

US EPA ARCHIVE DOCUMENT



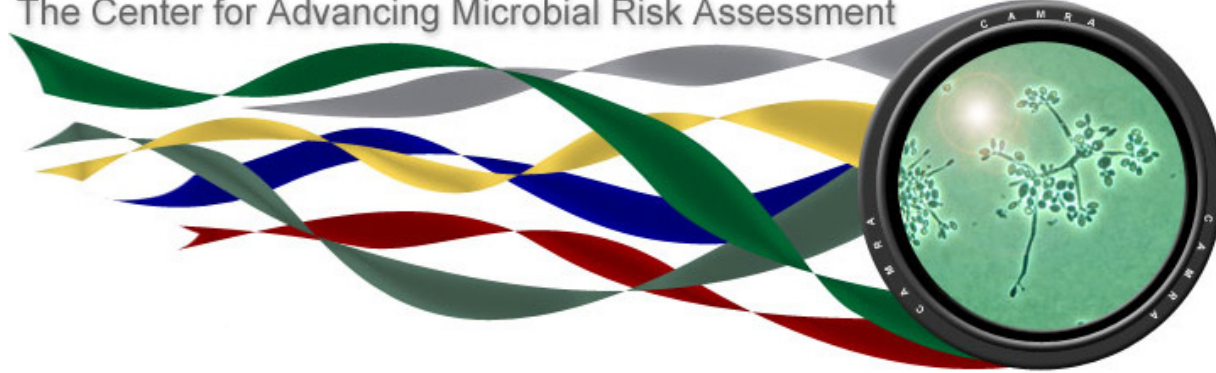
Improvement of Water Quality Modeling for Microbial Risk Assessment: Recent Advances

**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

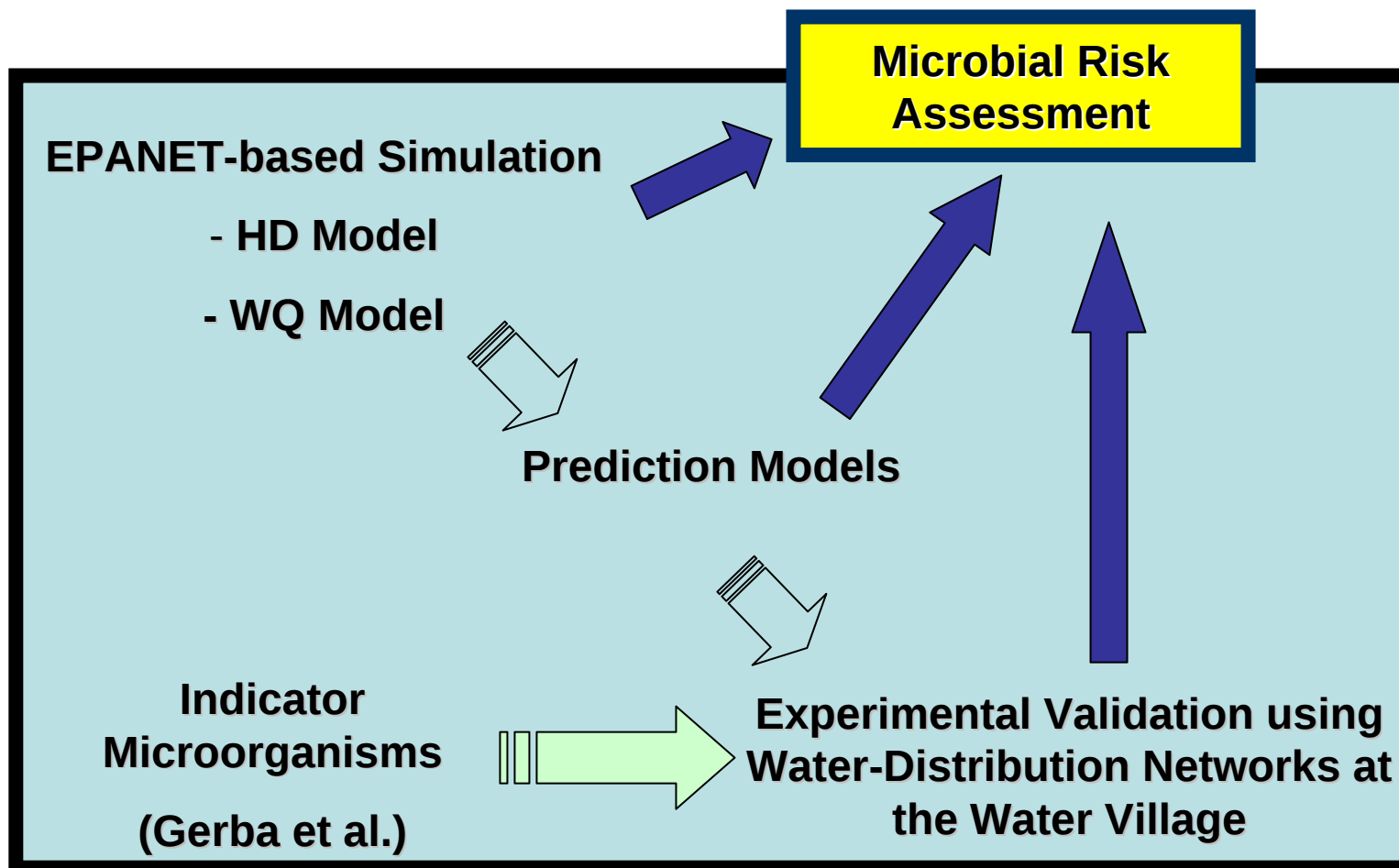
Carnegie Mellon University



The University of Arizona

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









Pilot Scale Experiments

Current Laboratories



Real-time
Water
Quality
Monitoring
Lab



Pumpin
Station



Main Building
Offices and
Computer Lab



50 m line

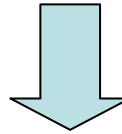
Water
Chemistry
Lab

Water
Treatment
Laboratory

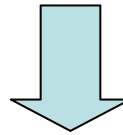
Trench

50 m line for dispersion experiments

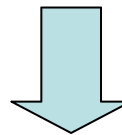
Step 1. Accurate Water Quality Modeling



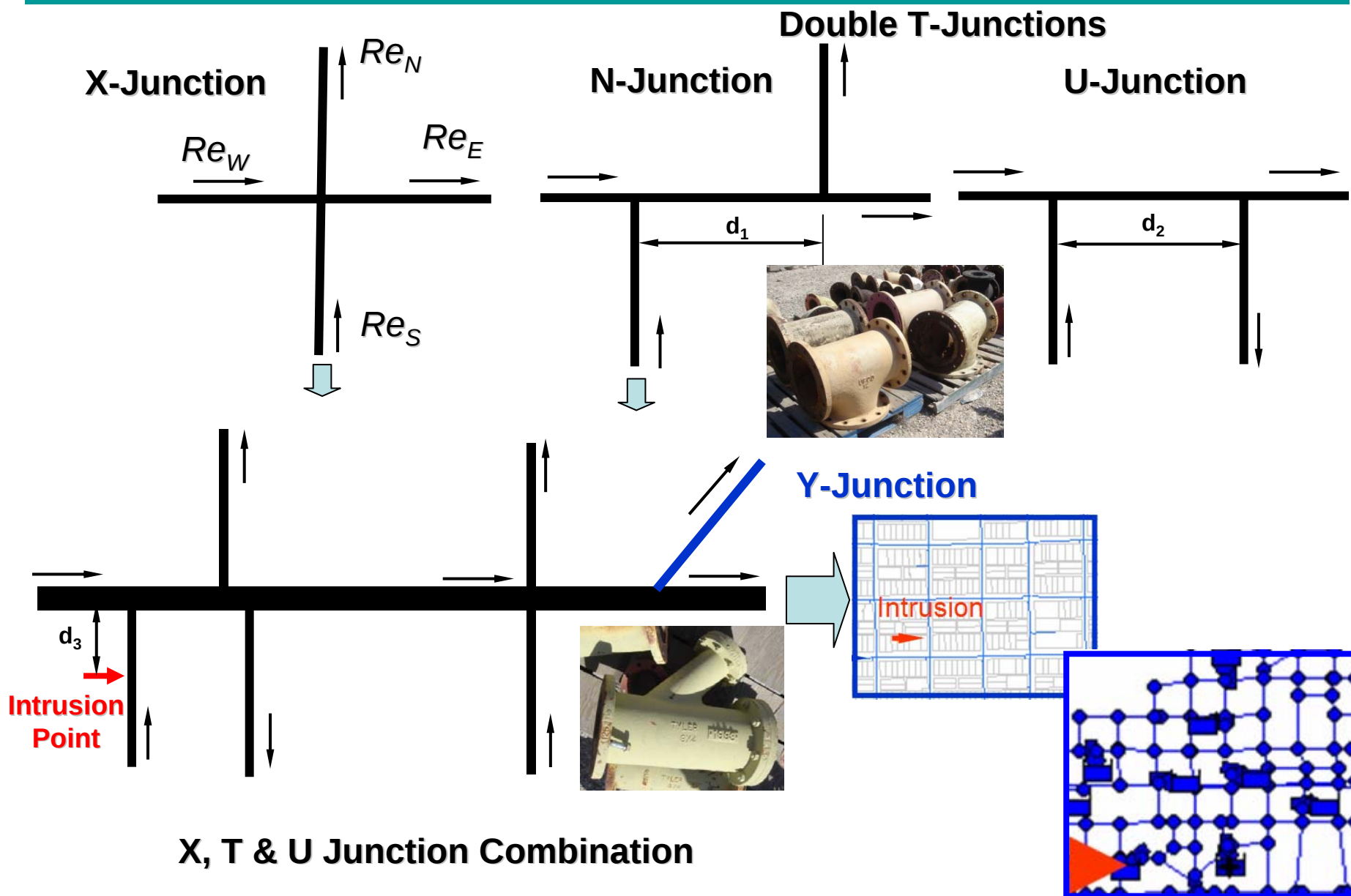
Step 2. Accurate Prediction of Dispersion of Microorganisms



Step 3. Use of Prediction Tools such as ANNs



Step 4. Accurate Quantitative Microbial Risk Assessment

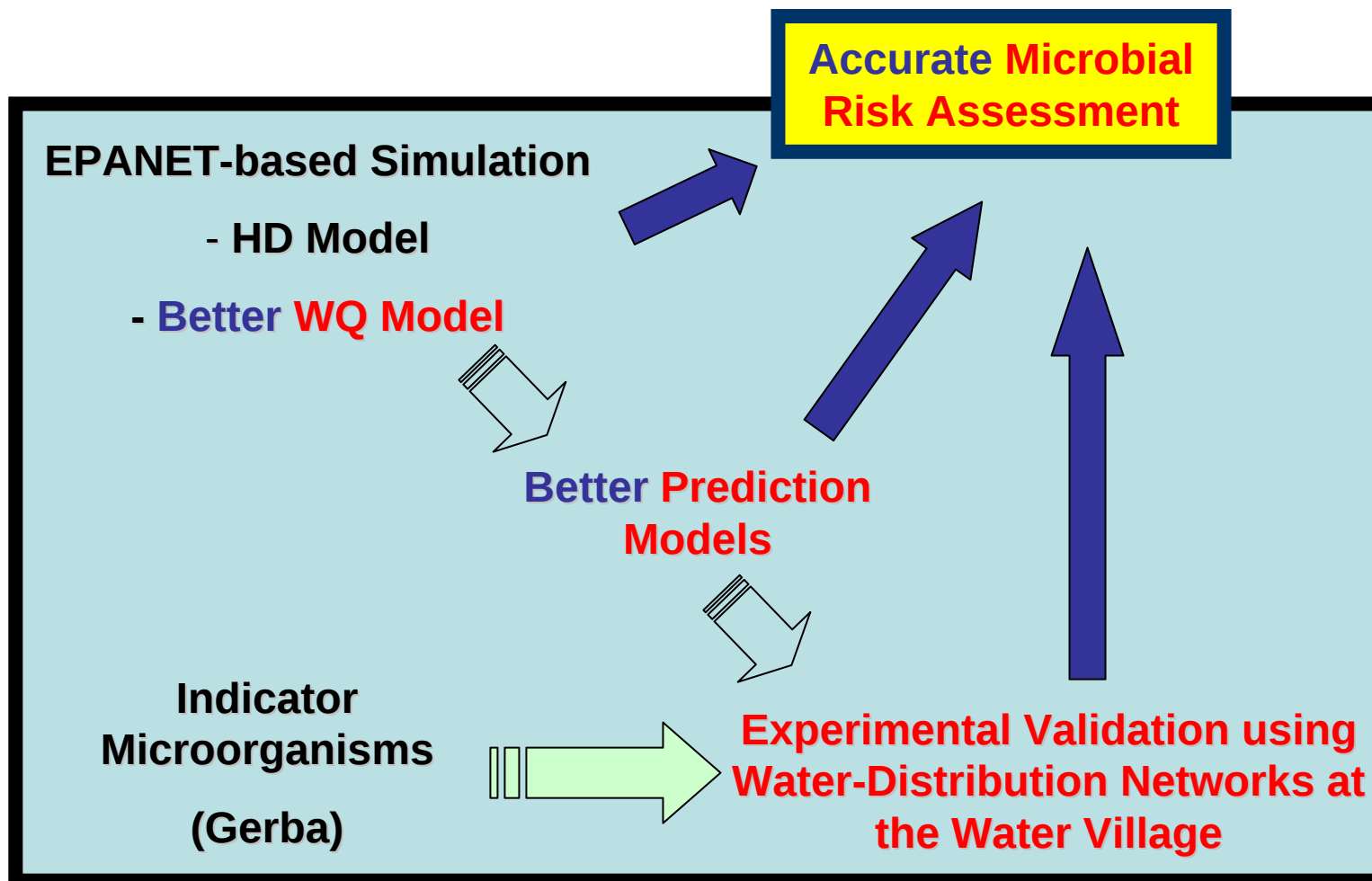


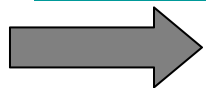
At present, prevailing network water quality models are based on two major simplifications.

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.

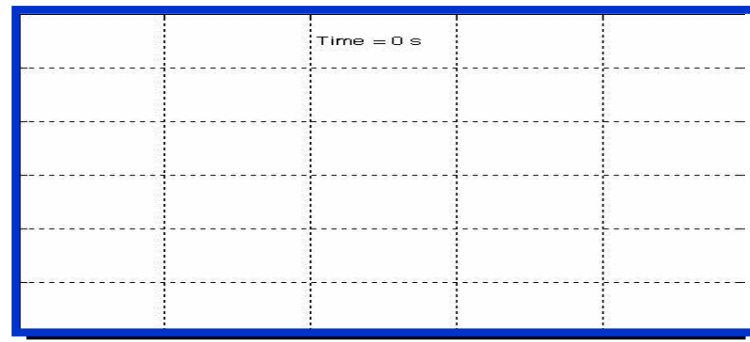
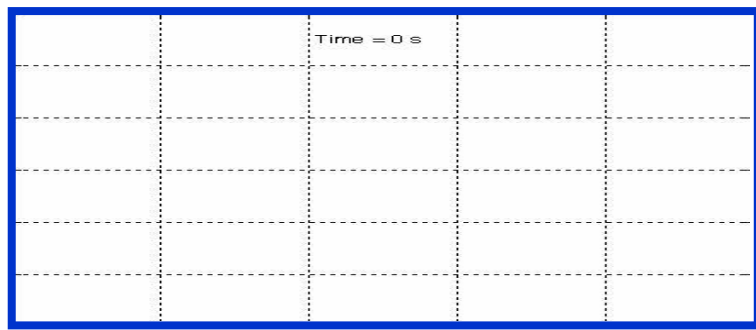
Recent investigations suggest that these assumptions **may NOT be valid** in real pipe networks.





$U = 0.635$ m/s $Re = 8,000$

$U = 1.27$ m/s $Re = 16,000$



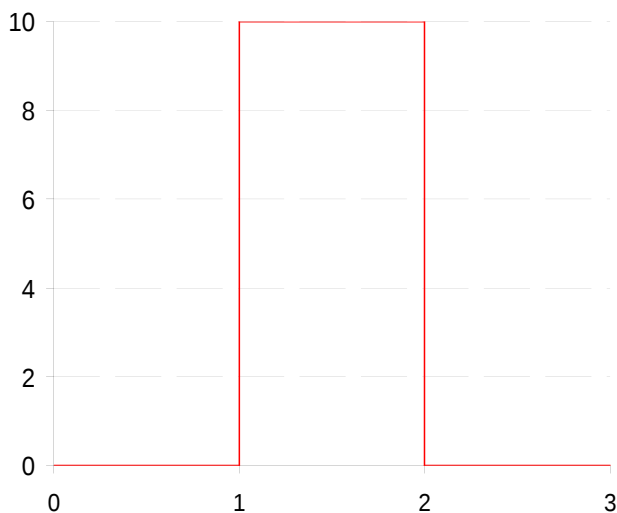
— $Re = 8,000$, $T_{max} = 6.5$ s

$Re = 8,000$, $T_{max} = 11.6$ s

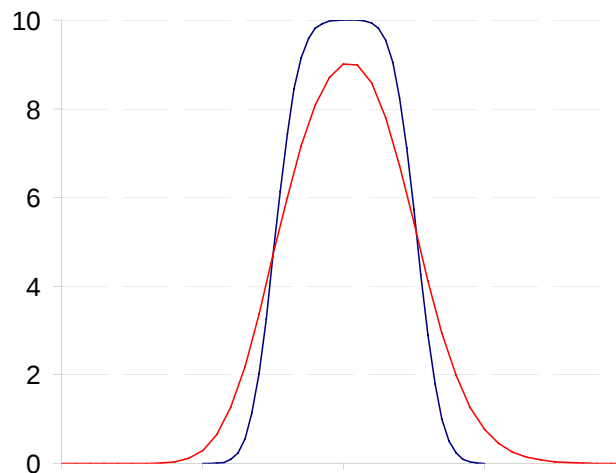
— $Re = 16,000$, $T_{max} = 4.05$ s

$Re = 16,000$, $T_{max} = 6.55$ s

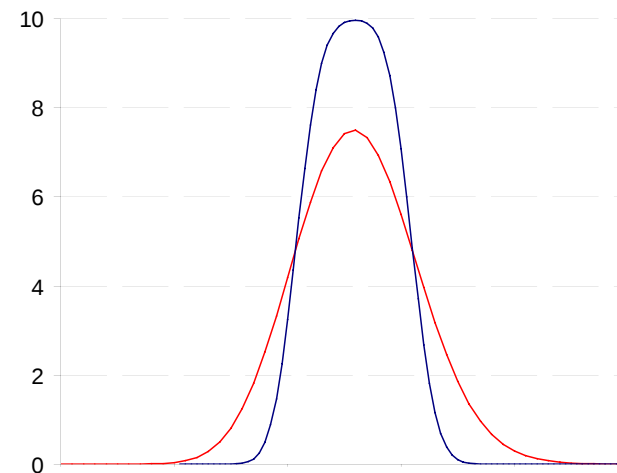
$X = 0$



$X = 250 D$

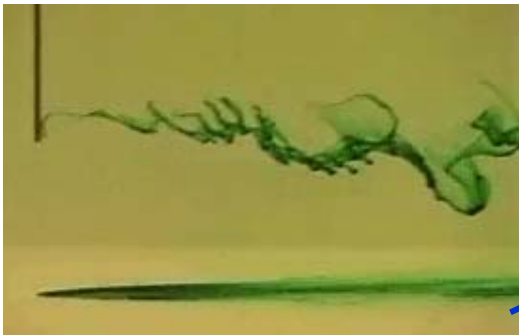
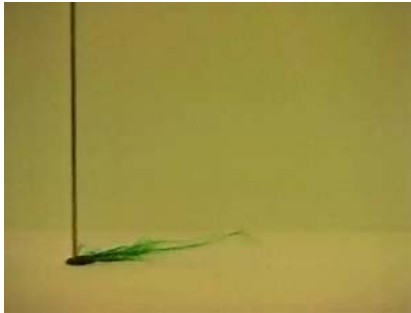


$X = 500 D$

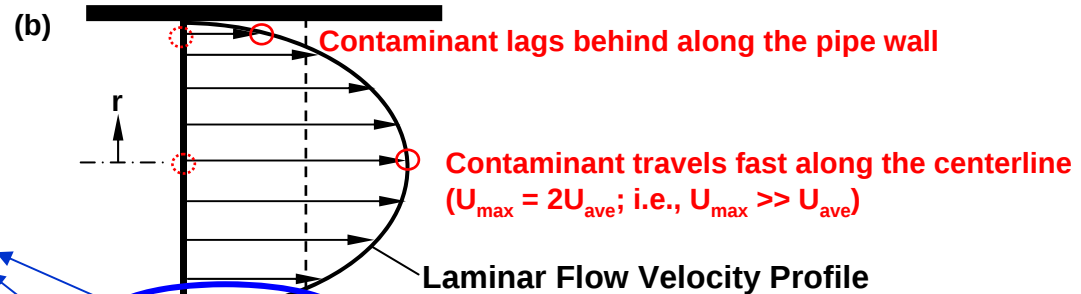
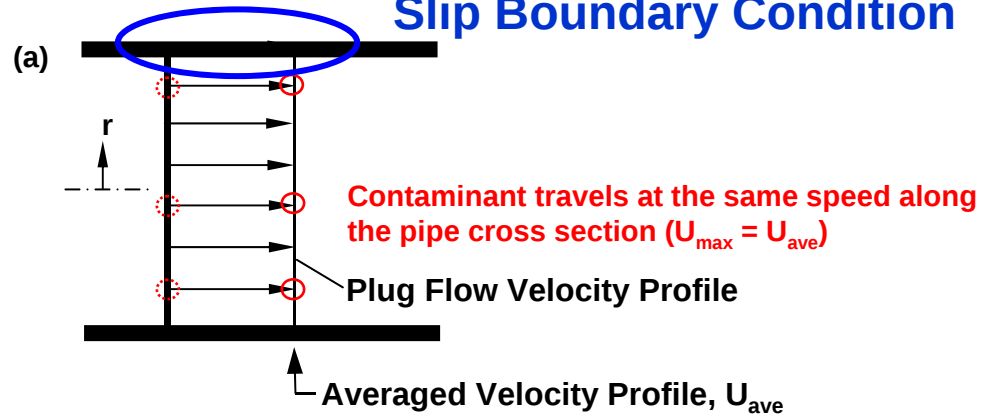


Why Axial Dispersion?

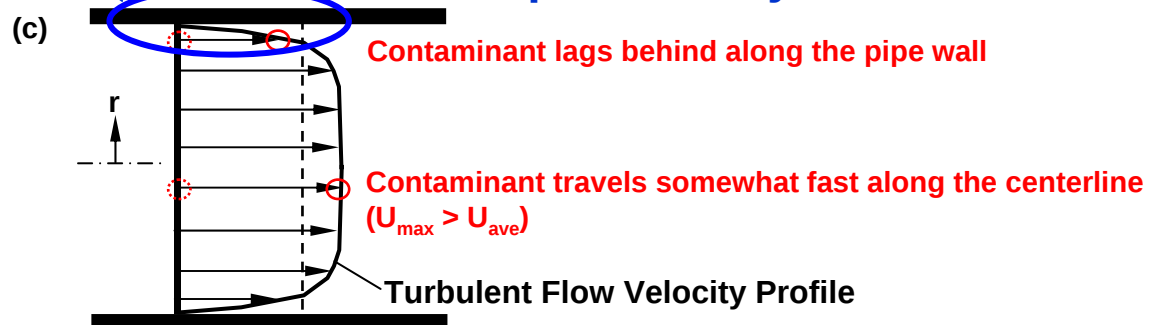
Typical tracer dispersion along the pipe wall

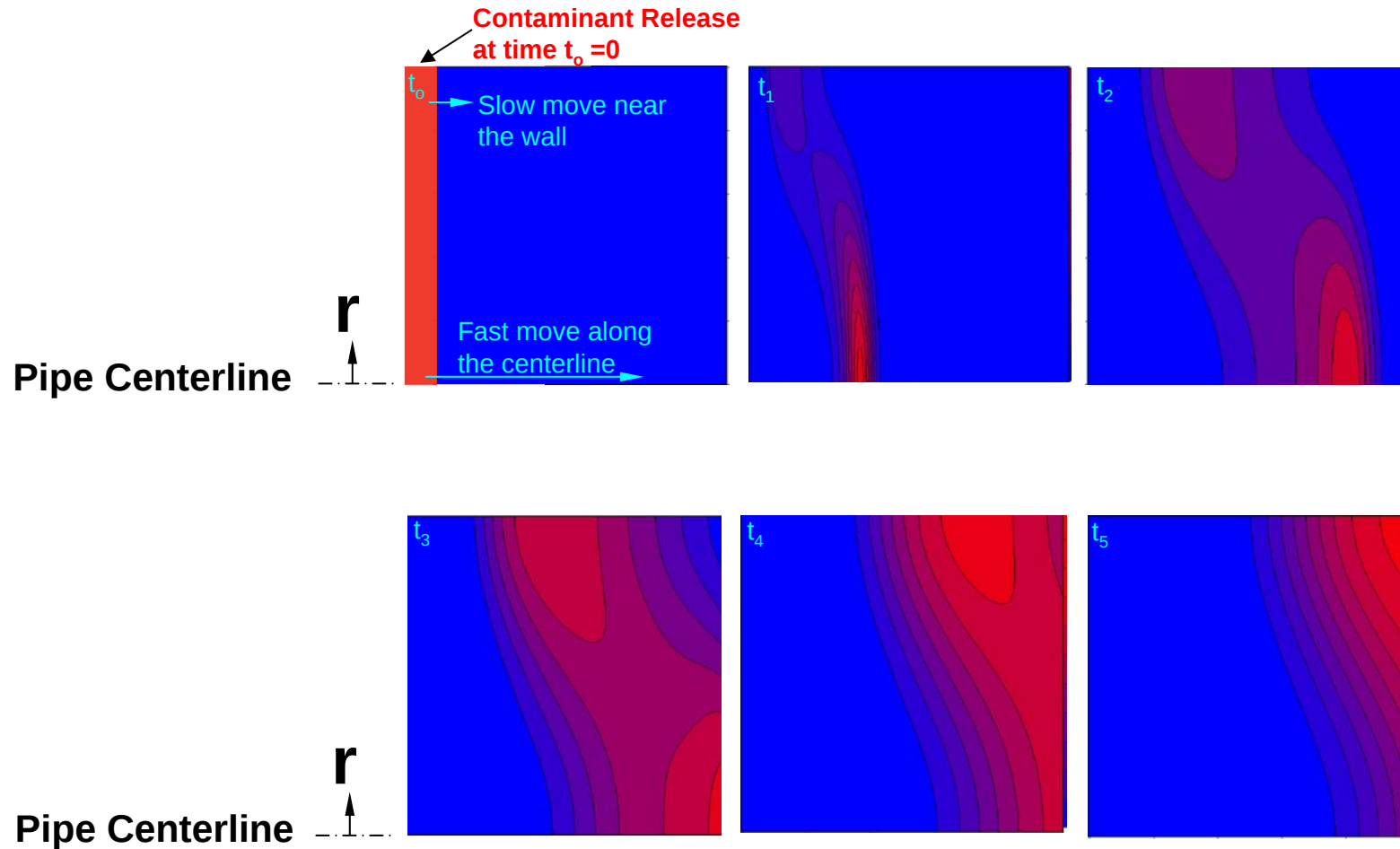


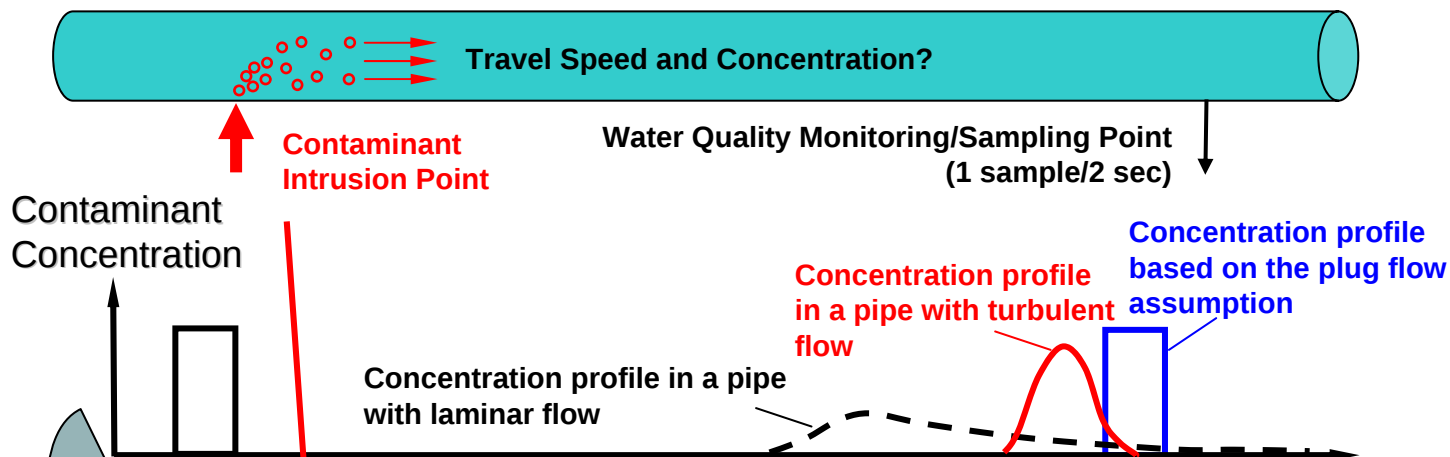
Slip Boundary Condition



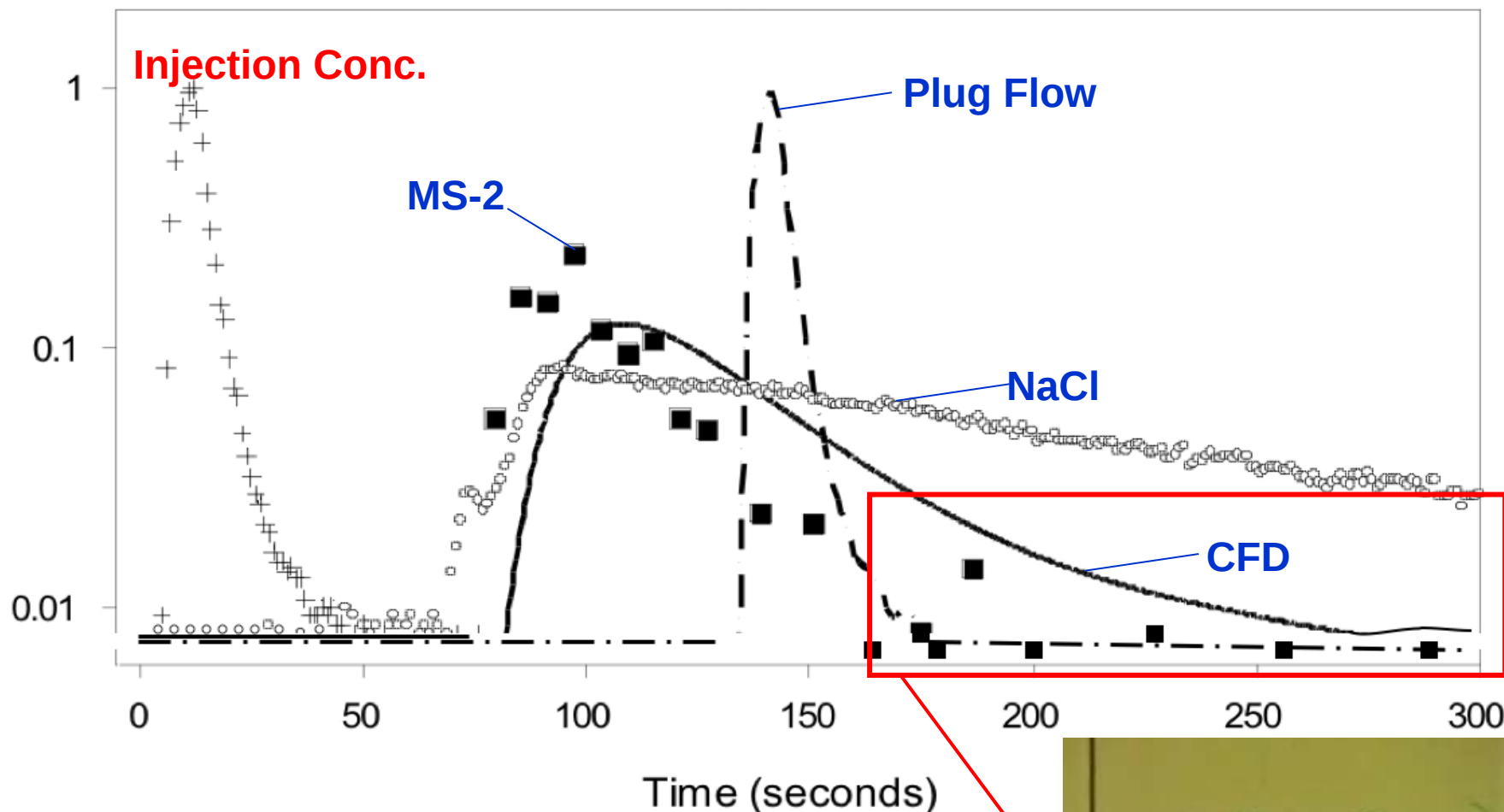
No-Slip Boundary Condition



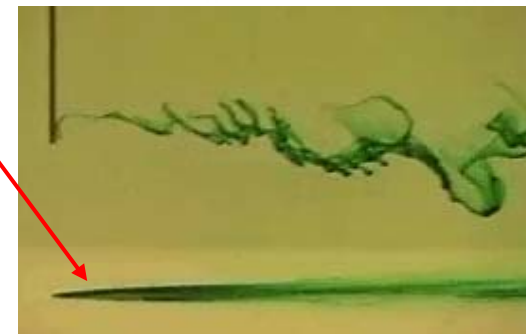


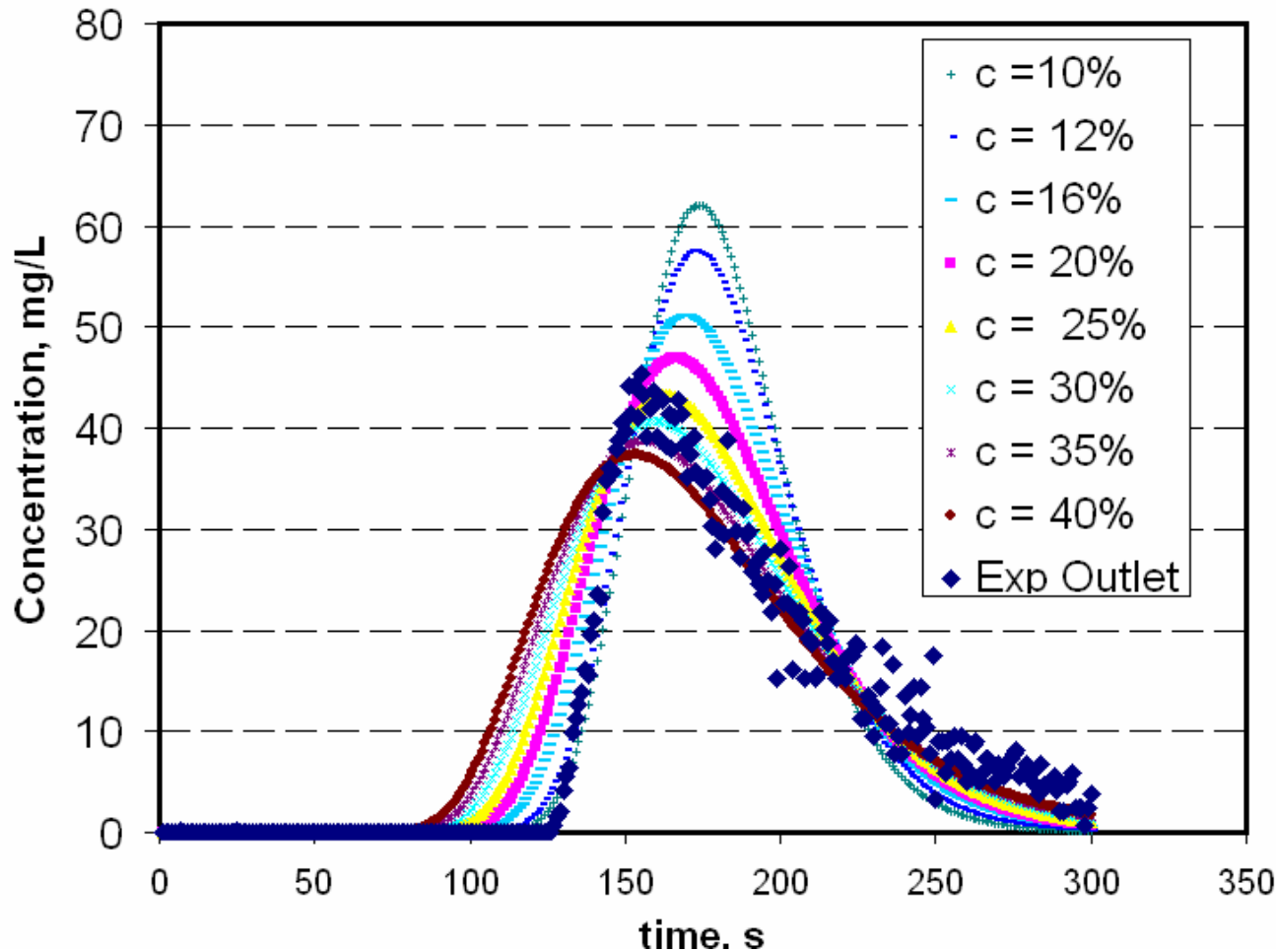


Injection of tracers Laminar flow

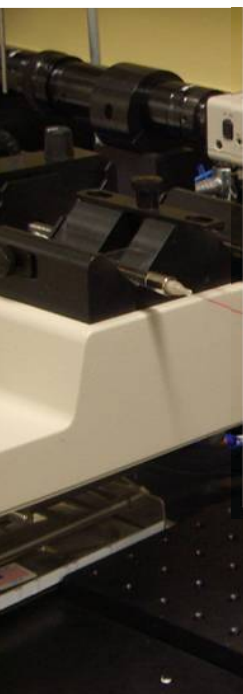
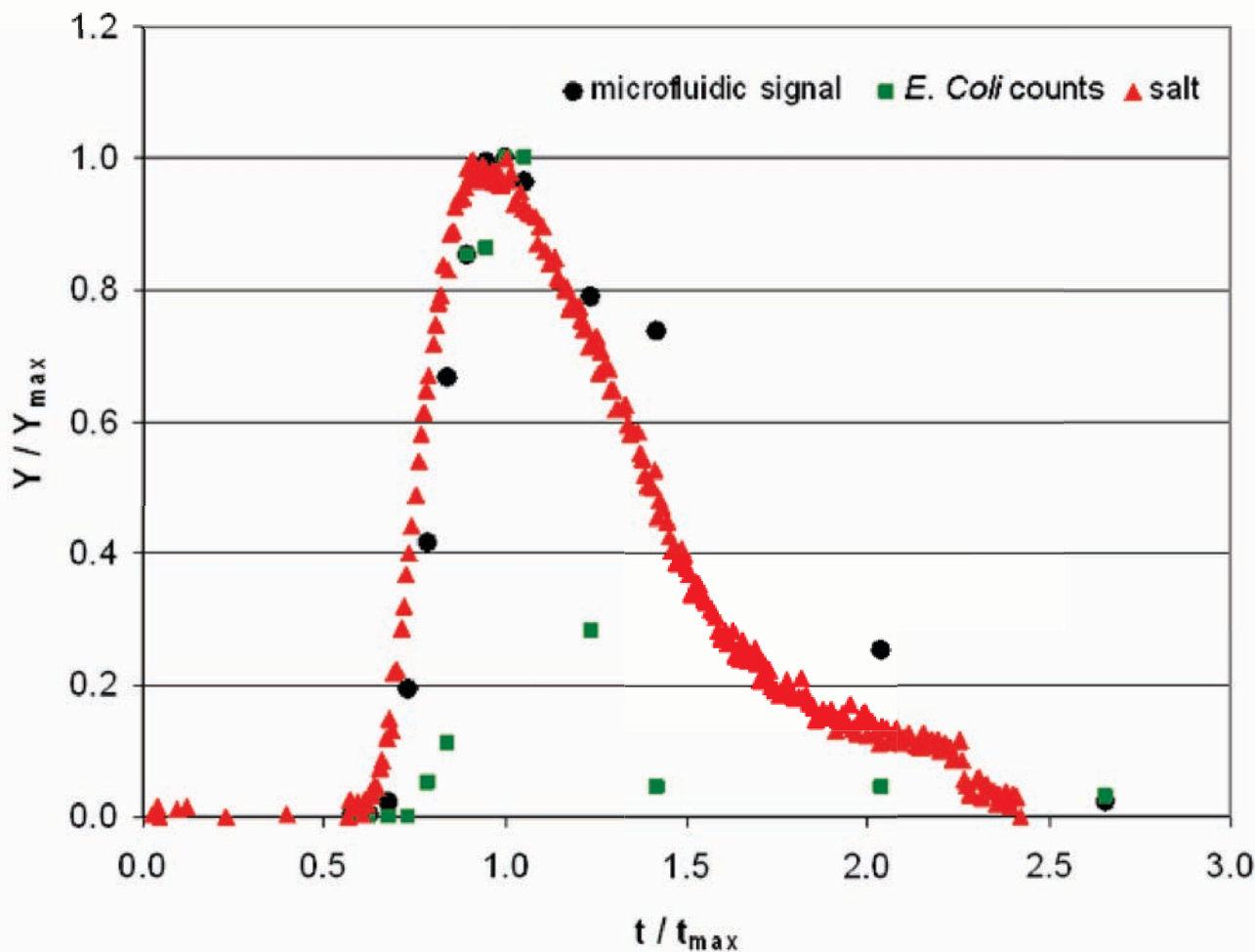


**Axial dispersion & tailing of microorganisms:
Detection limits, sampling frequency, sensor locations**





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



Courtesy of Professor Yoon
The Univ. of Arizona

Pumping
Station



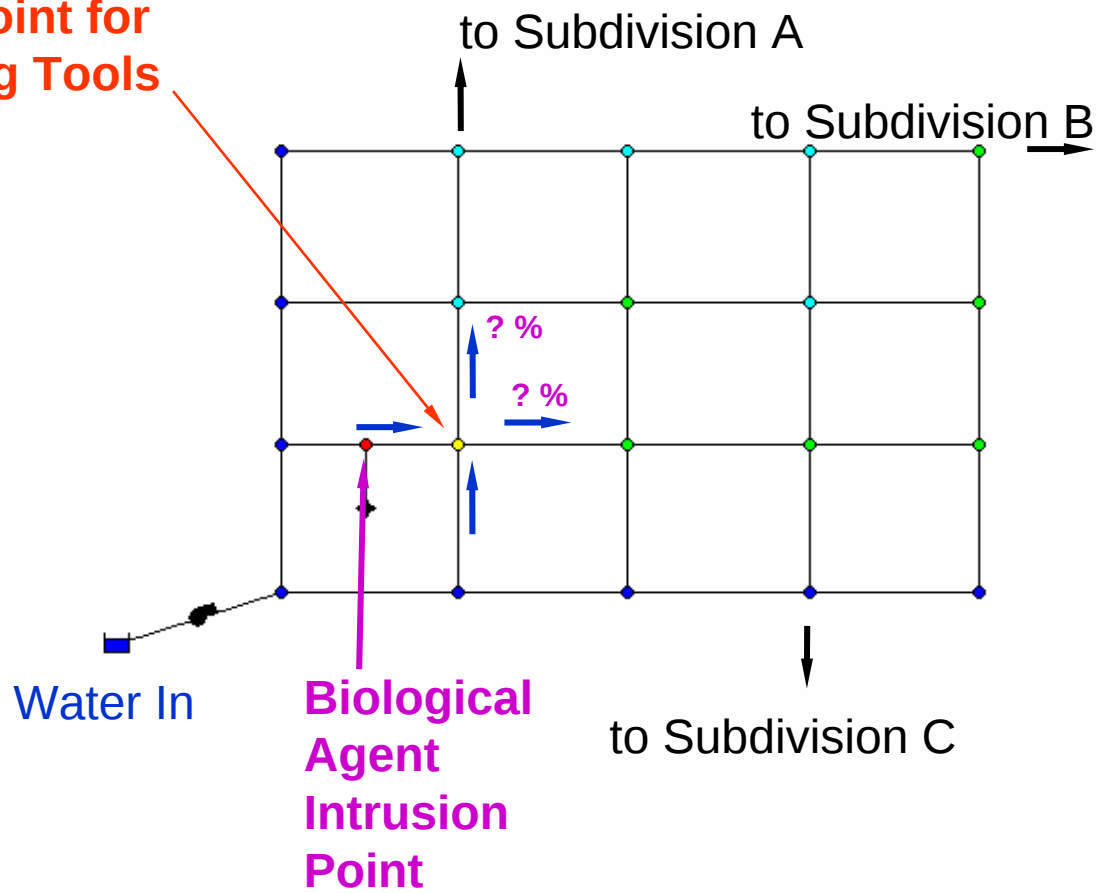
Main
Building

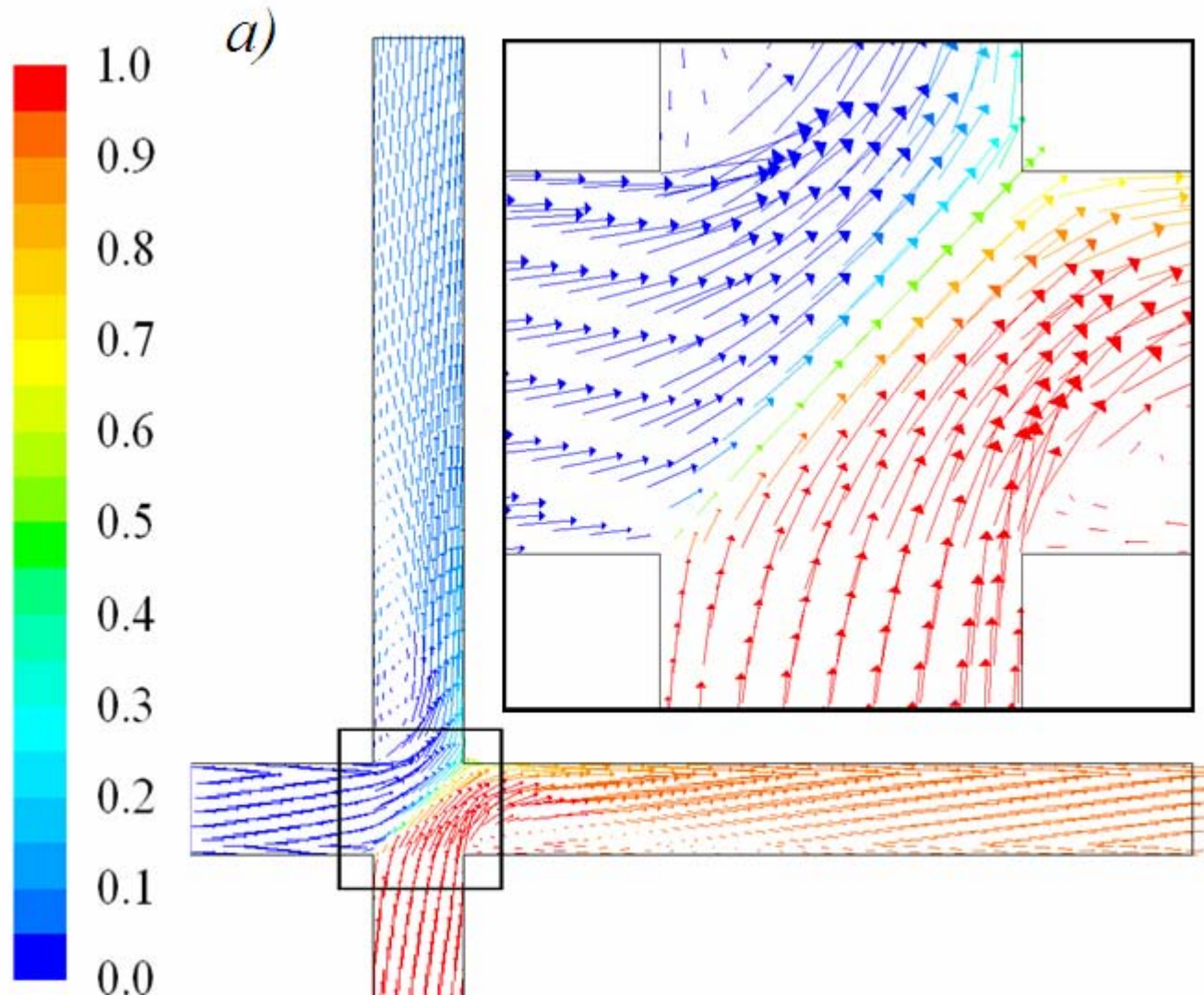


Trench

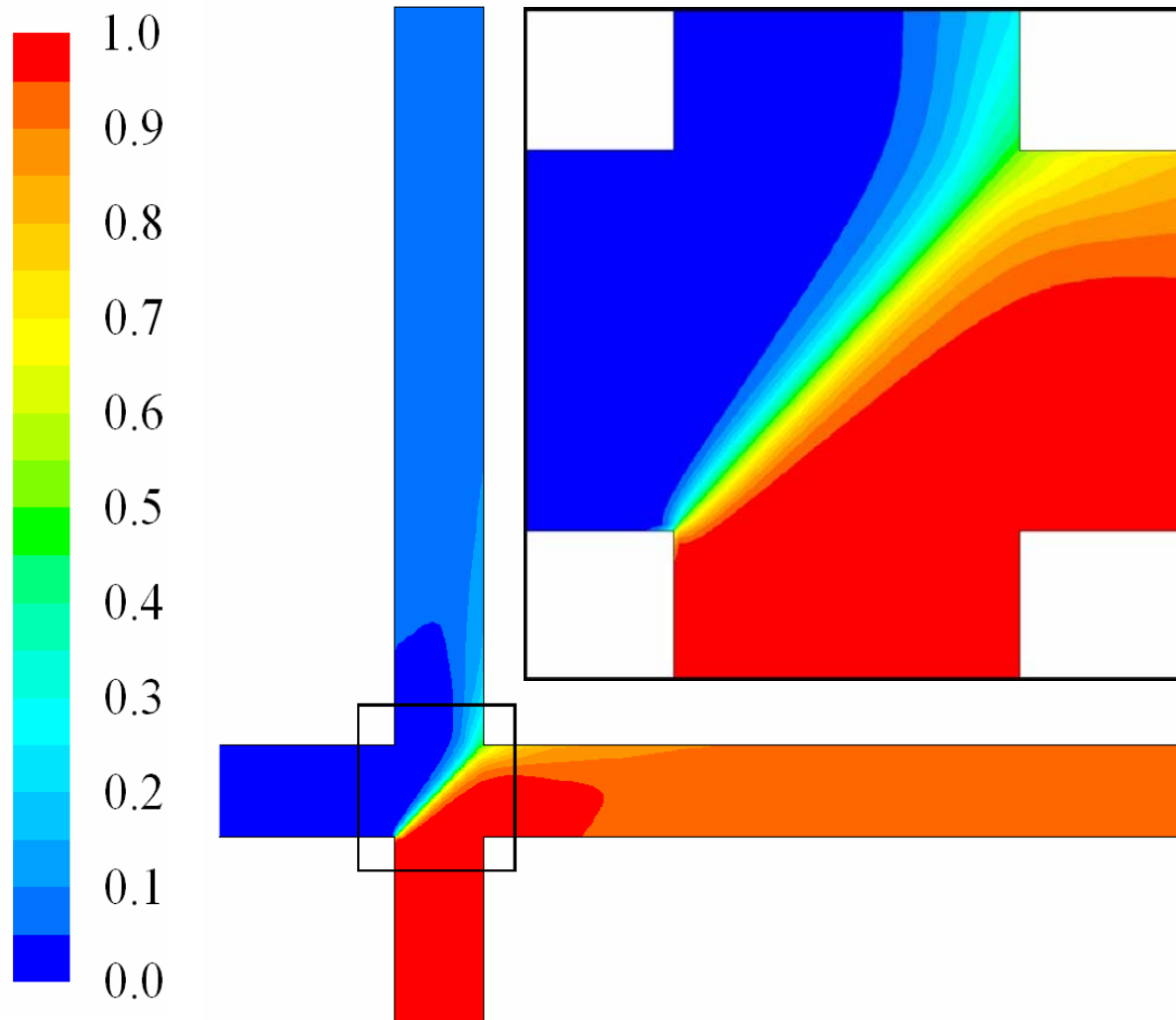
50 m line for dispersion experiments

**Perfect Mixing
Assumed at the
Cross Joint for
Modeling Tools**

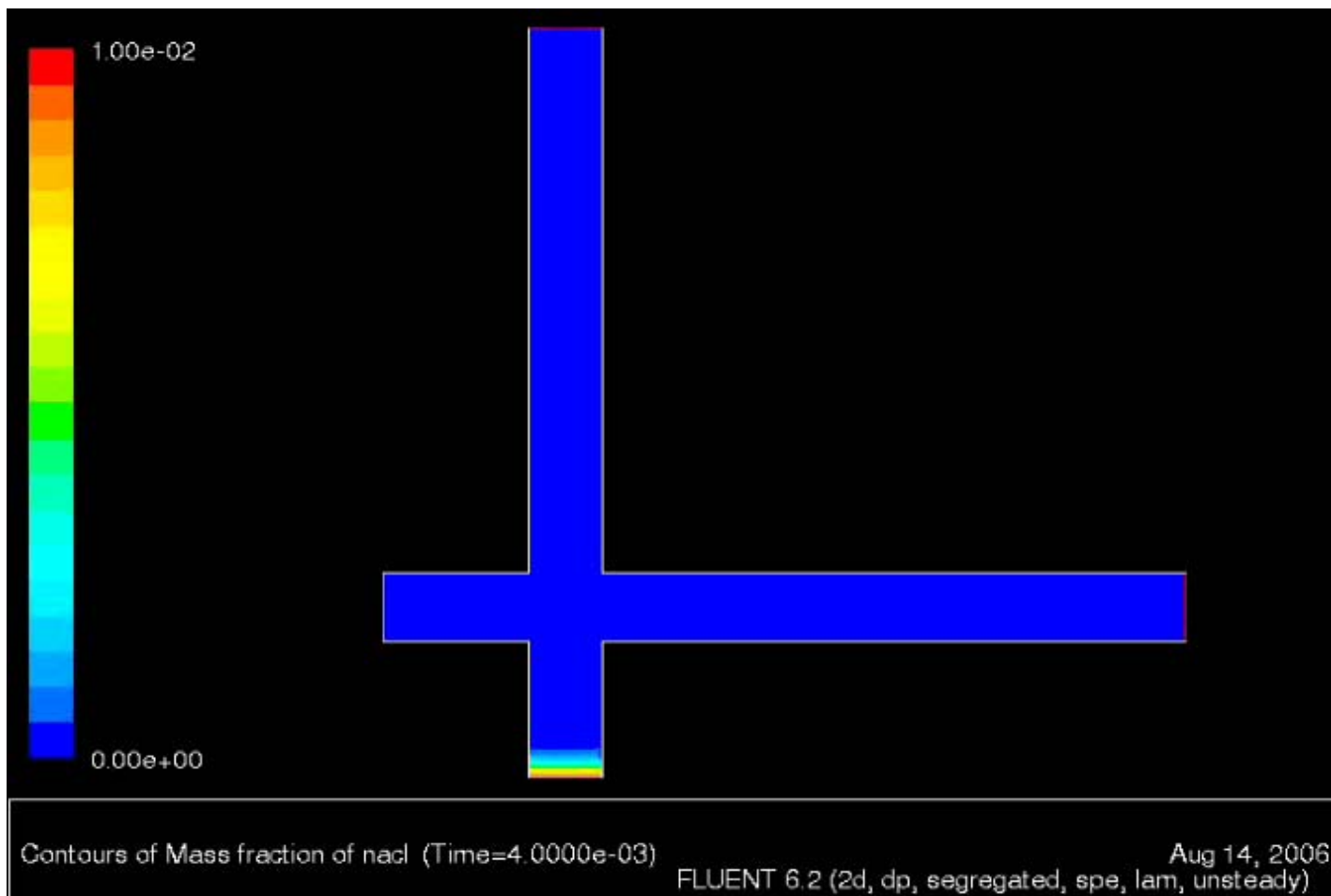




**Velocity vectors at a cross junction, when $Re_S = Re_W = Re_E = Re_N = 44,000$
($Re_{SW} = 1$, $Re_{E/N} = 1$), and $Sc_t = 0.1875$**

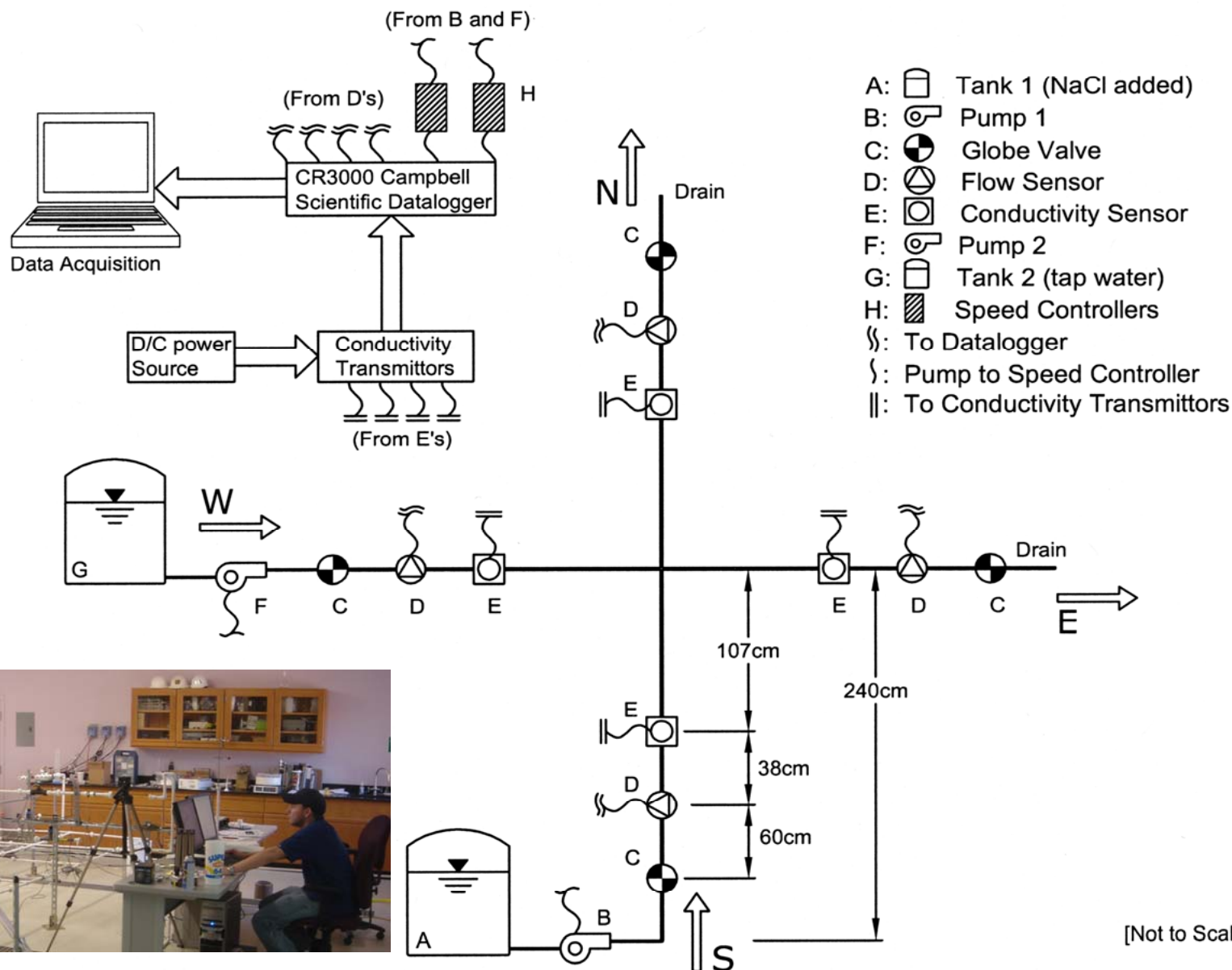


Dimensionless NaCl Concentration contours at a cross junction, when $Re_S = Re_W = Re_E = Re_N = 44,000$ ($Re_S/W = 1$, $Re_E/N=1$), and $Sct = 0.1875$



Simulation Video Clip

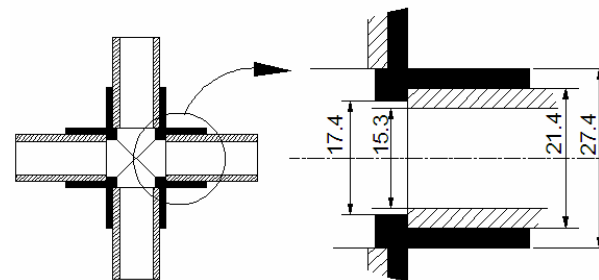
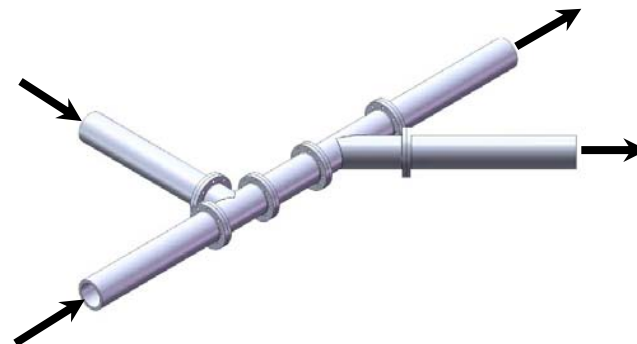
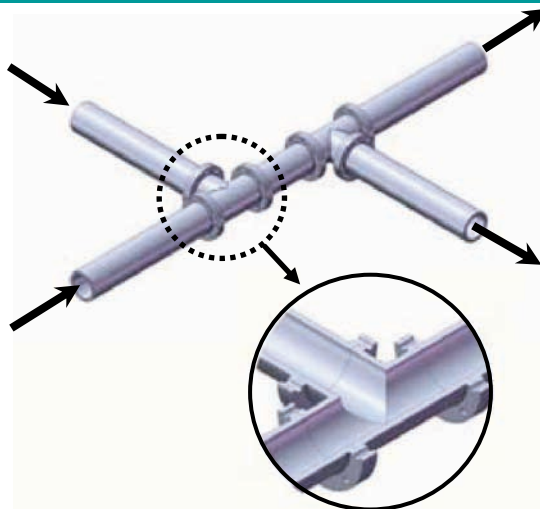
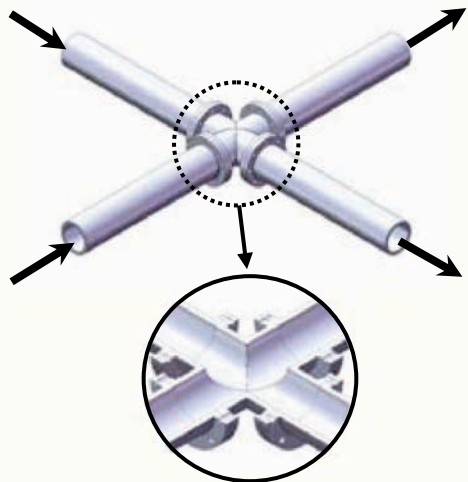
A schematic of the experimental setup

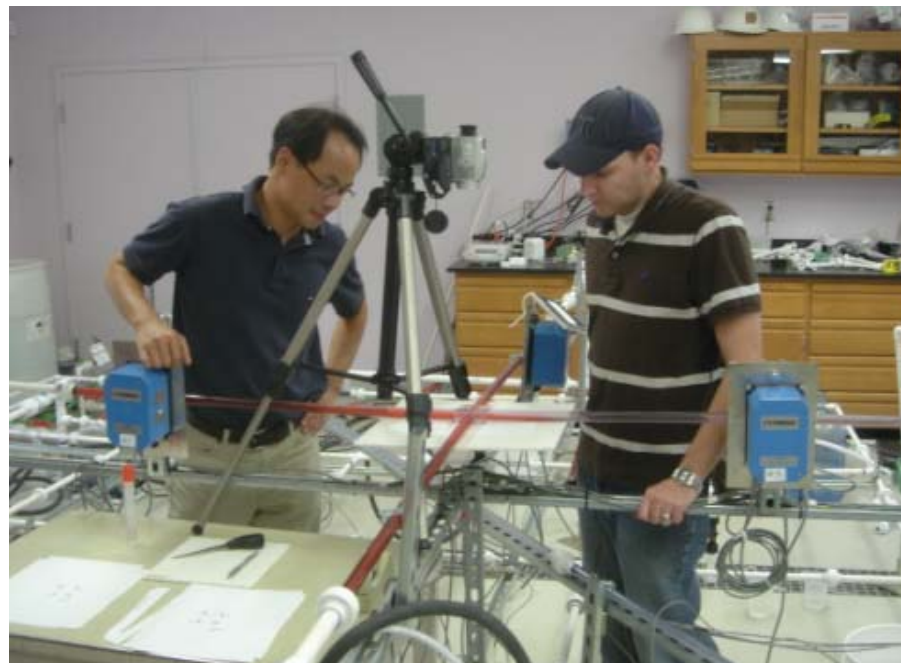
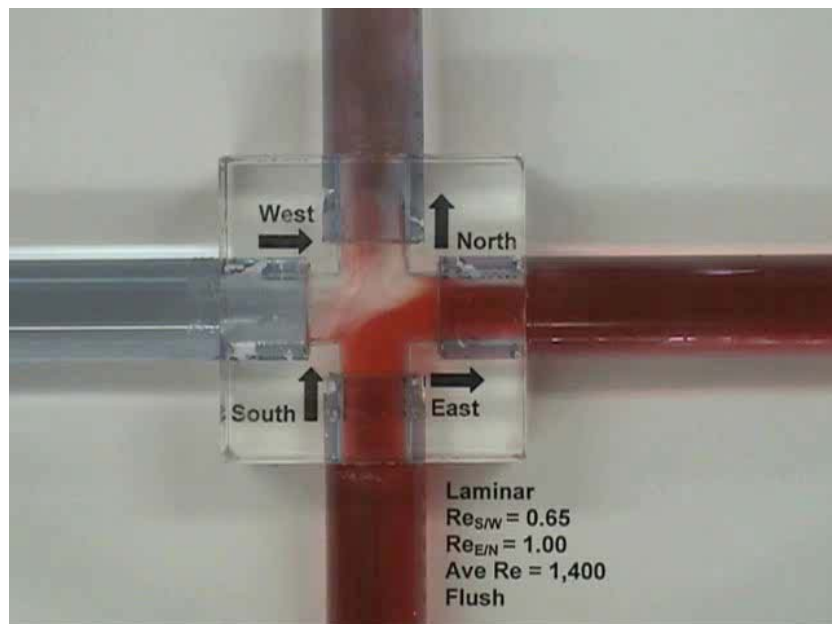


[Not to Scale]



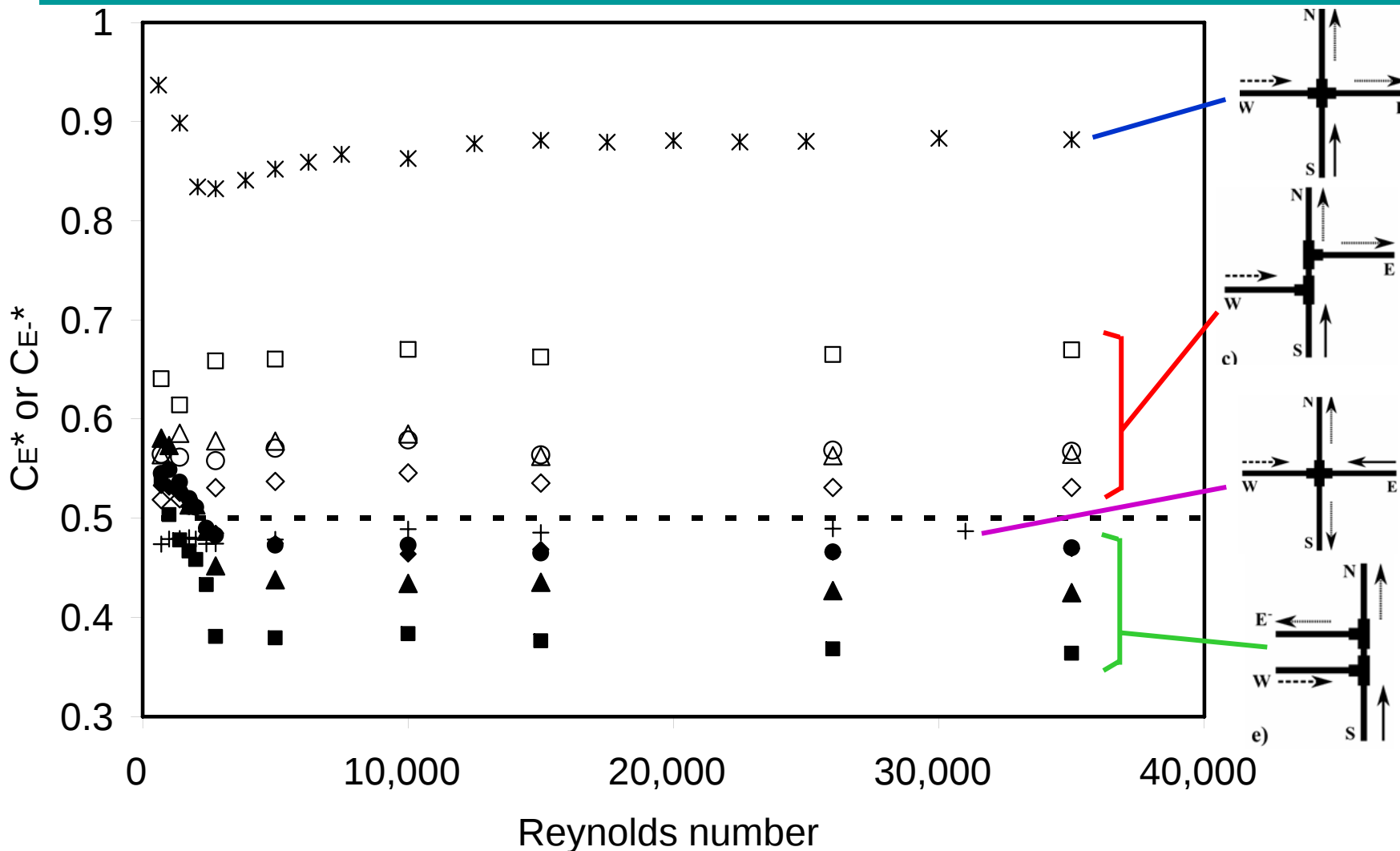
Representative Junction Types





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

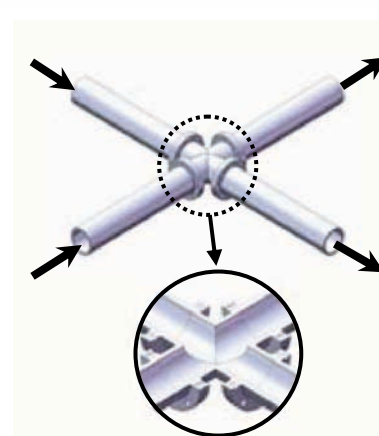
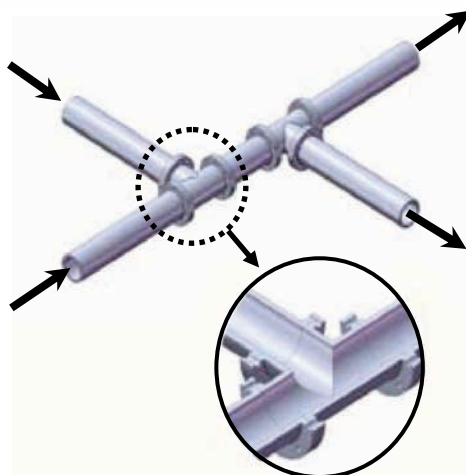
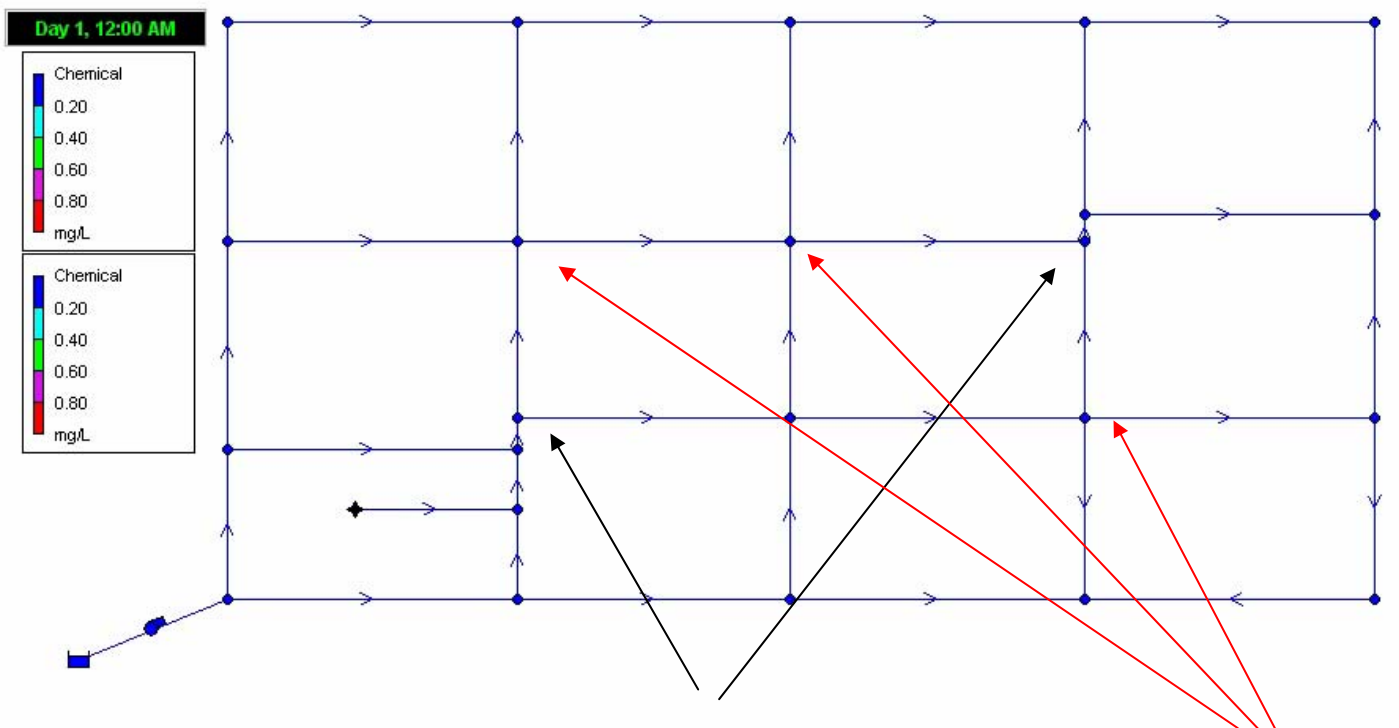
▲ D=5.0, U1

● D=7.5, U1

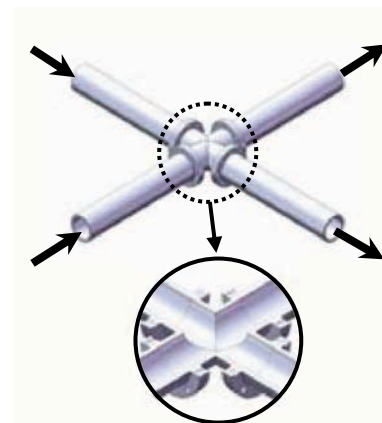
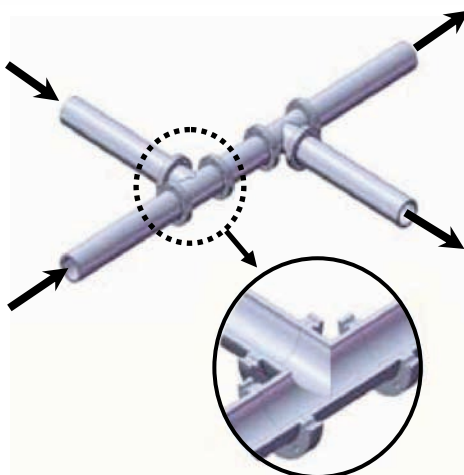
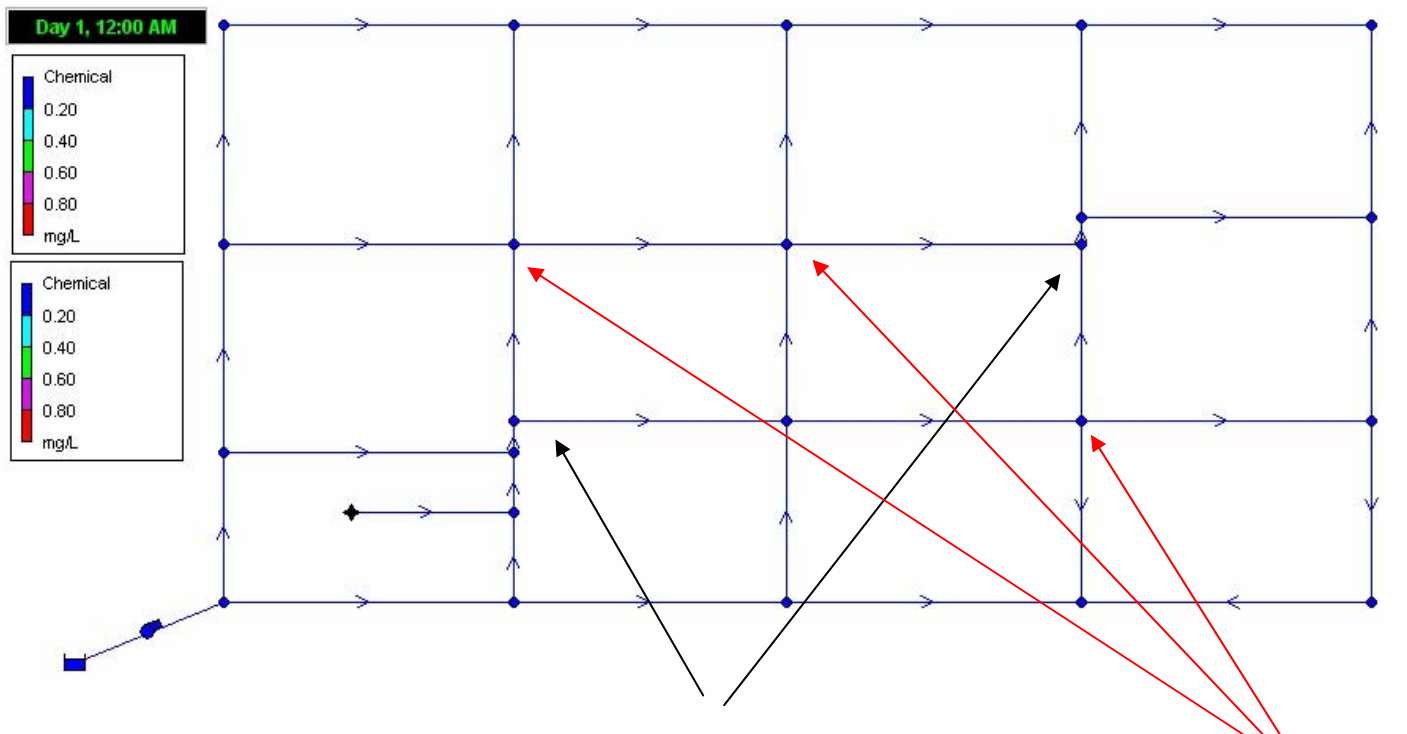
◆ D=10, U1

--- Complete Mixing

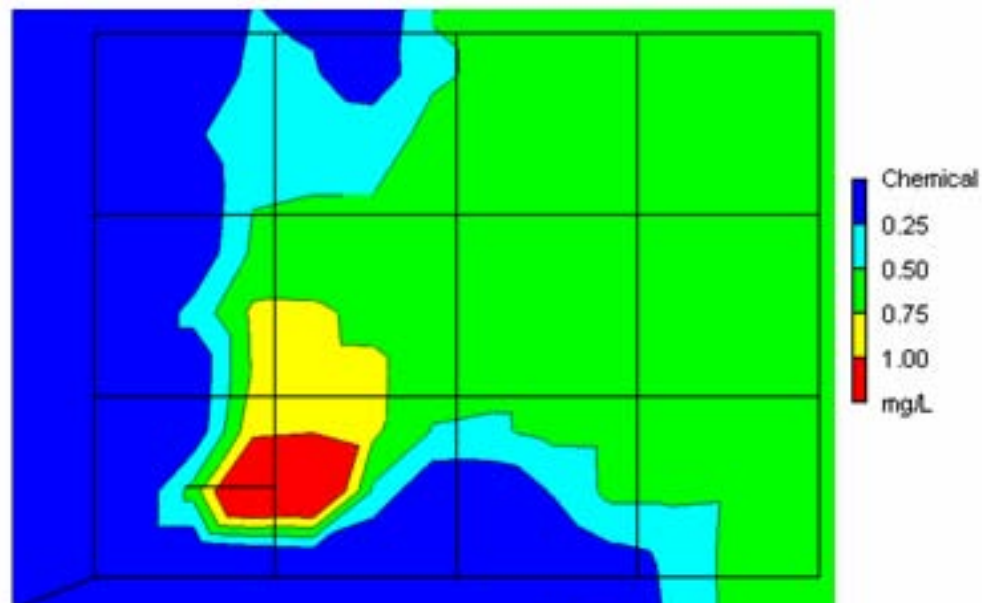
Development of **AZRED** (with Perfect Mixing Model)



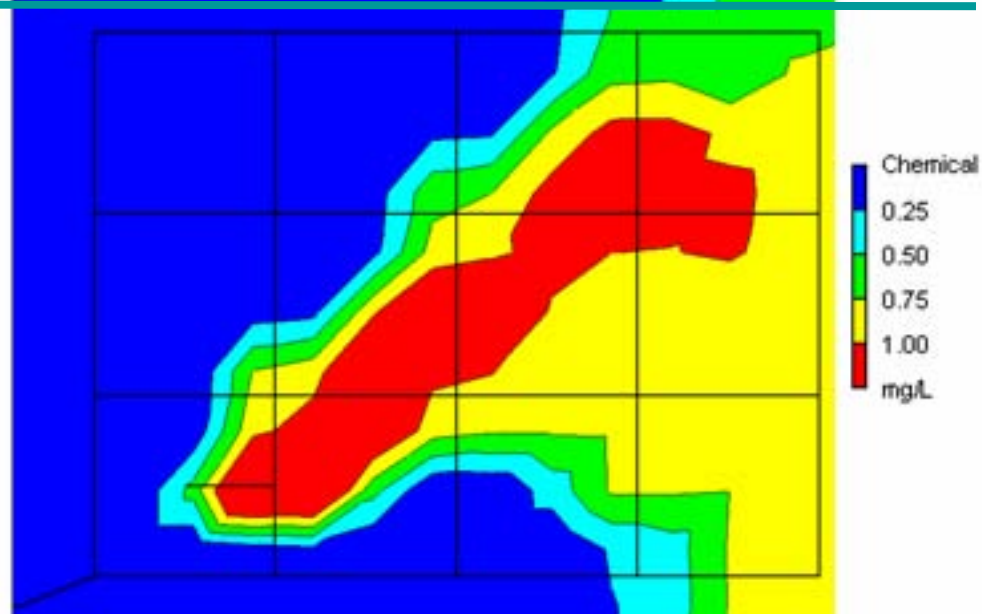
Development of **AZRED** (Improved Water Quality Model)



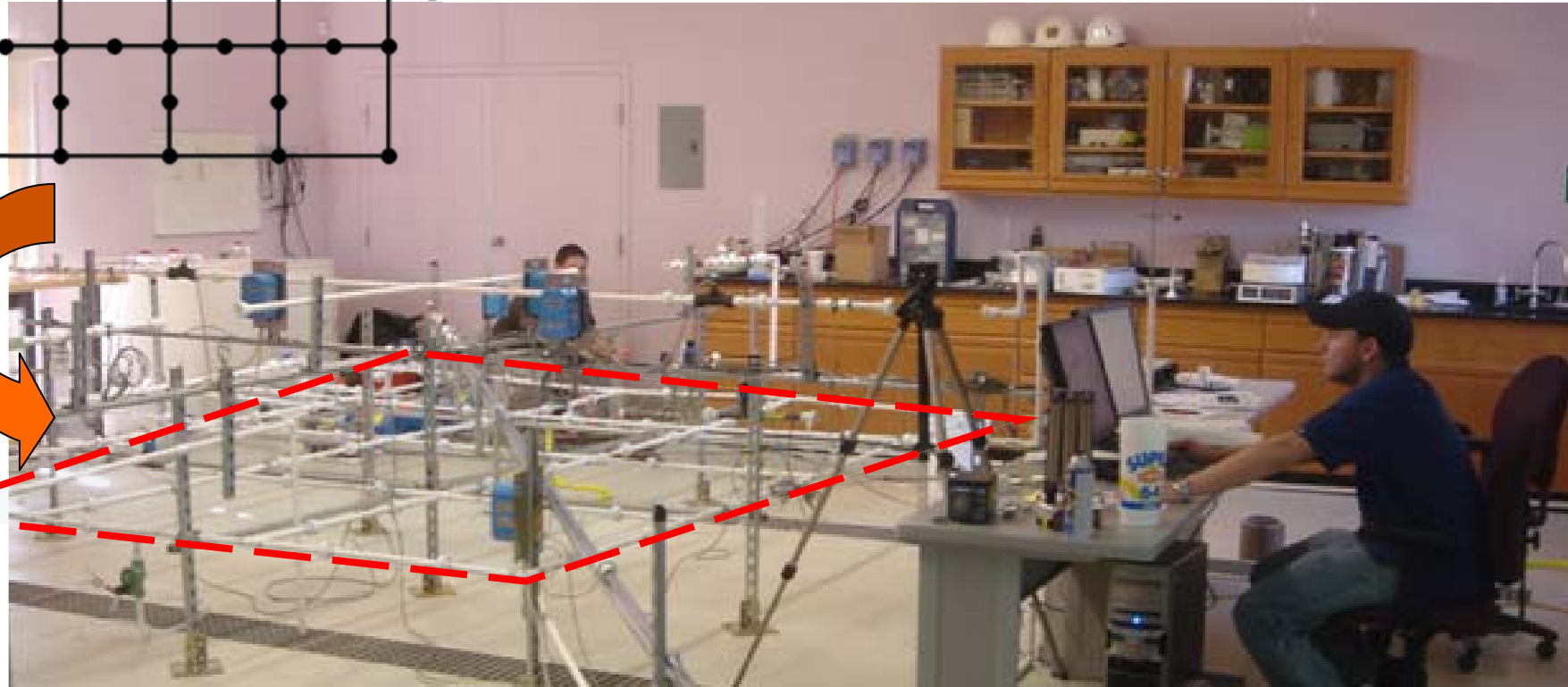
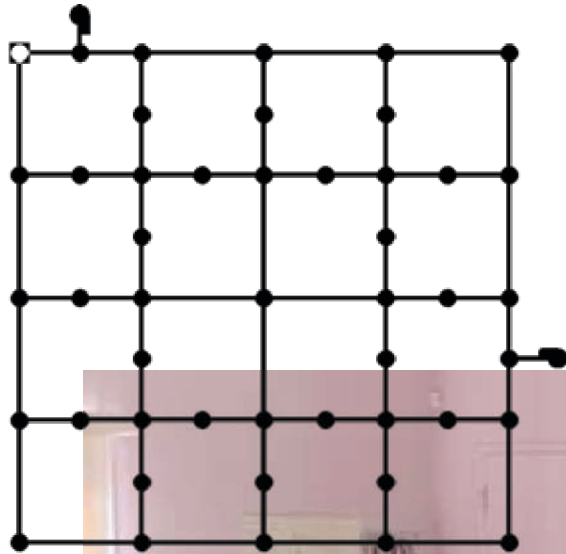
**Existing Model
based on
Perfect Mixing**

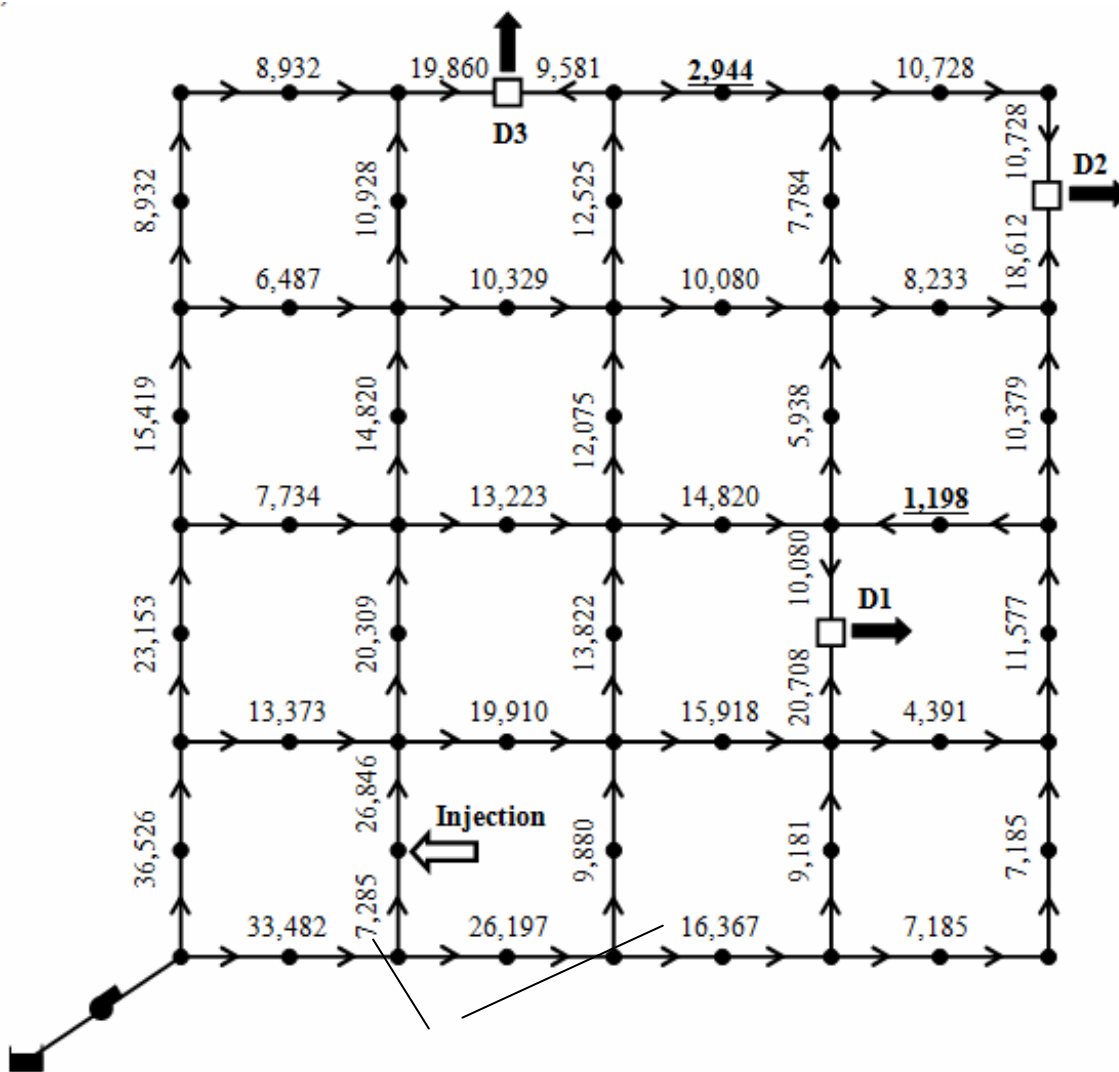


AZRED



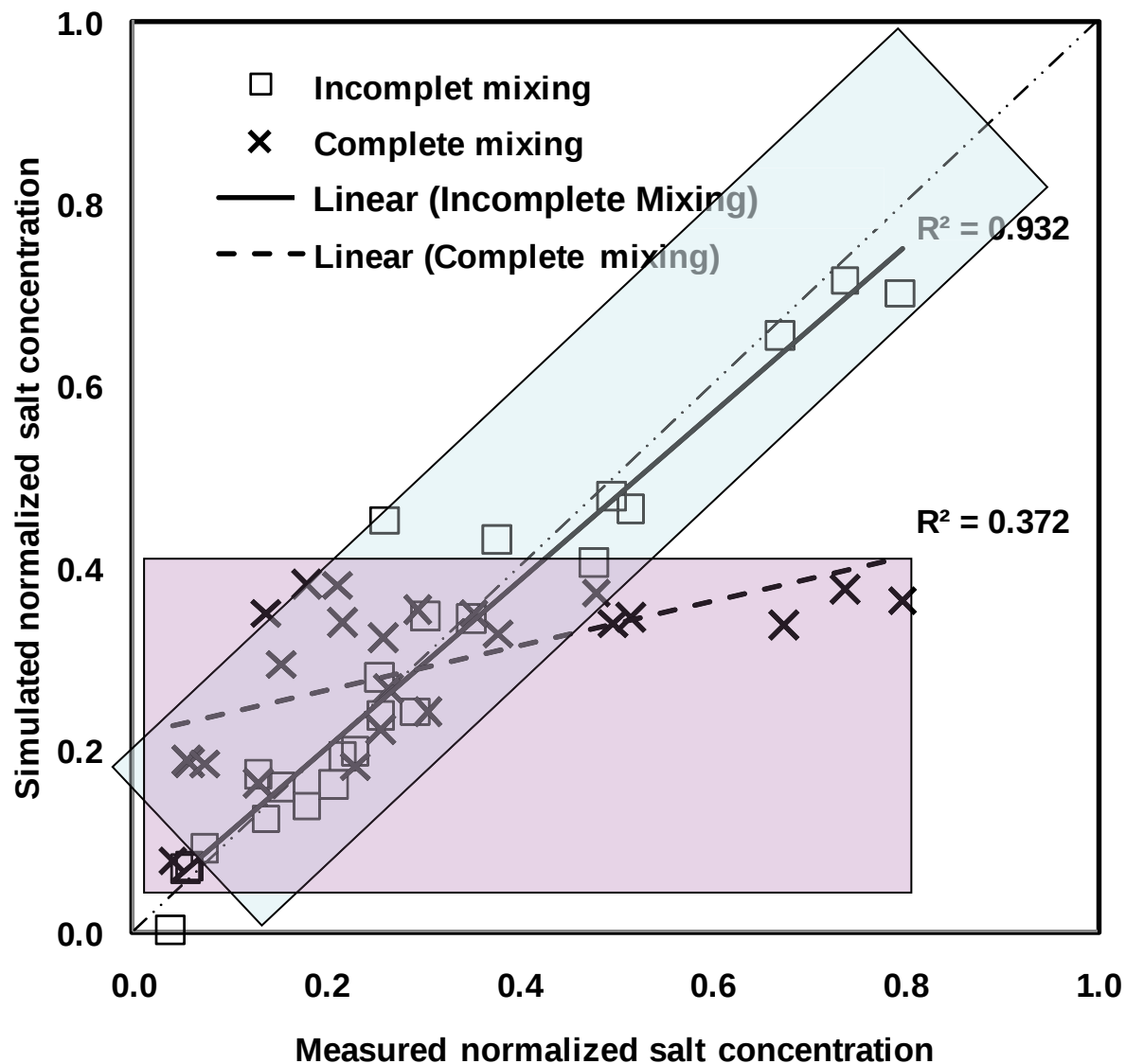
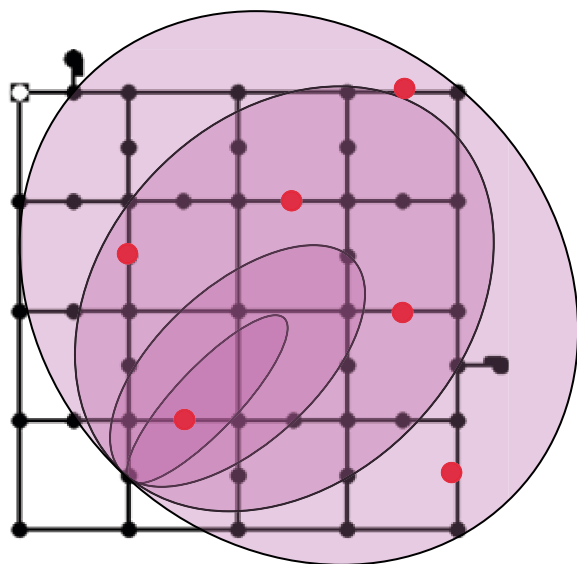
5x5 Network Establishment



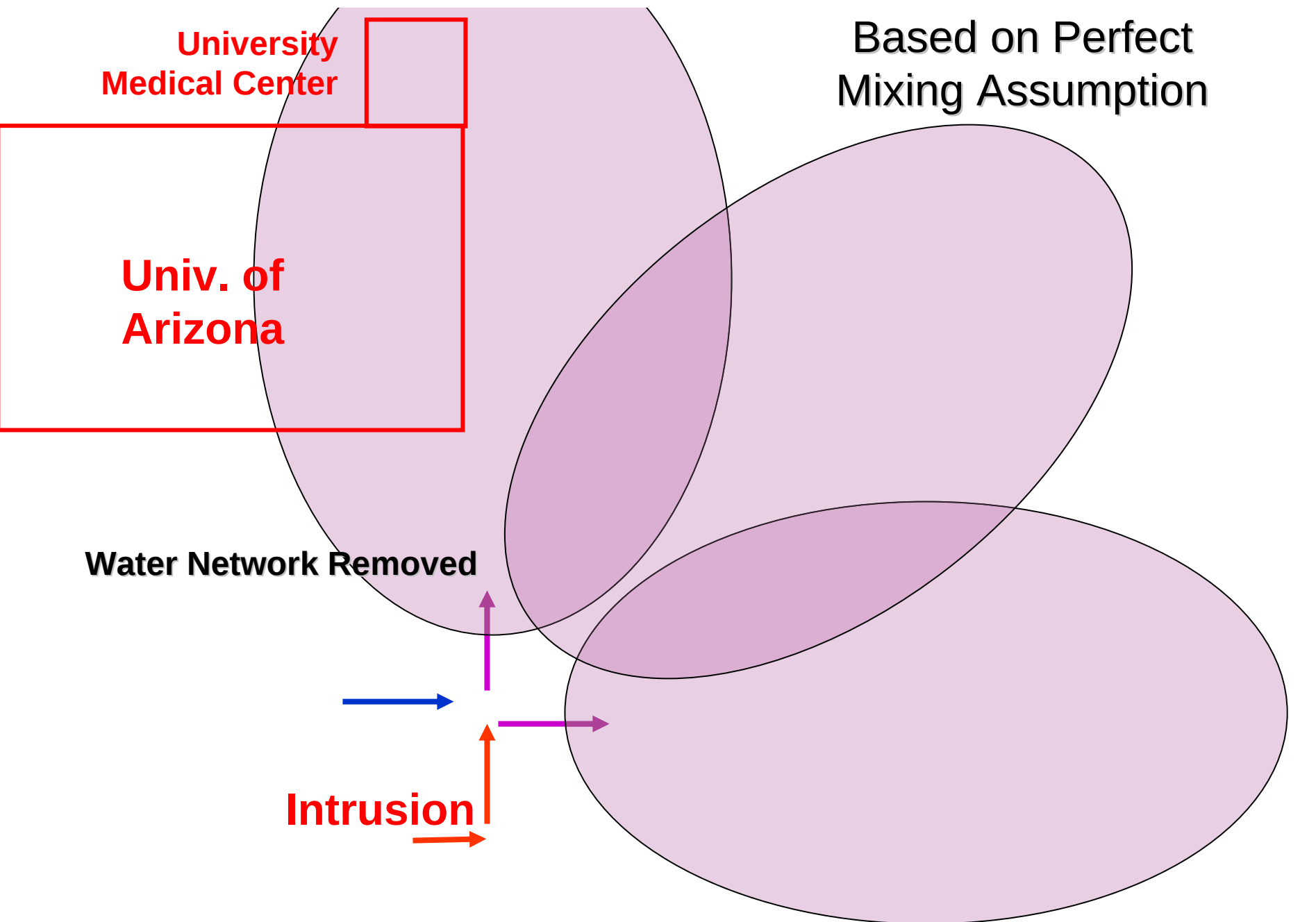


Reynolds Numbers

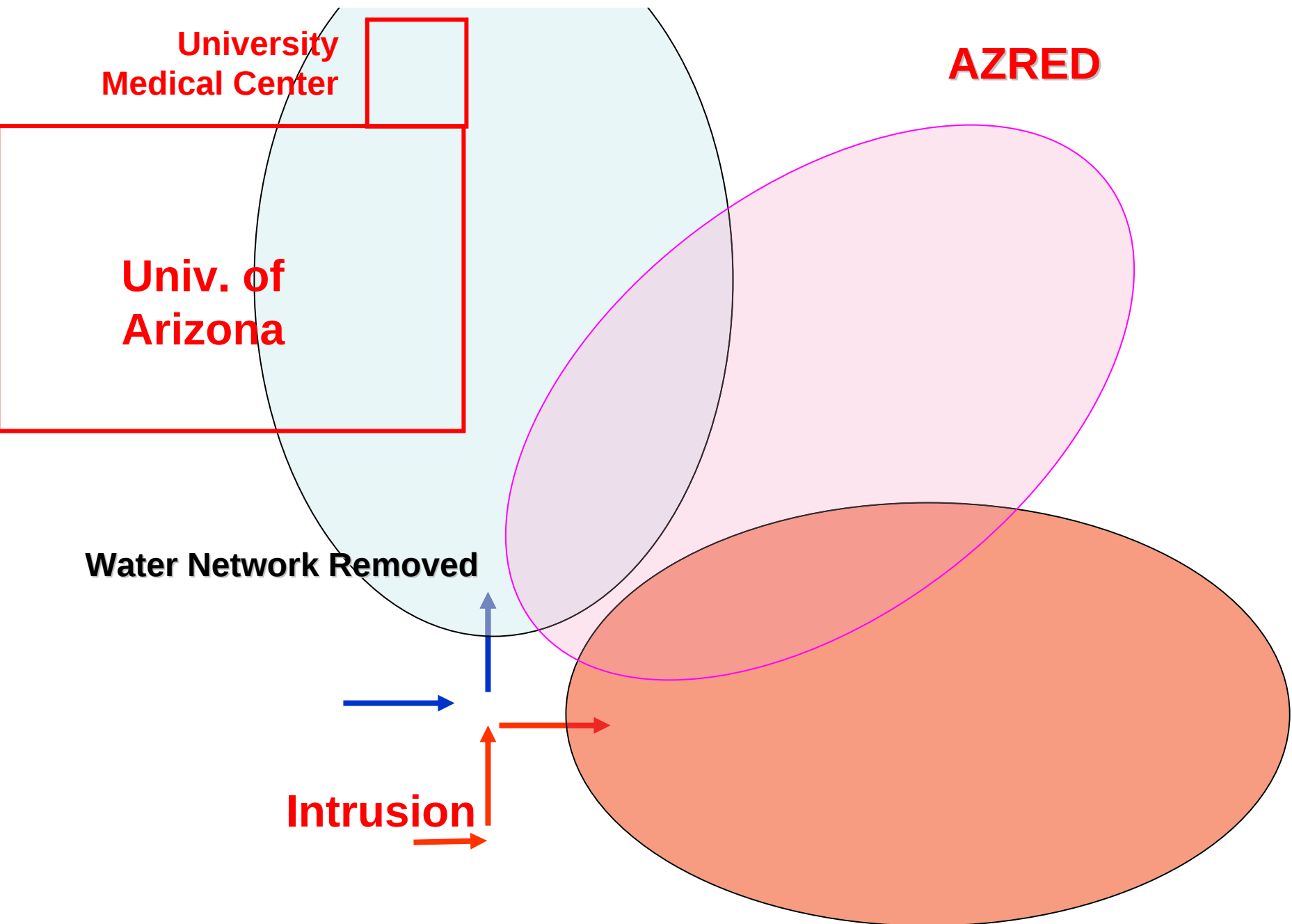
● Sensor Locations



Corresponding Risk Microbial Risk Assessment & Consequences



Corresponding Risk Microbial Risk Assessment & Consequences

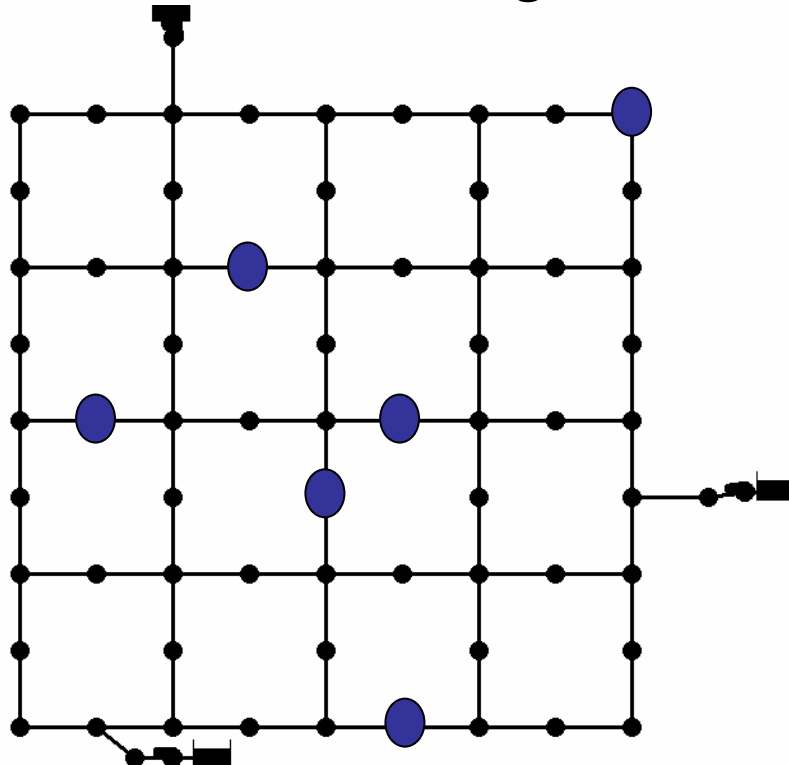


Minimum Hazard Levels (MHL) = 1 mg/L

Level of Services (LOS) = 25 m³

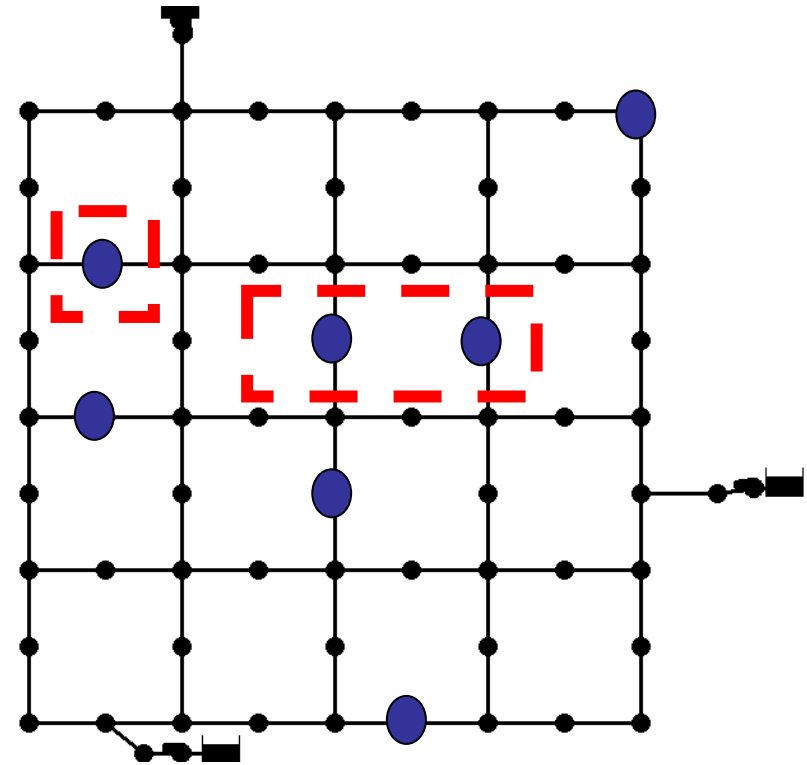
The total number of sensors and their locations change based on AZRED water quality data.

Perfect Mixing



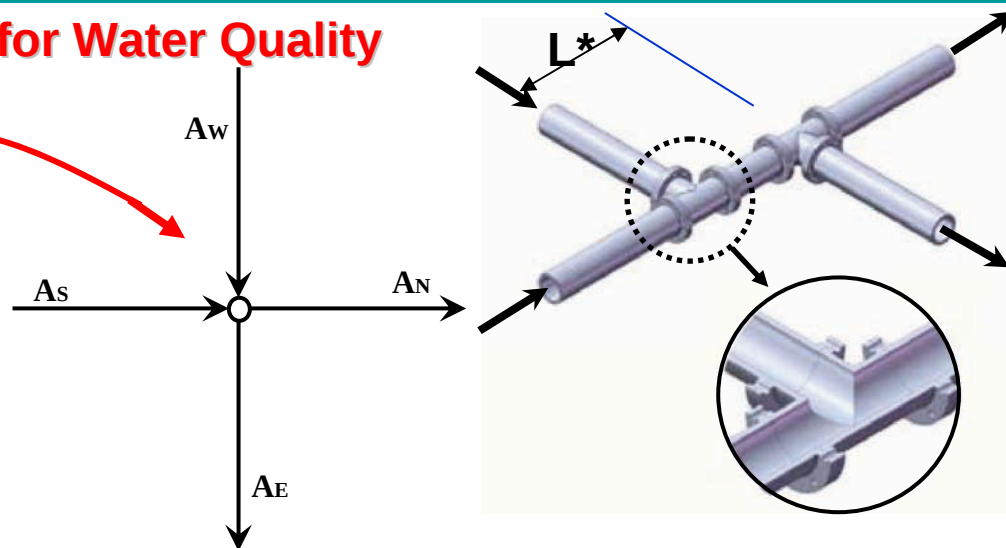
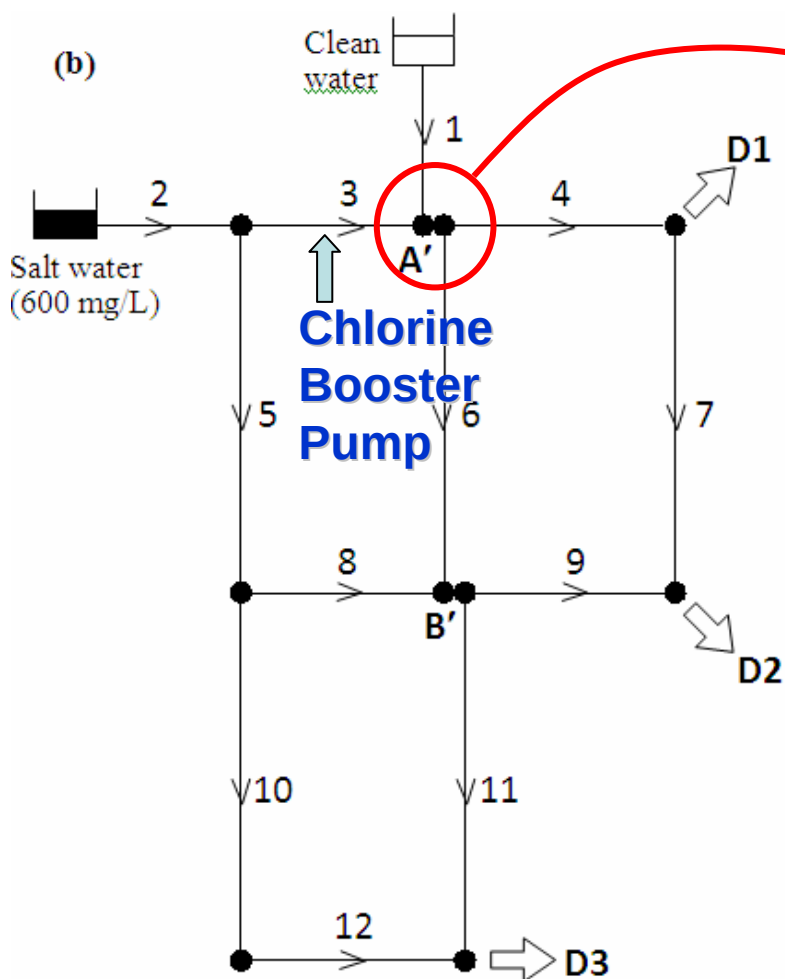
6 sensors

AZRED

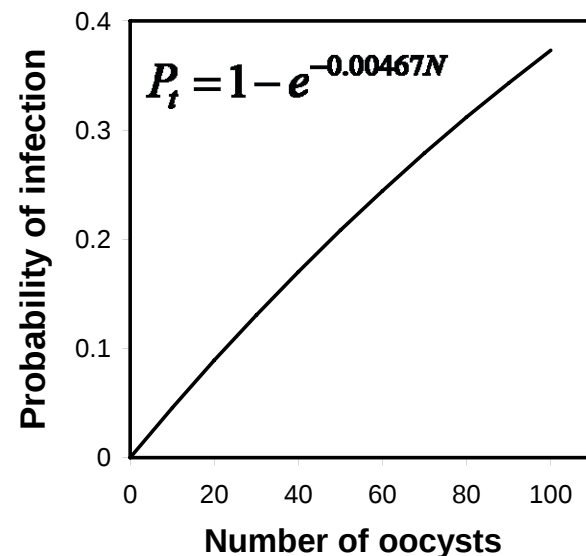
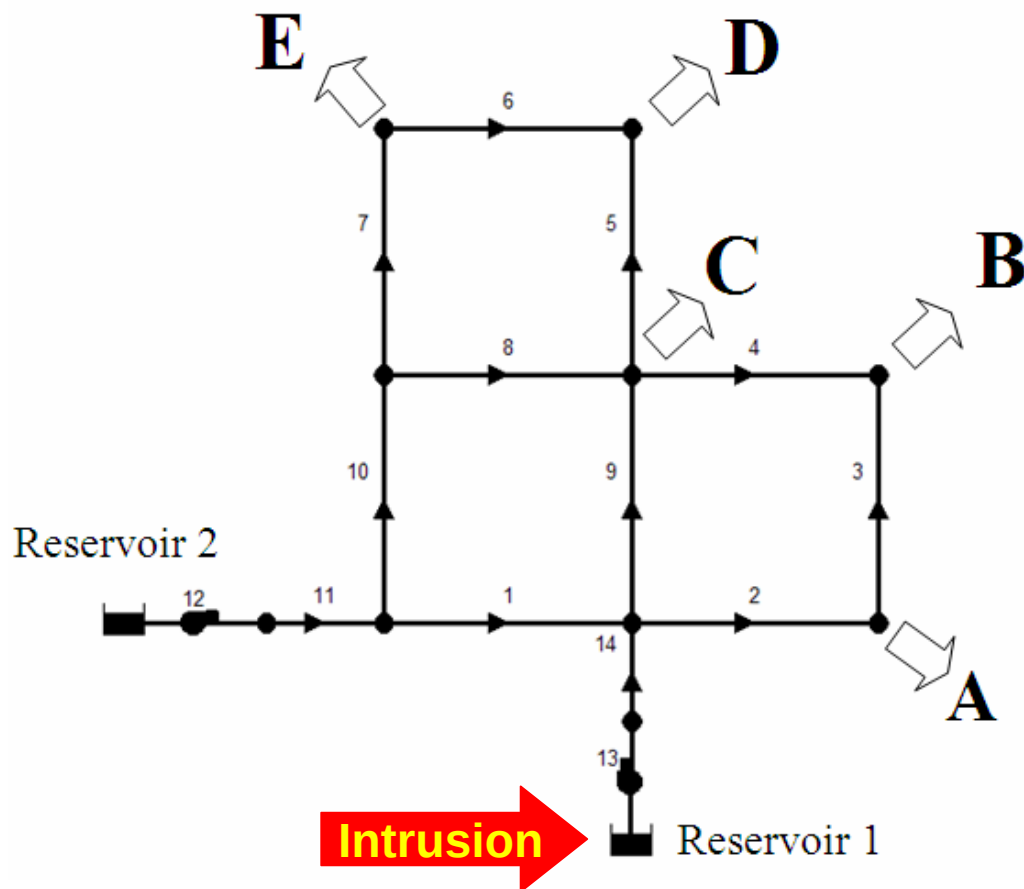


7 sensors

Water Age = Reliable Surrogate for Water Quality

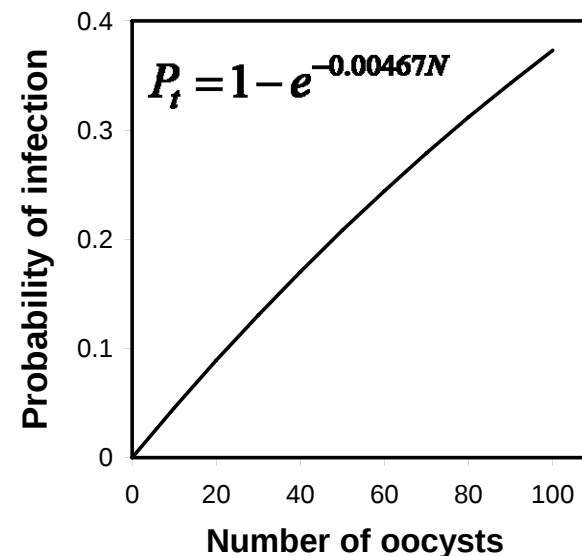
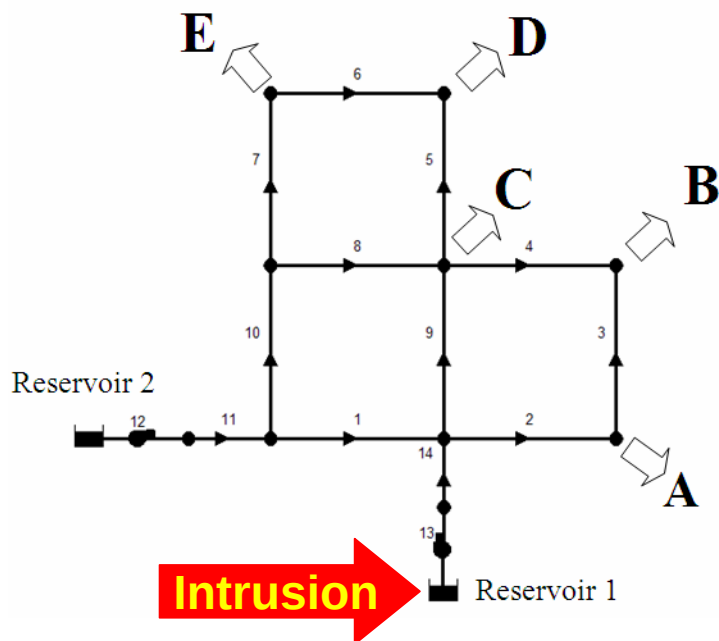


Assumption	L^*	Water age (hr)			
		A_S	A_W	A_N	A_E
Complete mixing (A_0)	-			2.27	2.27
Incomplete mixing (A_1)	0.0			2.76	1.62
	2.5	1.56	2.97	2.47	2.00
	5.0			2.35	2.15
	10.0			2.30	2.22



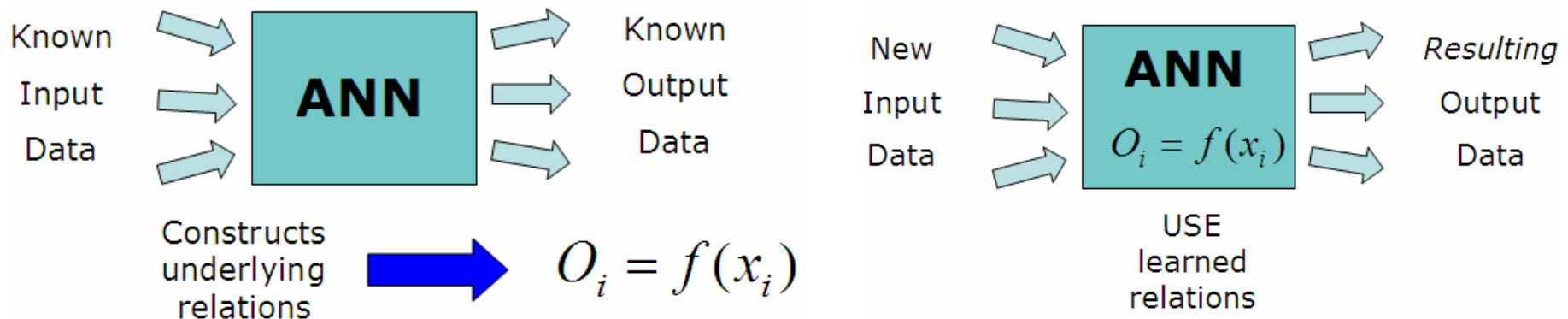
Dose-Response curve,
Cryptosporidium (Haas
et al., 1996)

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.

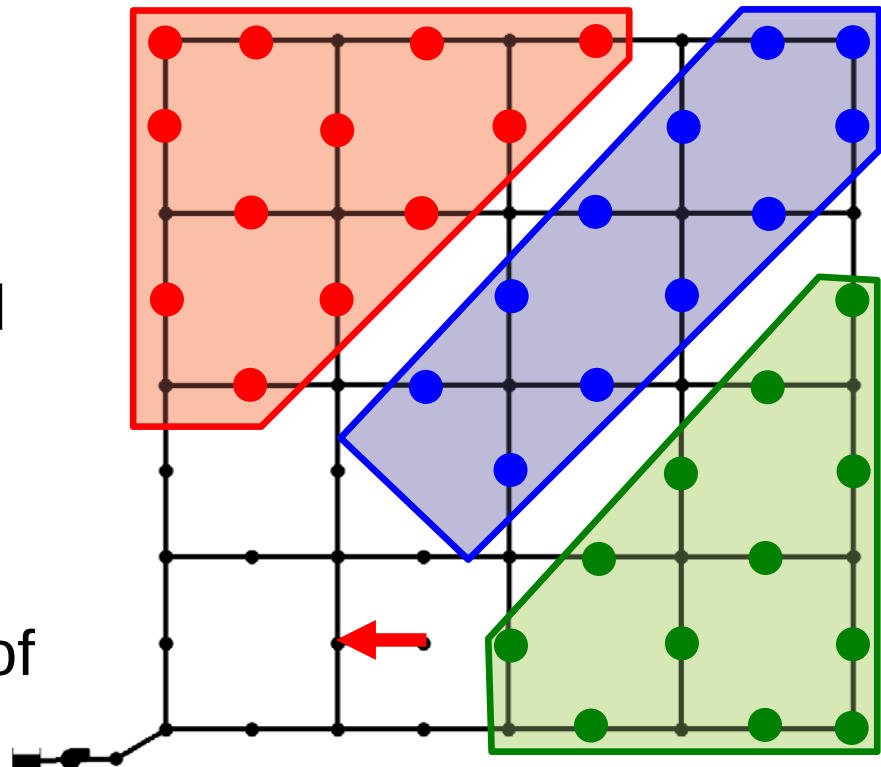


Dose-Response curve, *Cryptosporidium*
(Haas *et al.*, 1996)

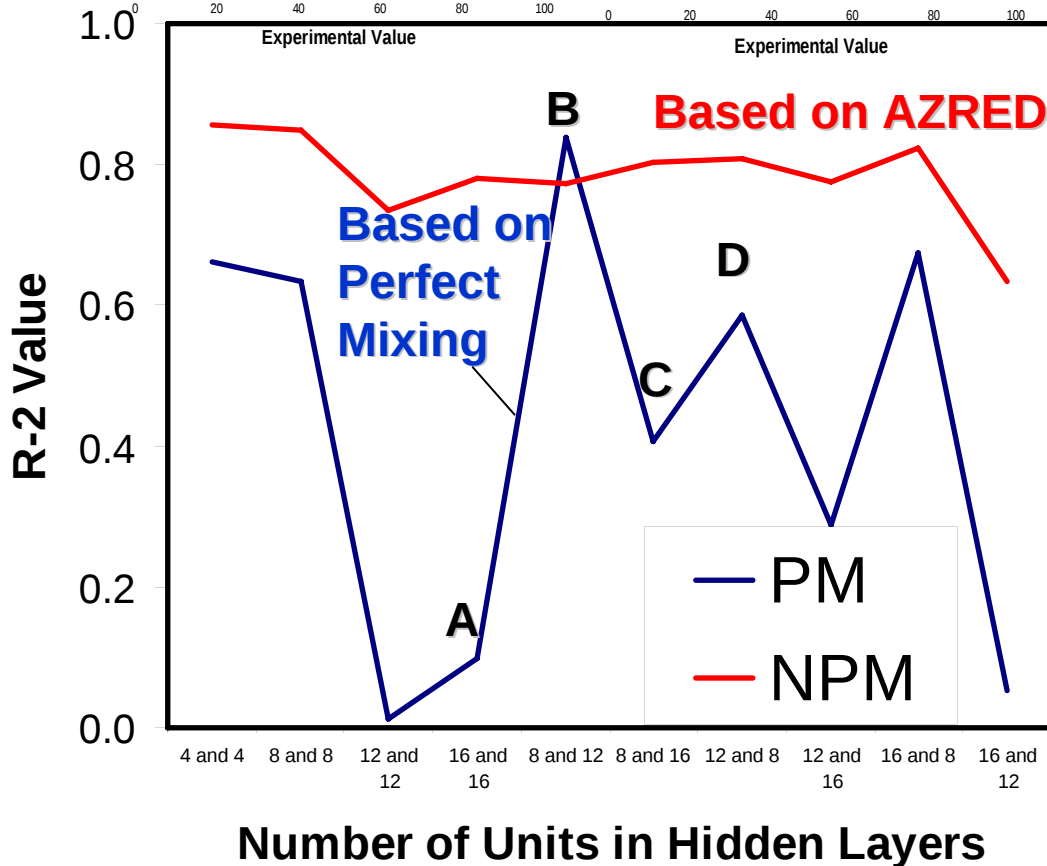
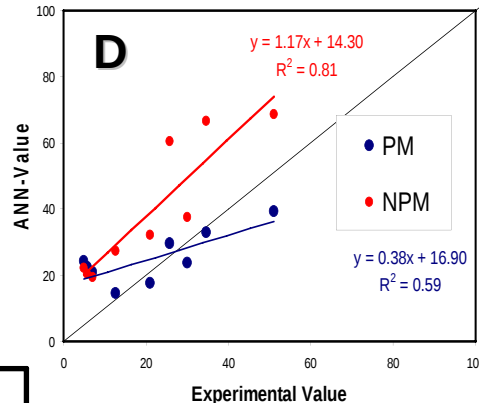
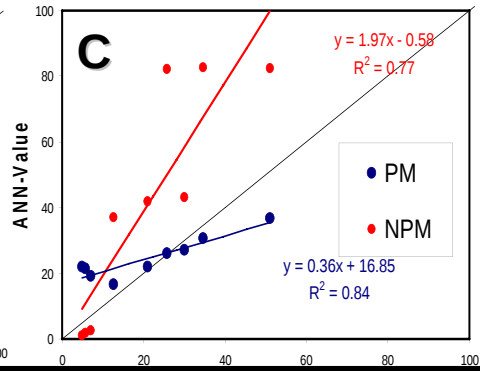
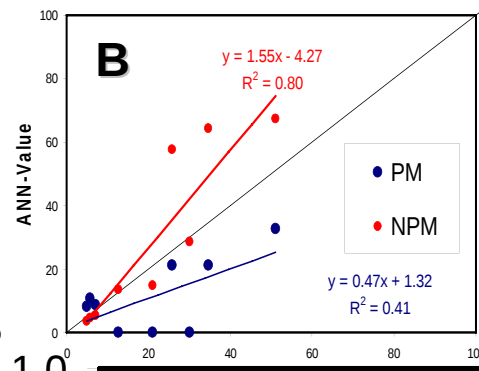
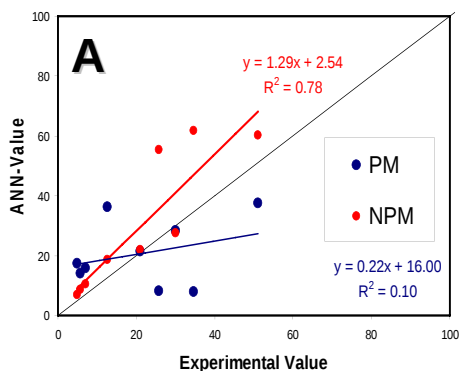
Neighborhood	Assumption	Oocyst concentration (oocyst/L)	Ingested oocyst per person (dose)	Probability of infection (exponential model)	Infected population
A	Complete Mixing	21.83	44	0.186	186
	Incomplete Mixing	37.35	75	0.294	294
D	Complete Mixing	7.14	14	0.063	158
	Incomplete Mixing	0.50	1	0.005	13



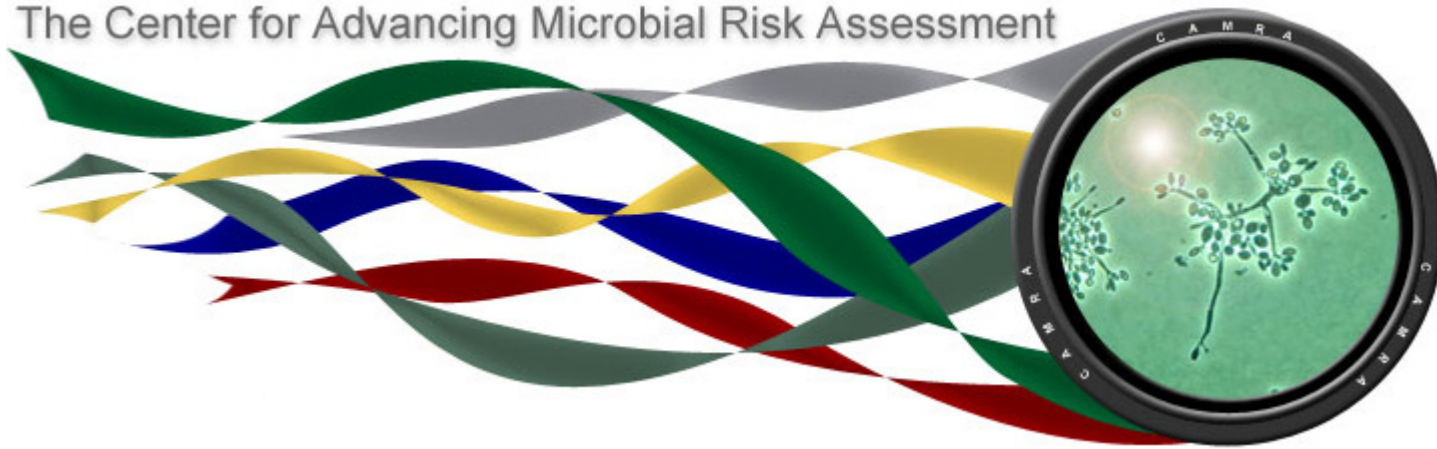
- Three random demands were set, one on each “region”
- AZRED simulations generated input data for ANN training and testing
- ANNs were trained on WQ models with both assumptions of mixing at junctions



ANN vs Experimental Results



The Center for Advancing Microbial Risk Assessment



Research Sponsors and Major Collaborators

