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# Antibody-based Biosensor



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Experiments: characterization of antibodies, determination of assay parameters, preparation and reactivation of Apo-glucose oxidase, synthesis.

**MD Simulations:** antigens binding to antibody, energy minimization, loop structures, sequence alignment.

**CFD Simulations**: flows in microchannels, complex geometry, property optimization, reaction-diffusion-transport of concentrations, parallelization.

**Manufacturing:** microsensor layer fabrication, micromixer fabrication and evaluation, nanoporous membrane.

## The immunosensor will use GOx mediated glucose oxidation for signal transduction

#### **Glucose Oxidation**



cose oxidase requires the cofactor for the catalysis of glucose to onic acid. This process involves initial reduction of FAD to FADH<sub>2</sub> consequent oxidation by ecular O<sub>2</sub> generating <u>H<sub>2</sub>O</u>2

#### **E-Chem Sensor**

Enzyme activity can be modulated the removal and introduction of th cofactor FAD. The cofactor can be efficiently dissociated under acidic conditions to yield apo glucose oxidase.



Thus the cofactor FAD can be conjugated to an analyte and utilized to modulate enzyme activity.

## General Strategy for E-chem Immunoassa





# **E-Chemical Immunoassay**



## **Reactivation of Apo-GOx**



## Antibody – Analyte Selection

ne Number	Ligand	K <sub>d</sub> (M)	Availability
20	Fluorescein	1.5 x 10 <sup>-9</sup>	Invitrogen
209	Fluorescein	3.6 x 10 <sup>-9</sup>	Fitzgerald International
6	2,9-dicarboxyl-1,10	7.5 x 10 <sup>-7</sup>	Blake et al., (2004) Bioconj.
	phenanthroline (DCP)		Chem. <b>15</b> :1125.
6	UO2 <sup>2+</sup> -DCP	9.1 x 10 <sup>-10</sup>	Ibid
3	EDTA	1.3 x 10 <sup>-8</sup>	Blake lab
3	Cu <sup>2+</sup> -EDTA	2.2 x 10 <sup>-9</sup>	Blake lab

### Analytes to be coupled to FAD





CC

C

## Synthesis of FAD Conjugate



## Summary

#### election and Characterization of Antibodies

Commercial and in-house antibodies have been characterized.

#### po-Glucose Oxidase has been prepared

Change in UV-VIS Spectra >300nm confirmed removal of FAD. Purification has been optimized to yield high quantity with low residual signal; storage conditions have been developed. Apo GOx has been transferred to LATech for sensor fabrication.

#### AD mediated Reactivation of Apo-Glucose Oxidase

Reactivation of Apo GOx was dependent on FAD concentration. Reactivated enzyme showed kinetics identical to native GO. Enzyme activity was not affected by components of the immunoassa

#### ynthesis of primary amine terminated FAD

N<sup>6</sup>-2-aminoethyl FAD has been synthesized and characterized. This intermediate was used to synthesize FAD-analyte conjugates. The apo enzyme could be reactivated with the FAD-FITC conjugate. A newly synthesized bifunctional crosslinker is also being tested for the preparation of FAD conjugates.

# **Antibody Bootcamp for Modelers**

### perimental Rotation in Blake Lab

Ashbaugh graduate student (Jain) spent one and half wee in Blake lab learning experimental protocols for antibody sensing.

Titer experiments performed to measure concentrations of antibody 5B2, Pb<sup>2+</sup>-DTPA-benzyl-BSA conjugate, metal chelator (DTPA), and Pb<sup>2+</sup>-DTPA.

Enzyme-Linked ImmunoSorbant Assay (ELISA) used for titer of monoclonal antibody 5B2 and Pb<sup>2+</sup> conjugate. Competitive inhibition ELISA used to infer the ability of DTPA and Pb<sup>2+</sup> to bind to 5B2.

# he Molecular Modeling Requires

- Creation of putative antibody models based on sequence Nodeler)
- Parameterization of the analytes that bind to the antibodie Gaussian)
- Docking analytes in different potential antibody binding ops; (PackMol)
- Optimization of the antibody-analyte interaction by *in silice* o*int mutations (Methods under development, REDS)*



# sensors: Computational Aspects MD

## ulations of 5B2 loop region (Test Cases)

nding of antigens to antibody occurs in loop domain. Aim to identify us nulations side chains in loop region that contribute to binding specifici ide antibody engineering.

vacuo energy minimizations of 5B2 LC and HC loops confirm previou entification of metal binding residue Lys<sup>58</sup>.

eplica Exchange Molecular Dynamic performed of 5B2 in vacuo and plicit solvent to generate families of loop structures for minimization to termine robustness of predictions and identify spatial and dynamic prrelations between key binding residues

tial findings: HC3 loop has more varied and xible structure than the other e antibody loops

LC1

HC<sub>3</sub>

## sensors: Computational Aspects MD

### Simulations of 5B2 loop region (continued II)

Replica Exchange (REMD):
*replica*: several simultaneous simulations
2 levels of parallelization
*exchange*: simulations swap information

### Simulation Characteristics

### Loops

~1000 atoms => 2CPUS/sim 10ns run time => Gb's data; Full REMD in 24hrs 64CPUS

### Full System

~10,000atoms => 4CPUS/sim 10ns run time => 10-100Gb data; Full REMD in 2wks 64CPUS

## **Biosensors: MD Fast Track Study**

high throughput simulation workflow

Bishop (CCS @ TU) Emir Embahsi & Tevik Kosar (CCT @ LSU)



retical and Computational Biophysics Group at UIUC

## yberTools Connections

1: Scheduling and Data Services.

e details of integrating our Molecular Modeling packages into WP 1 arong addressed by Drs. Thomas Bishop (Tulane) and Tevfik Kosar (LSU

#### 2: Information Services and Portals.

a. Thomas Bishop and Tevfik Kosar are collaborating to bring Bishop's A folding simulations on-line. The Workflow resulting from this effort care readily modified to investigate the antibody and analyte interactions.

#### <u>93: Visualization Services.</u>

rk is in progress to create modules that will permit all scientists involve project to visualize molecular models and other results via a commor er interface without the necessity of transferring data or installing softw local lab computers.

#### 4: Application Services and Toolkits.

Steven Rick (UNO) and Henry Ashbaugh are developing replica sulation techniques that will enable this group to efficiently identify anti p sequences that optimize the antibody-analyte interactions.

# Fluid Mechanics and Transport

**GOAL** → Computationally determine the optimal geometric configuration of the omega channel network to enhance mixing of two species.

Laminar flow field governed by continuity & Stokes equations:  $\nabla P = \mu \nabla^2 \mathbf{u}$ 

$$\nabla \cdot \mathbf{u} = 0$$

Boundary Element Method determines velocities and surface stresses



Omega channels developed by IfM

Streamlines resulting from constant pressure drop across model channel



# **Boundary Element Method**



Velocity **u** and stress  $\tau$  are approximated as quadratic polynomials, and at each node point, satisfy

$$\mathbf{C}_{ki}u_{i}(\mathbf{x}) + \sum_{m=1}^{N}\int_{s_{m}}\mathbf{T}_{ik}(\mathbf{x},\mathbf{y})u_{i}(\mathbf{y})dS_{m} = \frac{1}{\mu}\sum_{m=1}^{N}\int_{s_{m}}\mathbf{U}_{ik}(\mathbf{x},\mathbf{y})\tau_{i}(\mathbf{y})dS_{m}$$

Integral equation is expressed as system of linear equations:

$$H\mathbf{u} = G\tau$$

- Elements of H and G computed using Gaussian quadrature rules
- Optimization of simulation is being developed in conjunction with WP4 and will create a general purpose *CyberTool*.

# **Results: Particle Trajectory**

For more info see our poster!



Particles initially positioned along y-axis at x = 0

Path of each particle traced as it flows through the domain

Note inner particles travel more slowly than outer particles that migrate quickly across channel along outer walls

Results suggest domain modification is important to improve mixing

# licrosensor Mixing and Transpo

•Analyte-FAD conjugate and analyte from serum compete to bind with antibody

 Binding and release occur spontaneously as analytes and antibody are transported by fluid motion

ach analyte/antibody satisfy a reaction-diffusion equation:

$$\frac{\partial C_i}{\partial t} + \nabla \cdot (\mathbf{u} C_i) = D_i \nabla^2 C_i + R_i$$

- **u** fluid velocity
- $C_i$  concentration of each species
- $D_i$  diffusion coefficient
- $R_i(C)$  reaction term

# **Transport Methodology**

ansform equations into a boundary-fitted coordinate system



e the Finite Volume Method to solve for *concentration* 

ote: velocity field obtained from BEM code

ultiblock approach for omega channels

orking with Dr. Blake for reaction/diffusion rates

# CyberTools Connections

### <u> Current Work:</u>

- Parallelization of Stokes flow problem for use in the HPC environment (WP4: Mayank Tyagi, Shantenu Jha, Sanjay Kodiyalam)
  - OpenMP
- Visualization of model problem using TecPlot (with WP3)
- Generalization of code to develop a CyberTool package that solves Stokes flow equations



### uture Work:

- Parallelization of source code including transport

## Layer-by-Layer (LbL) Nanoporous Membrane for Immunoassay (sensor technology for enzyme deposition



manufacturing





Nanoparticle/polyion (or protein) bilayer, D = 5-50 nm

eme of the layer-by-layer bassembly by alternate orption of polycations and anions or nanoparticles SEM cross-section images of (glucos) oxidase/PAH)<sub>22</sub> multilayer on quartz (lef and (40 nm silica/PAH)<sub>6</sub> film on silv electrode (right).

## Polymer-based Electronic Microsensor Fabrication



## **Micromixer Fabrication**







**Fabrication** 

- Lithography
- ICP
- Bonding
- **Challenges**
- Connectors
- **Modifications**
- New set of connectors from Upchurch Scientific are being tested and evaluated

## **Micromixer Evaluation**



- difications
- esigned 'T' shape inlet and outlet for
- itial mixing
- antification
- nage analysis software
- sing fluorescent dyes for better signal/noise



New Micromixer Design

## Fluorescent dyes in the omega channels: /ISUALIZED WITH MICROSCOPY AND DIGITAL CAMERA



Water FITC

flow of fluids



# **Summary and Conclusions**

#### Microfluidic Component

- Fabricated and evaluated two sets of micromixers
- Designed new micromixer based on the results obtained (fabricated)

#### Nanoporous Membrane

LbL nanoassembly is being evaluated for fabricating nanoporous membrane; <u>New:</u> in collaboration with Dr. Scott Gold (Louisiana Tech.) for modeling of flow through porous membrane.

#### Reproducibility

 Evaluating PEDOT and carbon nanotube based microsensor for reproducibility, Selectivity, ar Life Time

#### Testing Procedures

- Currently testing florescent dyes and particles (proposed) for evaluating micromixers.
- Currently evaluating microscale sensor system based on carbon nanotubes
- Carbon-based Electrodes
- Under testing and fabrication
- CyberTools Connection
- Access Grid (AG) video conference with Tulane (23 July 2008): simulations and experimenta data. Evaluation of Cybertools link to visualization software: Vislt 1.9.1 (Windows version, DeCoster) thru the Cactus Code Link.
- VISIT OUR GRADUATE STUDENT POSTER! -- SENAKA KANAKAMEDALA-

## **Final Remarks**

#### eractions between the laboratory and the molecular dynamics groups

- rovided new models of protein structure that allow testing of hypotheses in sili or to time-consuming laboratory experiments.
- alidated in silico predictions based on laboratory experiments.
- entified methodological refinements based on laboratory results.

#### eractions between the laboratory and the micromanufacturing groups

- roadened the kinds of hardware and electronics that can used to construct the nsors.
- compted micromanufacturers to examine paradigms used to validate their dev g. they have modified the molecular identity and concentration ranges of reag ed to test their microscale mixers).

#### eractions between the fluid mechanics and the micromanufacturing gro

- rovided data to determine boundary conditions used in the simulations.
- stablished realistic geometries for fluid flow domains.
- elineated a plan to reduce the number of fabrication trials needed to optimize al device.