
THE SEDIFLOC PROJECT

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*RAASF Workshop, September 27, 2004
Ghent, Belgium*

Why sedifloc ?

- Settling is often still a bottleneck in WWTP
- Improve knowledge of AS flocculation
 - mechanism
 - environmental influences
- Model final clarifier
- Calibrate 1D-model

SBR (80L) *sludge breeding*

-STABILITY-

Monitoring:

On-line:

- DO concentration
- pH

Off-line:

Effluent:

- COD
- N_{total}, NO₂-N, NO₃-N, NH₃
- PO₄-P

Sludge

- MLSS, MLVSS, SVI
- DGGE (microbial stability)

Influent

- 7 days stability

Flocculation
Knowledge

Floc Unit (5L) *flocculation experiments*

-DYNAMICS-

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- DO concentration
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- Conductivity
- Shear stress
- PSD
- Image analysis
-

Off-line:

- Settling velocity
- SVI

1D-Clarifier Modelling

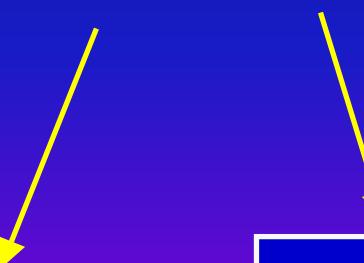
Virtual
Optimal
Experimental
Design
(Virtual OED)

2D-Clarifier Modelling
COMPUTATIONAL
FLUID DYNAMICS
(CFD)

Prediction of velocity, AS-conc.
pattern

Population
Balance Modelling
(PBM)

BIO-MATH



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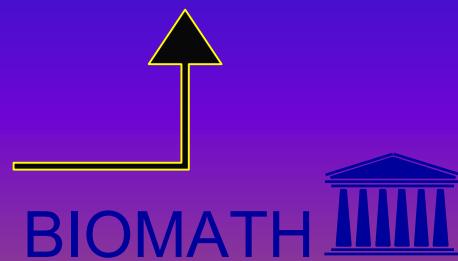
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Sequencing Batch Reactor



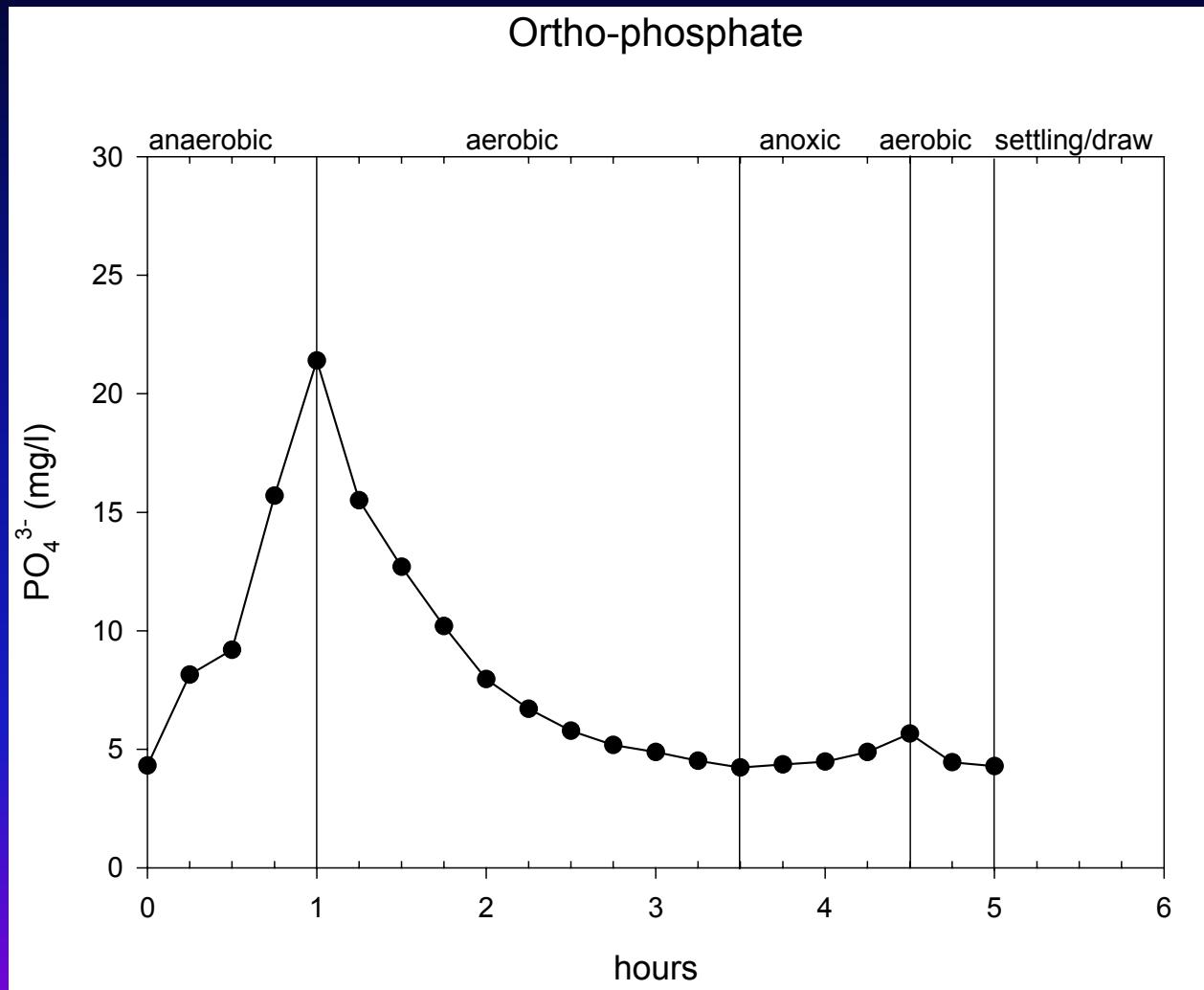
Sequencing Batch Reactor

- Aim → eliminate biological variability
- Characteristics
 - V= 80 l
 - SRT= 10 d
 - HRT = 12h
 - synthetic influent (COD/N/P = 100/13,7/2,14)
- Measurements
 - DO, pH, ORP, conductivity , weight (on-line)
 - COD, CODsol, Total-N, NH_4 , NO_3 , NO_2 , PO_4 (off-line)
 - MLSS (2-3 g/l), SVI (80-120 ml/g)
 - DGGE (microbial community)

SBR (2)

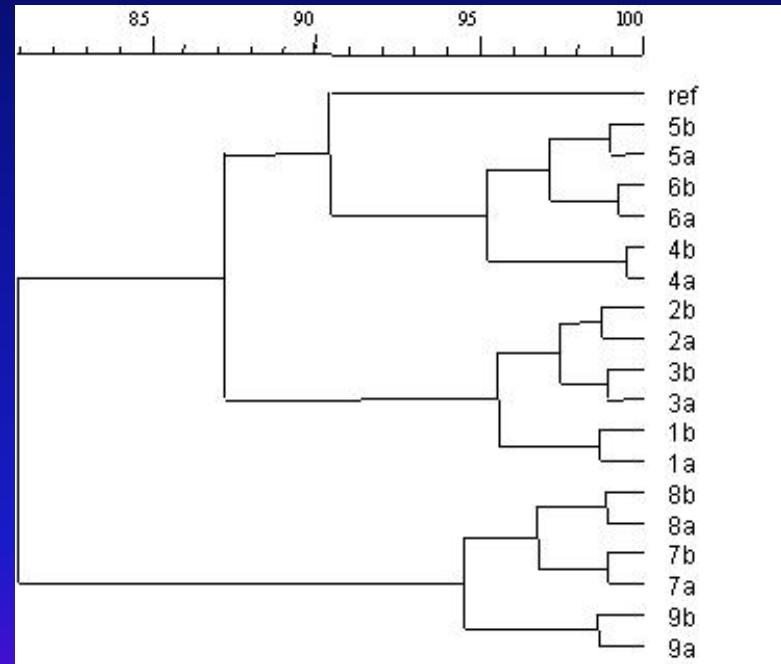
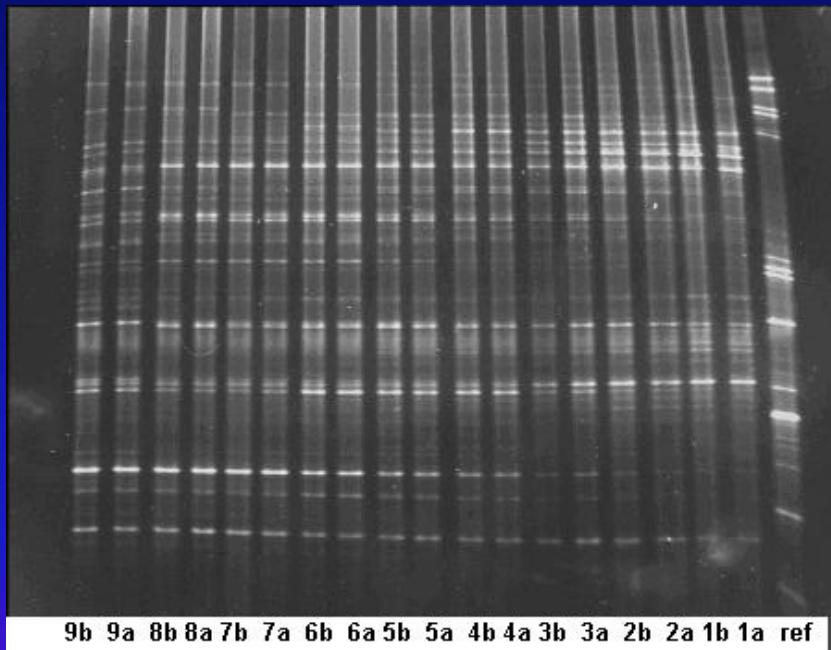
- Performance
 - 95 % COD-removal
 - 65 % N-removal
 - complete nitrification / incomplete denitrification
 - 65 % PO₄-removal

SBR (3)



SBR (4)

- DGGE analysis of 16S rRNA genes
examine temporal differences
within the activated sludge bacterial community



DGGE profile for replicate samples and cluster analysis

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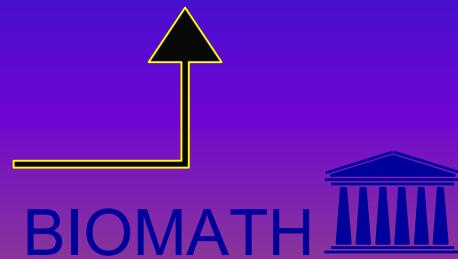
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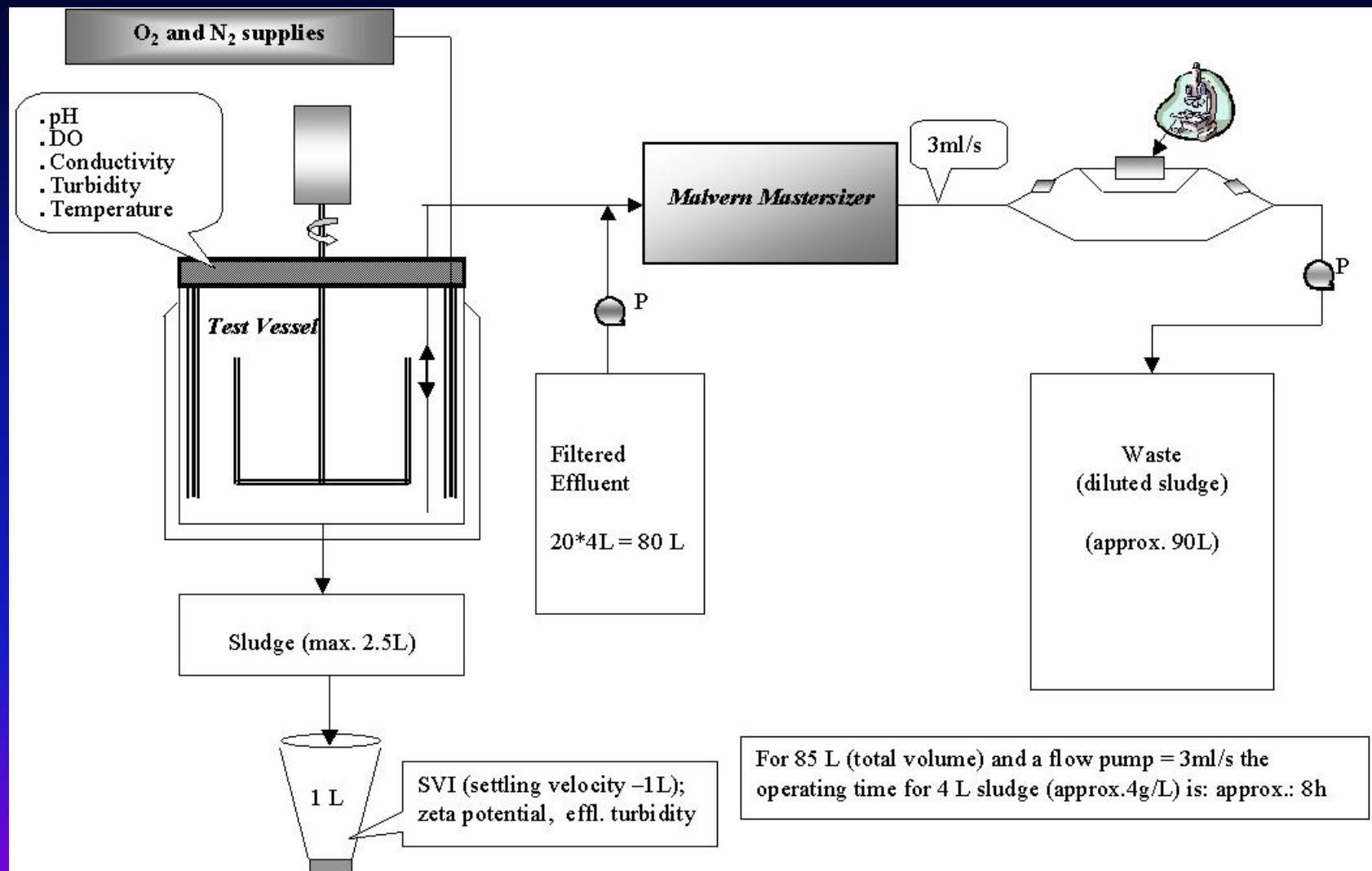
BIO-MATH



Floc Unit



Floc Unit



Floc Unit

- Features

- Mobile, T-controlled, bubbleless aeration
- Measurements:
 - DO, pH, conductivity, torque (on-line)
 - PSD (Malvern, Galai, image analysis) (on-line)
 - SS, SVI, Vs, zeta-potential (after flocculation experiment)
- Degrees of freedom
 - DO (O_2/N_2 -mixture)
 - Torque (viscometer)
 - Temperature
 - SS
 - Ionic strength (ion addition)
- Statistically based experimental design (full factorial)

Floc Unit

- Typical experiment
 - standardised starting condition
 - reflocculation in controlled environment
 - apply different environments
 - use orthogonal experimental design
 - flocculation = $f(?)$
 - implement in PBM

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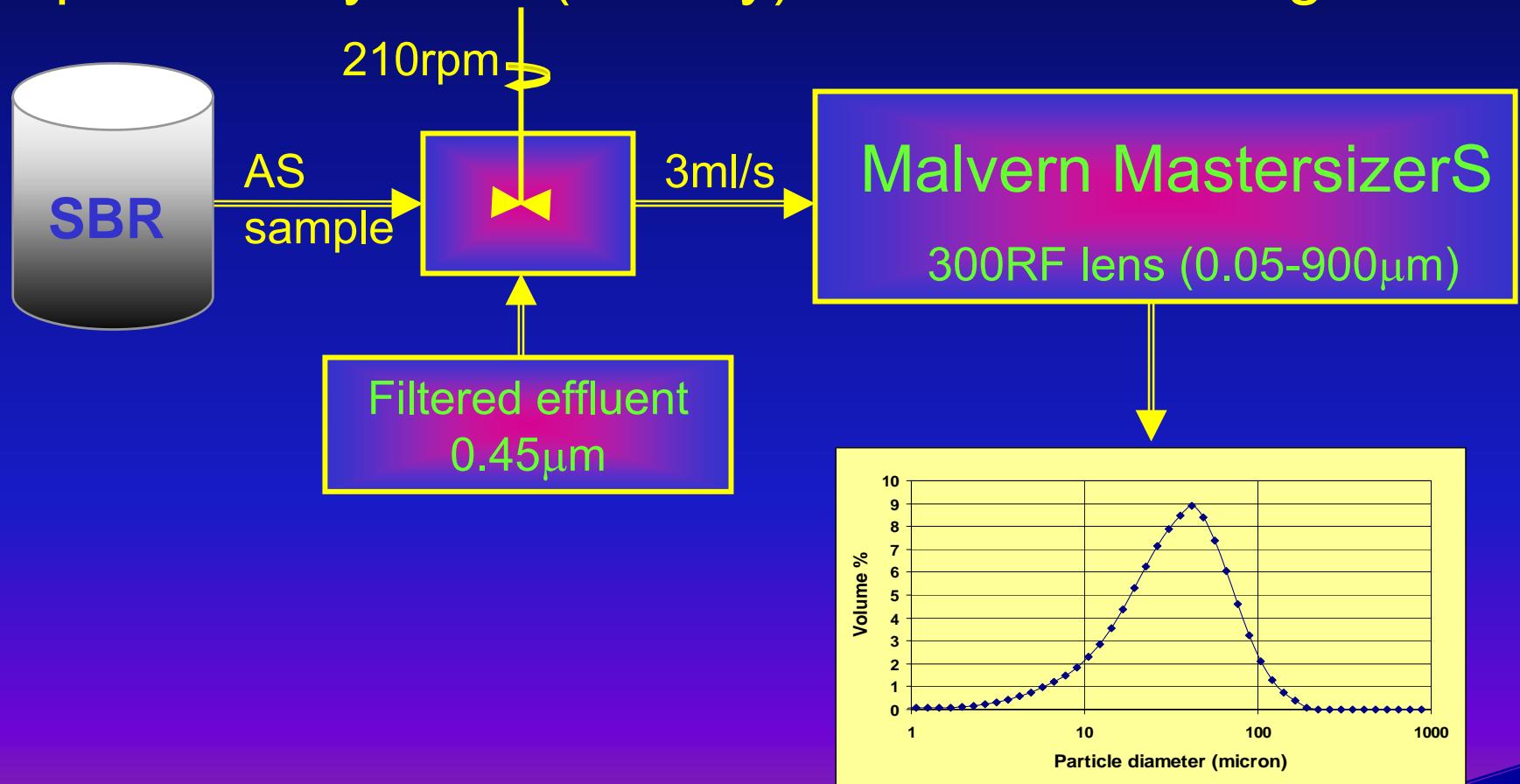
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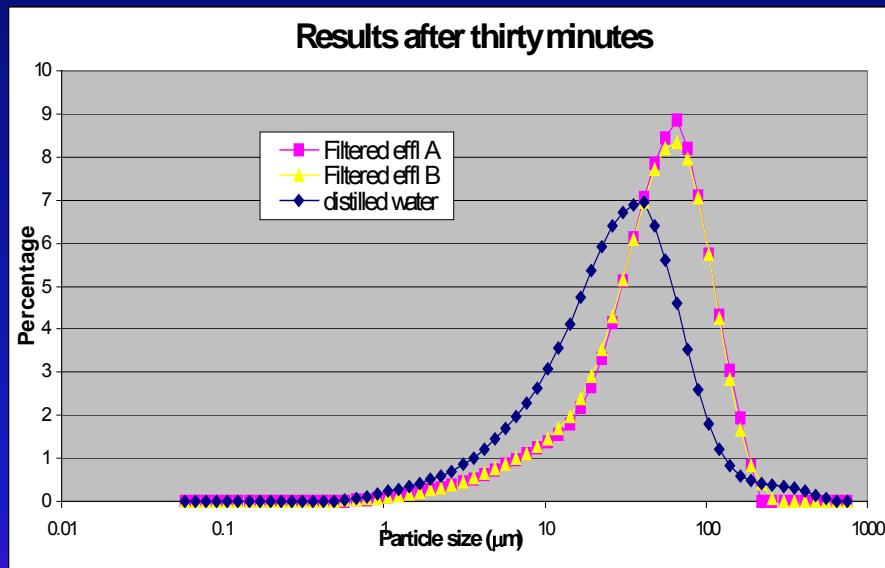
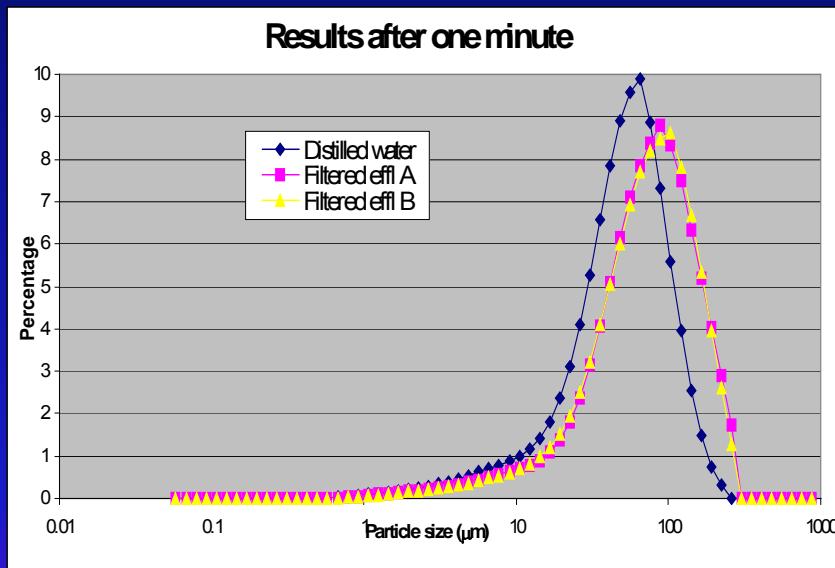
Flocculation knowledge

- On-line determination of PSD
- preliminary tests (weekly) – SBR Monitoring



Malvern PSD analysis

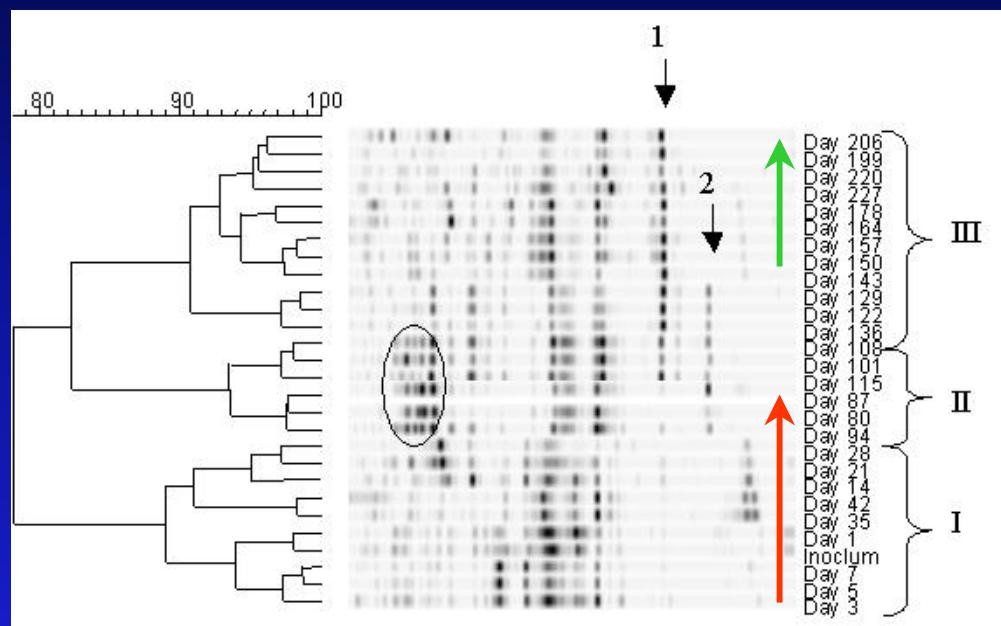
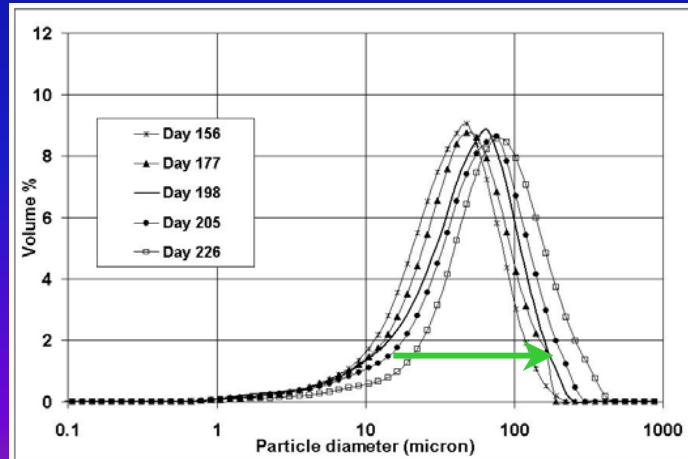
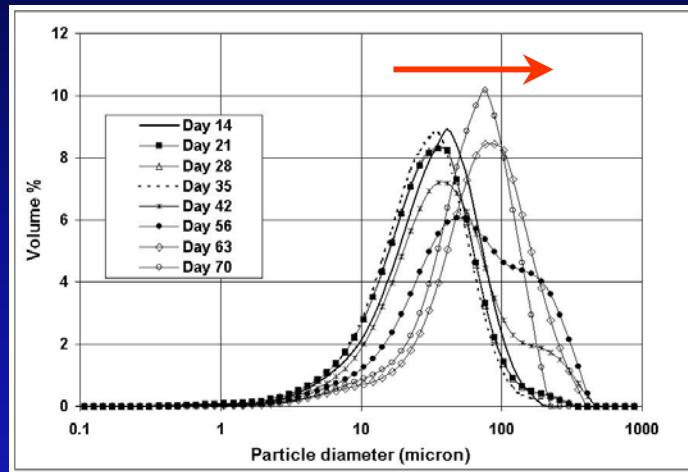
- Dilution Influences
 - Distilled water
 - Filter effluent ($0,45\mu\text{m}$ and $0.50\ \mu\text{m}$)



Diluting the sludge with distilled water caused an increase in the negative surface charge density (Mikkelsen et al. 1996)

Malvern PSD analysis

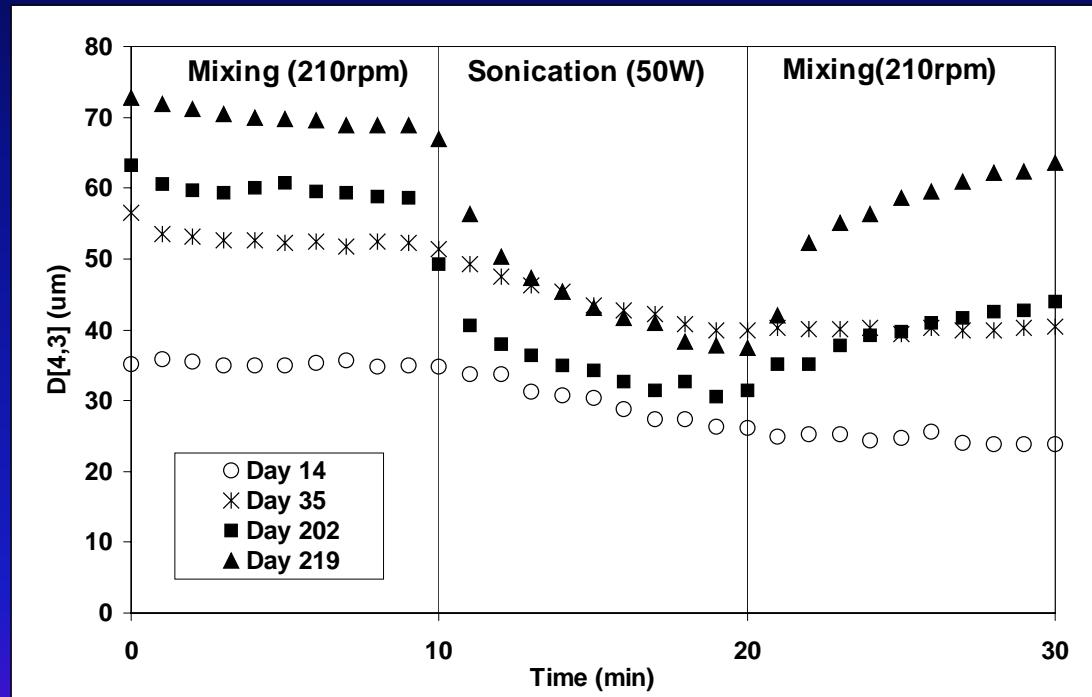
- Correlation with DGGE results



- I. Floc forming bacteria
- II. Floc forming and filamentous bacteria
- III. Filamentous bacteria

Malvern PSD analysis

- Evolution of D[4,3]



$$D[4,3] = \frac{\sum_{i=1}^n \Delta F_N(x)_i x_i^4}{\sum_{i=1}^n \Delta F_N(x)_i x_i^3}$$

- Non filamentous flocs (days 14 and 35)
- Filamentous flocs (days 202 and 219)

Malvern PSD analysis

- Important remarks
 - drawbacks of Malvern
 - assumption of sphericity
 - dilution is needed
 - Fraunhofer vs Mie theory
(refractive indices unknown)
 - combination with on-line image analysis
 - check for differences between PSDs
 - you see what is present
 - on-line image analysis is also possible

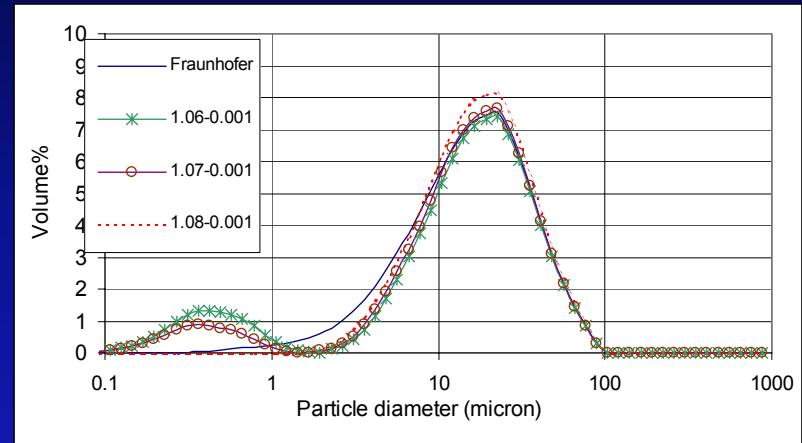


Image analysis (LabView)



Image analysis (LabView)

Press 'START' button to start image analysis

START

Configuration

Background storage path
Image storage path
Results storage path

CONTINUE

Background Threshold

CONTINUE

Image Acquiring

Press 'START' to start image-sampling

Particle filter:

Fill holes
no/yes

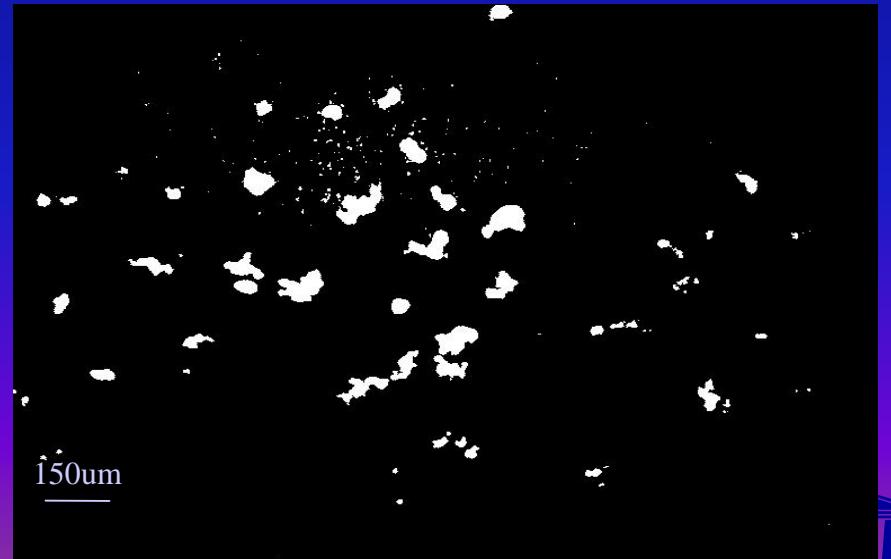
Parameter: Area (pixels)
Lower Value: 0.00
Upper Value: 0.00
Interval: 0.00
Include

Keep/Remove Particles
False: Keep

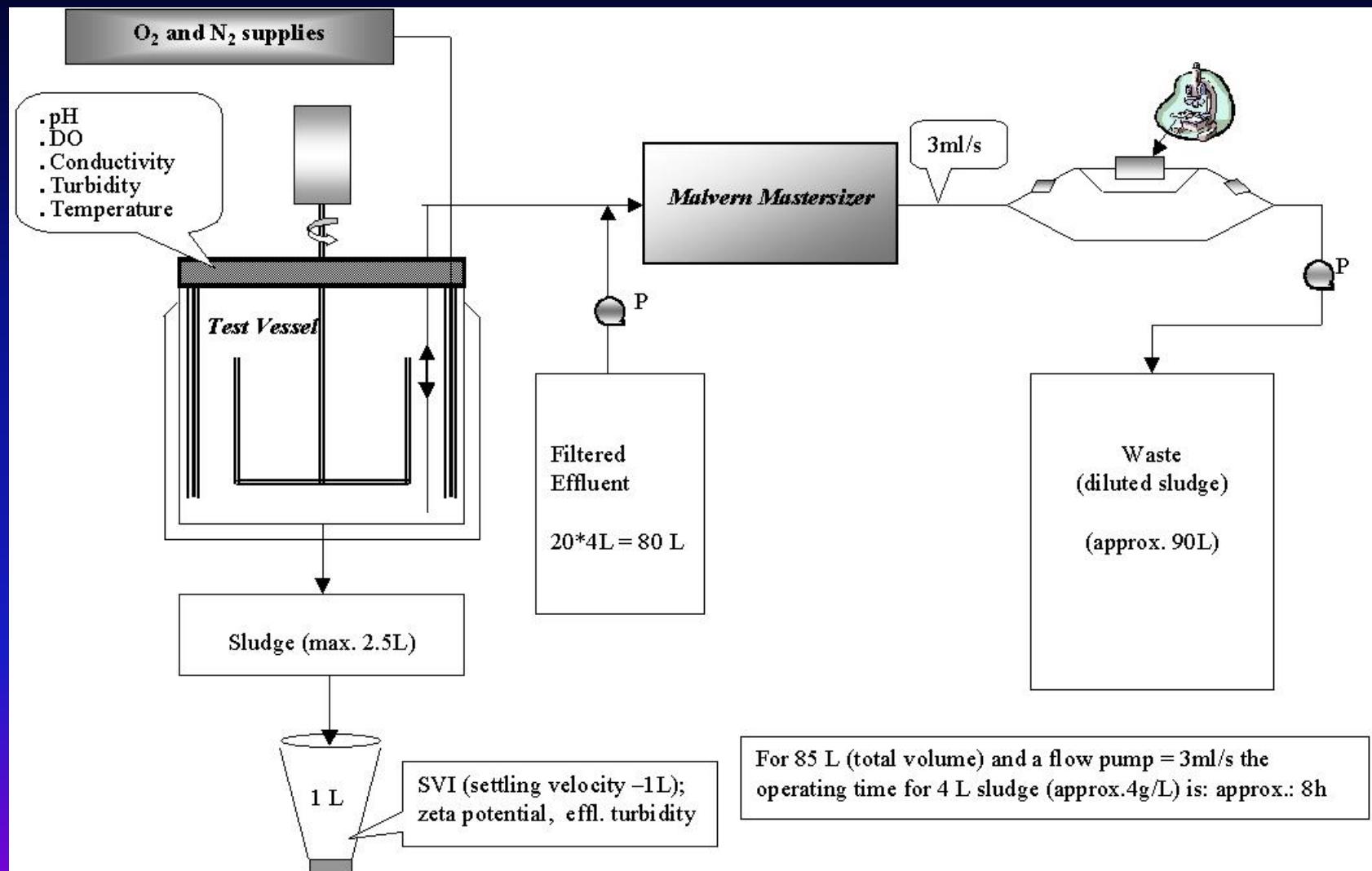
CONTINUE

Image Analysis

Processing Images ...



Floc Unit



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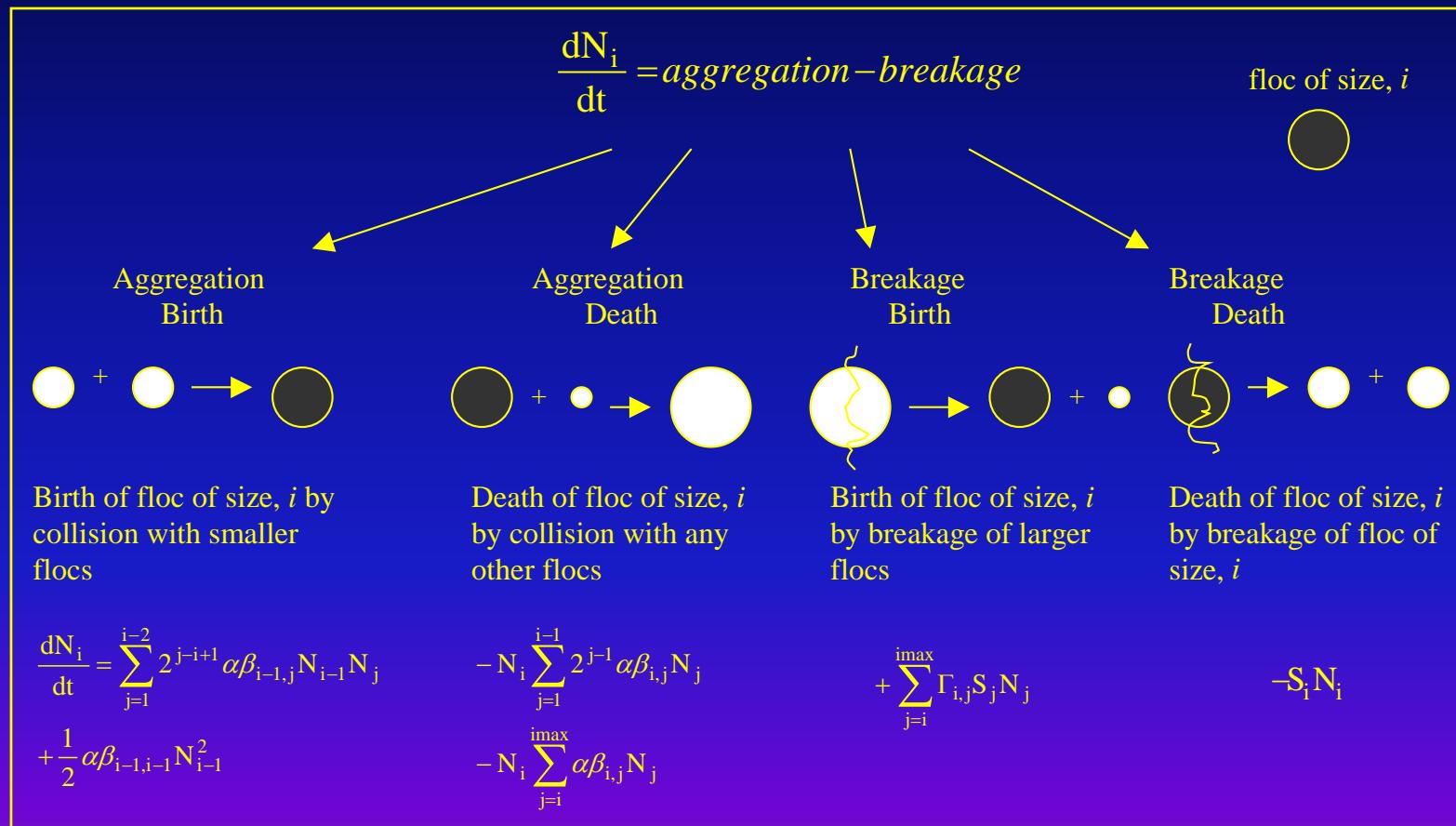
Flocculation
Knowledge

Population
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BIO-MATH

Population Balance Modelling

Hounslow (1988), Biggs (2000)



PBM: Model structure - Aggregation

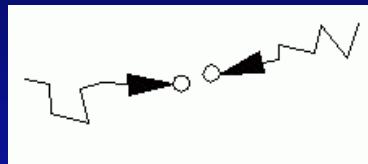
Aggregation kernels (summary)

Collision frequency, $\beta(i,j)$

- Brownian motion

$$\beta(i,j) = f(T, G, A_H, d(i), d(j), \eta)$$

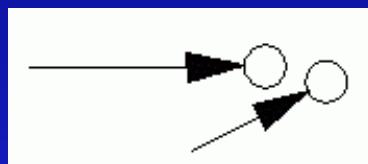
Smoluchowski (1917)



- Shear flocculation

$$\beta(i,j) = f(G, d(i), d(j))$$

Camp & Stein (1943)

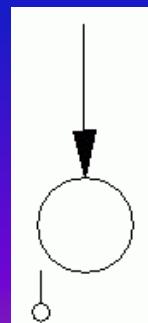


$$\beta(i,j) = f(G, d(i), d(j), D_f)$$

Lee et al. (1996)

- Differential sedimentation

Lee et al. (1996)



Collision efficiency, $\alpha(i,j)$

- $\alpha(i,j) = \text{constant}$

$$\alpha(i,j) = f(\eta, G, A_H, d(i), d(j))$$

Adler (1981)

$$\alpha(i,j) = f(\eta, G, A_H, d(i), d(j), e, \xi)$$

Kusters (1997)

Ducoste (2002)

G – shear rate

e - porosity

η – viscosity

ξ - permeability

A_H – Hamaker

T - temperature

d – particle diameter

D_f – fractal dimension

PBM: Model structure - Breakage

Breakage kernels (summary)

Breakage frequency, $S(i)$

- $S(i) = f(d(i))$
Spicer & Pratsinis (1996)
- $S(i) = f(d(i), \varepsilon, k, d(i) \dots)$
Kusters (1991)

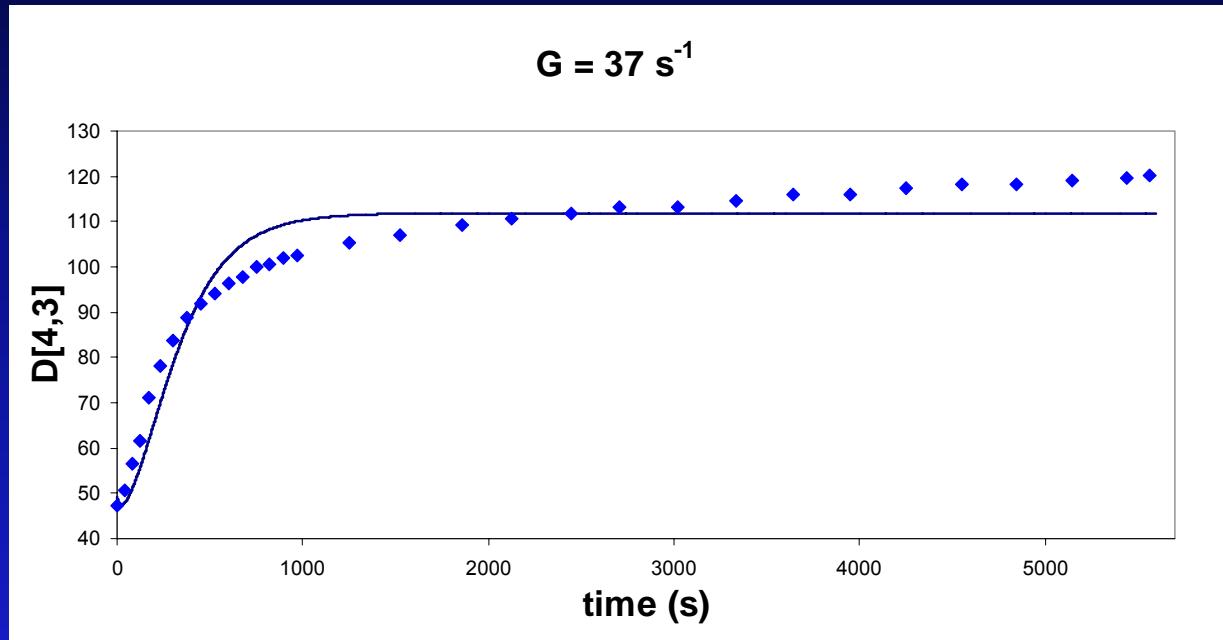
Fragment distribution function, $\Gamma(i,j)$

Binary breakage

Different combinations of aggregation and breakage kernels are implemented, simulated and compared to experimental results

PBM: Simulation study

- Model as used by Biggs (2000)
- model fitted
on $D[4,3]$

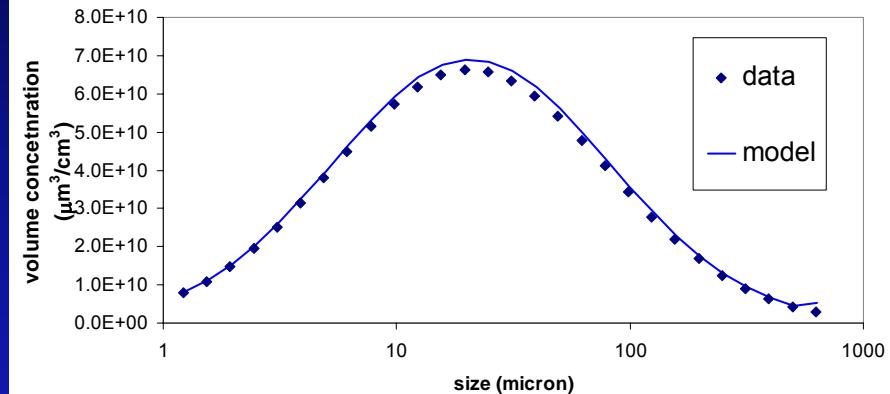


- too fast / lower steady-state mass mean
- model lacks flexibility

PBM: Simulation study

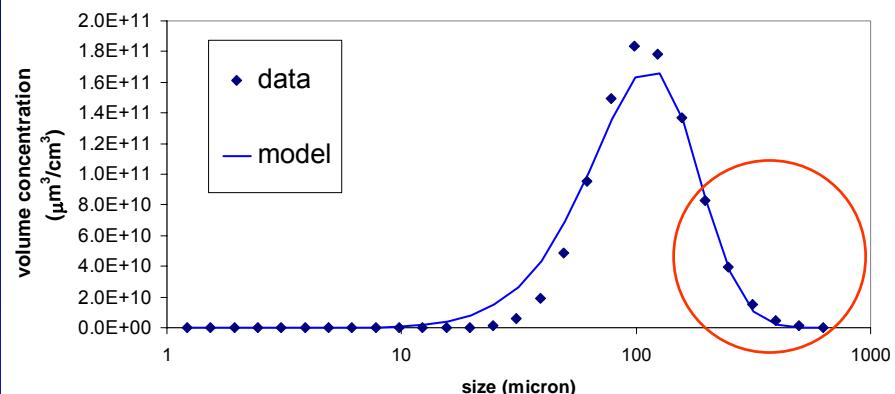
- inaccurate estimates of PSDs

volume distribution time 0



Volume distribution
at time zero

volume distribution end



Number distribution
at end of experiment

- D[4,3] volume based



fit on PSDs

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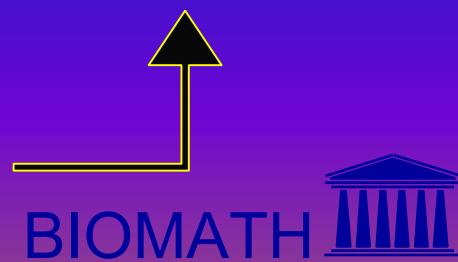
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2D-clarifier model

- **Aim**

- Modeling of sludge transport in clarifier, including PBM
- Computational Fluid Dynamics in Fluent software
- 2-dimensional
- Validation of CFD-model by
 - tracer test
 - solids concentration
 - particle size distributions
 - velocities

2D-clarifier model (cont'd)

- Model equations for CFD:

Continuity equation:

$$\frac{\partial \rho}{\partial t} + \frac{\partial}{\partial x_i} (\rho u_i) = 0$$

Momentum equation:

$$\frac{\partial}{\partial t} (\rho u_i) + \frac{\partial}{\partial x_j} (\rho u_i u_j) = - \frac{\partial p}{\partial x_i} + \frac{\partial \tau_{ij}}{\partial x_j} + \rho g_i$$

$$\tau_{ij} = (\mu + \mu_t) \left(\frac{\partial u_i}{\partial x_j} + \frac{\partial u_j}{\partial x_i} \right) - \frac{2}{3} (\mu + \mu_t) \frac{\partial u_1}{\partial x_1} \delta_{ij}$$

Sludge transport equation:

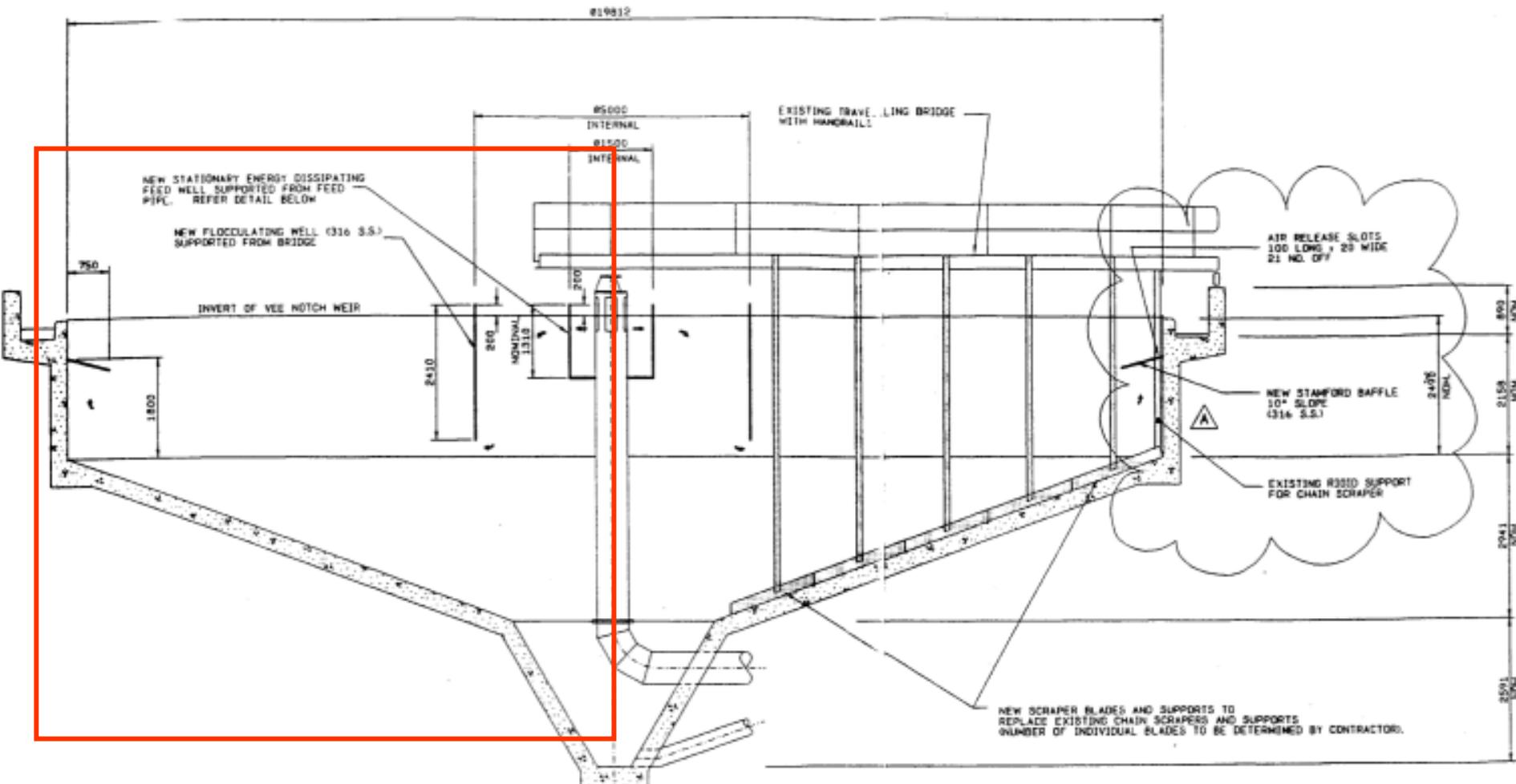
$$\frac{\partial}{\partial t} (\rho \phi) + \frac{\partial}{\partial x_i} \left[\rho \phi (u_i + v_i) - \Gamma \frac{\partial \phi}{\partial x_i} \right] = 0$$

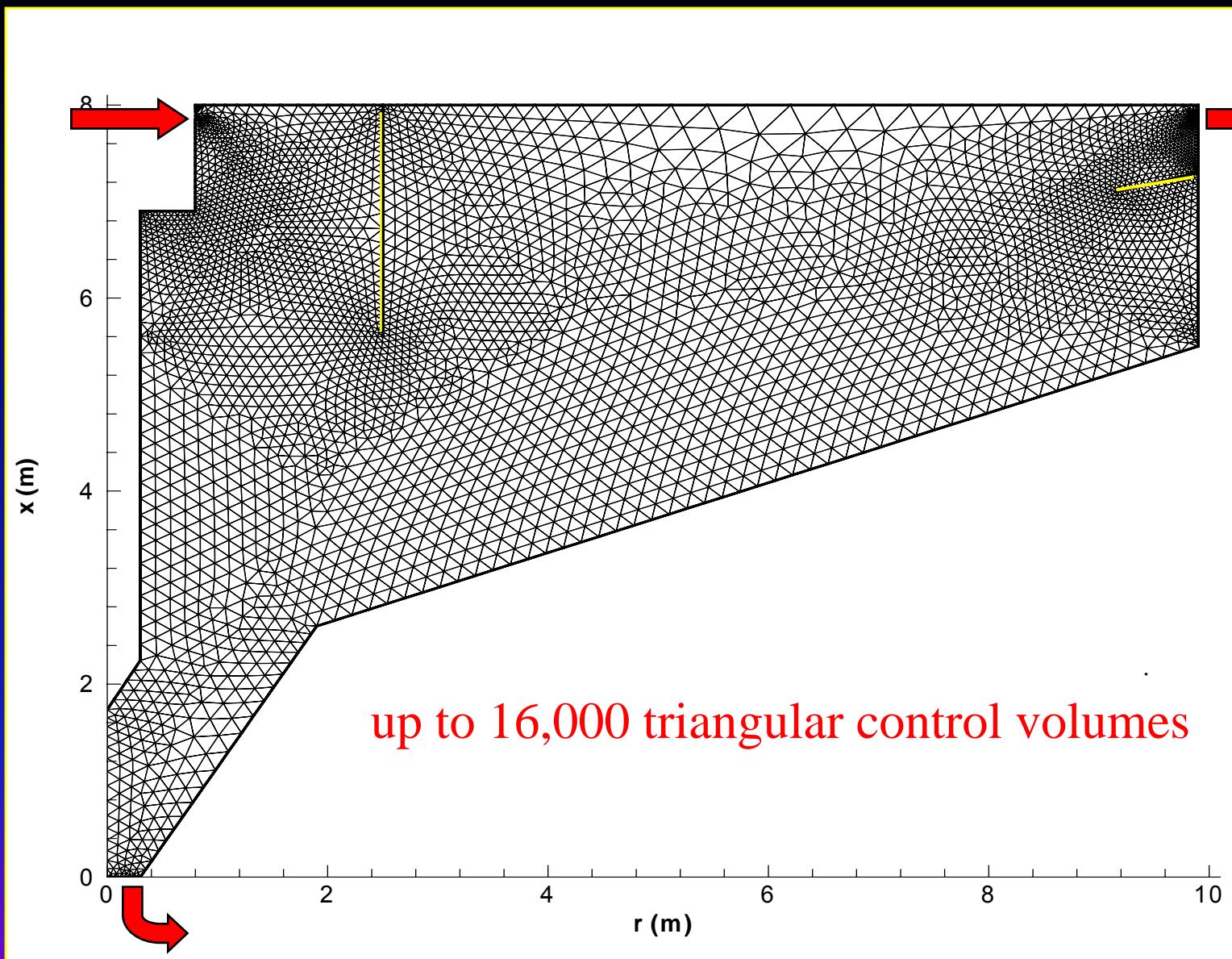
Density:

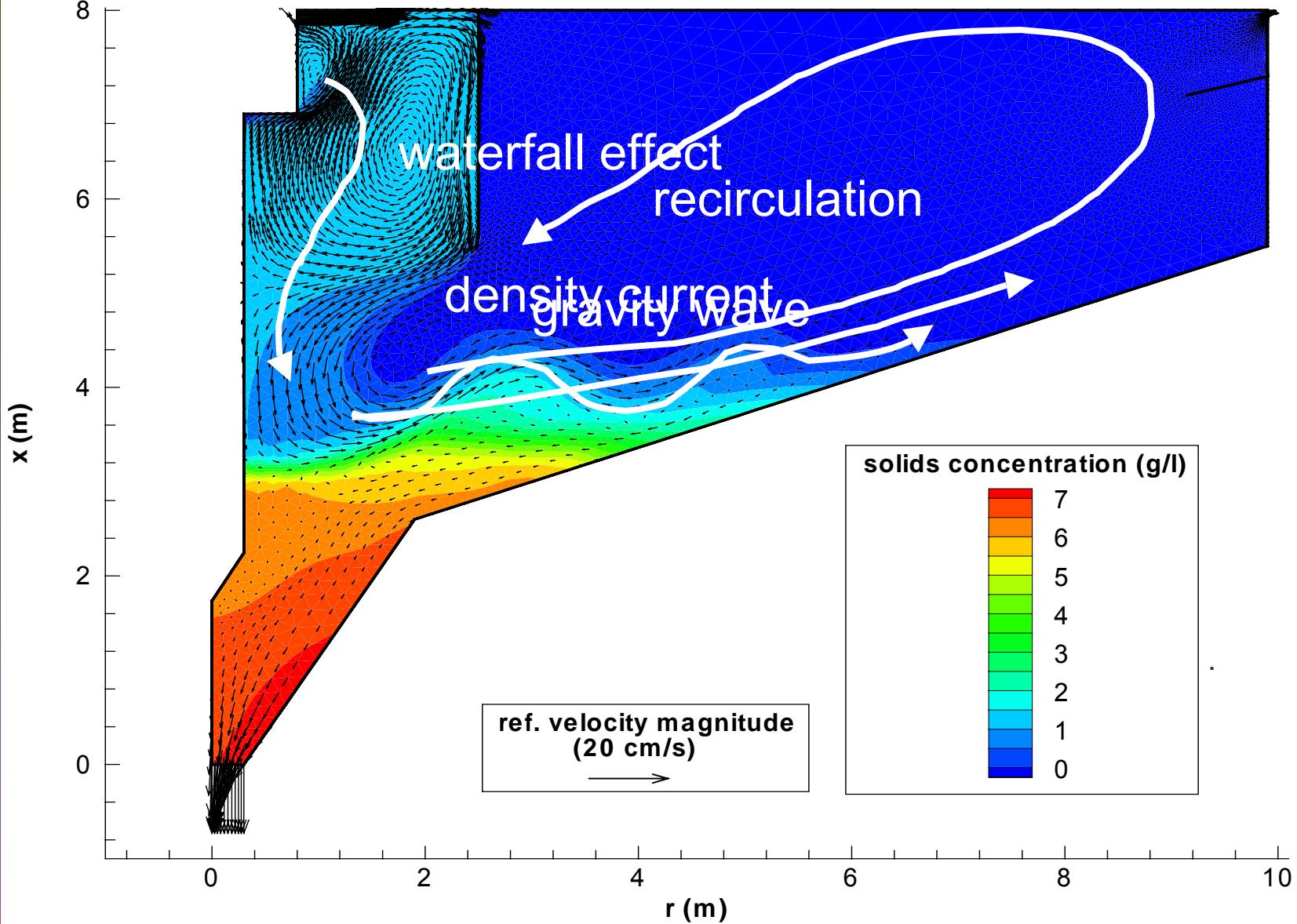
$$\rho = \frac{\rho_w}{1 - \phi \left(1 - \frac{\rho_w}{\rho_s} \right)}$$

2D-clarifier model (cont'd)

- Oxley Creek settling tank

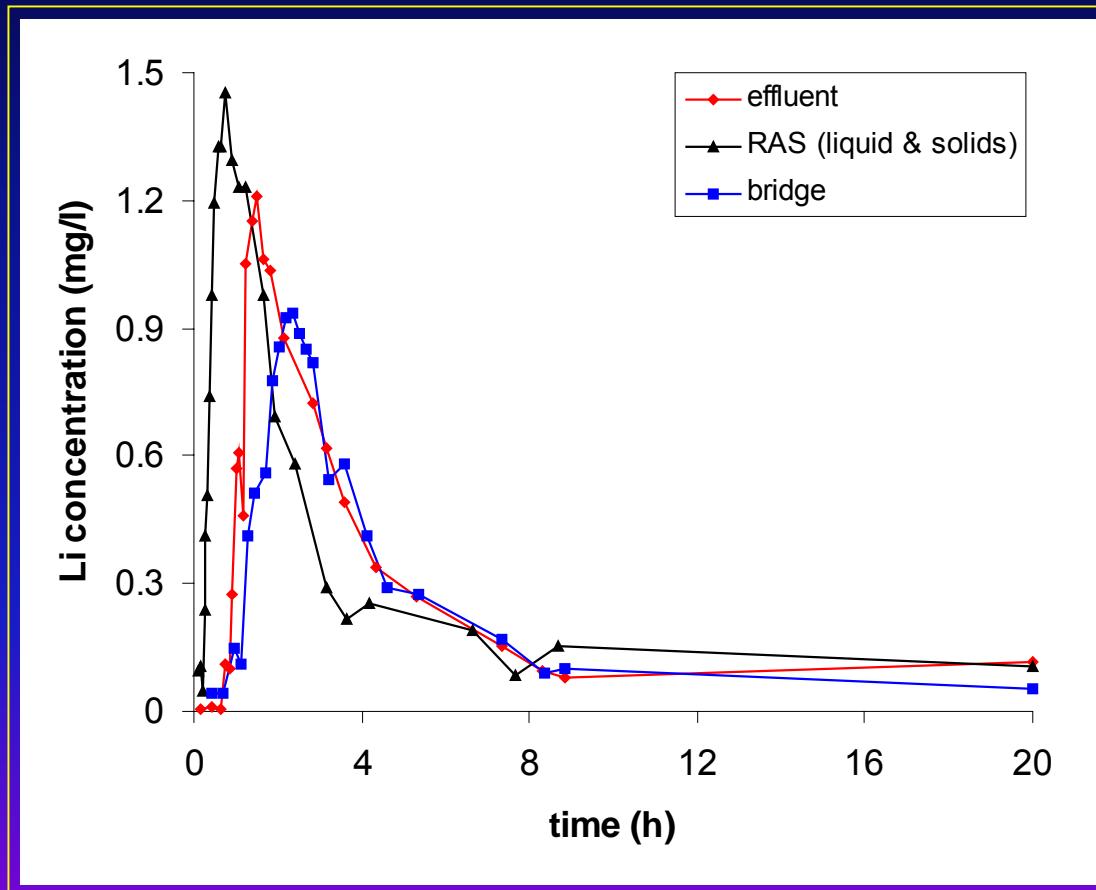




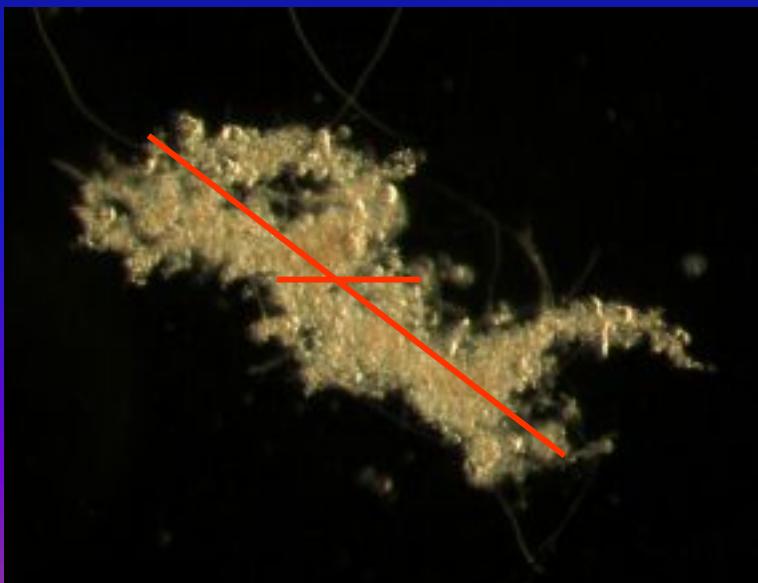
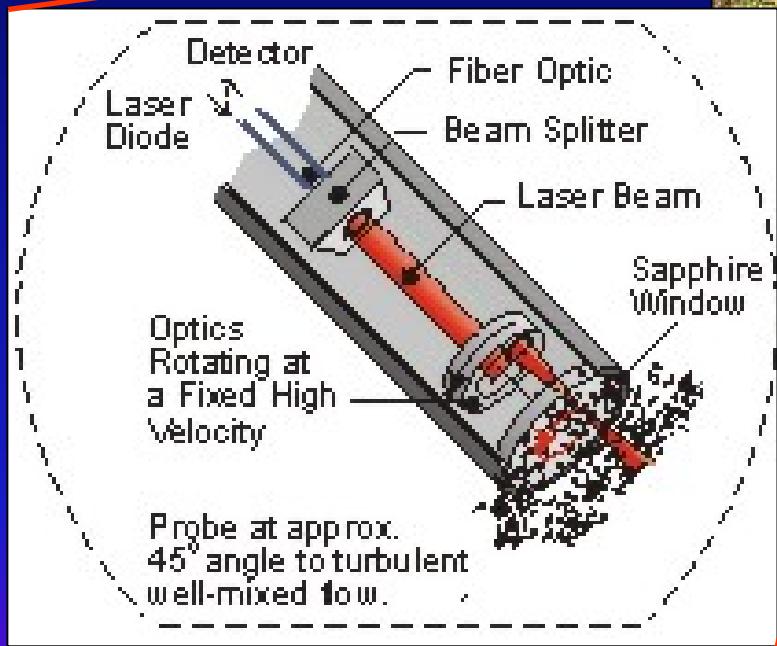


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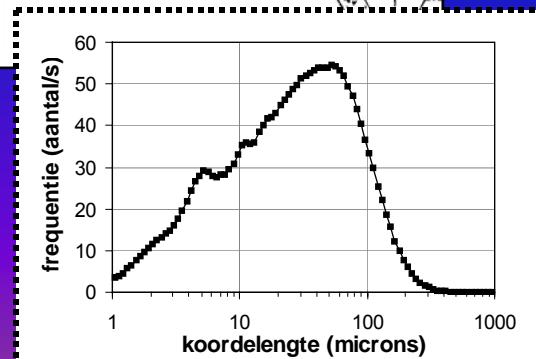
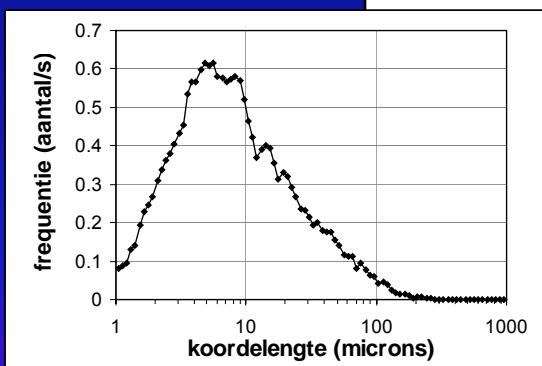
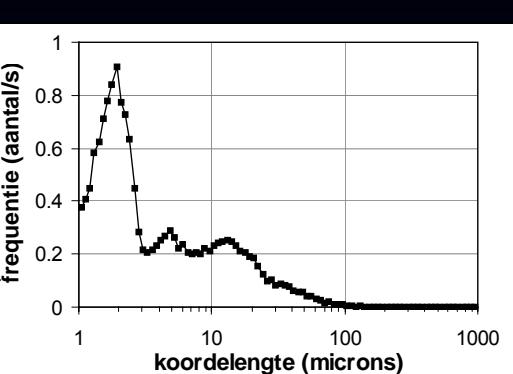
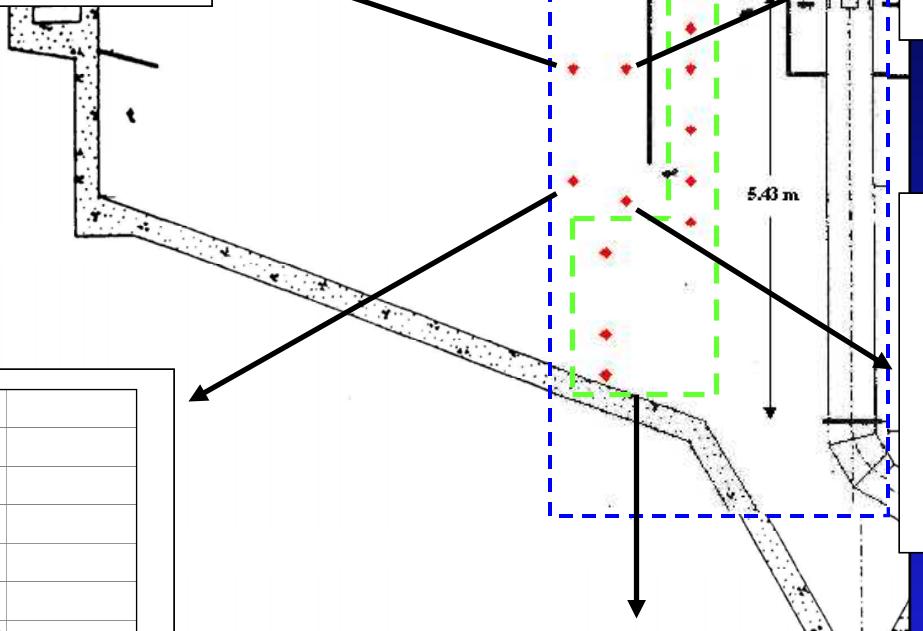
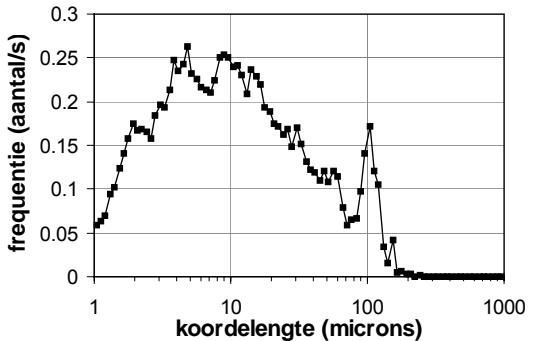
- Results of tracer test (LiCl)



In situ PSD (Lasentech)



PSD (Lasentech)



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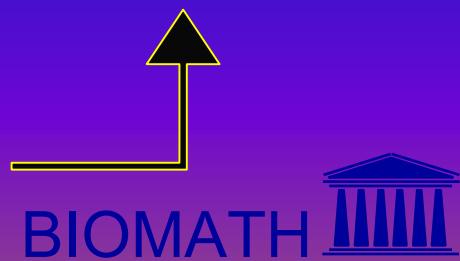
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1D-clarifier model with dispersion

- Continuity equation:

$$\frac{\partial C(z,t)}{\partial t} = -\frac{\partial(F(C(z,t),z,t)}{\partial z} + \frac{\partial}{\partial z}\left(D(z,t) * \frac{\partial C(z,t)}{\partial z}\right) + s(z,t)$$

- Dispersion for clarification and thickening zone:

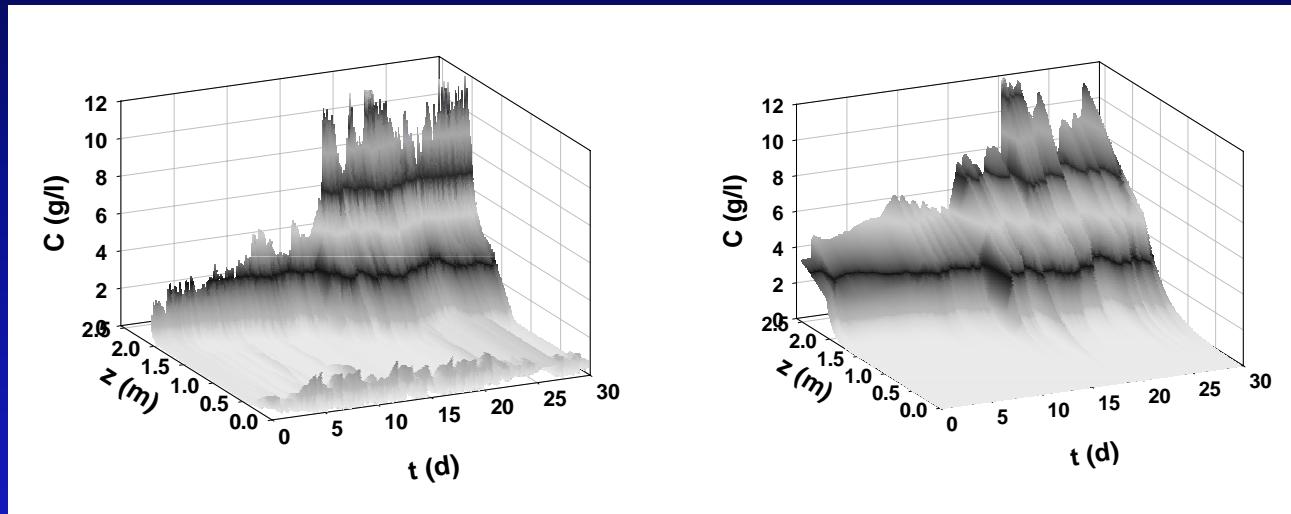
$$D_1(t) = D_{11} * e^{\alpha * \frac{Q_e(t)}{Q_f(t)}}$$

$$D_2(t) = D_{22} * e^{\beta * \frac{Q_u(t)}{Q_f(t)}}$$

- Variable location of feed layer to consider density currents
- Numerical integration
 - differencing of spatial derivatives of PDE (100 layers)
 - LSODA for numerical integration of (stiff) ODE system

1D-clarifier model (3)

- Measured (left) and predicted (right) sludge profile vs time



- Dynamics are described well
- Discrepancy at concentrations higher than 6 g/l, but good results for C_{recycle}

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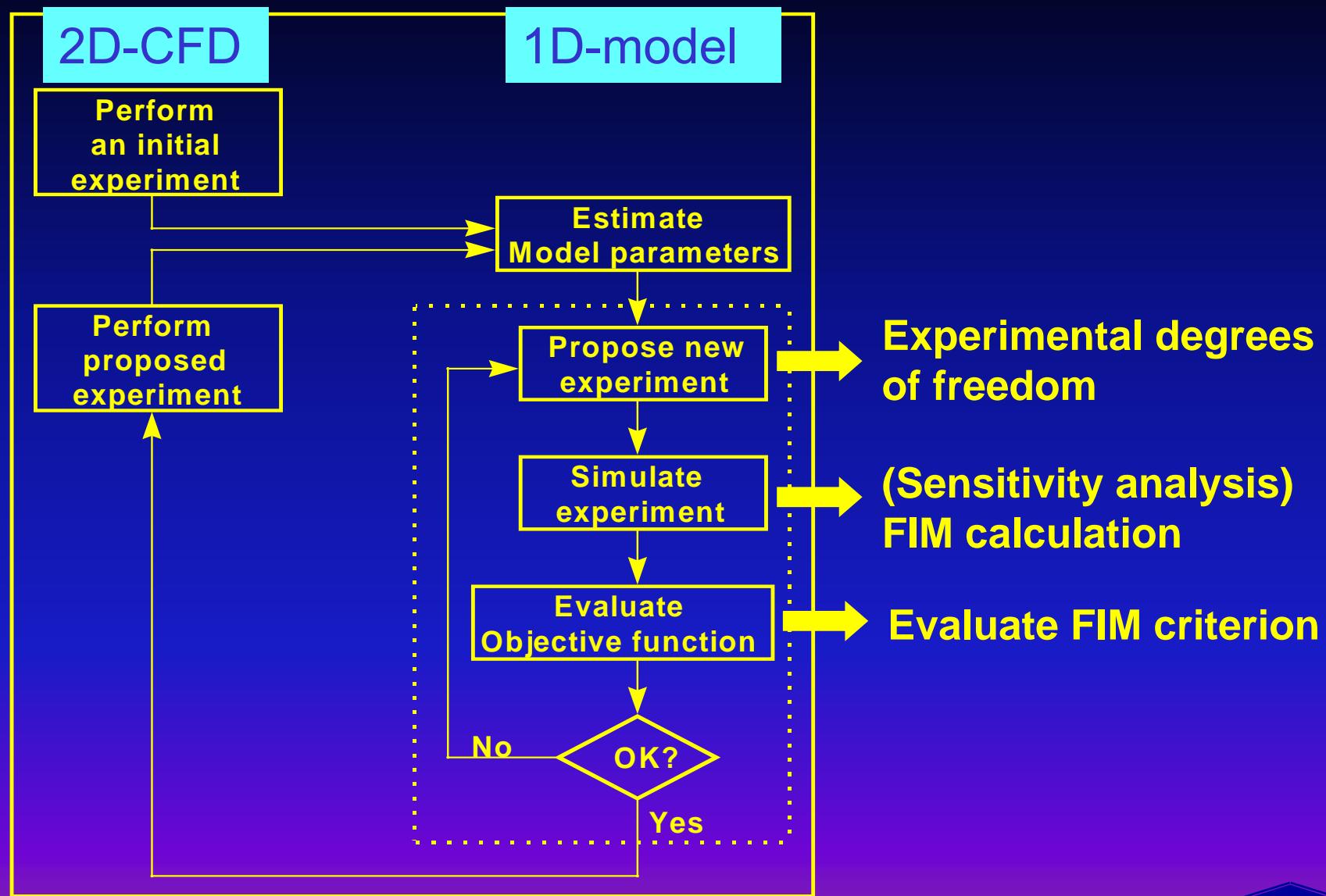
Population
Balance Modelling
(PBM)

BIO-MATH

Virtual Optimal Experimental Design

- Bridge between 1D and 2D clarifier models
=Knowledge transfer mechanism
- Propose simulation experiments (measure where you like !) to be performed with 2D CFD clarifier model for optimal calibration (low errors on parameters) of the 1D clarifier model
- Tool = WEST
- Implementations performed in WEST:
 - Sensitivity analysis
 - Fisher Information Matrix evaluation (information content)
 - Optimal Experimental Design criteria

Experimental design methodology



SBR (80L) *sludge breeding*

-STABILITY-

Monitoring:

On-line:

- DO concentration
- pH

Off-line:

Effluent:

- COD
- N_{total}, NO₂-N, NO₃-N, NH₃
- PO₄-P

Sludge

- MLSS, MLVSS, SVI
- DGGE (microbial stability)

Influent

- 7 days stability

Flocculation
Knowledge

Floc Unit (5L) *flocculation experiments*

-DYNAMICS-

Monitoring:

On-line:

- DO concentration
- pH
- Conductivity
- Shear stress
- PSD
- Image analysis
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Off-line:

- Settling velocity
- SVI

1D-Clarifier Modelling

Virtual
Optimal
Experimental
Design
(Virtual OED)

2D-Clarifier Modelling
COMPUTATIONAL
FLUID DYNAMICS
(CFD)

Prediction of velocity, AS-conc.
pattern

Population
Balance Modelling
(PBM)

BIO-MATH