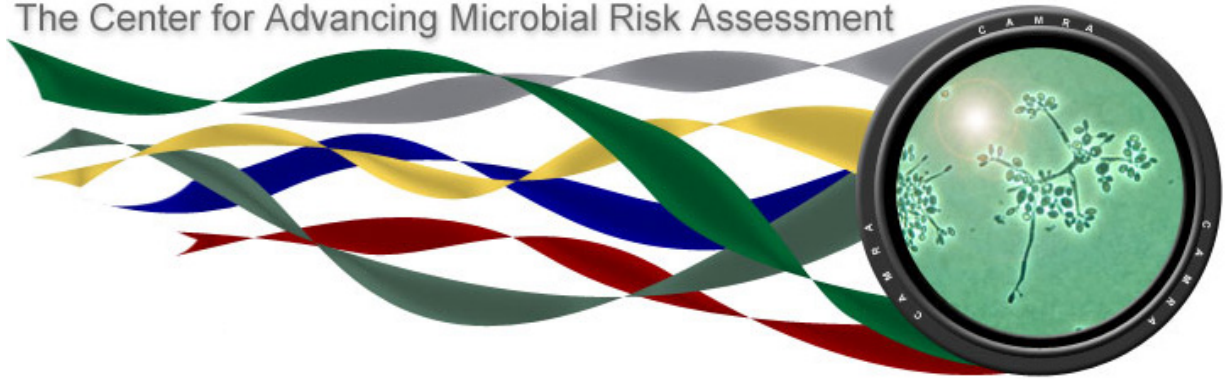
Improvement of Water Quality Modeling for Microbial Risk Assessment: Recent Advances

Chris Choi, Professor

CAMRA/The University of Arizona

CAMRA – National Homeland Security Center

The Center for Advancing Microbial Risk Assessment



Michigan State University

The University of Michigan

The University of California, Berkeley

Drexel University

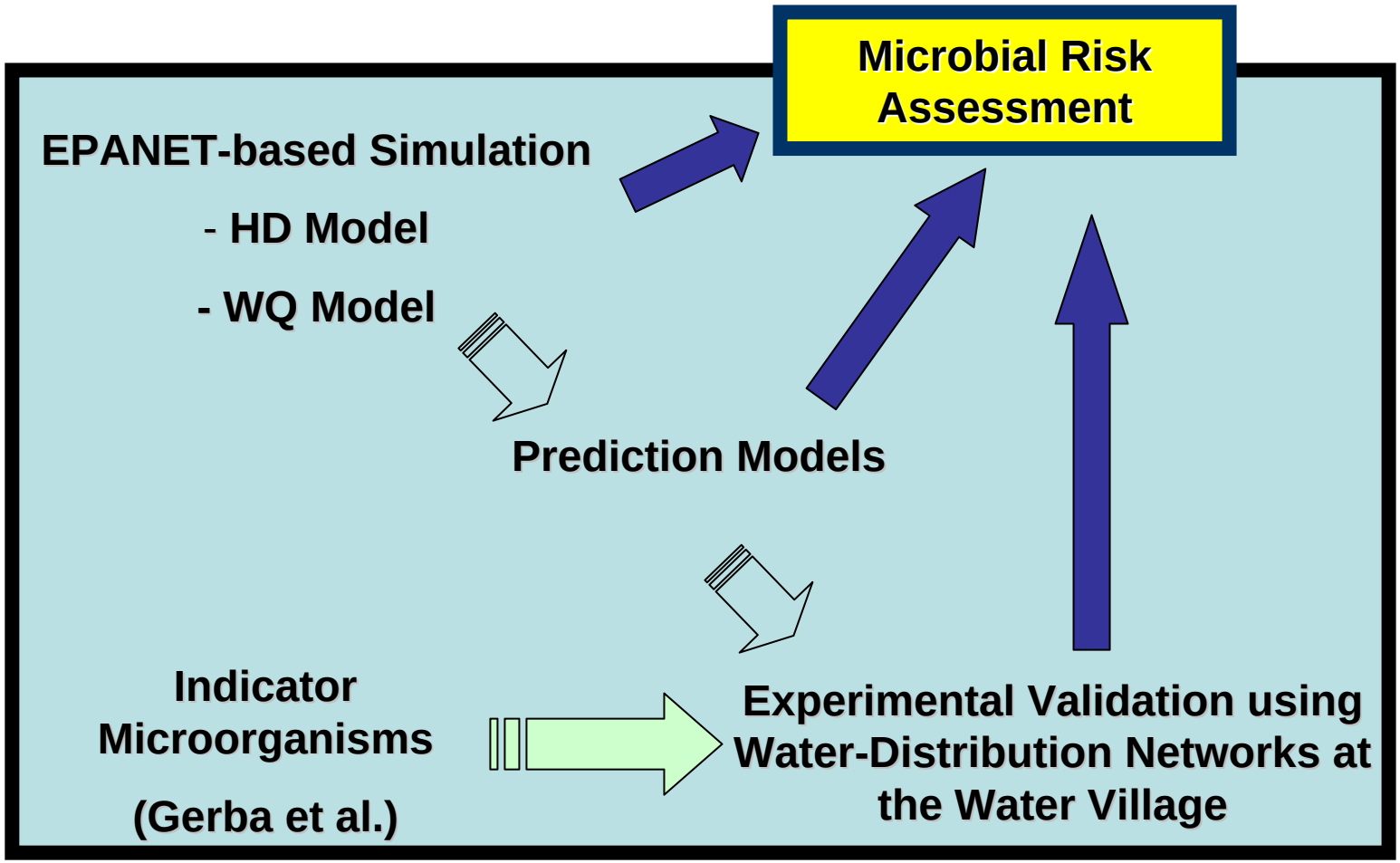
Northern Arizona University

Carnegie Mellon University

The University of Arizona

Main Research Focus of Choi's Group: Water Distribution Systems

Research Plan



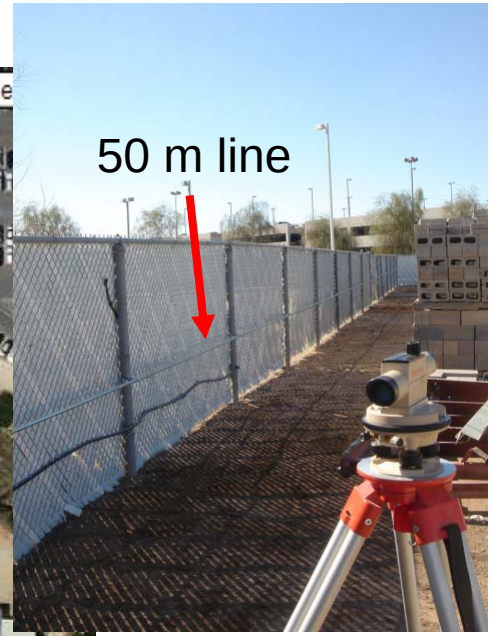
Point of Use & Microbiology Laboratory

Water Village

Water Distribution Systems Laboratory

Pilot Scale Experiments

Current Laboratories



Real-time Water Quality Monitoring Lab

Water Chemistry Lab

Water Treatment Laboratory

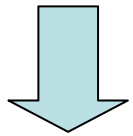
Pumpin Station

Trench

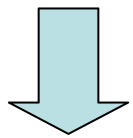
50 m line for dispersion experiments

Research Plan

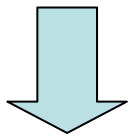
Step 1. Accurate Water Quality Modeling



Step 2. Accurate Prediction of Dispersion of Microorganisms

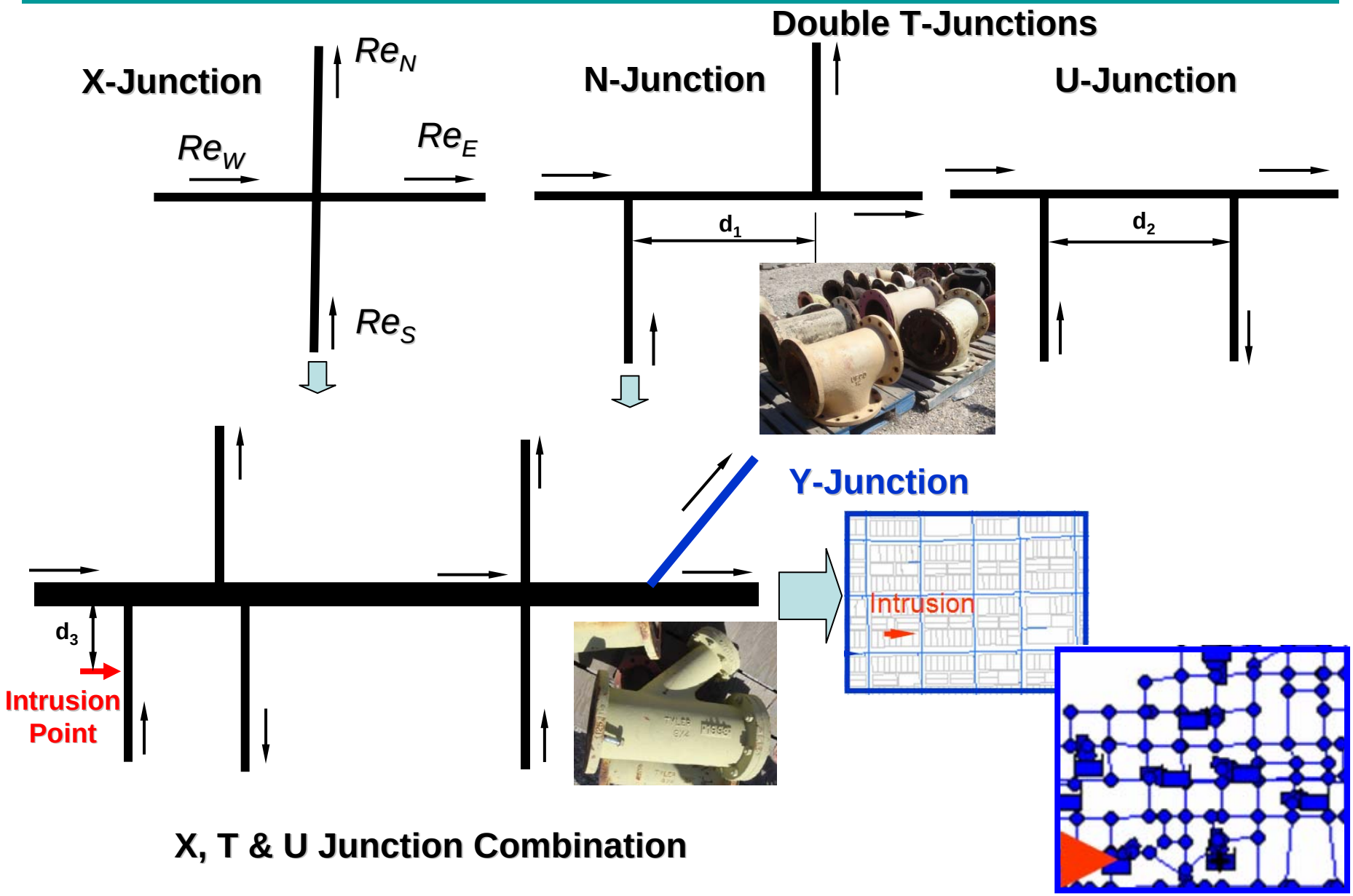


Step 3. Use of Prediction Tools such as ANNs



Step 4. Accurate Quantitative Microbial Risk Assessment

Building a Real-World Network Model



Two Significant Assumptions for EPANET Water Quality Model

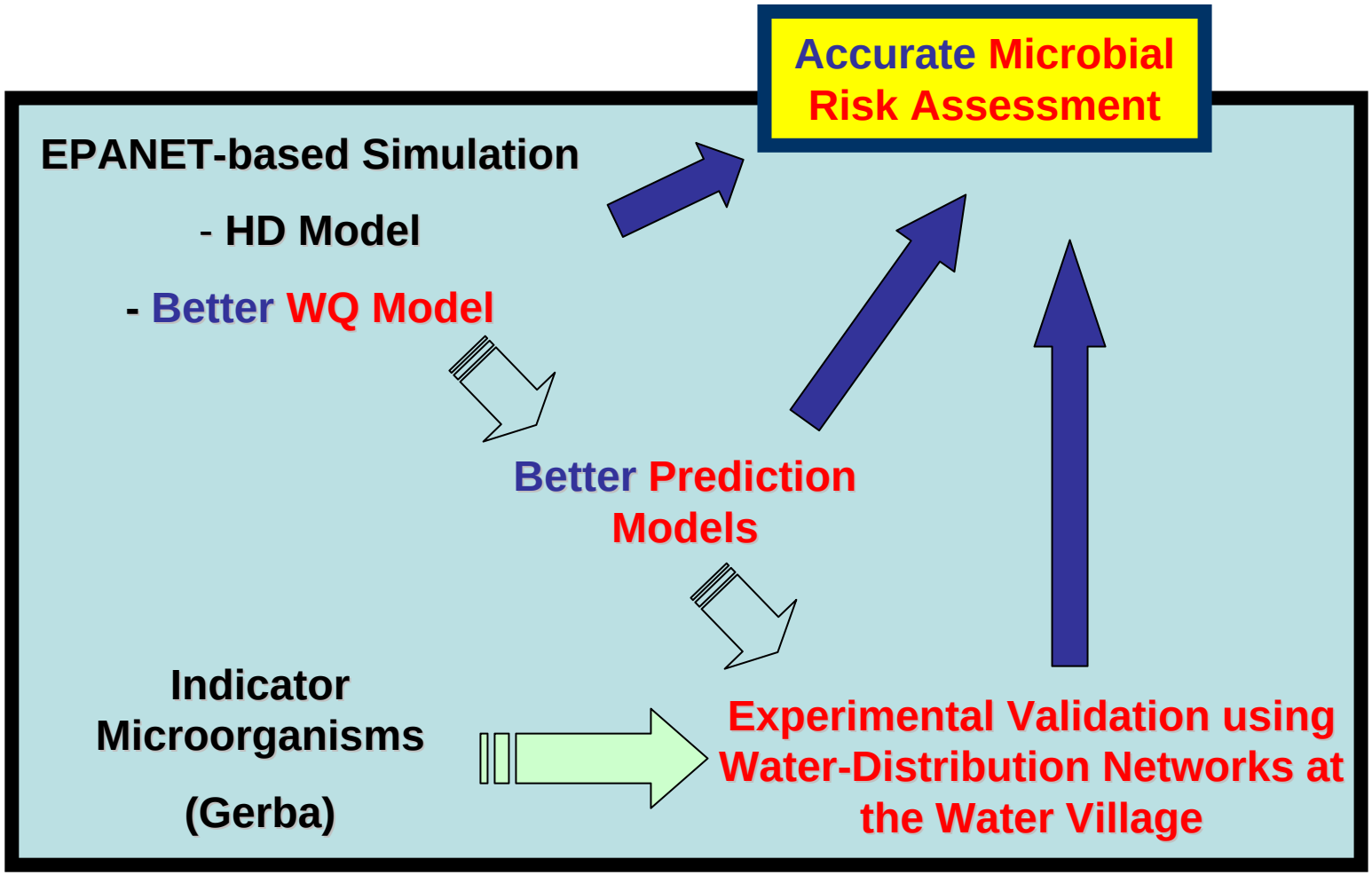
At present, prevailing network water quality models are based on two major simplifications.

First, longitudinal dispersion of the solute mass along the pipe axis is ignored, and "plug flow" is assumed to prevail.

Second, solute mixing is assumed to be "complete and instantaneous" at the pipe junctions.

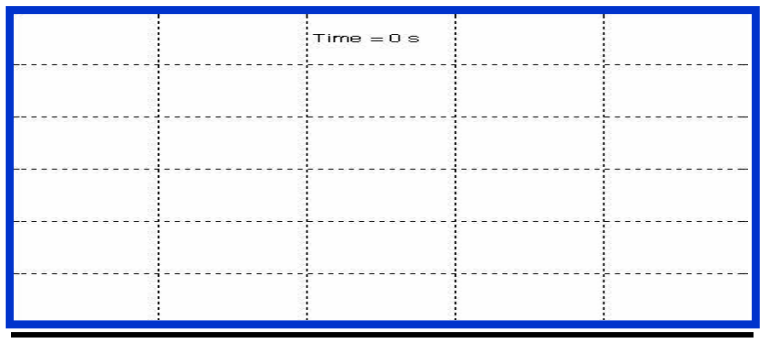
Recent investigations suggest that these assumptions may NOT be valid in real pipe networks.

Potential Improvement

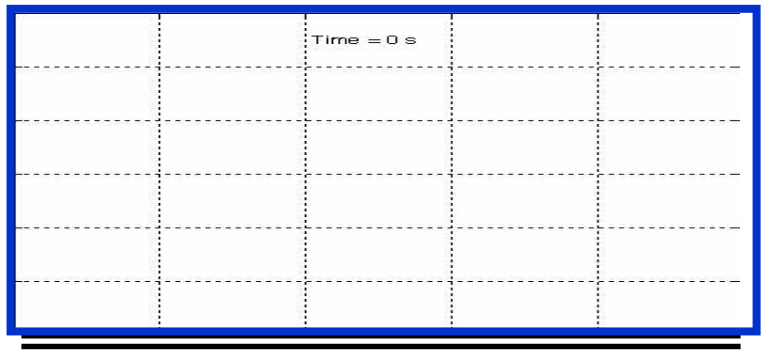


CFD Simulation for Pipe length = 6.35 m, D = ½ in.

U = 0.635 m/s Re = 8,000



U = 1.27 m/s Re = 16,000



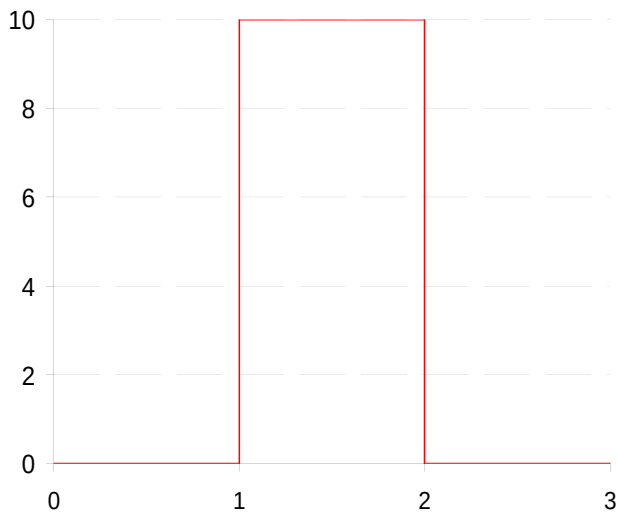
Re = 8,000 , Tmax = 6.5 s

Re = 16,000 , Tmax = 4.05 s

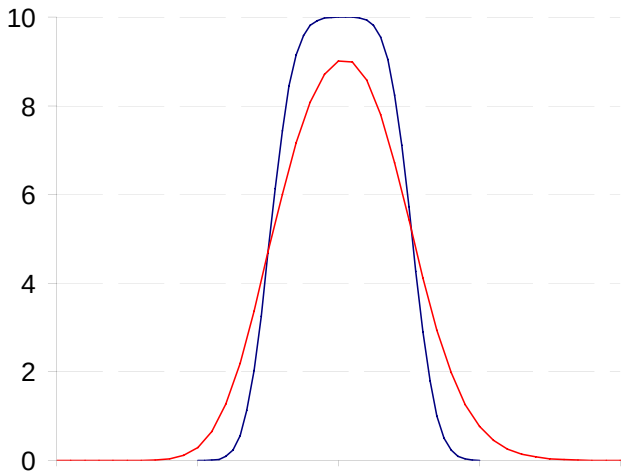
Re = 8,000 , Tmax = 11.6 s

Re = 16,000 , Tmax = 6.55 s

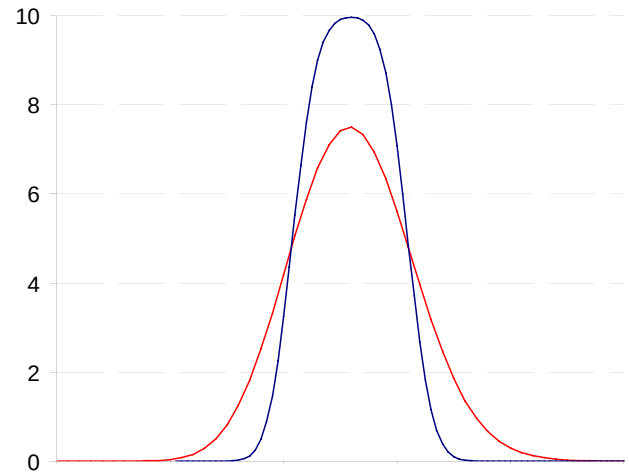
X = 0



X = 250 D

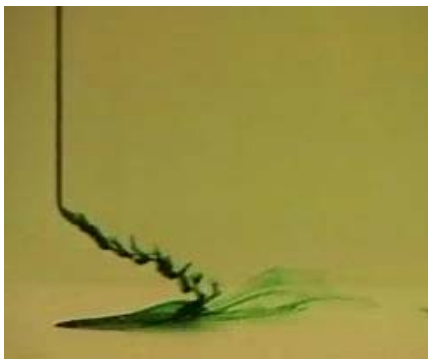


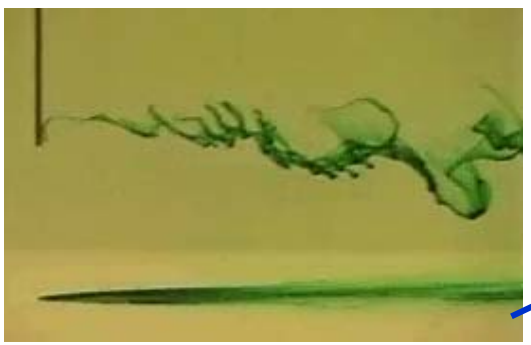
X = 500 D



Why Axial Dispersion?

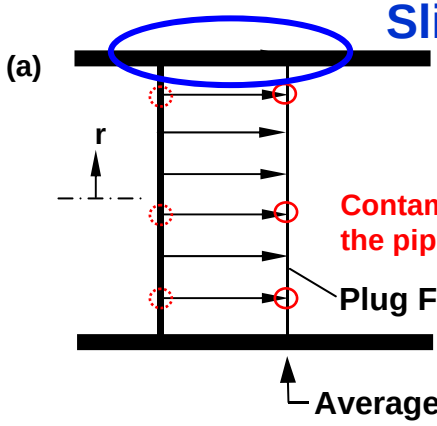
Typical tracer dispersion along the pipe wall

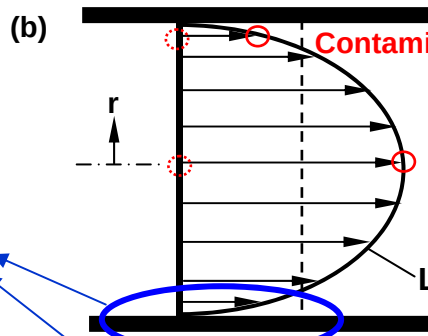




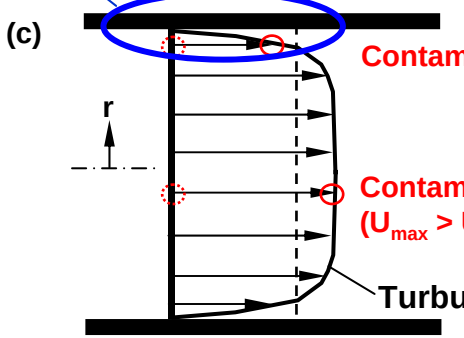
(Moran et al. 2003)

Slip Boundary Condition

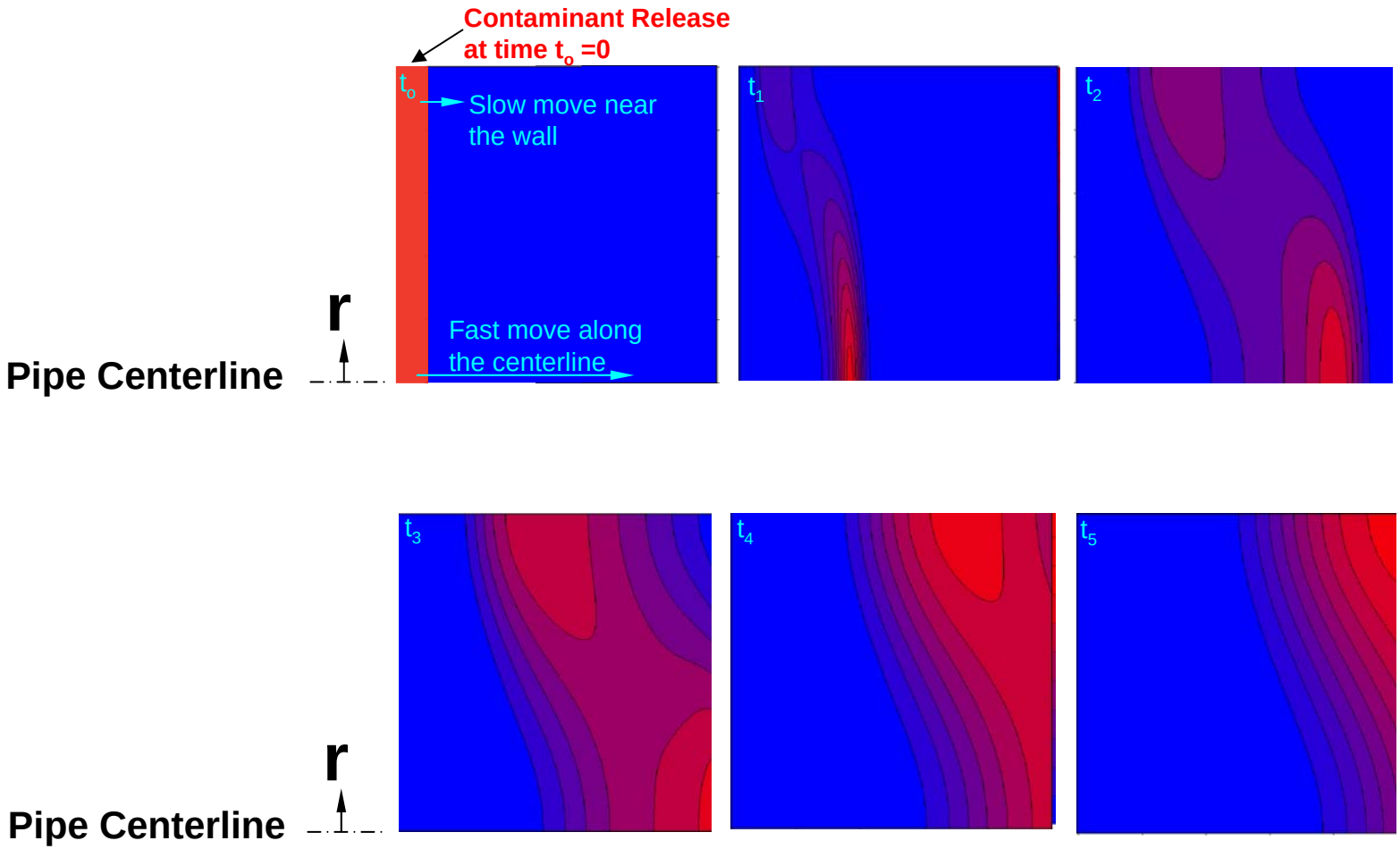




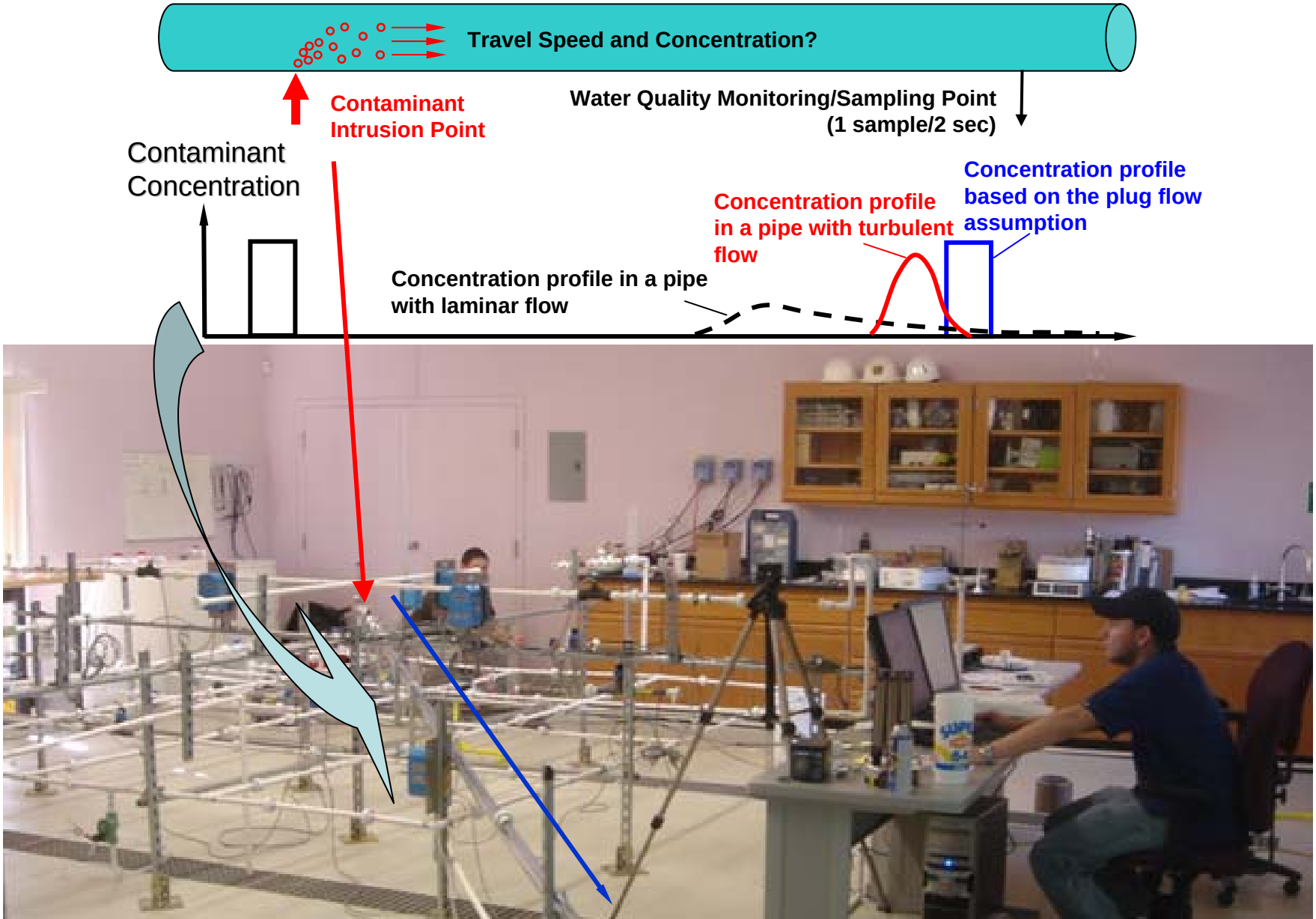
No-Slip Boundary Condition



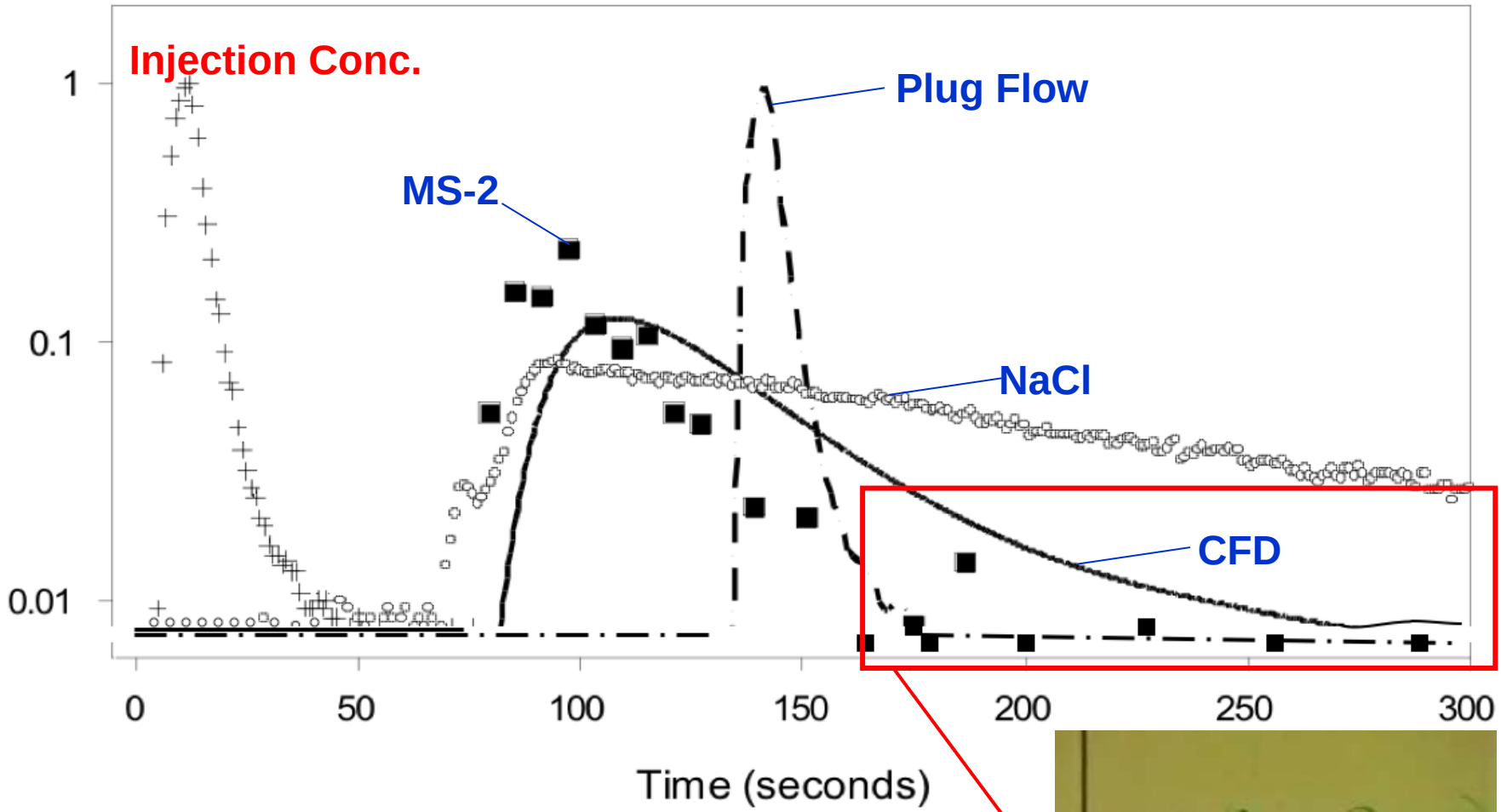
Computational Fluid Dynamics Simulations

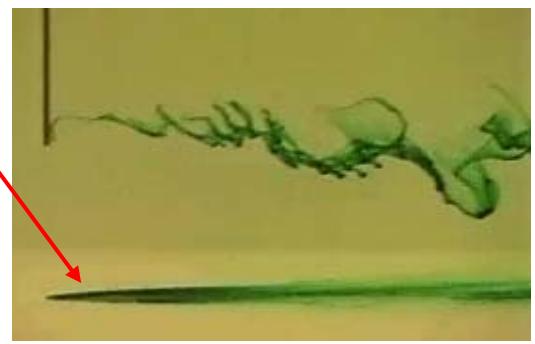


Experimental Setup



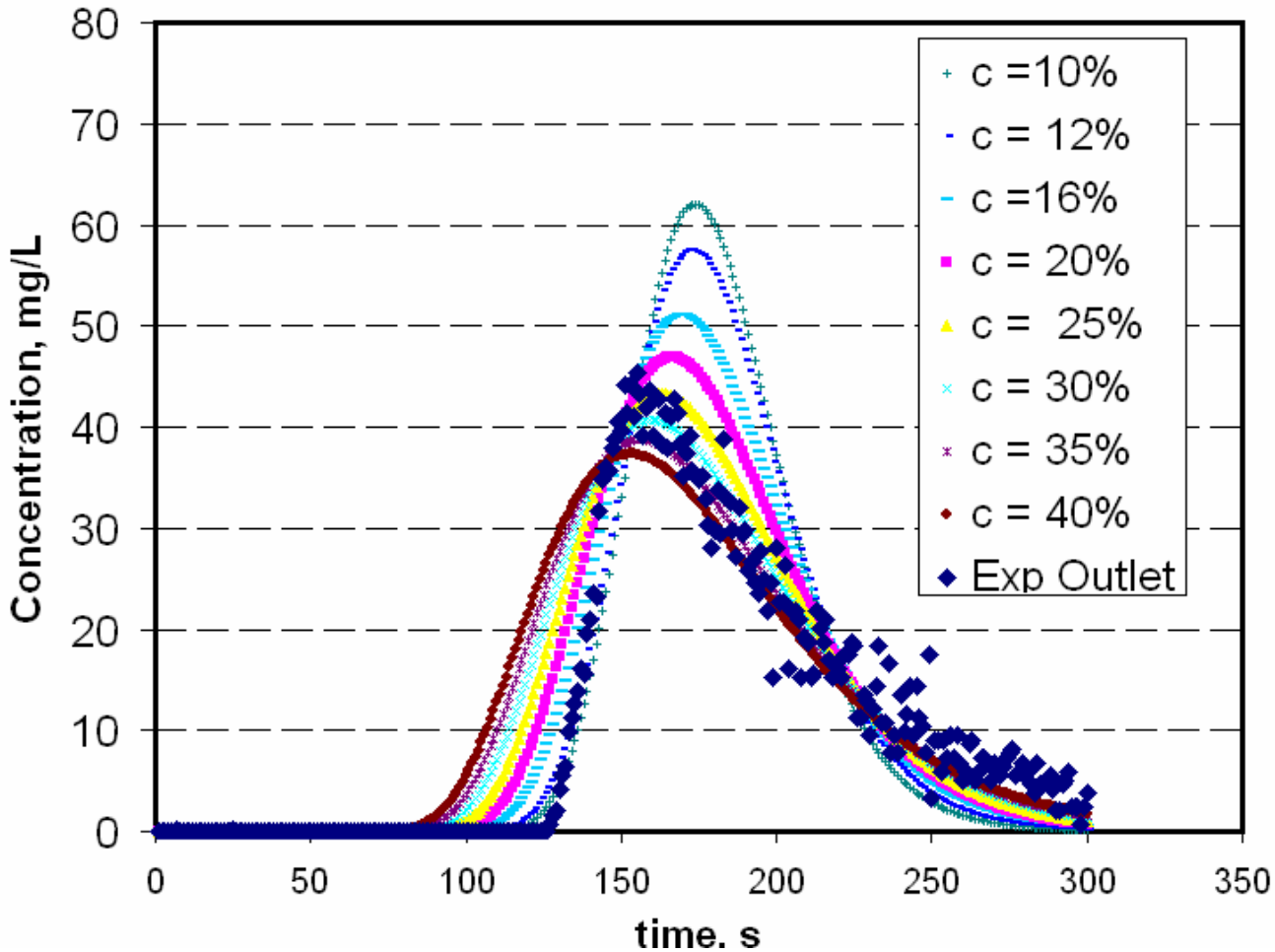
Injection of tracers
Laminar flow





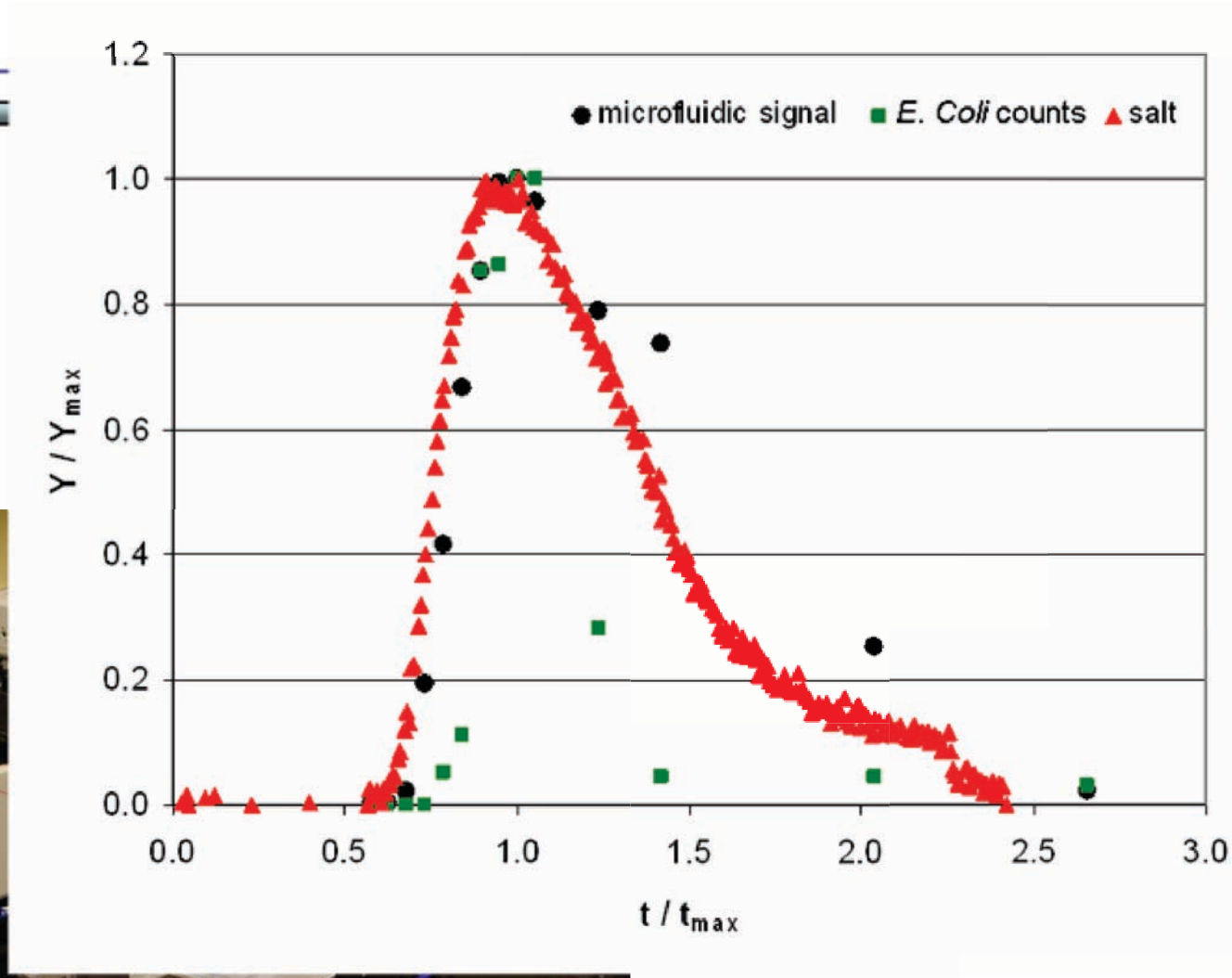
Axial dispersion & tailing of microorganisms:
Detection limits, sampling frequency, sensor locations

Examination of Existing Models



ADRNET (Buchberger and Li)/IMTARED (Tzatchkov)/In-House Code (Kang and Lansey)/Experimental Data (Choi)

Real Time Biosensor Research

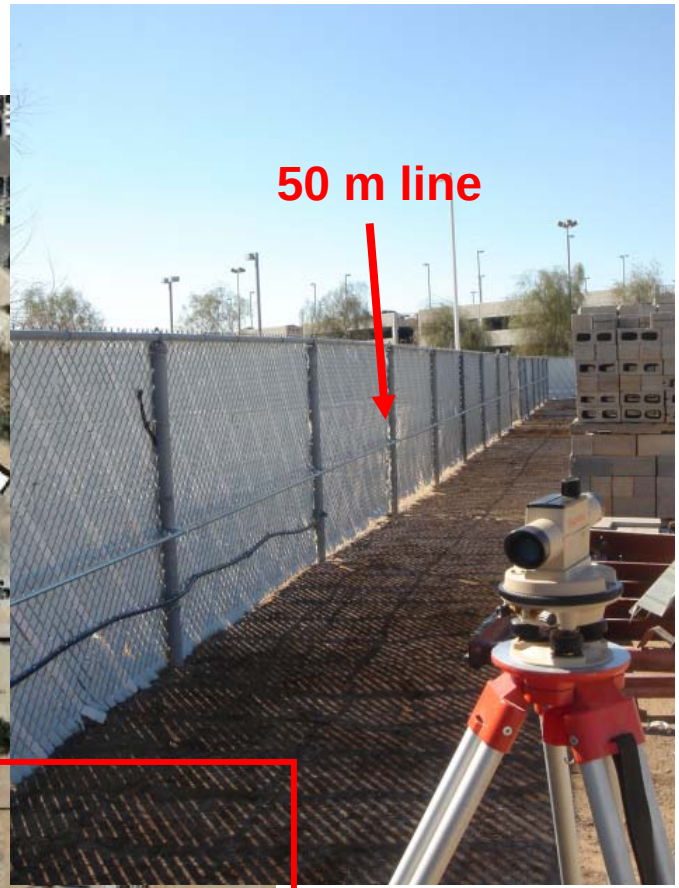


Syringe pump

Courtesy of Professor Yoon
The Univ. of Arizona

Lab Scale ⇨ Pilot Scale Experiments

Pumping Station

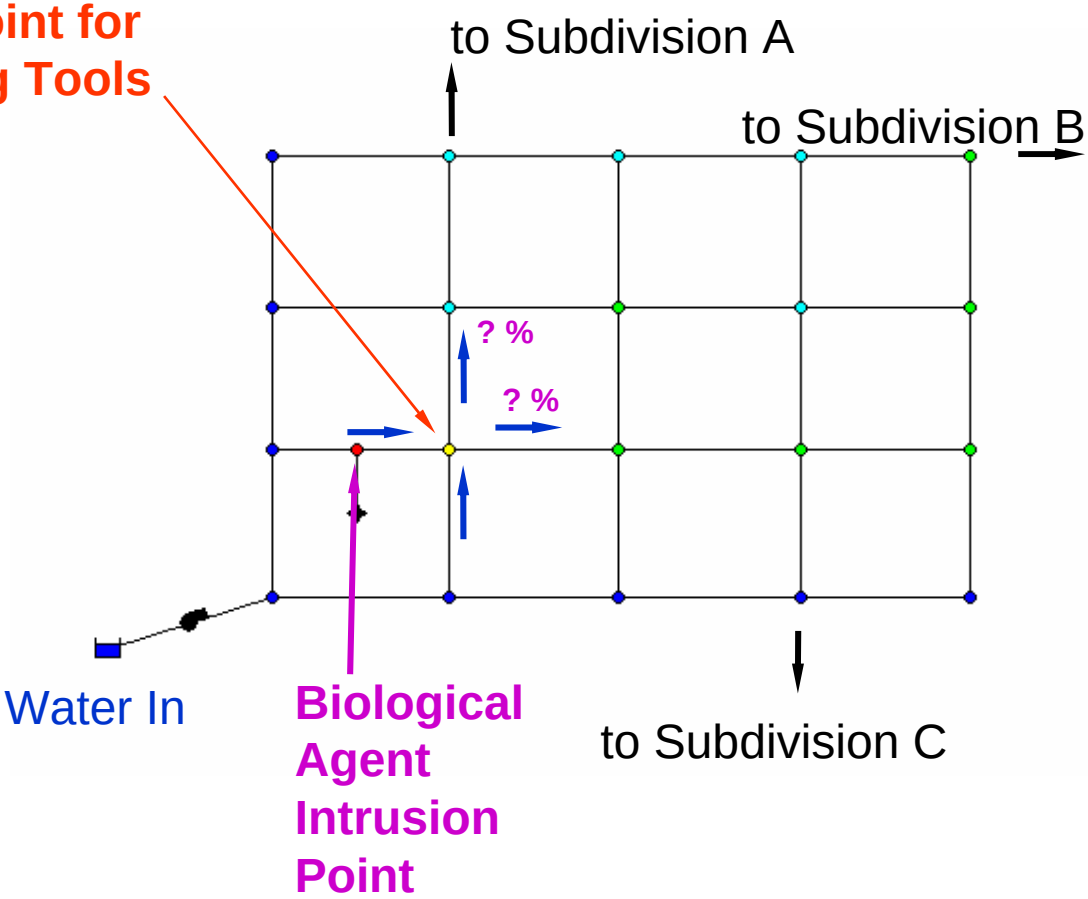


Trench

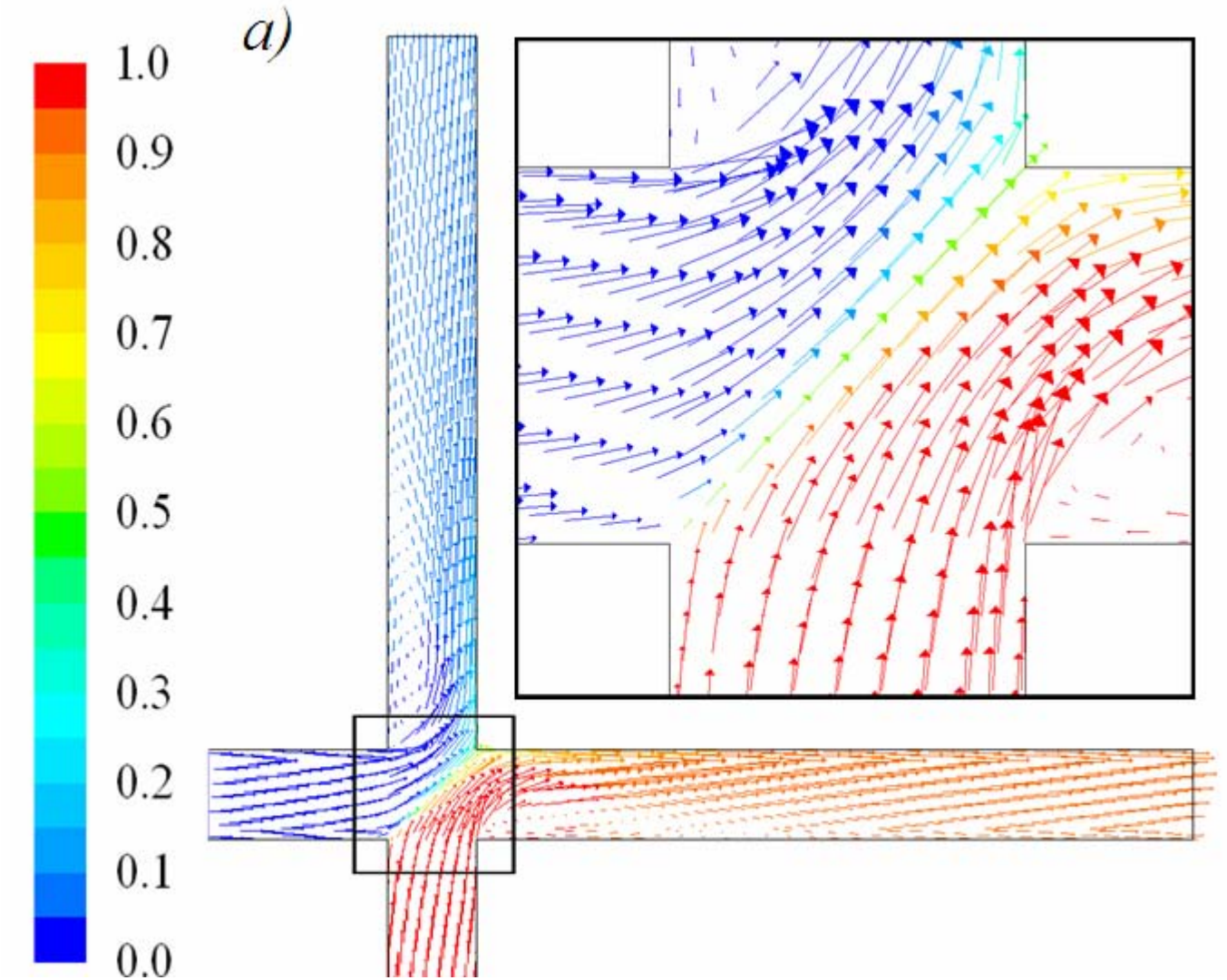
50 m line for dispersion experiments

Mixing at Junctions

Perfect Mixing Assumed at the Cross Joint for Modeling Tools

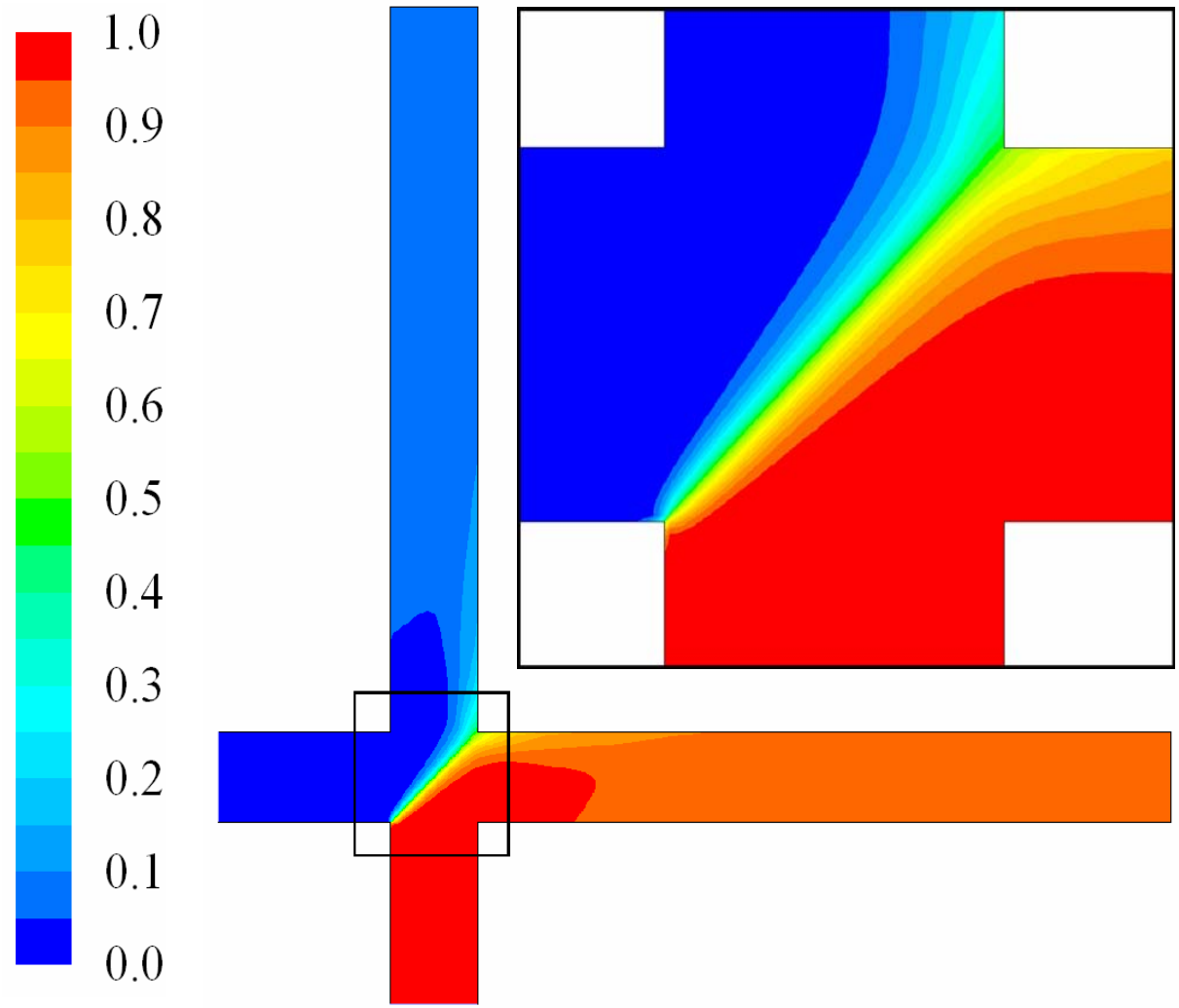


Mixing patterns along the interface



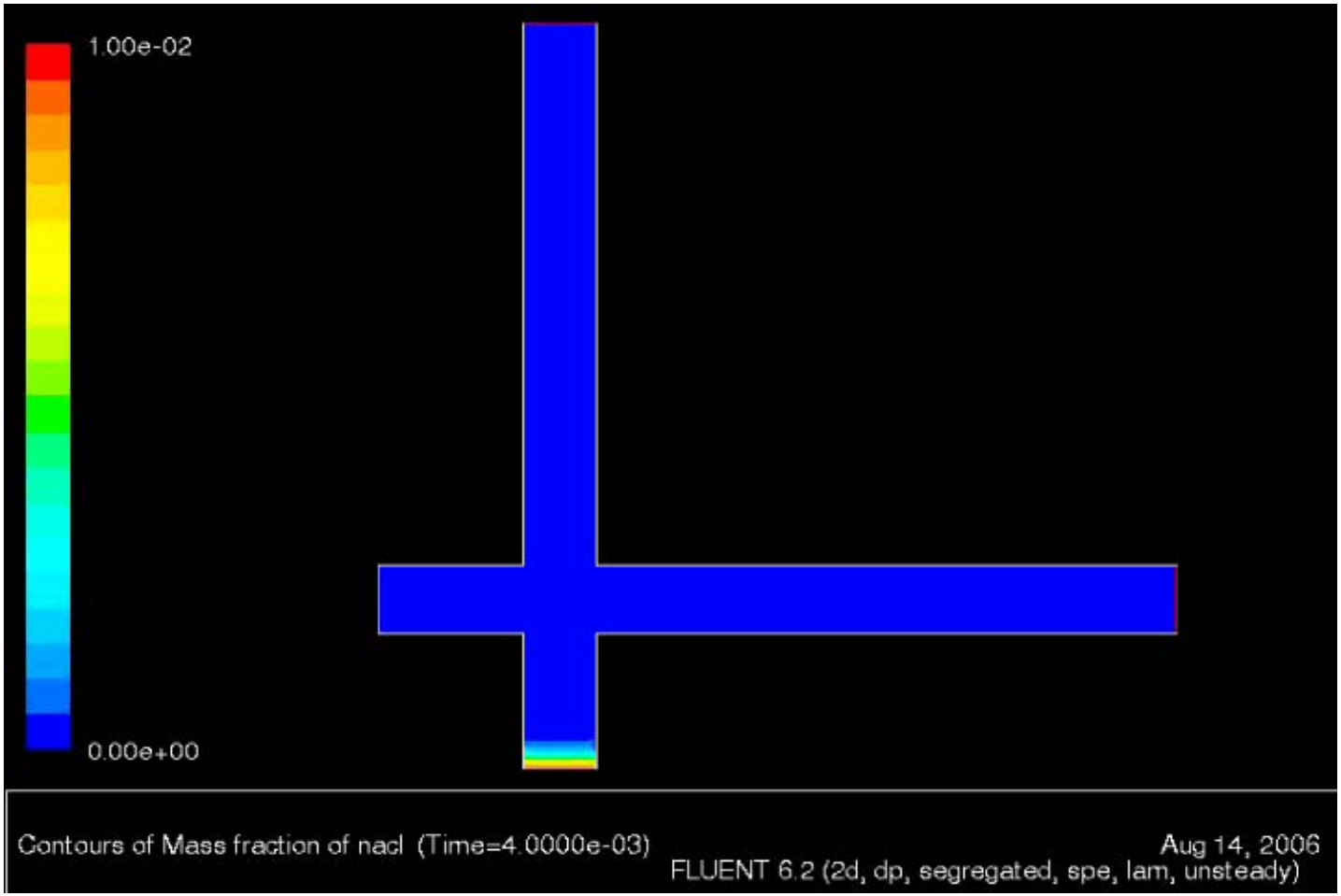
Velocity vectors at a cross junction, when $Re_S = Re_W = Re_E = Re_N = 44{,}000$ ($Re_{S/W} = 1$, $Re_{E/N}=1$), and $Sc_t = 0.1875$

Mixing patterns along the interface



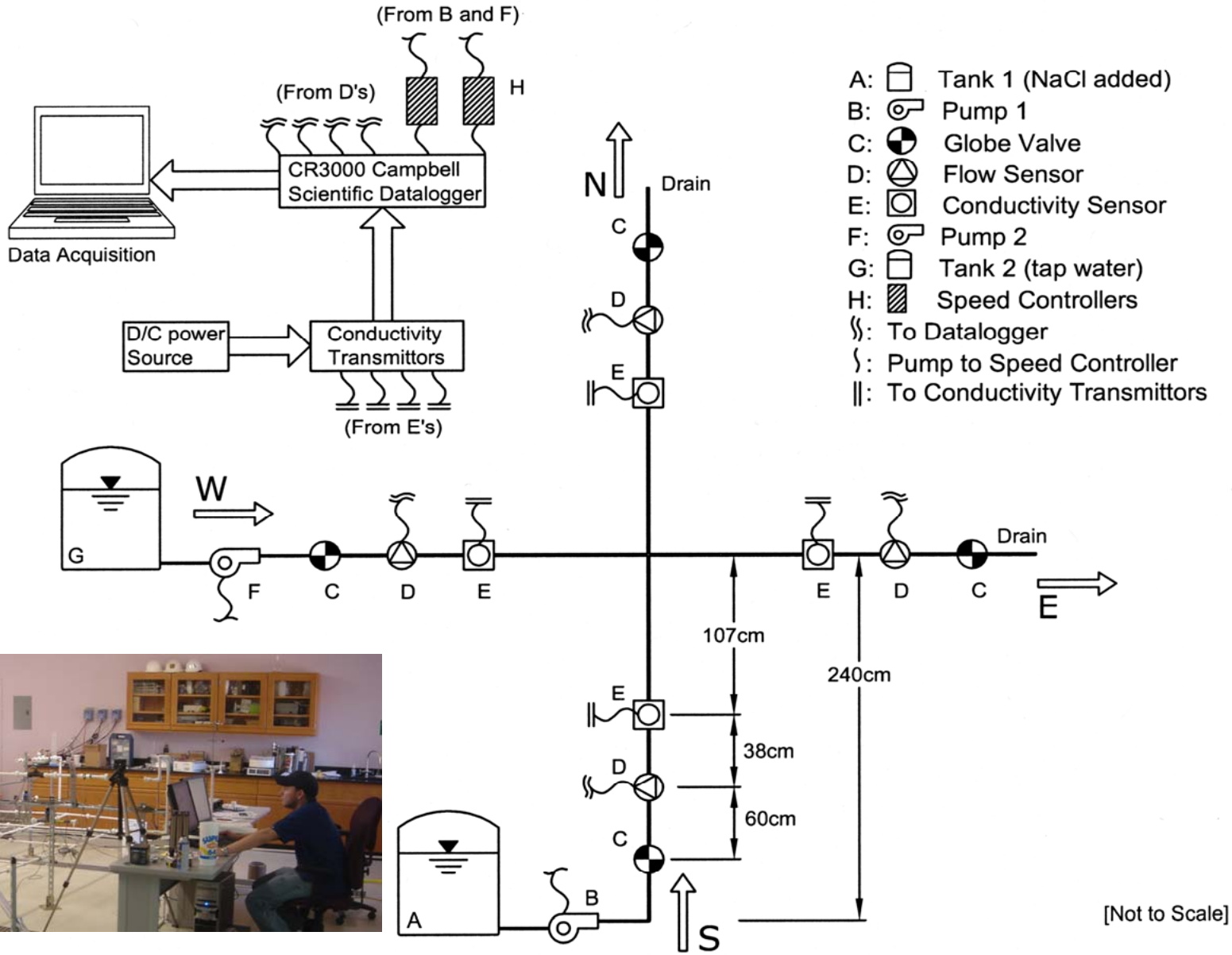
Dimensionless NaCl Concentration contours at a cross junction, when ReS = ReW = ReE = ReN = 44,000 (ReS/W = 1, ReE/N=1), and Sct = 0.1875

Mixing patterns along the interface

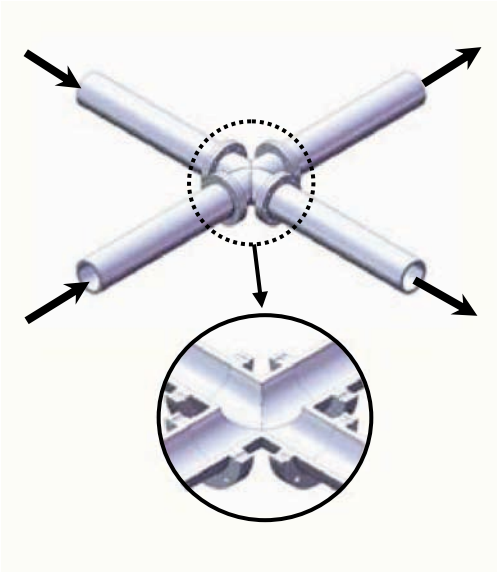


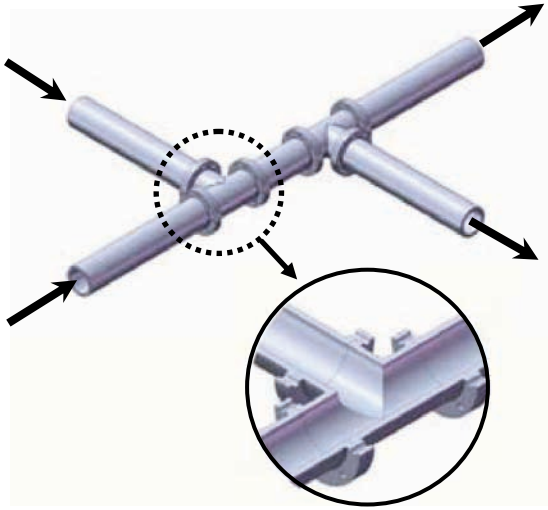
Simulation Video Clip

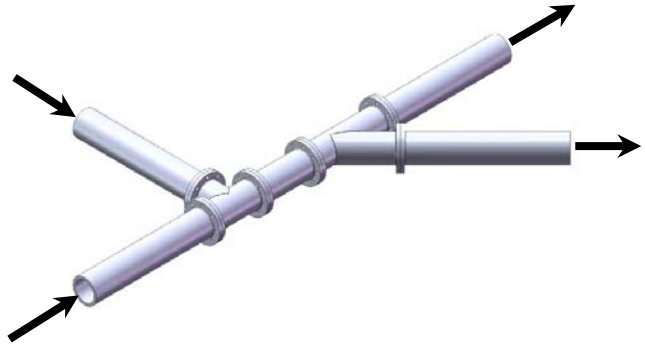
A schematic of the experimental setup

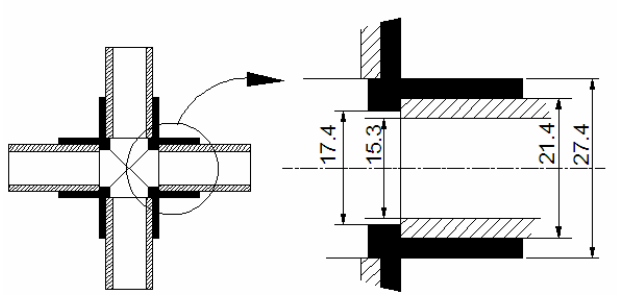


Representative Junction Types

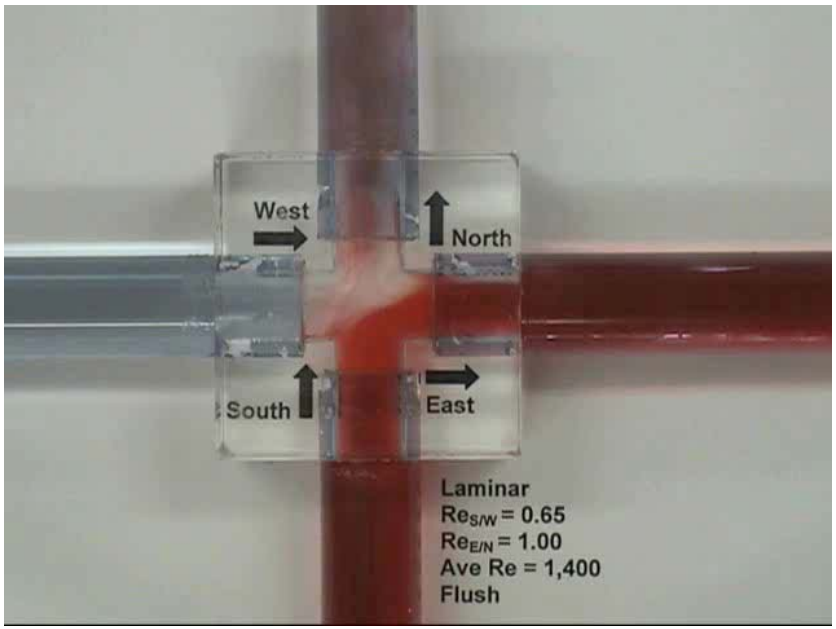


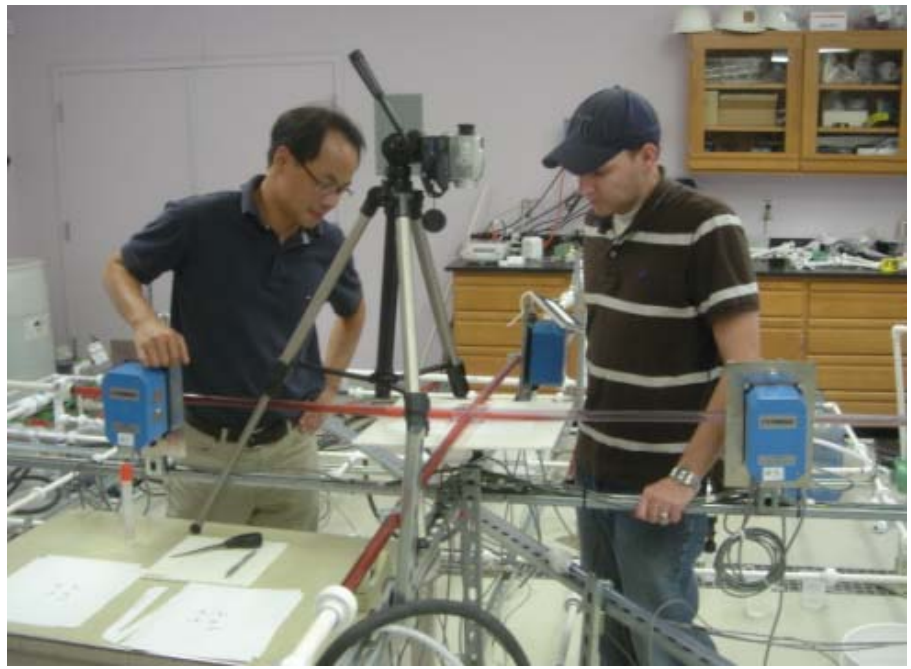






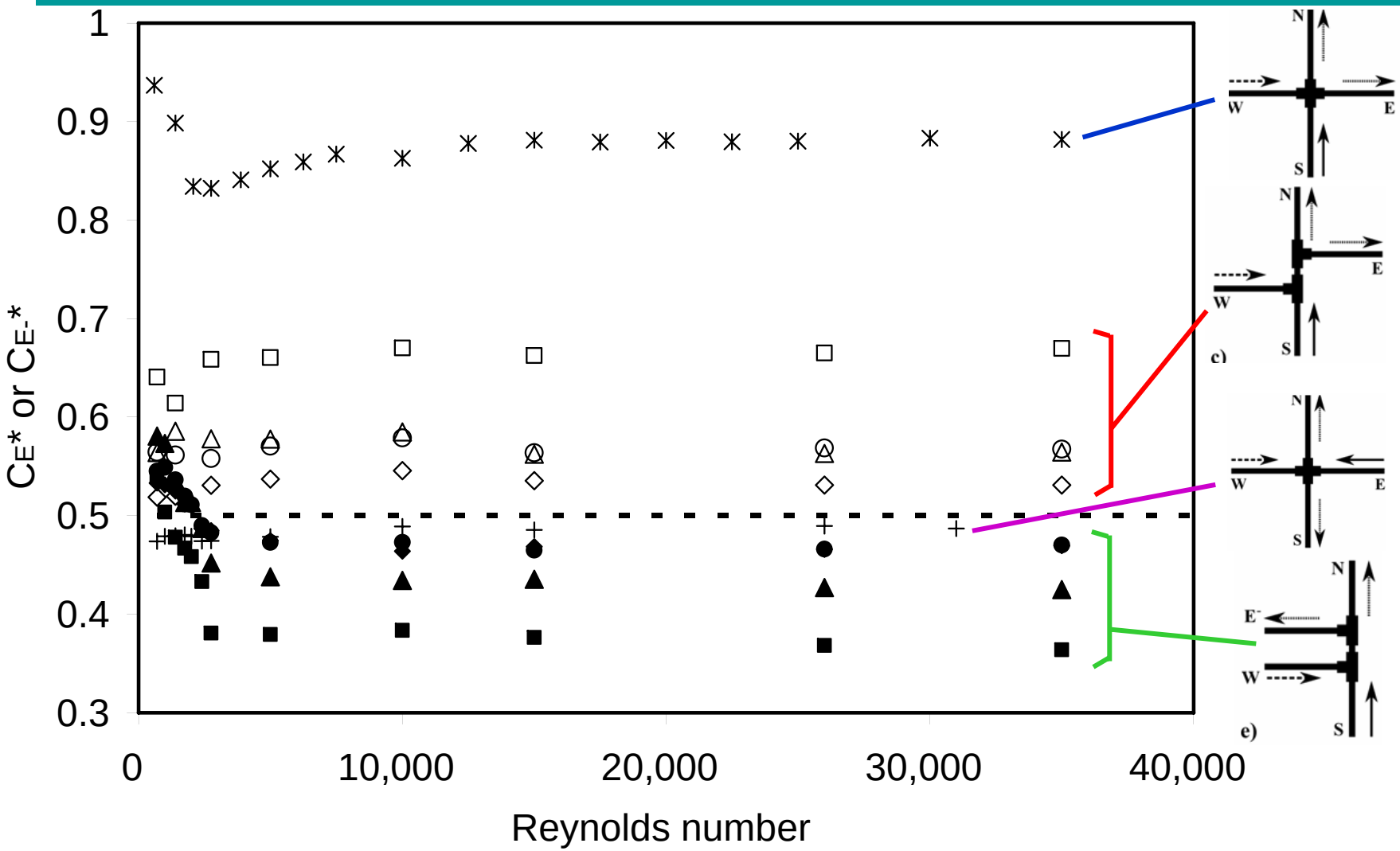
Flow Visualization





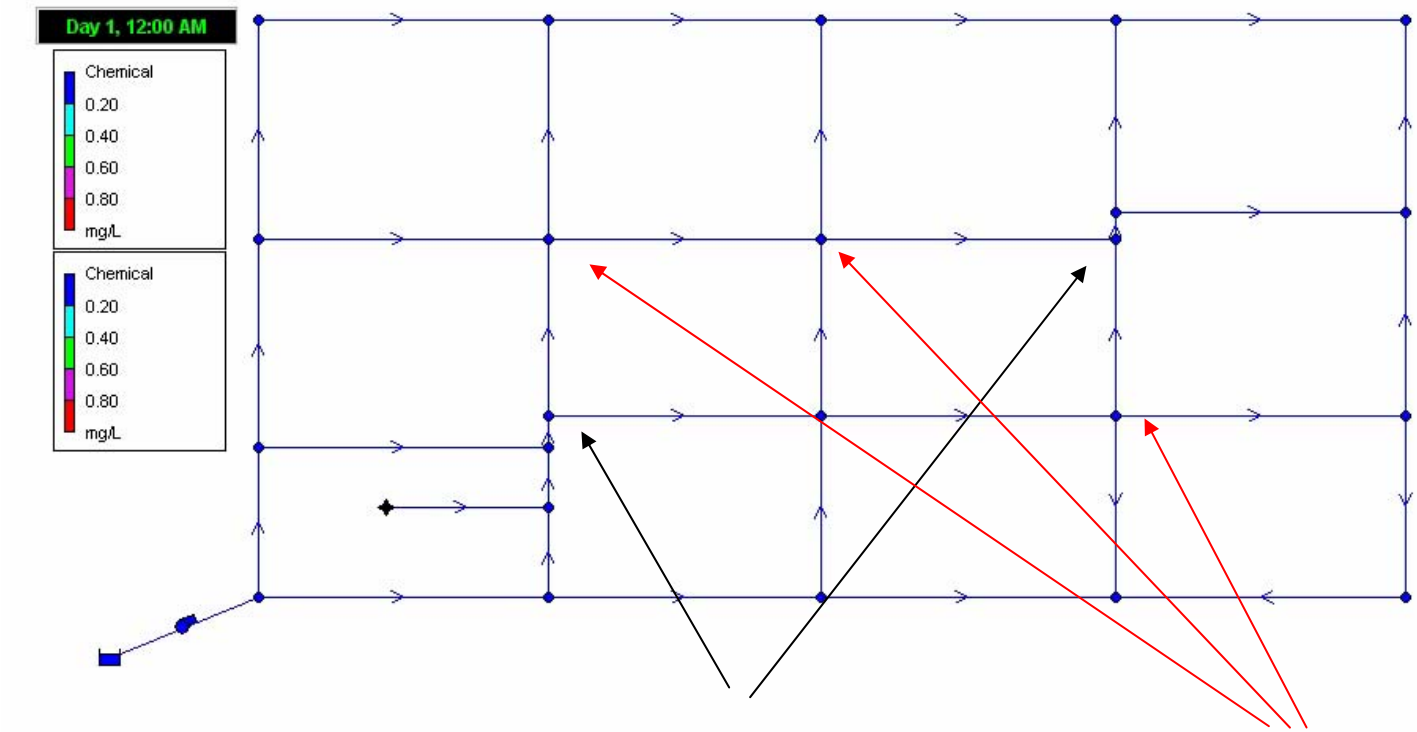
X, T, Y junctions for Scenario 1

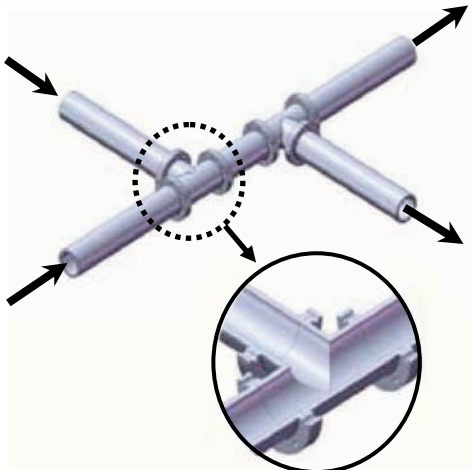
$Re_E = Re_W = Re_S = Re_N$

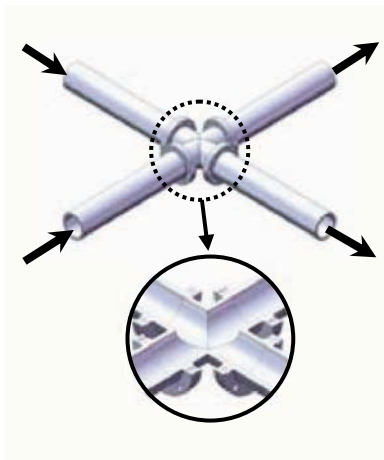


✱ D=0, Cross, X1	+ D=0, Cross, X2	□ D=2.5, N1	△ D=5.0, N1
○ D=7.5, N1	◇ D=10, N1	■ D=2.5, U1	▲ D=5.0, U1
● D=7.5, U1	◆ D=10, U1	– – – Complete Mixing	

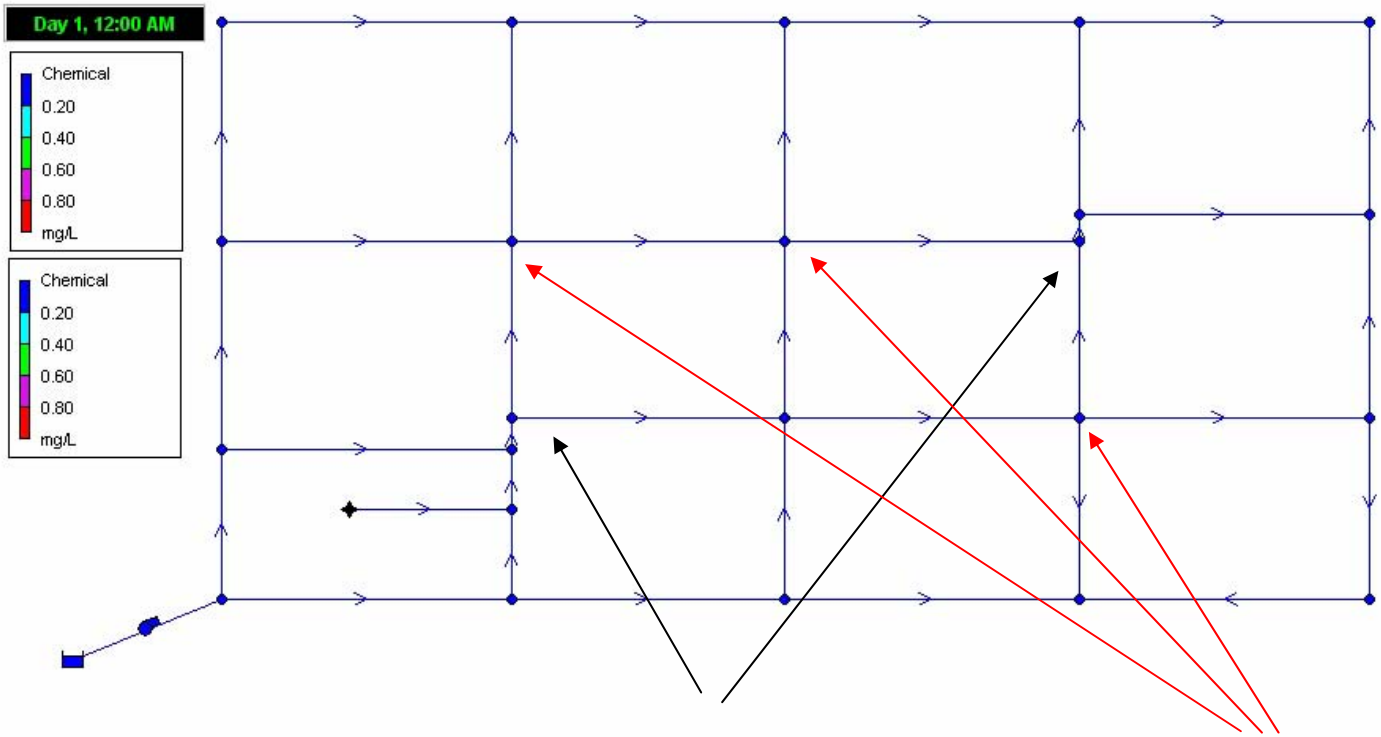
Development of AZRED (with Perfect Mixing Model)

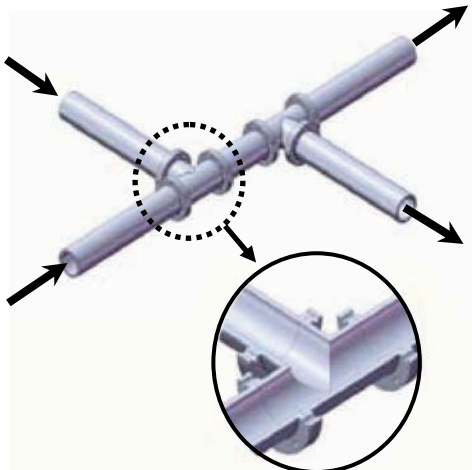


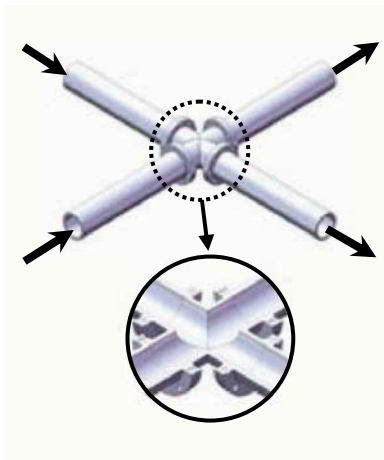




Development of AZRED (Improved Water Quality Model)

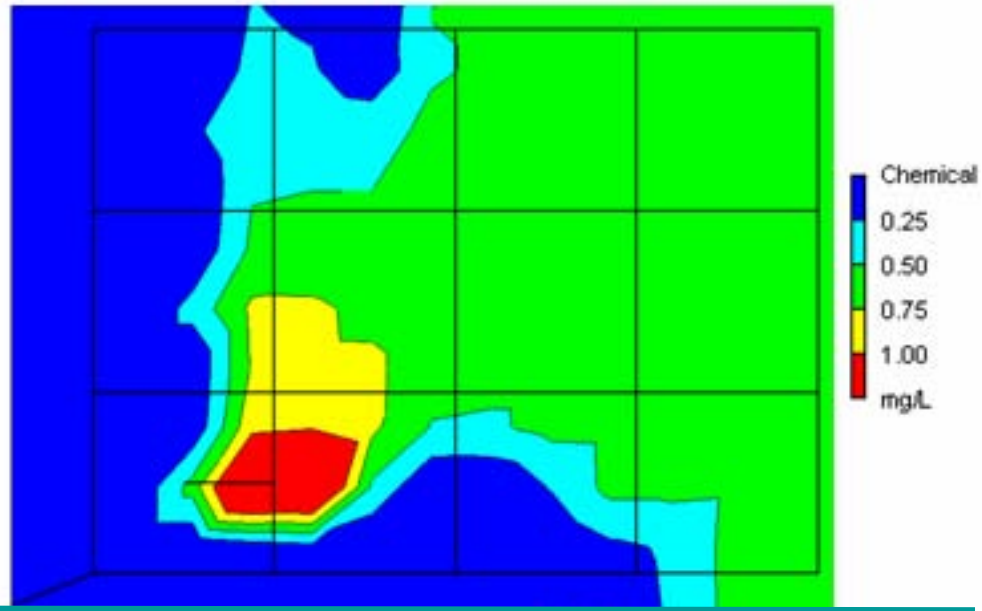




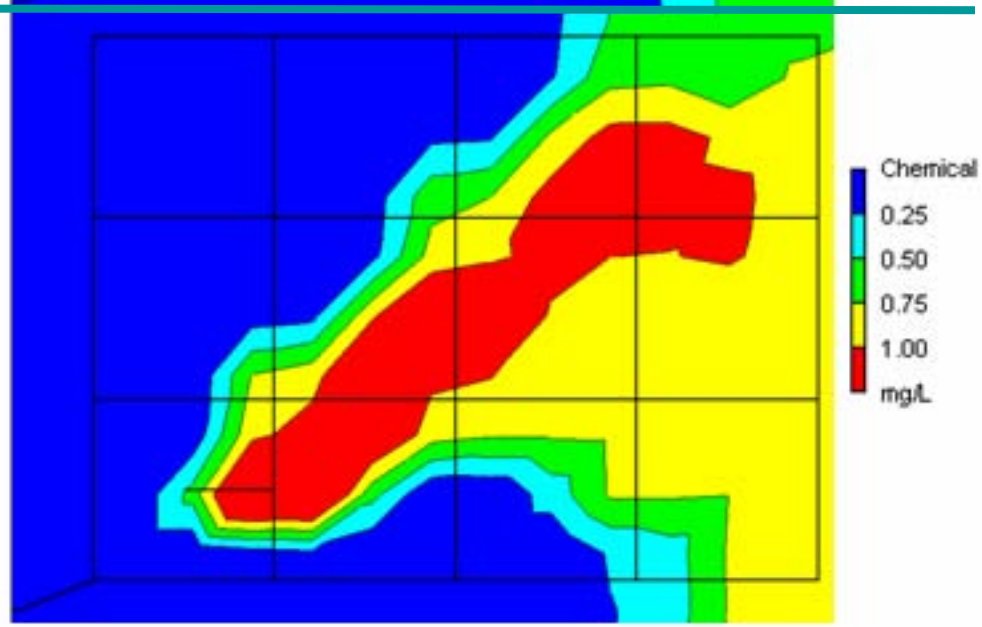


Comparison

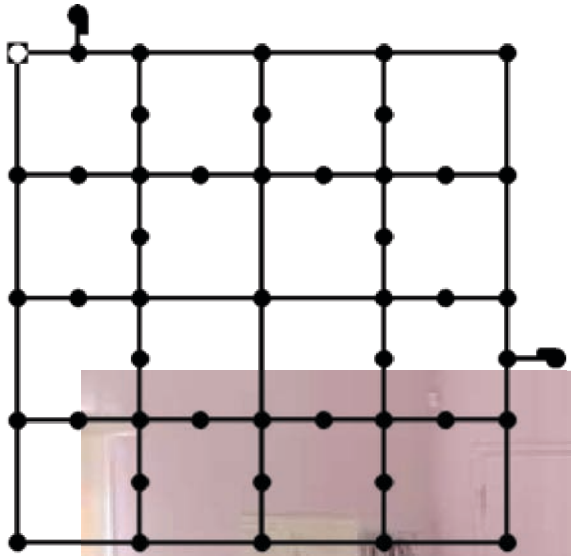
Existing Model based on Perfect Mixing

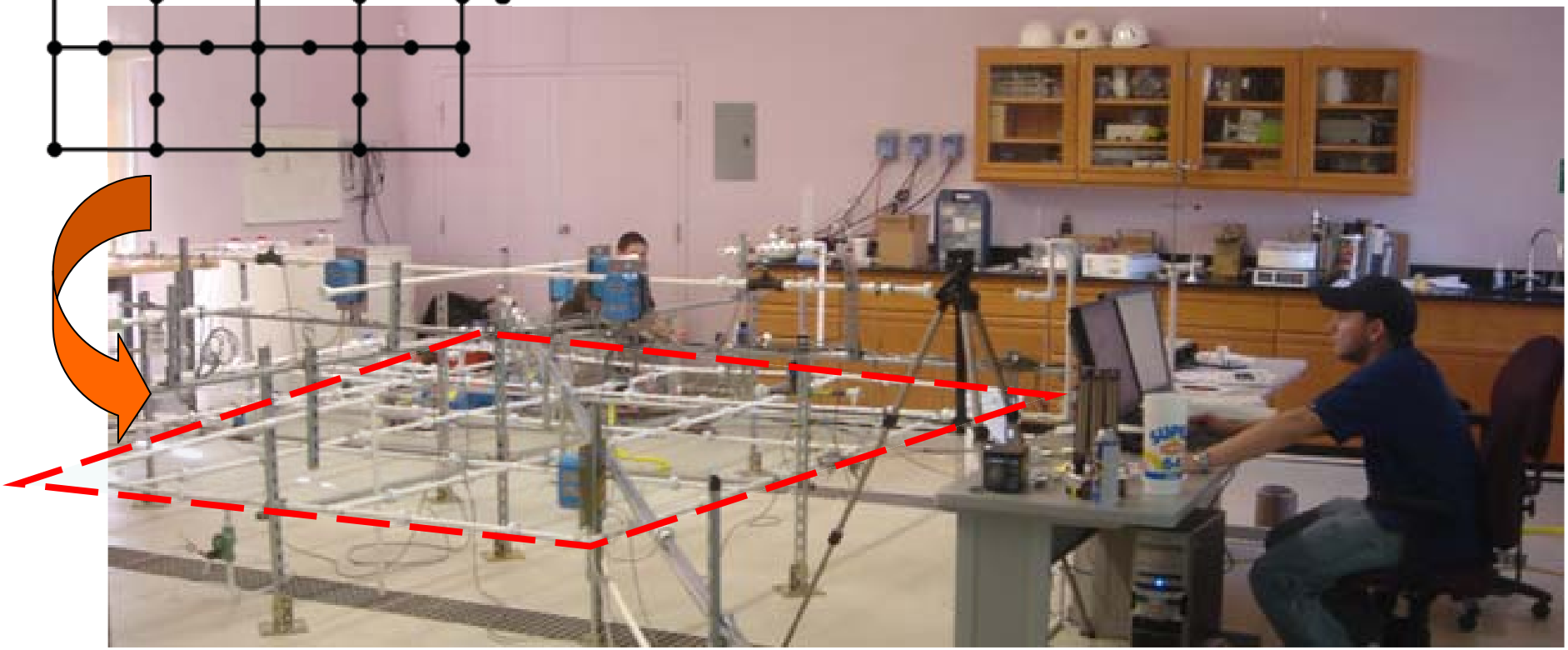


AZRED

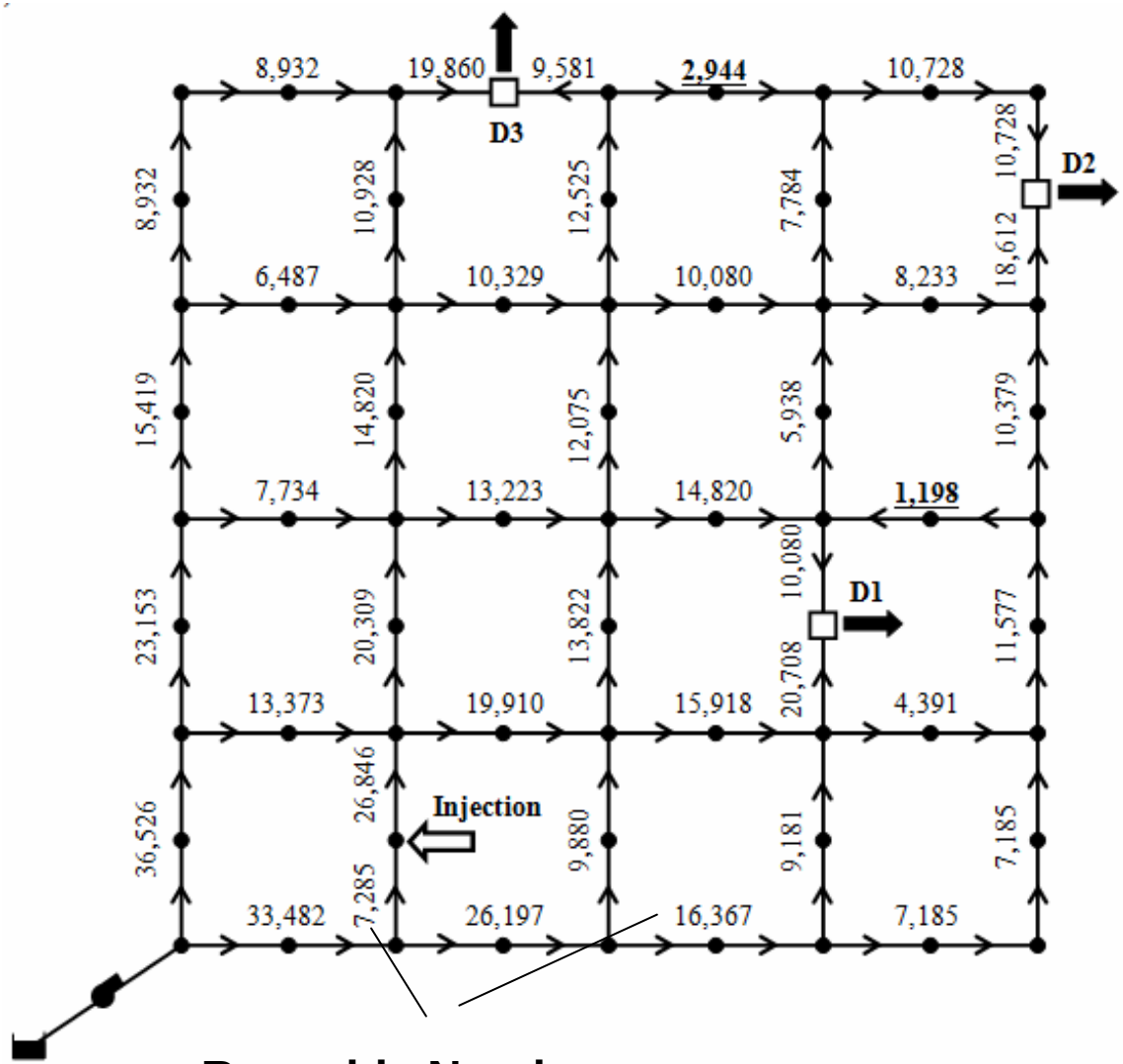


5x5 Network Establishment





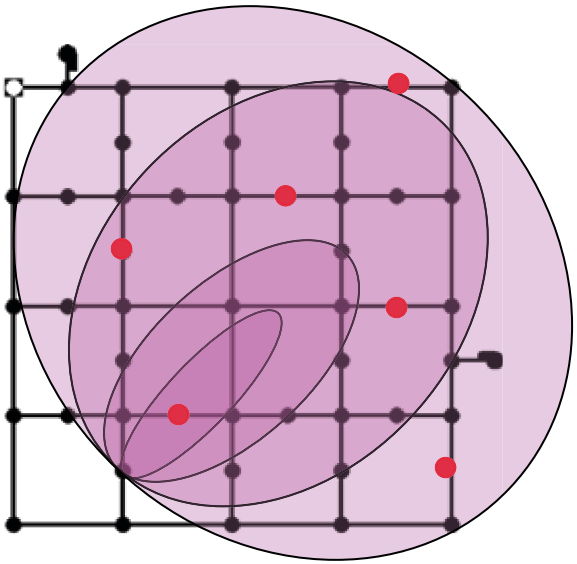
EPANET Hydraulic Model (Experimentally Validated)

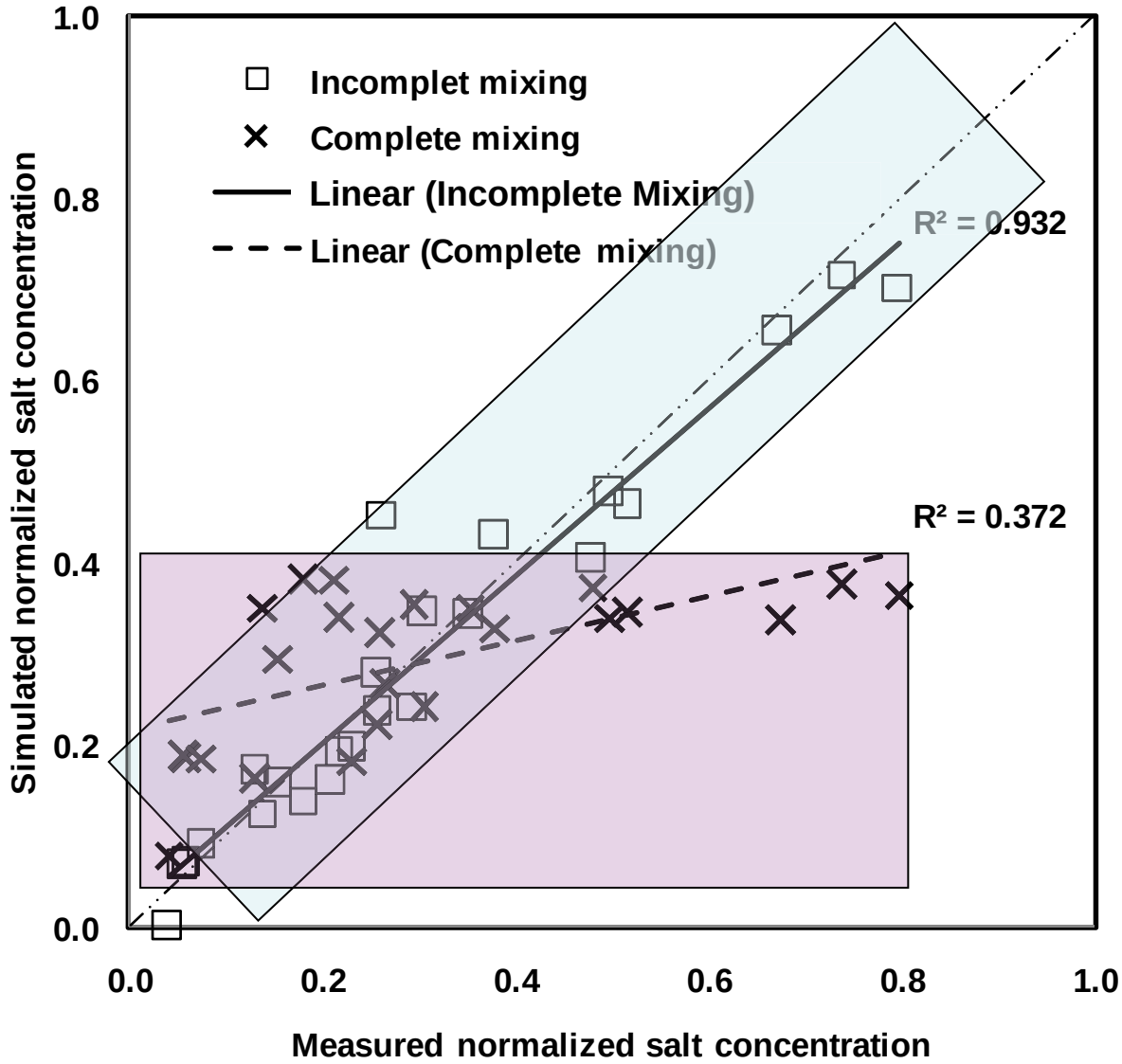


Reynolds Numbers

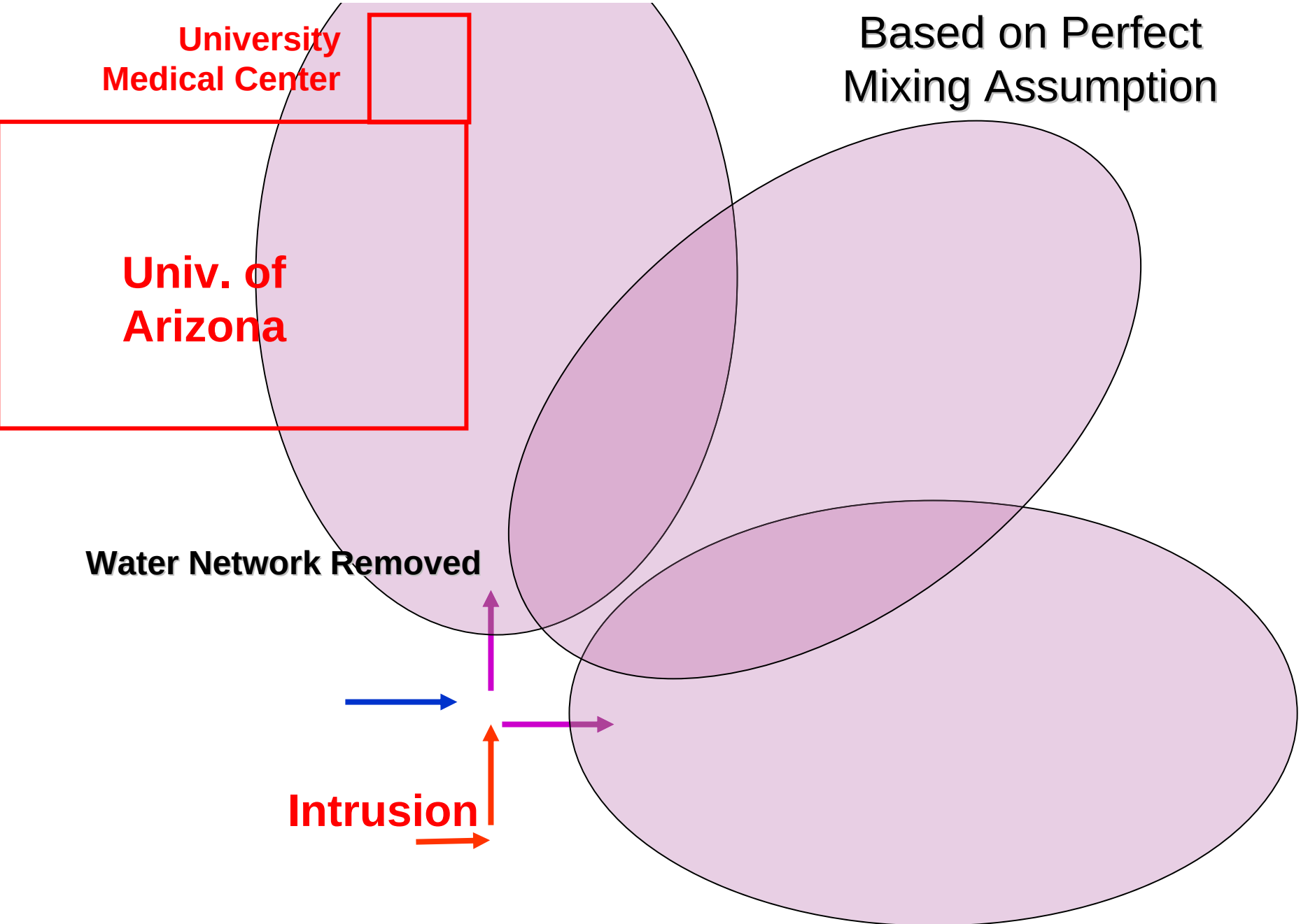
AZRED Water Quality Model vs. Experimental Results

- Sensor Locations

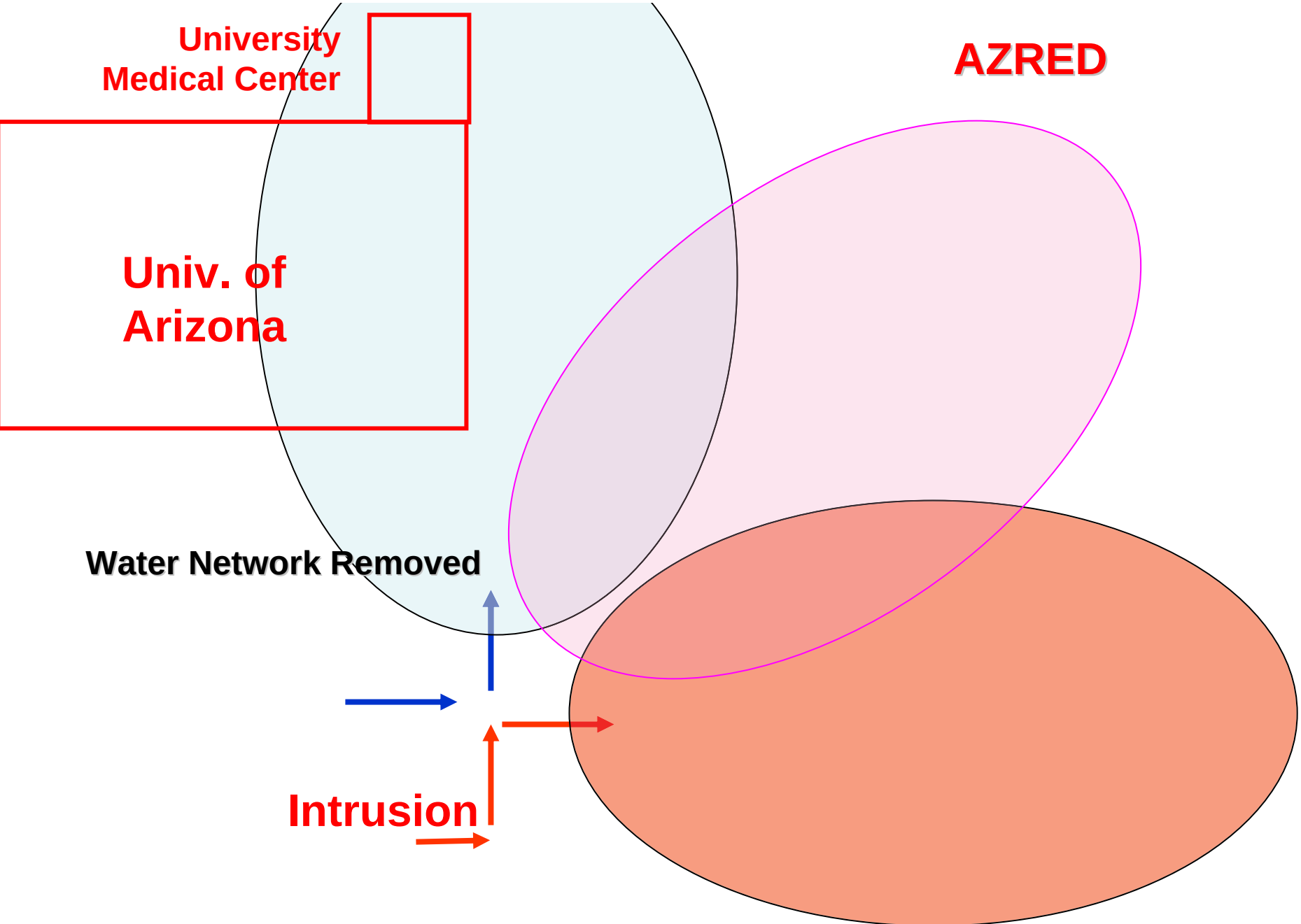




Corresponding Risk Microbial Risk Assessment & Consequences



Corresponding Risk Microbial Risk Assessment & Consequences

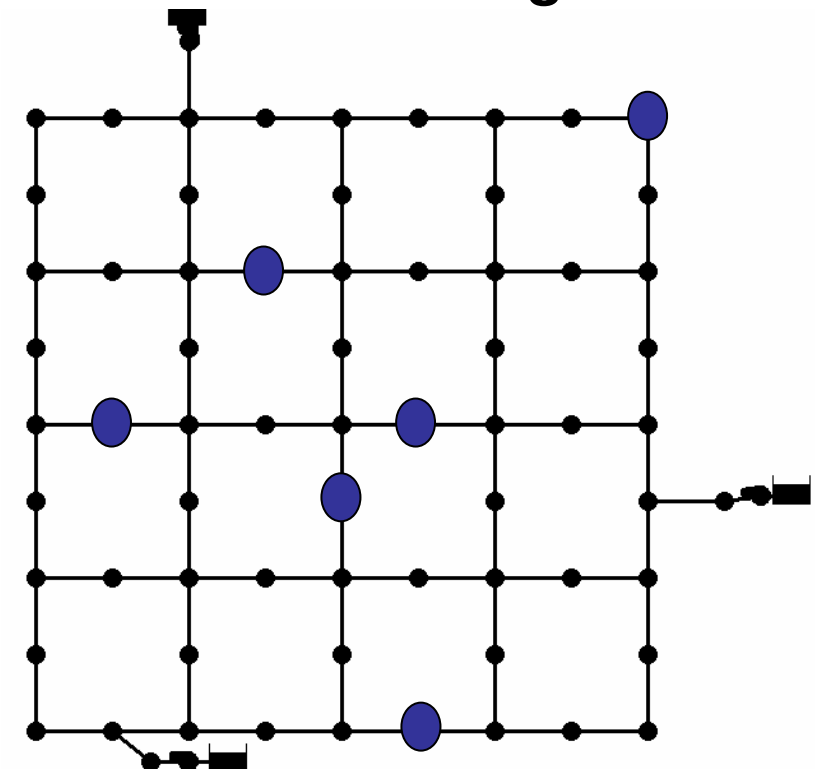


Optimal Sensor Placement

Minimum Hazard Levels (MHL) = 1 mg/L
Level of Services (LOS) = 25 m^3

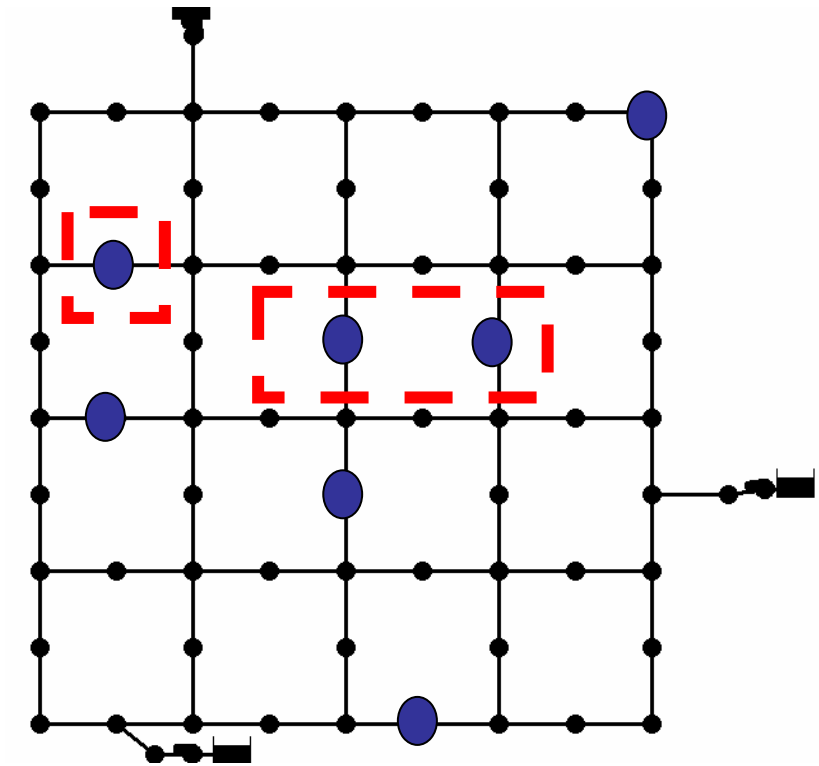
The total number of sensors and their locations change based on AZRED water quality data.

Perfect Mixing



6 sensors

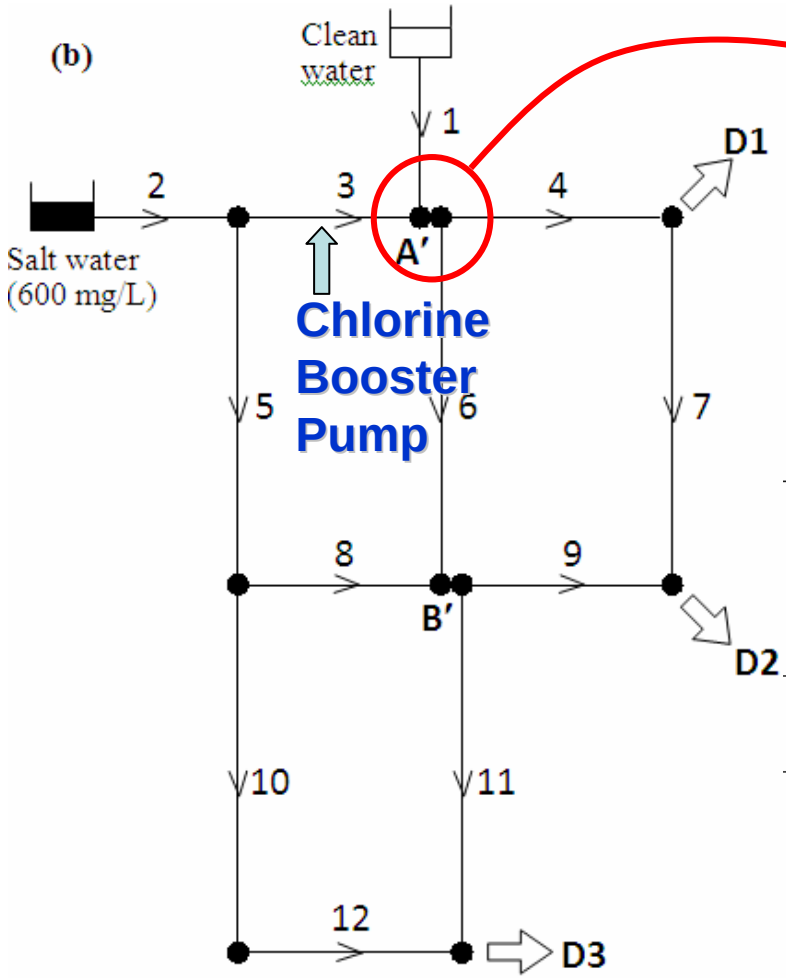
AZRED

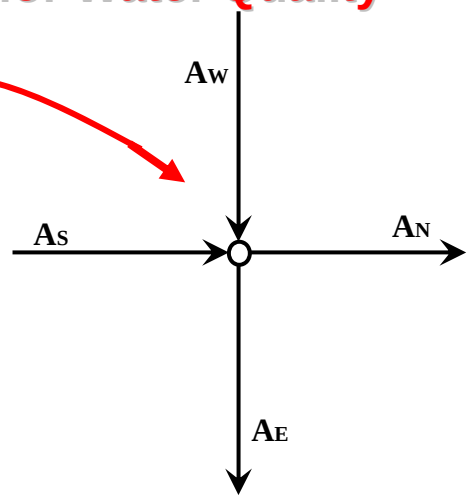


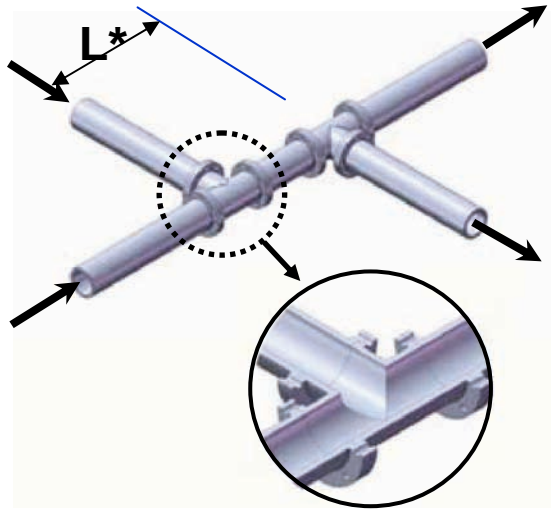
7 sensors

Beyond Water Security: Water Safety and System Design

Water Age = Reliable Surrogate for Water Quality

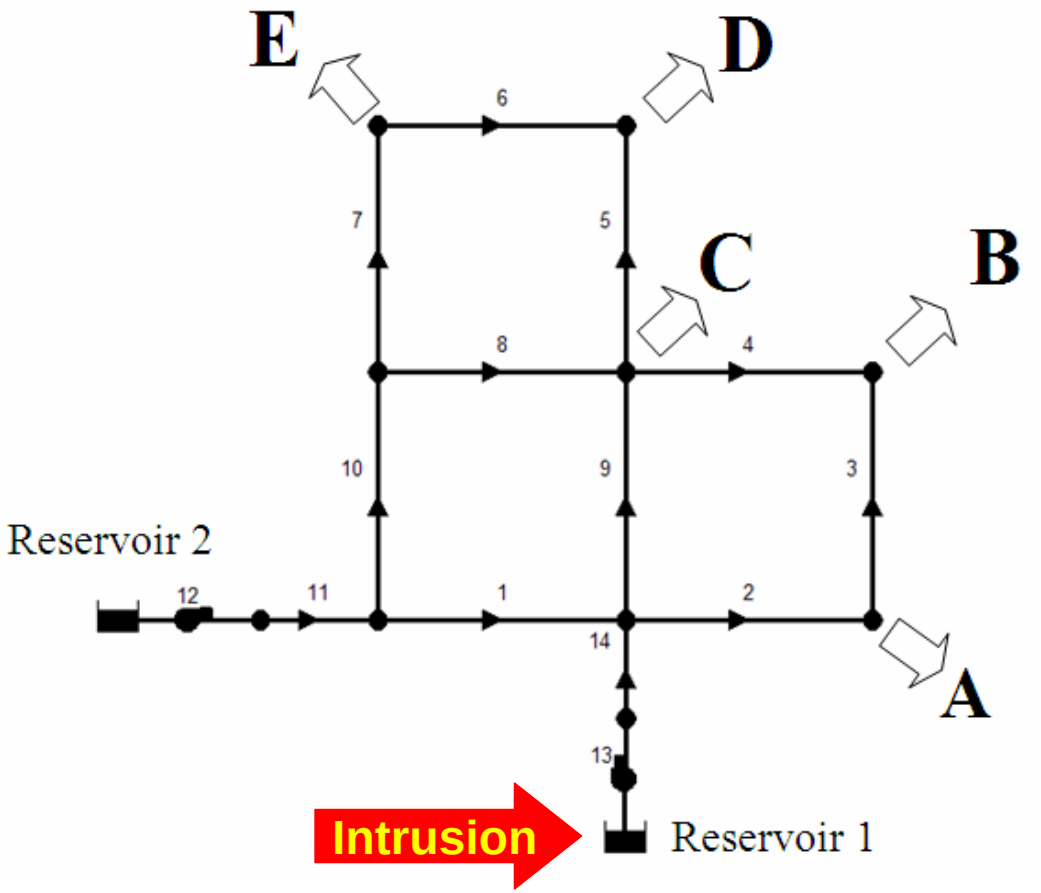


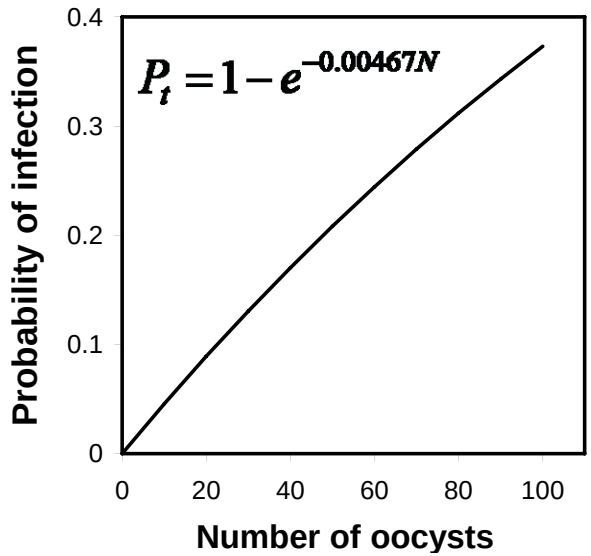




Assumption	L*	Water age (hr)			
		A_S	A_W	A_N	A_E
Complete mixing (A0)	-	1.56	2.97	2.27	2.27
Incomplete mixing (A1)	0.0			2.76	1.62
	2.5			2.47	2.00
	5.0			2.35	2.15
	10.0			2.30	2.22

Microbial Risk Assessment

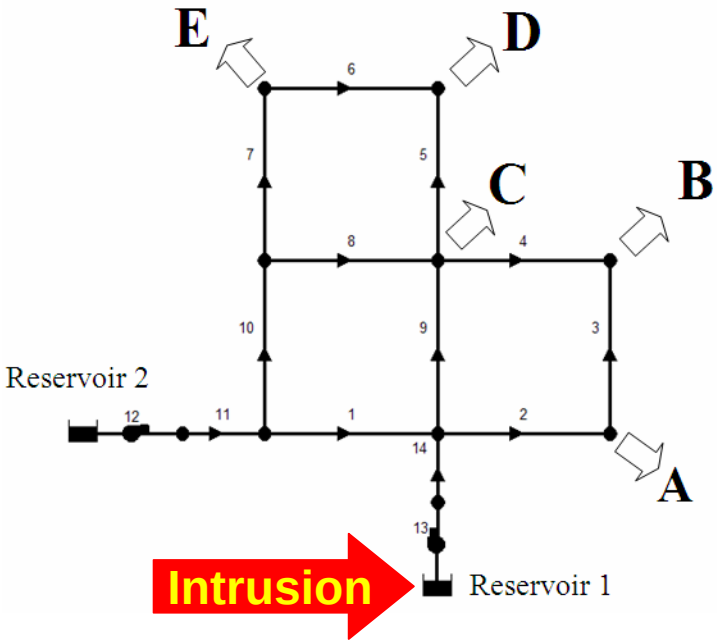


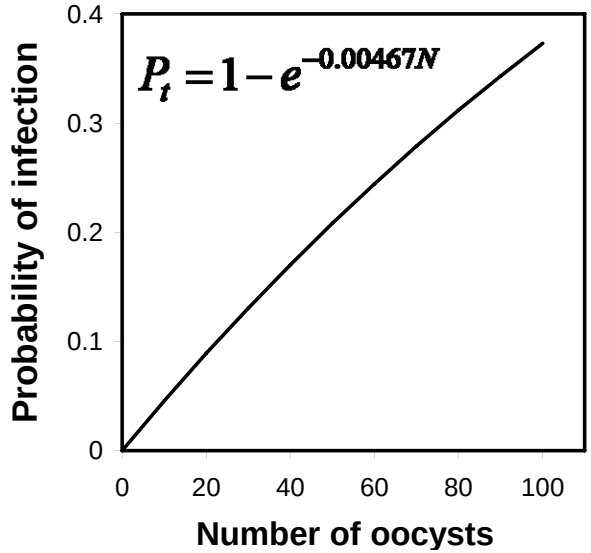


Dose-Response curve, Cryptosporidium (Haas *et al*., 1996)

Based on **'complete' and 'incomplete'** mixing assumptions, **calculate the number of infected people during this event**, when the water consumption of tap water is assumed equal to 2 liters per person per day, and an exposure time of one day.

Significance of Modeling for QMRA

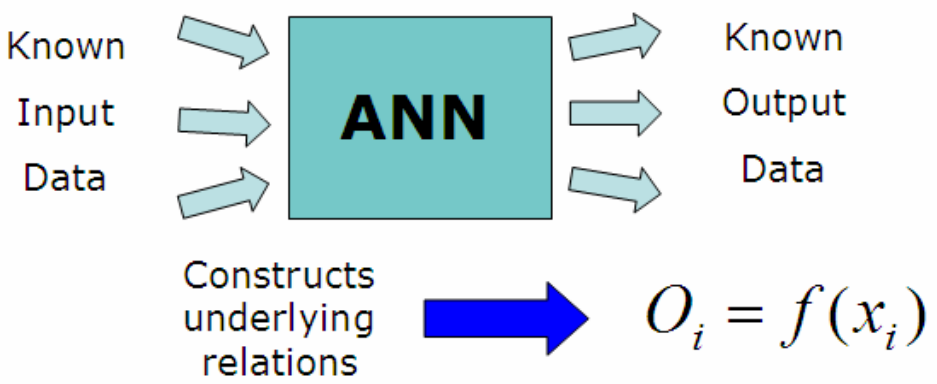




Dose-Response curve, Cryptosporidium (Haas *et al*., 1996)

Neighborhood	Assumption	Oocyst concentration (oocyst/L)	Ingested oocyst per person (dose)	Probability of infection (exponential model)	Infected population
A	Complete Mixing	21.83	44	0.186	186
	Incomplete Mixing	37.35	75	0.294	294
D	Complete Mixing	7.14	14	0.063	158
	Incomplete Mixing	0.50	1	0.005	13

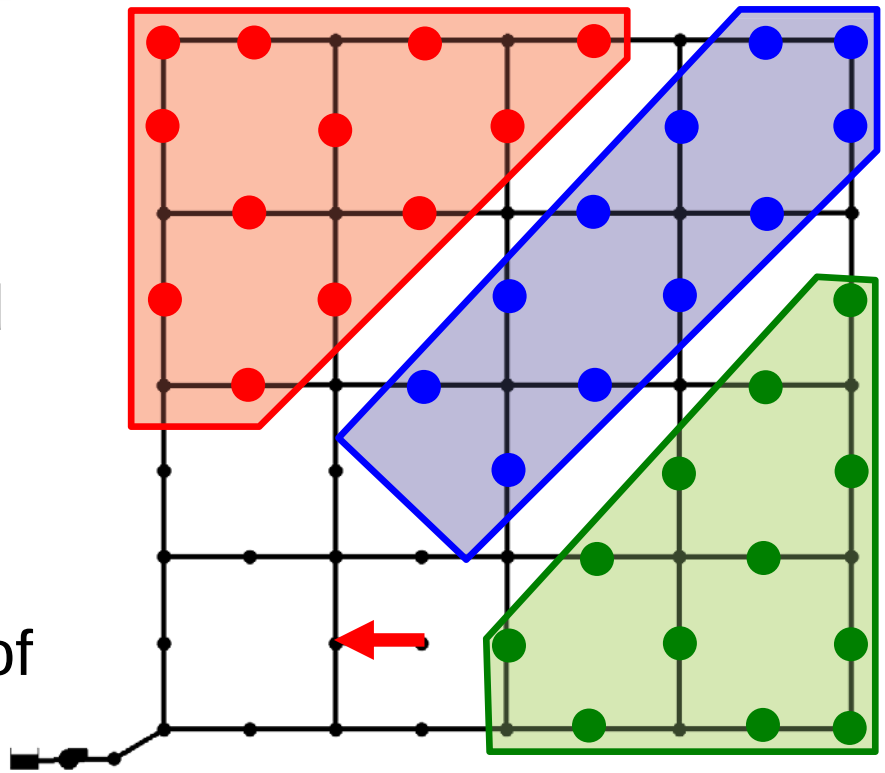
Artificial Neural Networks with AZRED



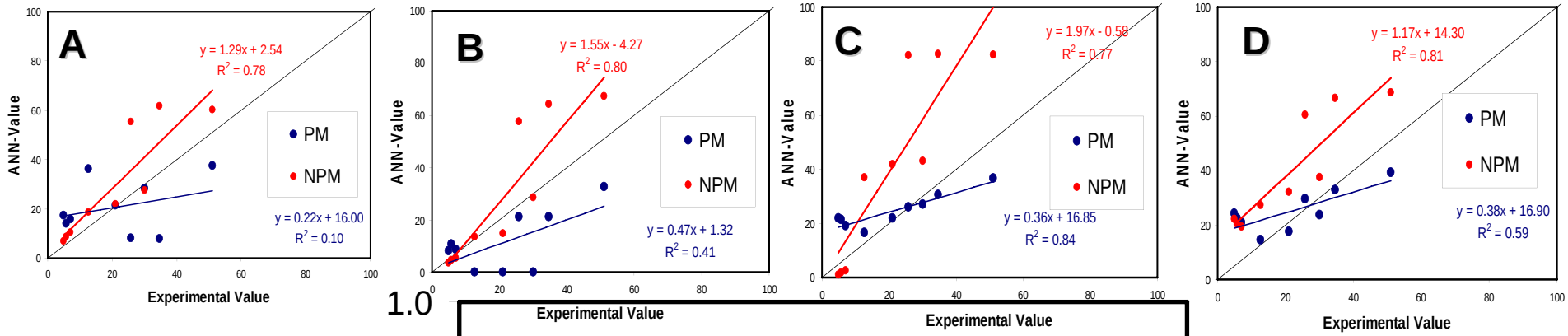
-Three random demands were set, one on each “region”

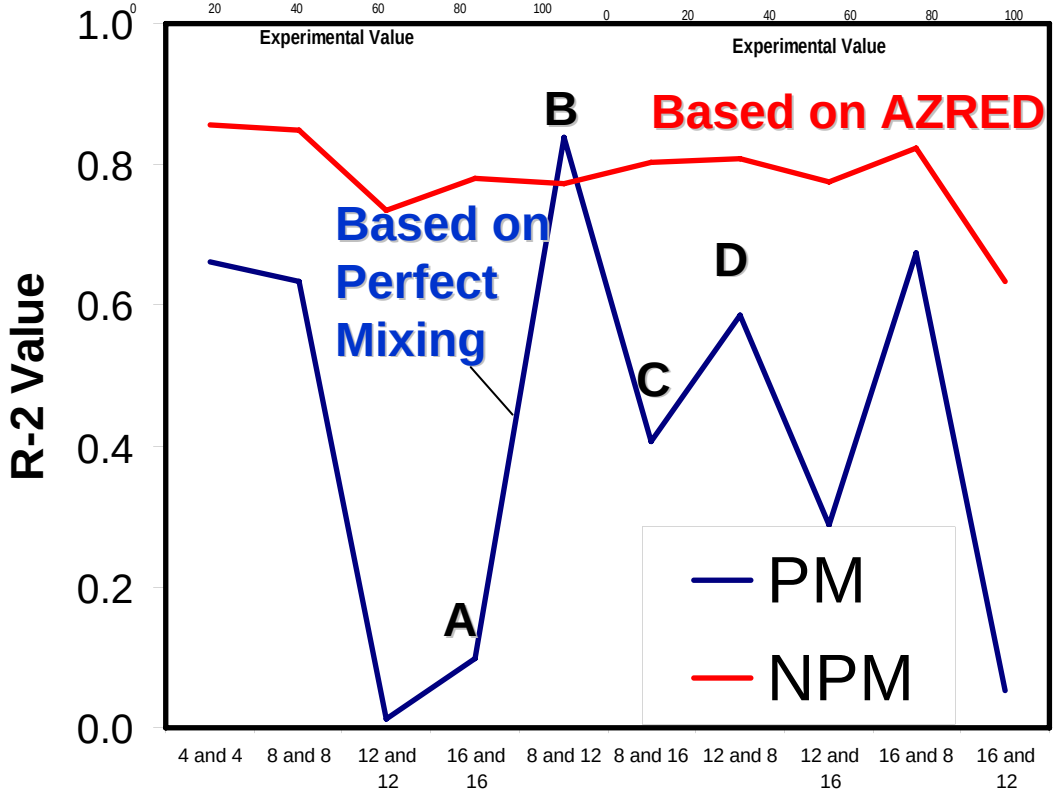
- AZRED simulations generated input data for ANN training and testing

- ANNs were trained on WQ models with both assumptions of mixing at junctions

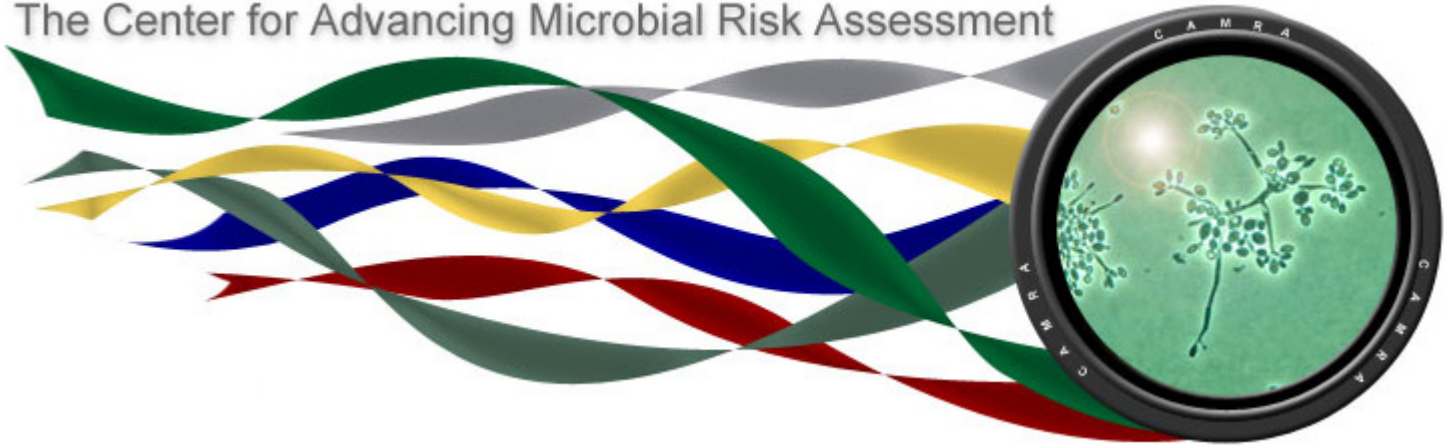


ANN vs Experimental Results





The Center for Advancing Microbial Risk Assessment



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