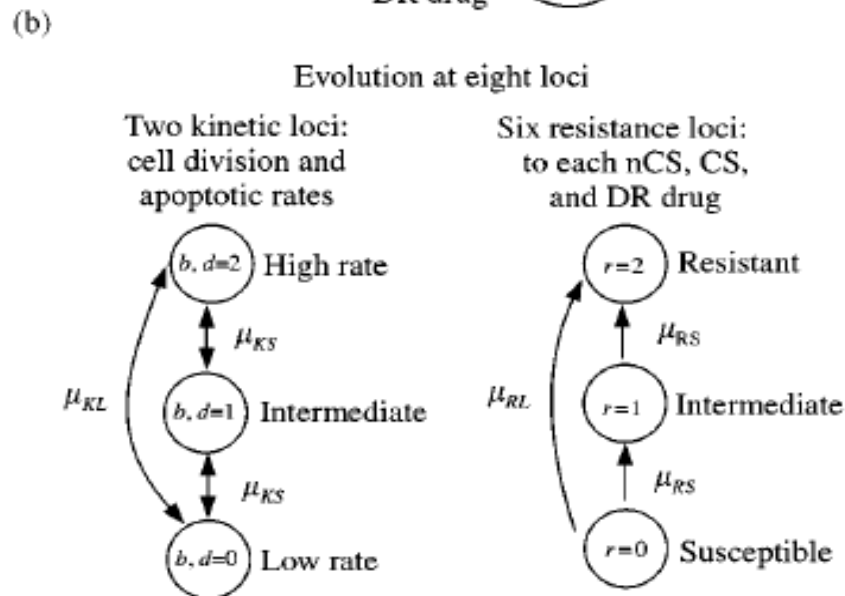
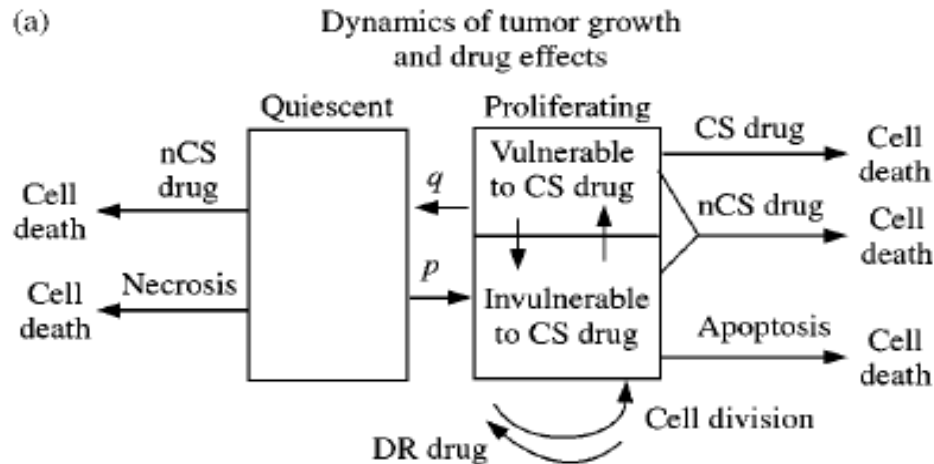
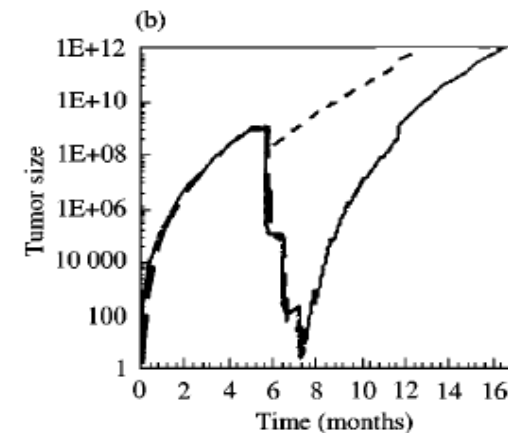


FIG. 1. (a) Diagram of the cellular dynamics of the model. Cells in the proliferating state may divide, die via apoptosis, or enter the quiescent state. Cells in the resting state may die via necrosis or transfer back to the proliferating state. nCS drugs kill cells in both the resting and the proliferating states, while CS drugs kill only a fraction of proliferating cells in a particular phase of the cell cycle. DR drugs slow the progression of cells through cell division. (b) Diagram of mutations in the model that generate intra-tumor heterogeneity. Mutations may occur at loci affecting the cell division and apoptotic rates, as well as at six loci each controlling drug resistance to a particular drug. Mutations of large effect occur with probability  $\mu_{RL} = \mu_{KL} = 10^{-6}$ , and those of small effect with probability  $\mu_{RS} = \mu_{KS} = 10^{-4}$ .



# Prognosis Tree

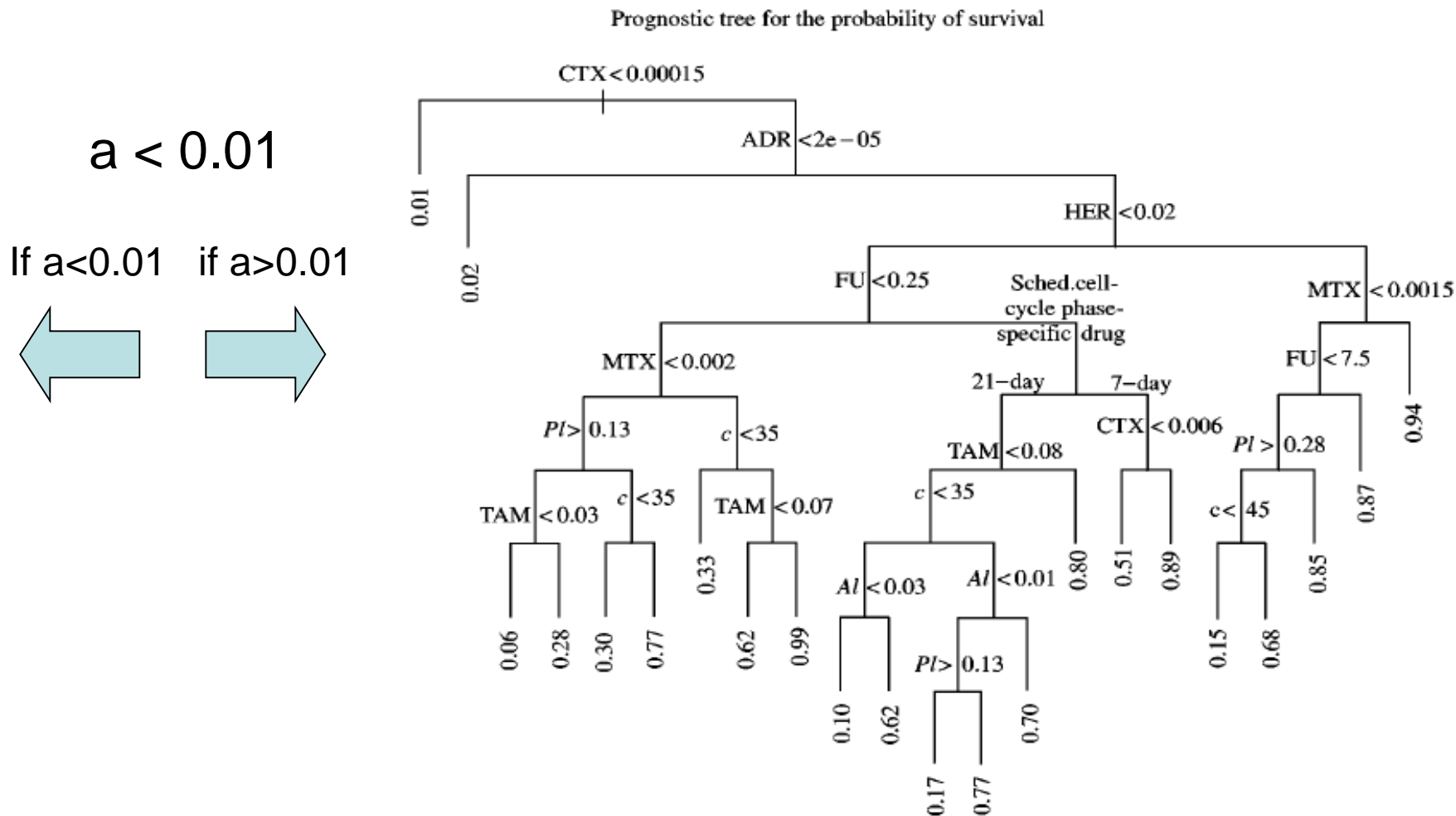
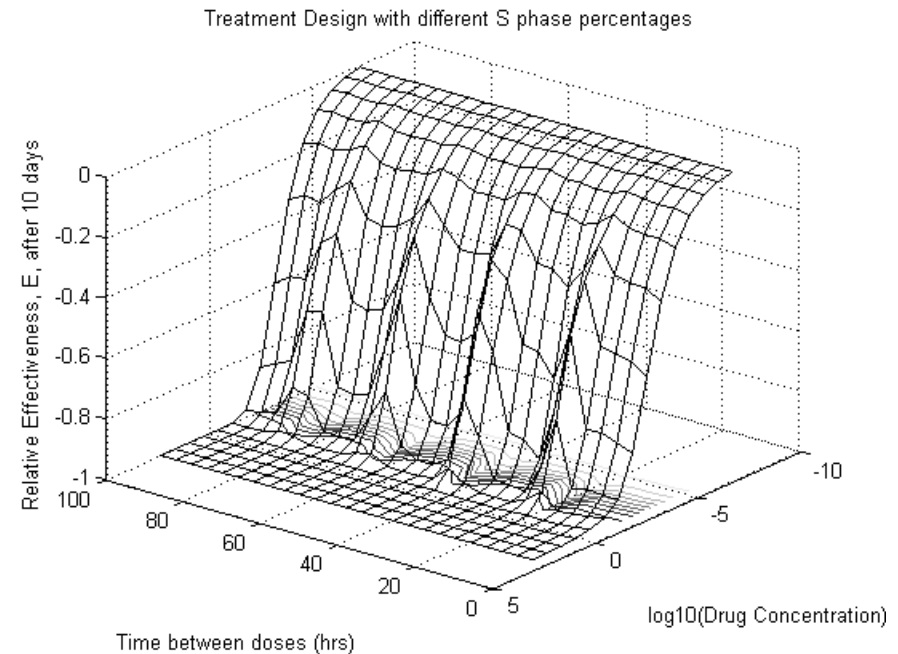
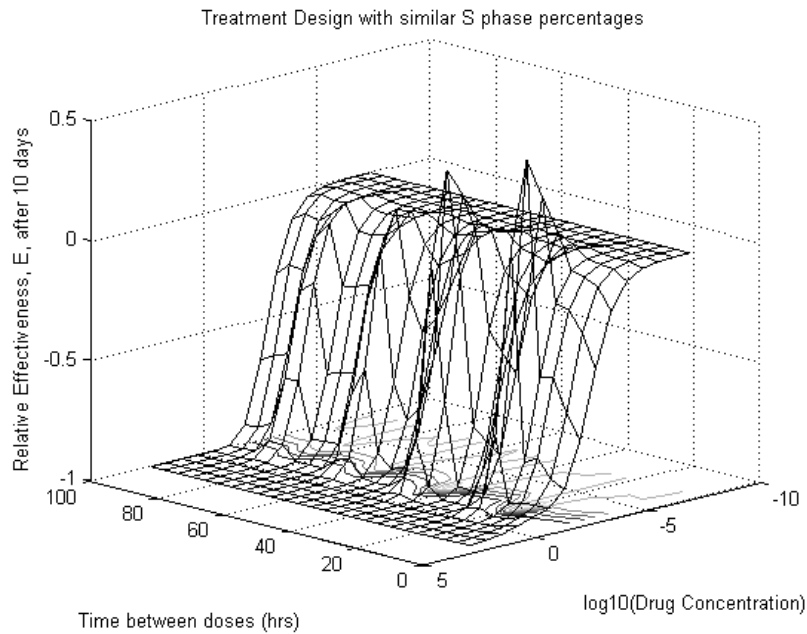


FIG. 6. Decision tree constructed from the simulation of 26 896 tumors. The physician-controlled factors of drug dose and schedule are illustrated in bold, and the tumor-specific factors of *AI*, *PI*, and *c* are in italics. (a) Probability of survival. (b) Duration in months from tumor detection to patient death.

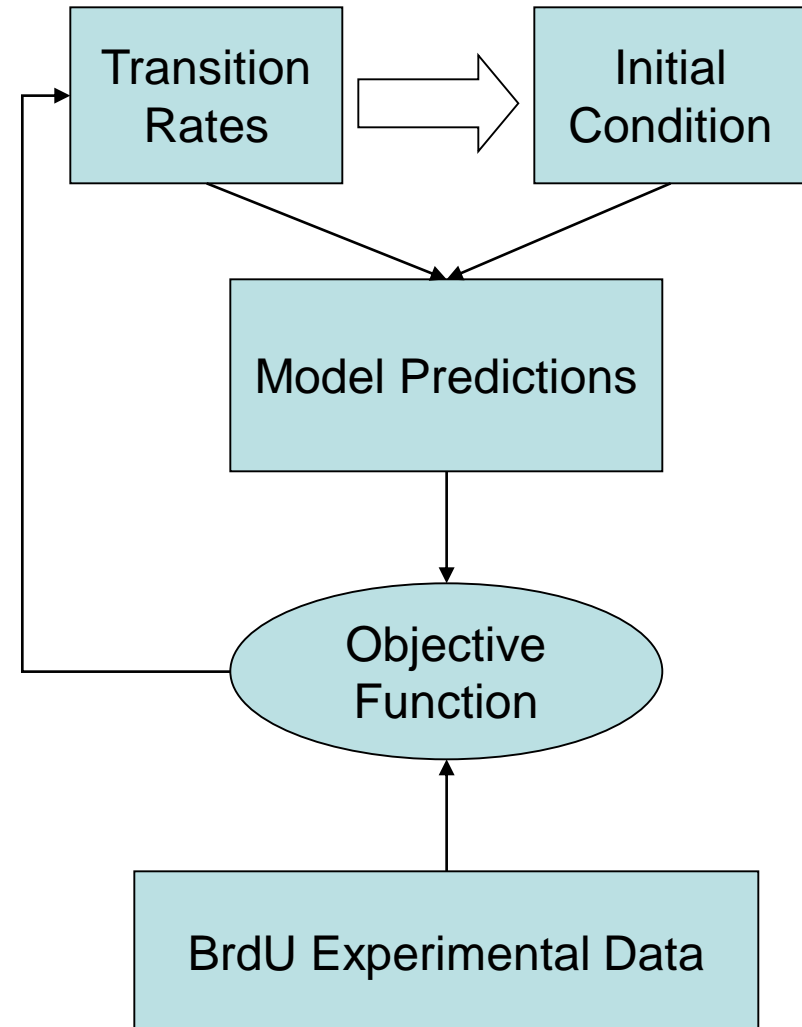
# Ramkrishna's resonance chemotherapy model



- Hypothesis testing (potential protocols)
- Patient specific treatments
- Treatment strength necessary or desired

# Cell Cycle Transitions $\Gamma(1, \tau)$

- Cannot measure ages
- Balanced growth
  - $\Gamma \Rightarrow$  age distributions
  - Predict dynamics
- BrdU
  - Labels S subpopulation
  - Phase transient amidst balanced growth
- Match transition rates



# Initial Condition

- Balanced growth

$$n_i(\tau, t) = N_i(t) f_i(\tau) \quad \text{where} \quad \int_0^{\infty} f_i(\tau) d\tau = 1$$

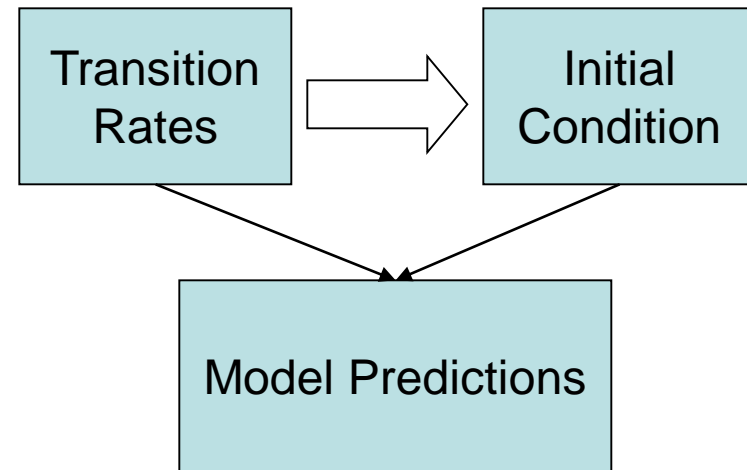
- Population increase

$$\frac{d\mathbf{N}(t)}{dt} = \mathbf{D}\mathbf{N}(t)$$

$$D_{ij} = \int_0^{\infty} [D_{ij}^{in}(\tau) - D_{ij}^{out}(\tau)] f_j(\tau) d\tau$$

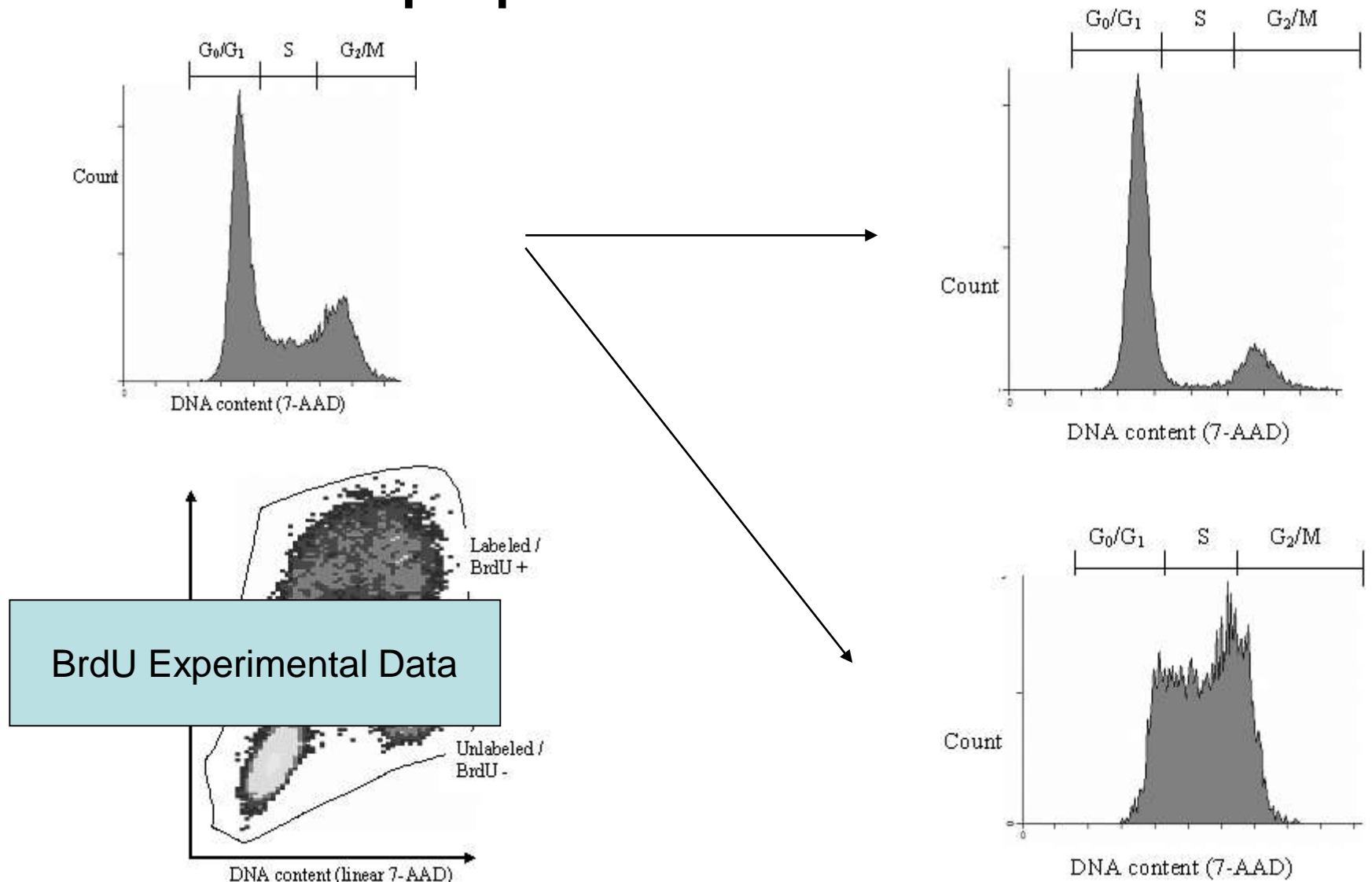
- Eigen analysis

- Balanced growth age-distribution



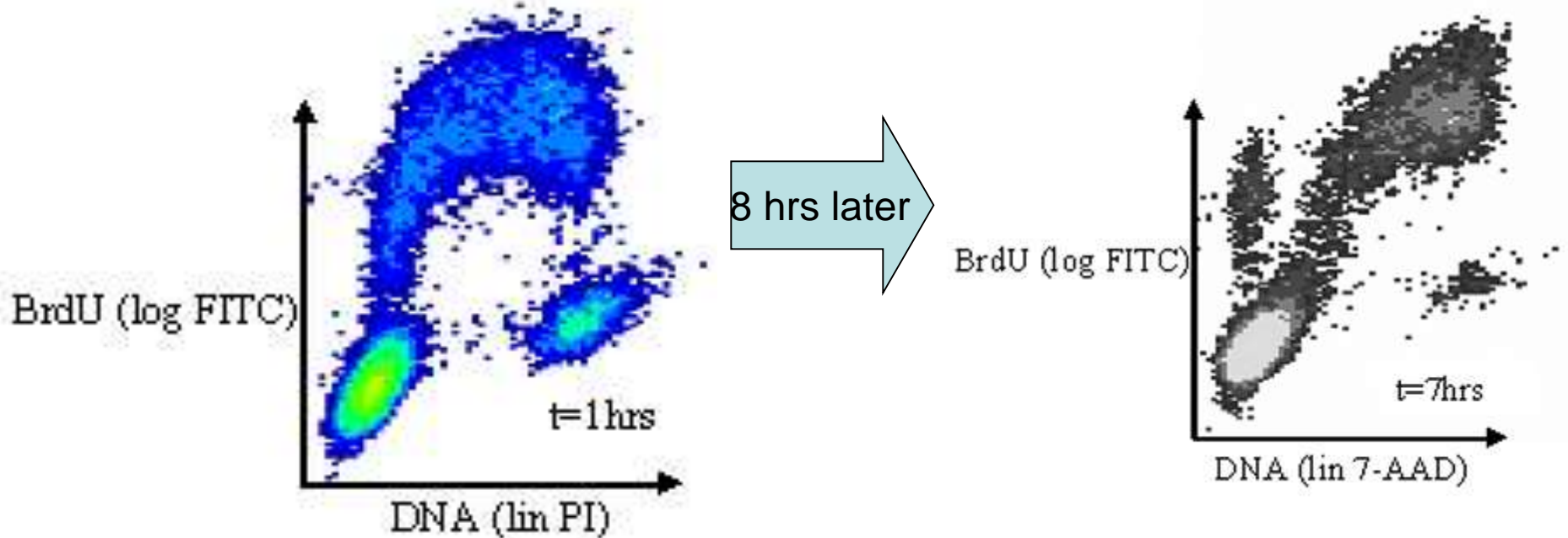
$$f_i(\tau) = \left\{ \int_0^{\infty} \exp \left[ - \int_0^{\tau'} (D_{ii}^{out}(\tau'') + \mu) d\tau'' \right] d\tau' \right\}^{-1} \exp \left[ - \int_0^{\tau} (D_{ii}^{out}(\tau') + \mu) d\tau' \right]$$

# Unbalanced subpopulation growth amidst total population balance growth

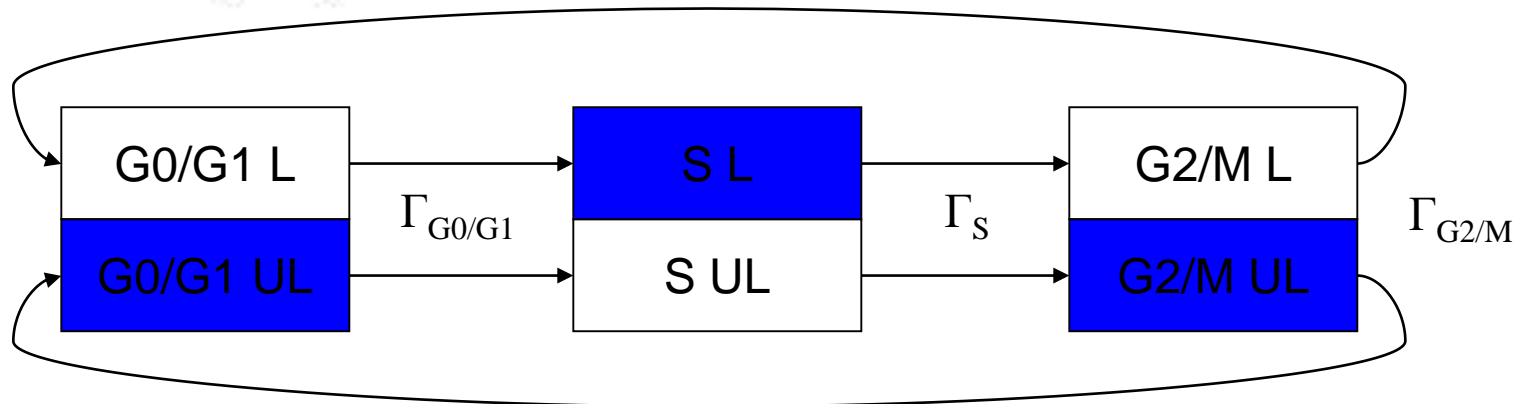




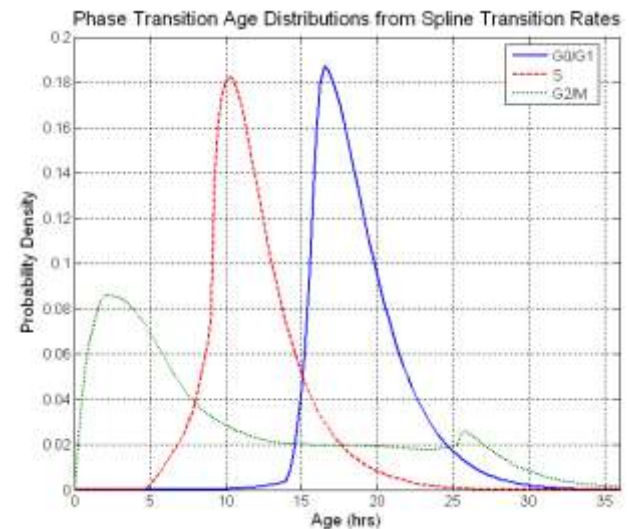
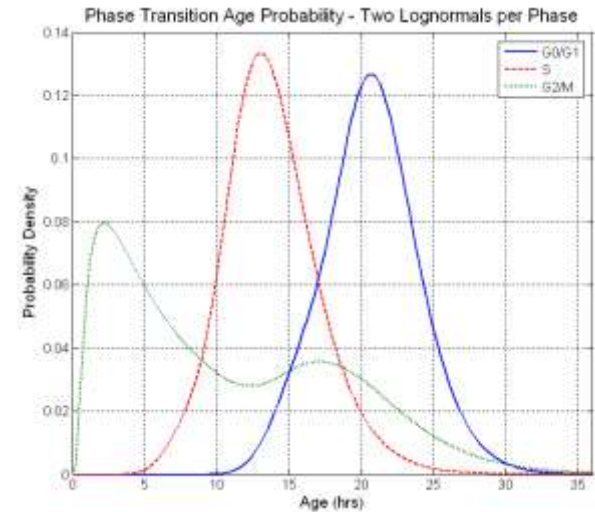
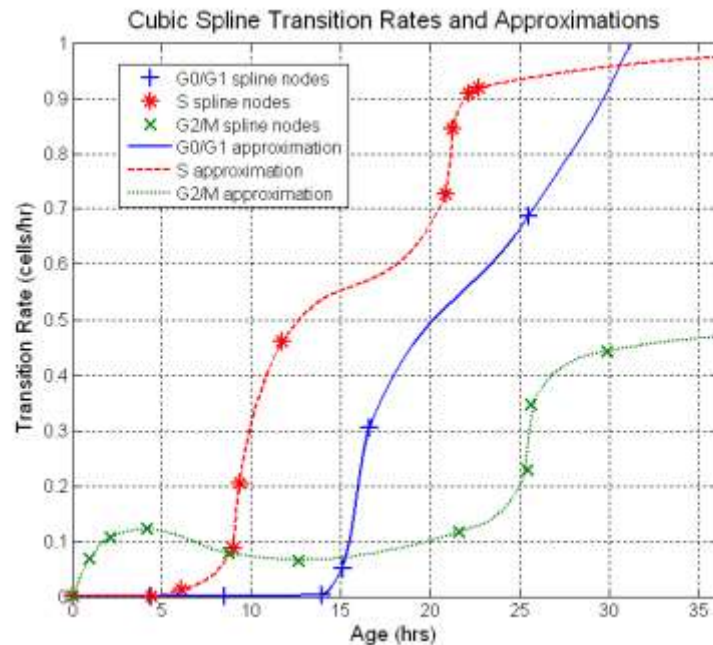
# Cell Cycle Transition Rates



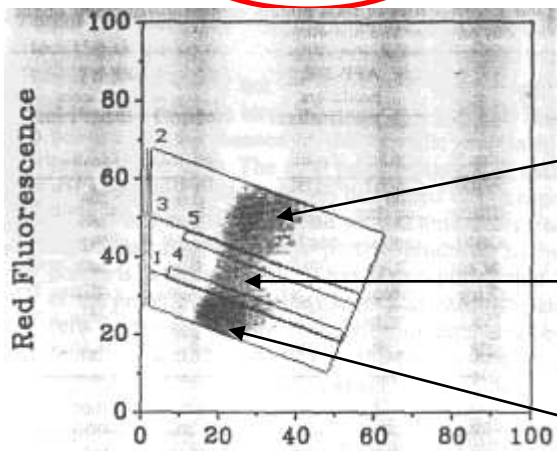
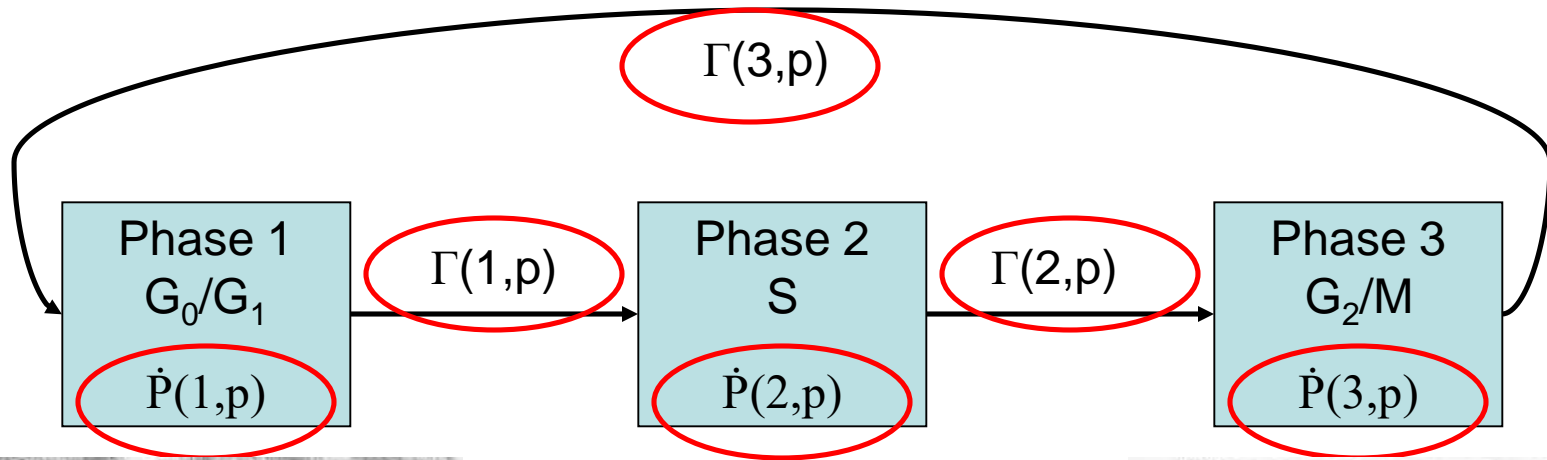
**BrdU  
Labeling  
(cell cycle)**



# Transition Rates $\Leftrightarrow$ Residence Time Distribution



# Example – protein (p) structured cell cycle rates



**Forward Angle Light Scatter**

Figure 8. Cytogram of red fluorescence intensity vs. forward-angle light scattering intensity for AFP-27E cells which were stained with propidium iodide and FITC after receiving the described treatment. Labeled regions represent gates that were used to generate protein content distributions on the condition that cell properties fell within the indicated region. This cytogram was constructed with data acquired for  $5 \times 10^5$  cells.

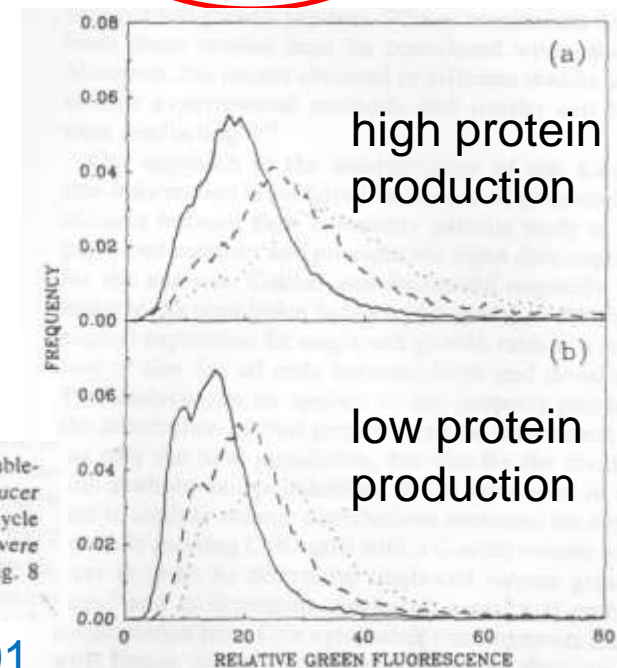
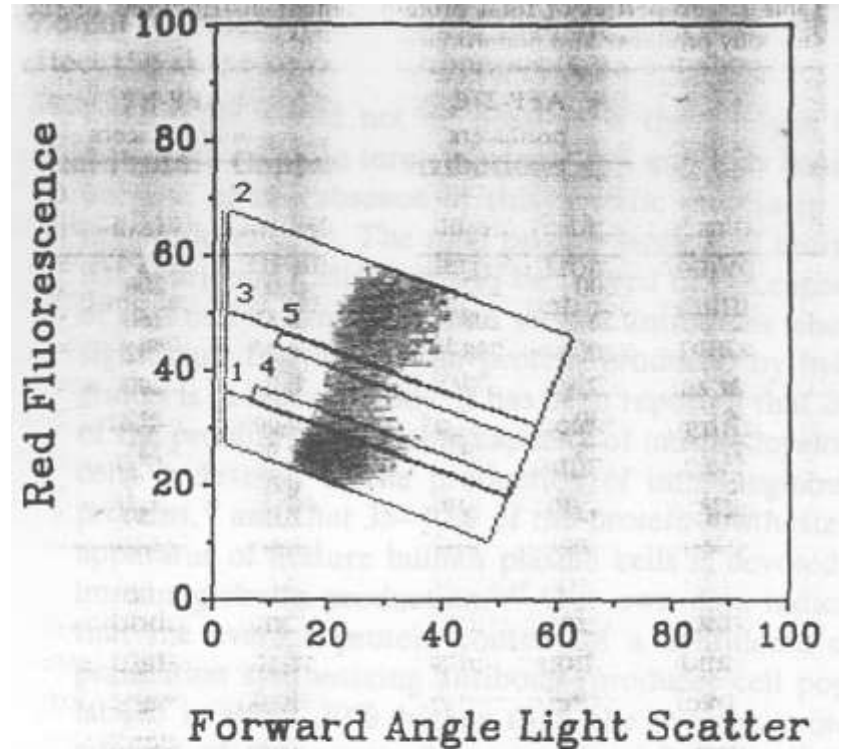


Figure 9. Distributions of green fluorescence intensity for double-stained (a) AFP-27E producer cells and (b) AFP-NP nonproducer cells from the  $G_1$  (—),  $S$  (----), and  $G_2 + M$  (····) cell cycle phases. These distributions are frequency functions which were obtained according to the described gating procedure (see Fig. 8 and text).

# Rate Identification Procedure

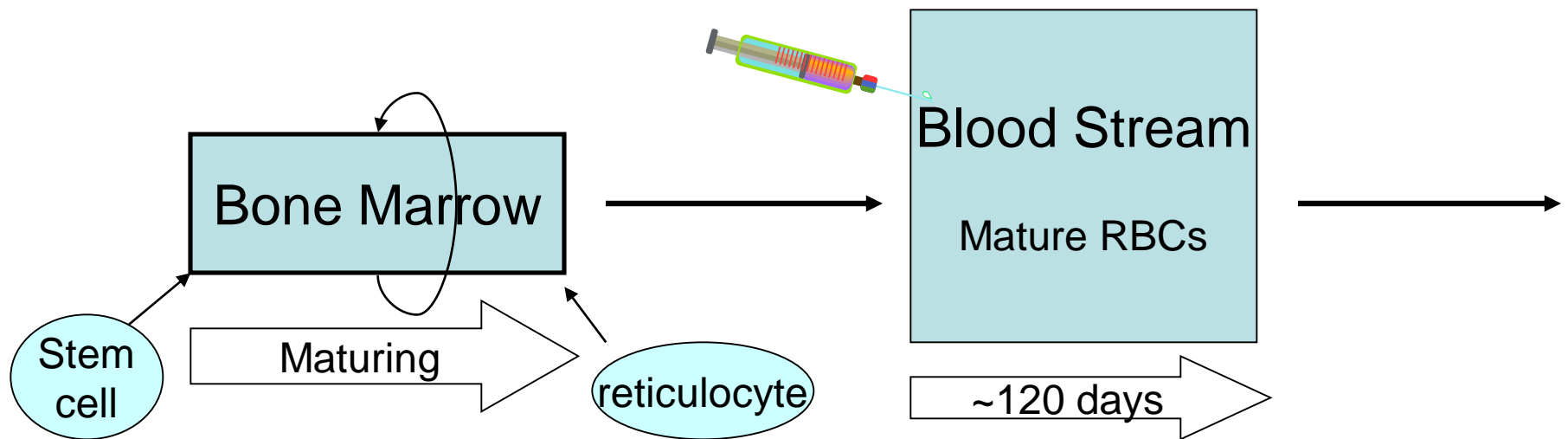
- Assume balanced growth
  - Specific growth rate
- Measure
  - Stable cell cycle phase protein distributions
  - Protein distributions at transition
- Inverse model
  - Phase transition rates
  - Protein synthesis rates



**Figure 8.** Cytogram of red fluorescence intensity vs. forward-angle light scattering intensity for AFP-27E cells which were stained with propidium iodide and FITC after receiving the described treatment. Labeled regions represent gates that were used to generate protein content distributions on the condition that cell properties fell within the indicated region. This cytogram was constructed with data acquired for  $5 \times 10^4$  cells.

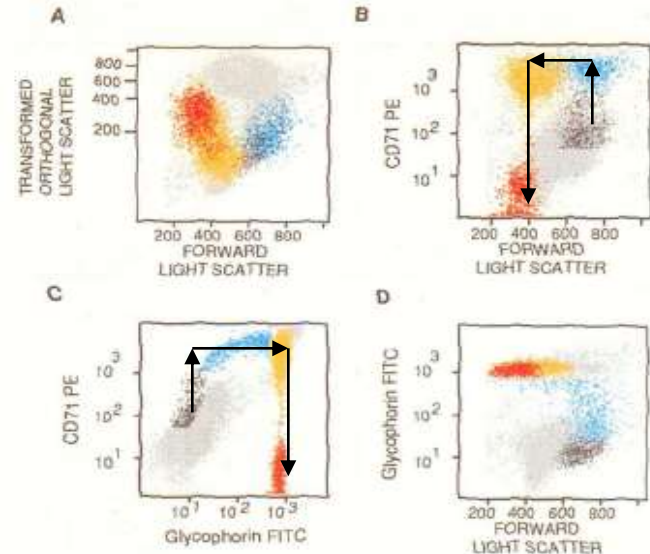
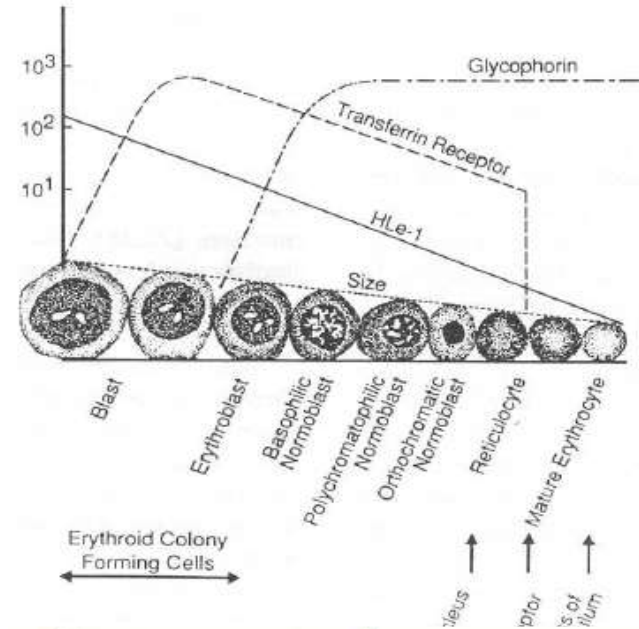
# Modeling a surrogate marker for the drug 6MP

- Bone marrow
  - 6MP inhibits DNA synthesis
- Blood stream
  - Red blood cells (RBCs) become larger
  - RBC size correlates with steady-state 6MP level



# RBC Maturation

- Cell volume
- DNA inhibition
  - Time since division
  - *DNA synthesis rate*
- Maturation
  - Discrete state
    - Transferrin and glycoporphin A
  - *Continuous*
    - Time spent in state





# Maturation + division + growth rates

- Maturation
  - Tag cells in 1<sup>st</sup> stage with pkh26
  - Track movement
- Division
  - Cell generations
- Growth
  - Satisfy volume distributions

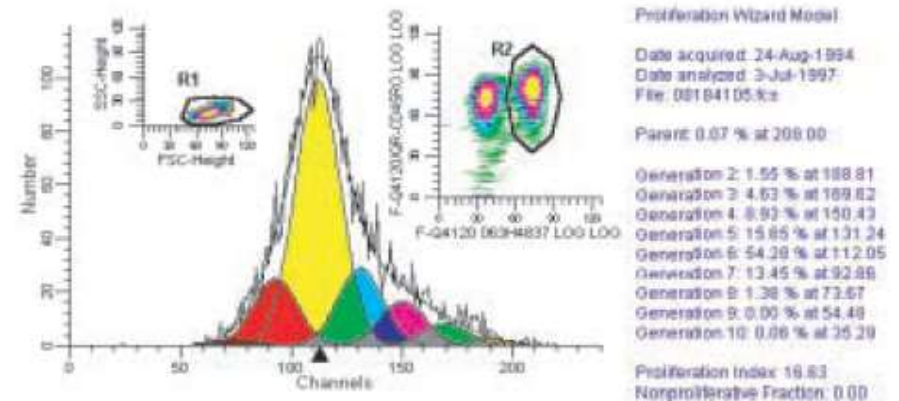
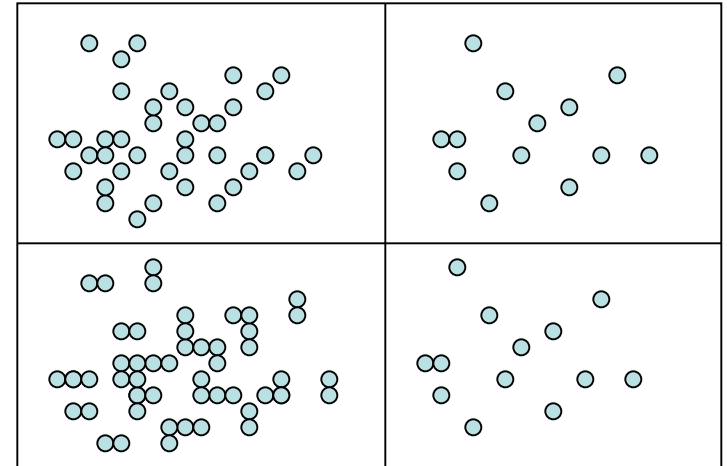


Figure 4. Deconvoluted histogram of CD4<sup>+</sup>/CD45RO<sup>+</sup> subset in PKH26-labeled PBL activated with 2.5  $\mu$ g/ml PHA for 6 days. Inset shows gates set by light scatter and immunophenotype.





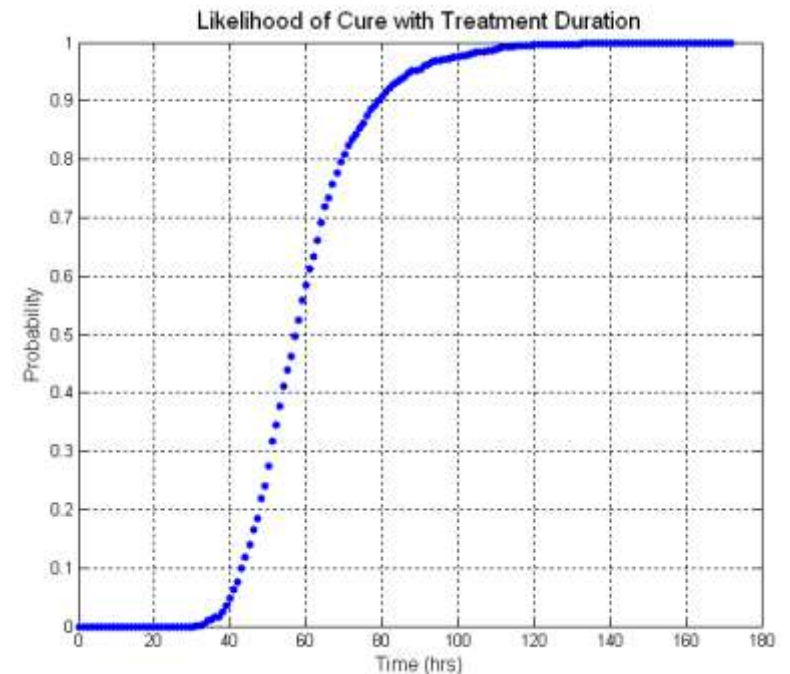
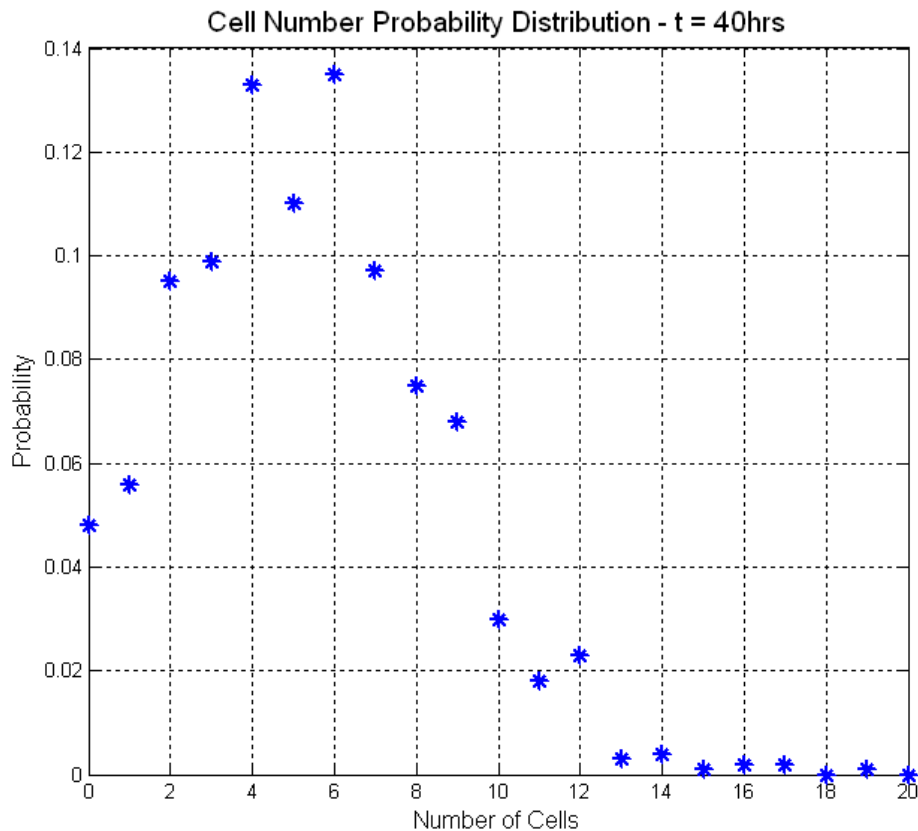
# Treatment Evaluation

- Actual cell behavior may be unobservable
  - Remission
  - Predictive models
- Model parameters known
- Treatment constantly adjusted
  - Immune system (Neutrophil count)
  - Toxic side effects
- Objective
  - Increase cure “rate”
  - Increase quality of life
- Quantitative comparison
  - Expected population
  - Likelihood of cure

# Small number of cells

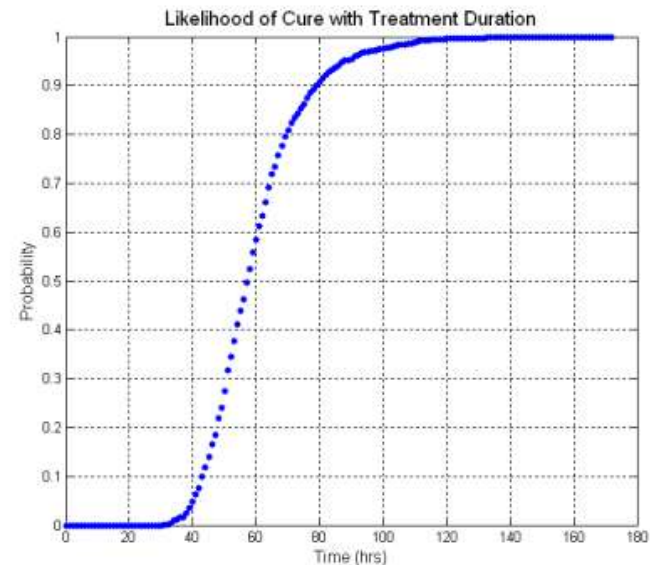
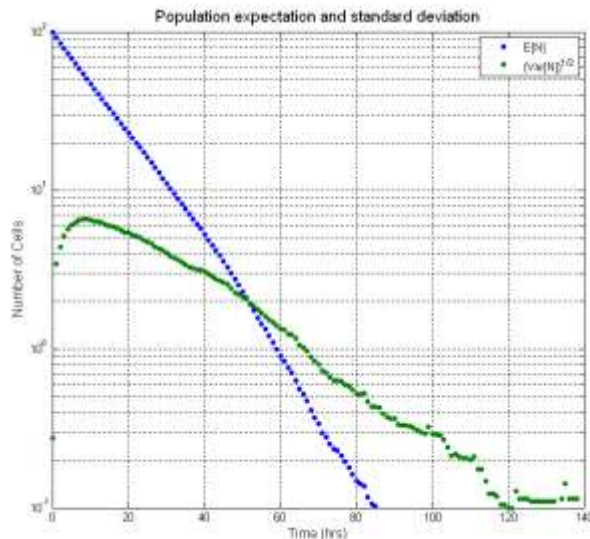
- Uncertainty in timing of events
  - Extrinsically stochastic
  - Average out if large number of cells
  - Greater variations if small number of cells
- Master probability density
  - Monte Carlo simulations
  - Cell number probability distribution
- Seminal cancerous cells
- Nearing “cure”
- **Dependent upon transition rate functions**

# Cell number probability distribution & likelihood of cure

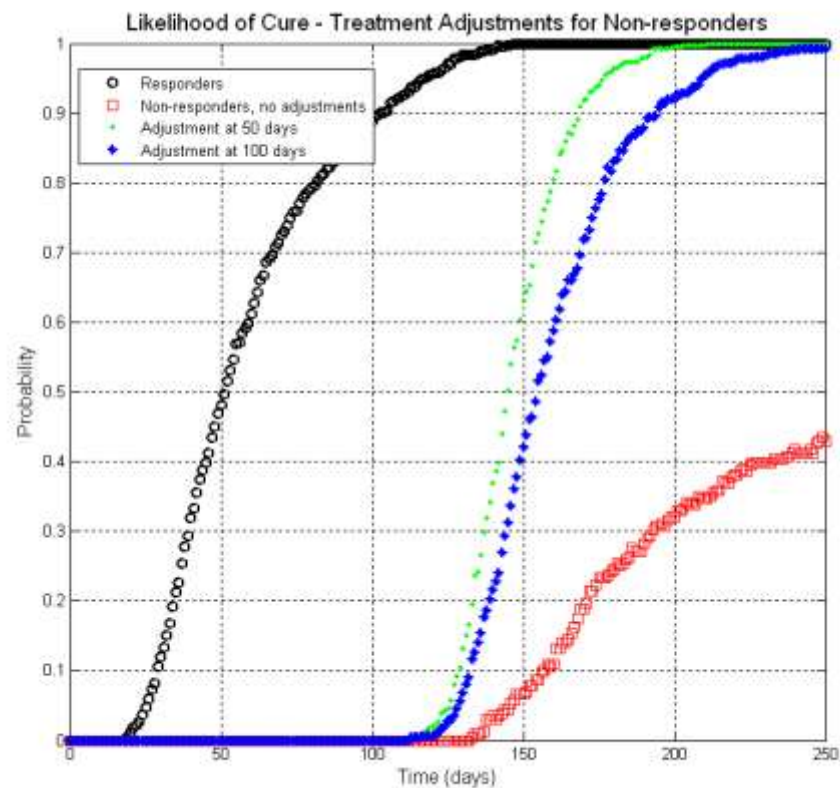
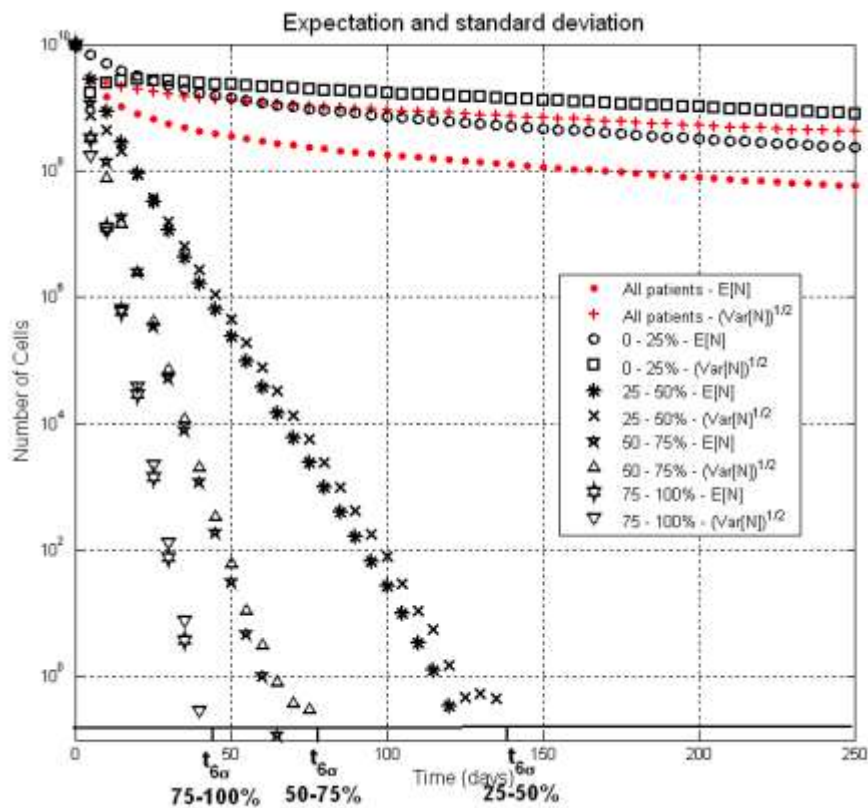


# Approximate treatment necessary to nearly ensure cure

- Small expected population
  - Small population mean
- Be certain that the population is small
  - Small standard deviation in number of cells



# Patient Variability



# Quiescence – importance of structure of transition rates

