

Spectroscopy Systems for Biodetection

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Presentation overview

- **Overview of Spectroscopy Methods**
- **Chemical Markers**
- **Physicochemical Markers**
- **Recent Advances**
- **Example: Uv-vis Spectroscopy**
- **Vision for the Future**

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Applications of Spectrometer Systems for Biodetection

- **Detection of Chemical Markers**
 - Invasive (molecular)
 - Non-invasive (molecular-phenotype)
- **Physicochemical Markers**
 - Invasive (labelling)
 - Non-invasive (phenotype)
- **Culture Methods (amplification)**
- **Non-Culture Methods (direct)**
- **Laboratory Applications**
- **Field (in-situ) Applications**

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Detection of Chemical Markers

- **GC-MS**
 - Carbohydrates
 - Polysaccharides, Lipopolysaccharides
 - Fatty Acids
- **FT-IR**
 - Total chemical and biochemical composition
- **UV Resonance Raman Spectroscopy**
 - Aminoacids, proteins, nucleic acids, pigments
- **Multidimensional Fluorescence**
 - Natural Fluorescence
 - Fluorescence Labelling
- **Fluorescence Lifetimes**
- **UV-Vis Spectroscopy**

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Detection of Chemical Markers

- **Advantages:**
 - High Sensitivity and Specificity to the identified markers (Analytical Chemistry)
 - Trace Analysis Possible
 - Profiling of : carbohydrates, sugars, fatty acids, etc.
- **Issues:**
 - Marker Identification
 - Sampling (cultures, environmental, clinical)
 - Sample preparation (derivatization, amplification)
 - Interferences: require fractionation GC, HPLC

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Detection of Physicochemical Markers

- **Angular Light Scattering**
 - Static LS
 - Dynamic LS
- **Microscopy**
 - Fluorescence
 - FT-IR
- **Imaging Methods**
- **Hybrid Methods**
 - UV-Vis Spectroscopy

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Detection of Physicochemical Markers

- **Advantages:**
 - High Sensitivity
 - Portable Instrumentation
 - Non-invasive
- **Issues:**
 - Marker Identification (size, shape, polarization)
 - Non-Specific markers
 - Sampling
 - Sample preparation
 - Interferences: require fractionation (Flow Cytometry)

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Analytical/Interpretation Methods

- **Theoretical**
 - Band assignment
 - Quantification
 - Linear Models (Beer-Lambert)
 - Non-Linear Models (Mie theory)
- **Statistical**
 - Multivariate Statistical Models
 - Factor Analysis
 - Principal Components
 - Genetic Algorithms
 - Classification Algorithms
 - Pattern Recognition

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Recent Advances

- **Miniaturized Instrumentation**
 - Detectors (MS, IR, Uv-vis, Raman)
 - Sampling and Sample preparation systems
 - Microfluidics
 - Nano-technology
- **Interpretation-Analysis**
 - Improved Scattering Algorithms (T-Matrix)
 - Enhanced Computation Capabilities
 - Improved Multivariate Statistical Models
 - Factor Analysis
 - Partial Least Squares
 - Pattern Recognition

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Biophotonics Analyzer Requirements

- **Portable laboratories**
 - Autonomous
 - Miniaturized
- **Sophisticated sensors & instrumentation**
- **Multipurpose or multi-sensor capabilities**
- **Immobilized reagent technologies**
- **Rapid screening & diagnosis capabilities**
- **Efficient communication subsystems**
- **Local high performance processing capability**

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Implications of Miniaturized Fiber-Optics Spectrometry

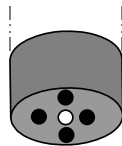
- Measurements (spectrometer systems) can be taken to the process
- Sensors (fiber-optics) can be configured
- Sensors (fiber-distal ends) can be designed for specific applications (polymer & sol-gel immobilized reagents)
- Detectors (spectrometers) can be configured
- Design sampling-measurement systems with the objective of increasing the information content of a measurement (detector and sensor bundles)

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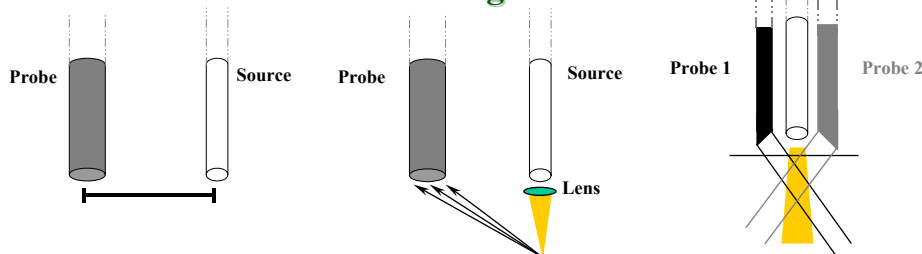
Reflectance/Backscattering

Probe



○ Source
● Detectors

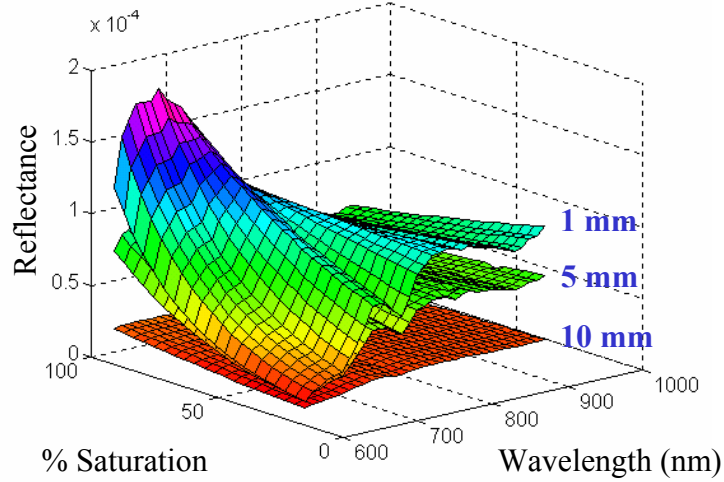
Probe Configurations



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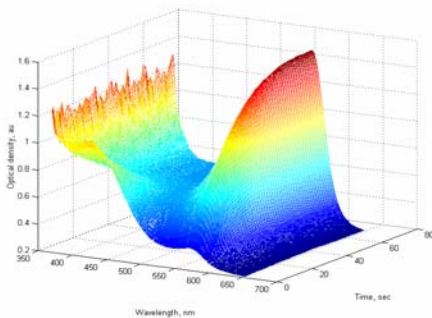
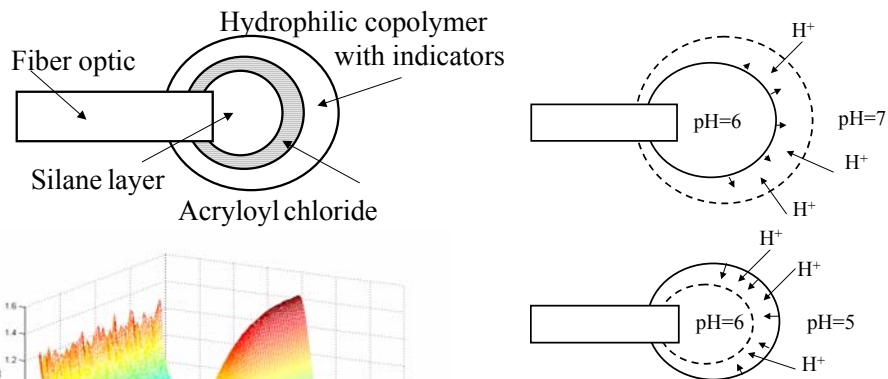
Reflectance of Optically Dense Media for Three Penetration Depths



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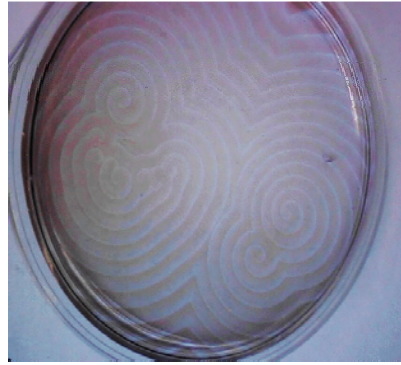
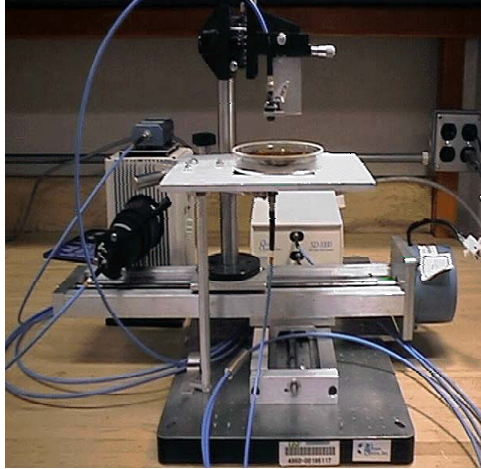
Fiber Optic Configuration



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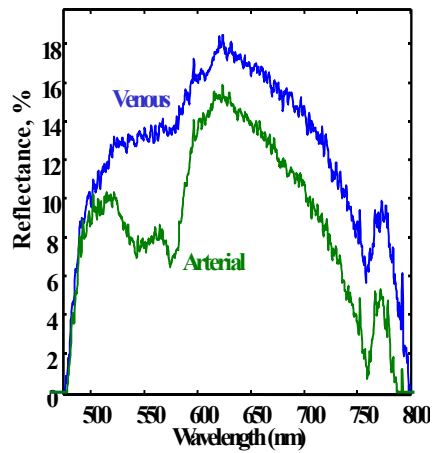
Spatial Patterns: the Belousov-Zhabotinsky Reaction



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Endoscopy: application to fetal surgery



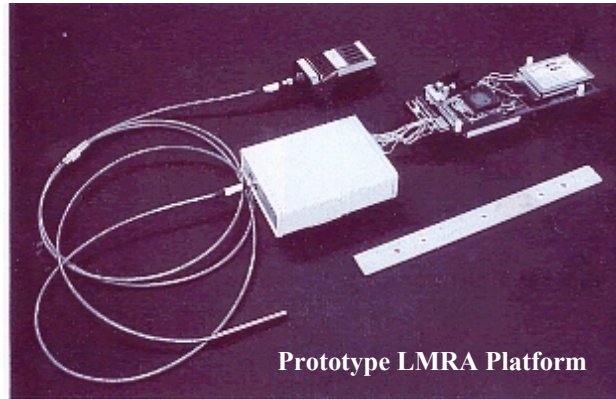
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Measurement Platform

Subsystems

Detector-Sensor Array
Electronics Pack
Communication Unit
Global Positioning System



Prototype LMRA Platform

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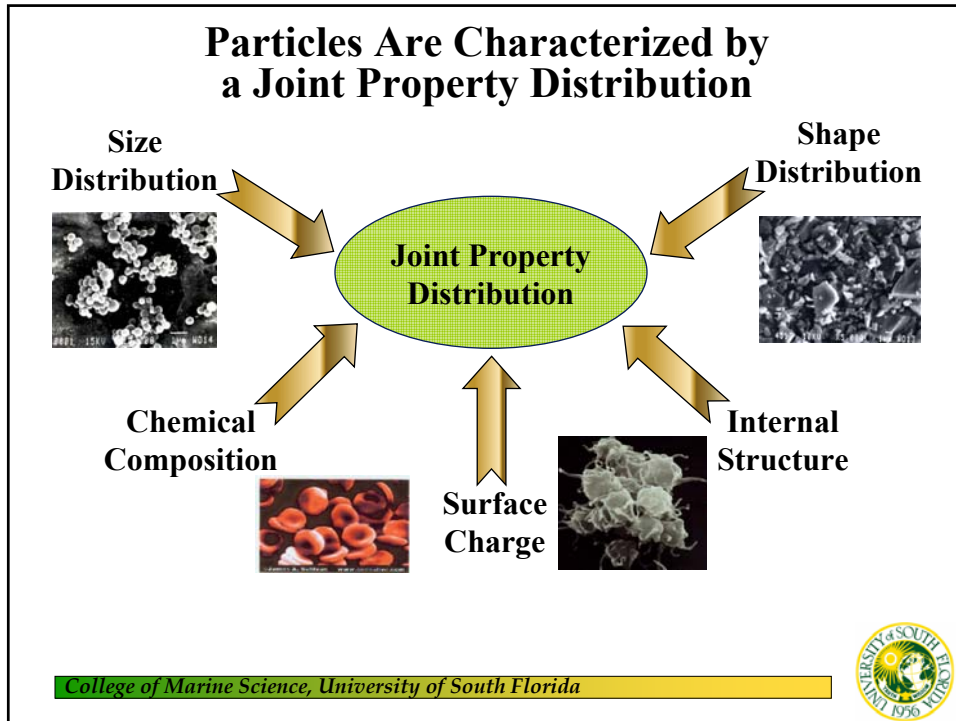



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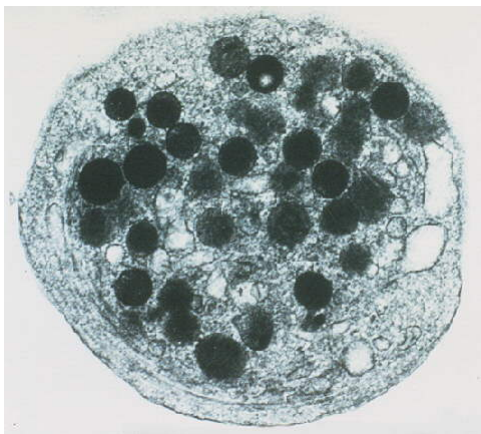
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- ### Issues in Bio-particle Characterization
- | | |
|---|--|
| Dilute Dispersions <ul style="list-style-type: none">▪ Enumeration▪ Shape▪ Composition▪ Internal Structure▪ Size▪ Charge▪ Orientation▪ Mobility▪ Optical properties▪ Identification▪ Limits of Detection | Concentrated Dispersions <p>Dense Media: skin, Suspensions, blood, cell cultures</p> <ul style="list-style-type: none">▪ Fluid structure<ul style="list-style-type: none">▪ Particle-Particle Interactions▪ Hydrodynamic Interactions (colony forming)▪ Sampling<ul style="list-style-type: none">▪ Sample Integrity▪ Representative Sampling▪ Background<ul style="list-style-type: none">▪ Growth Media▪ Detection Limits (contrast) |
|---|--|
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- 

Spectrometry of Complex Particles: *Platelets*



Dense Bodies:

ADP/ATP (~1 M total)

Calcium (2 M)

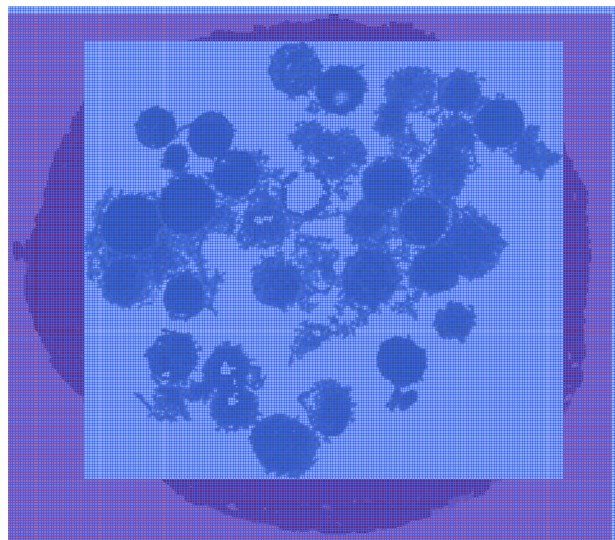
Serotonin (0.065 M)

Pyrophosphate (0.3 M)

Granule Diameter: 20 to 30 nm

Platelet Diameter: 1.5 to 2.5 μm

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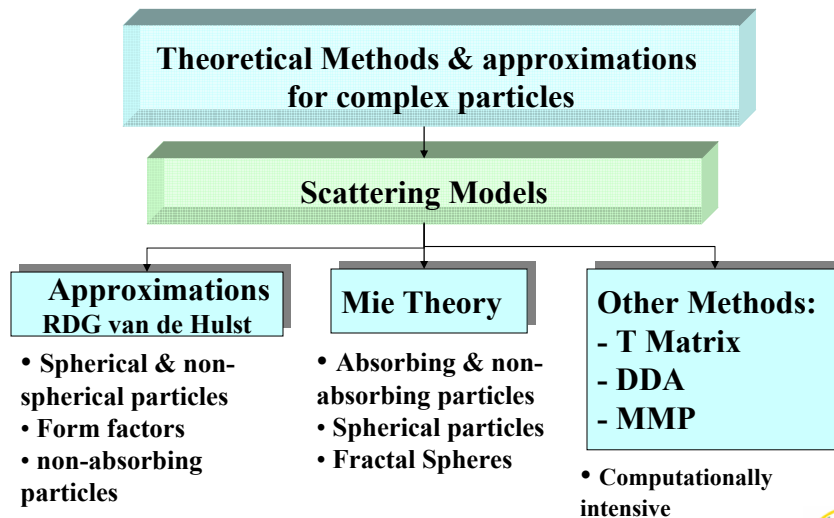
Advantages of Multiwavelength Spectroscopy

- **large dynamic range (resolution: α/λ)**
- **composition is accessible through absorption**
- **real time measurements (sample times < 0.01 sec)**
- **large signal/noise**
- **suitable for in situ, at-line and in-line applications**

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Interpretation of spectral data



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Single Scattering Models (Mie)

Optical Properties

Particle Size Distribution

$$\tau(\lambda_o) = \ell N_p \frac{\pi}{4} \int_0^\infty Q(m, \alpha) D^2 f(D) dD$$

$$m(\lambda) = \frac{n_1(\lambda)}{n_o} + i \frac{k_1(\lambda)}{n_o} \quad \alpha = \frac{\pi D}{\lambda}$$

$$\varepsilon(\lambda)_{obs} = \sum_j^M \frac{4\pi k_1(\lambda, j)}{\lambda} p(j): \quad n_1(\lambda)_{obs} = \sum_j^M n_1(\lambda, j) p(j) Q_{sca}$$

$$n(\omega) - 1 = \frac{2}{\pi} \int_0^\infty \frac{\Omega k(\Omega)}{\Omega^2 - \omega^2} d\Omega: \quad k(\omega) = \frac{-2\omega}{\pi} \int_0^\infty \frac{n(\Omega)}{\Omega^2 - \omega^2} d\Omega$$

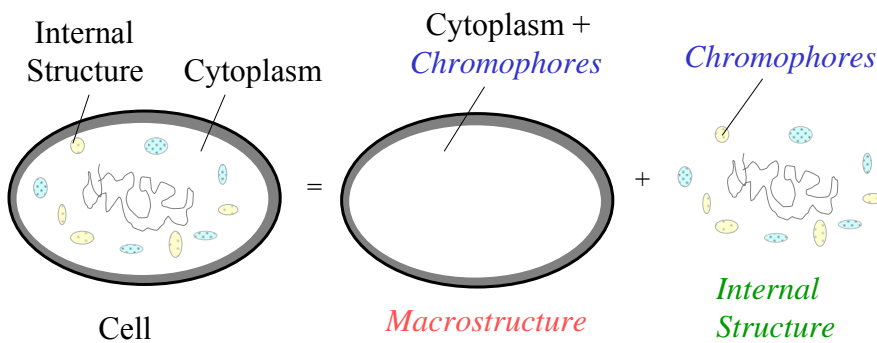
Macroscopic concentration n
 $N_p, Q_{abs}, & Q_{sca}$

Particle D
 $N_p, Q_{abs}, & Q_{sca}$

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Theoretical interpretation model

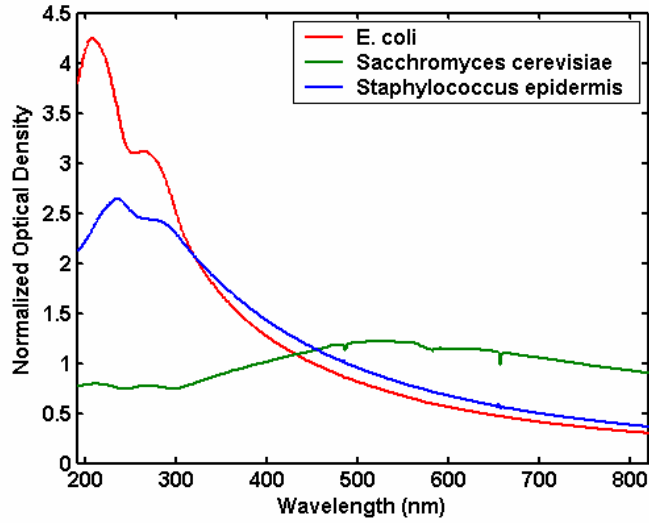


$$\tau(\lambda_o) = N_p \ell \left(\frac{\pi}{4} \right) \int_0^\infty Q_{ext}(m(\lambda_o), D) D^2 f(D) dD$$

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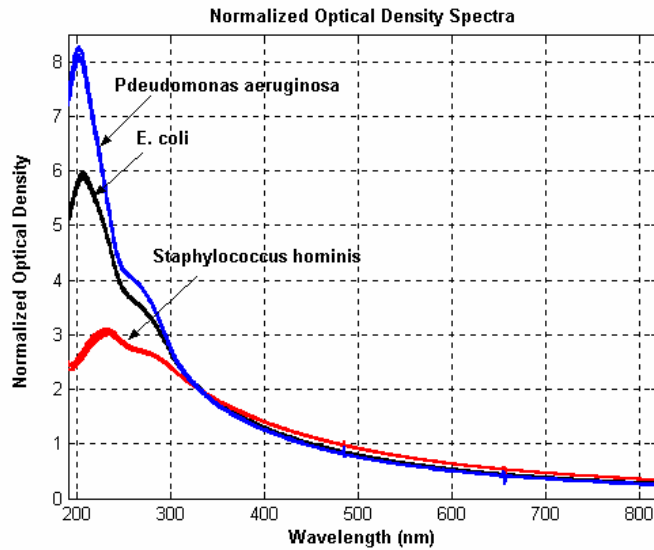
Typical Normalized Optical Density Spectra Measured from Suspensions of *Microorganisms*



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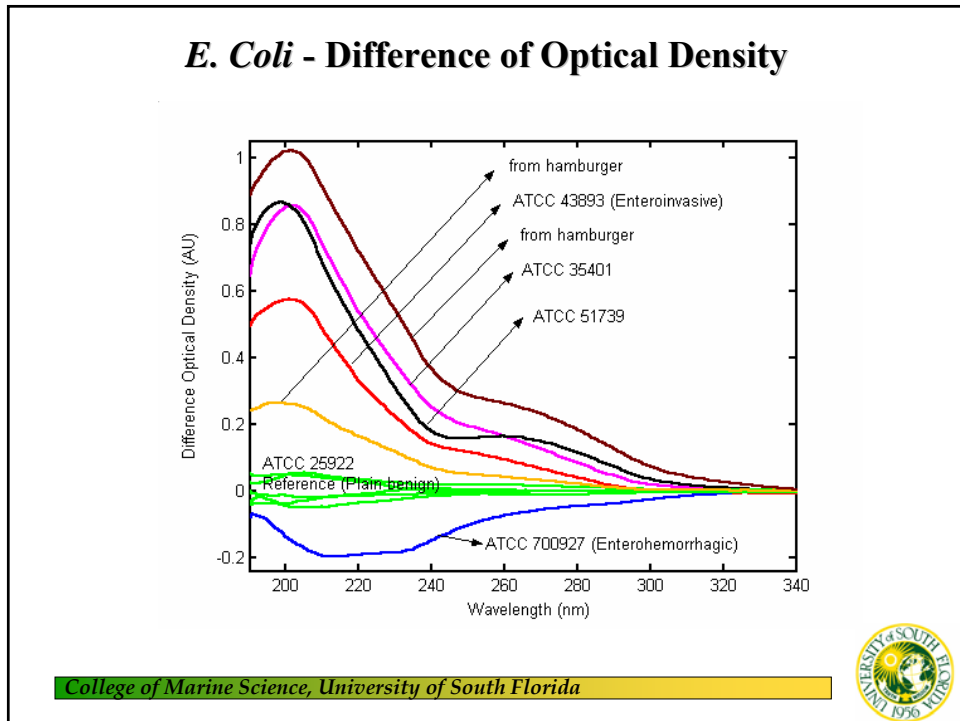
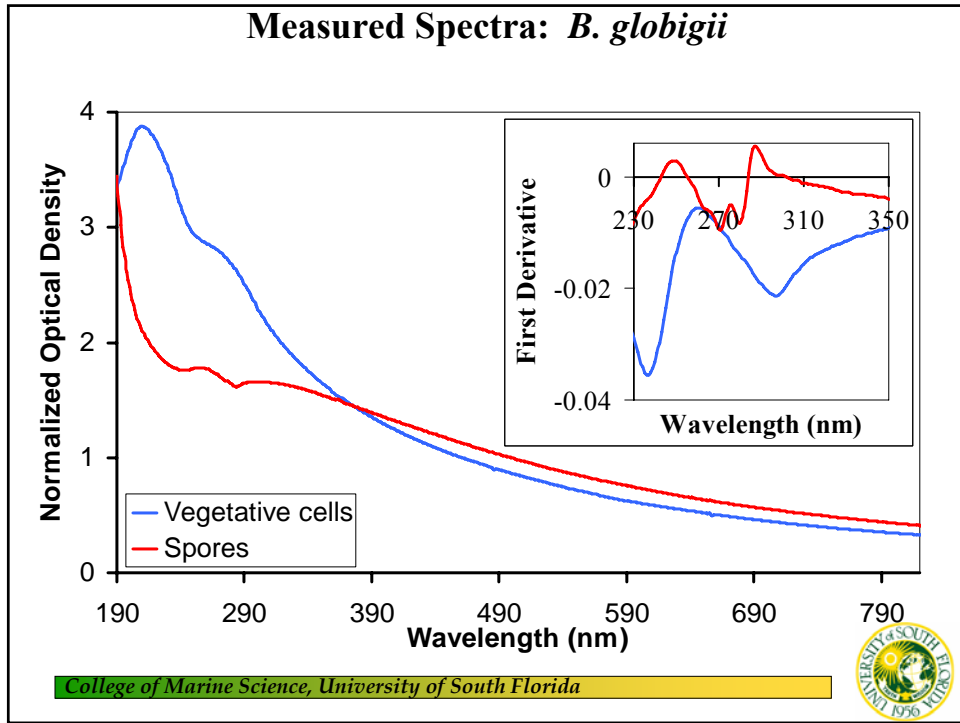


Comparison of Vegetative Cells

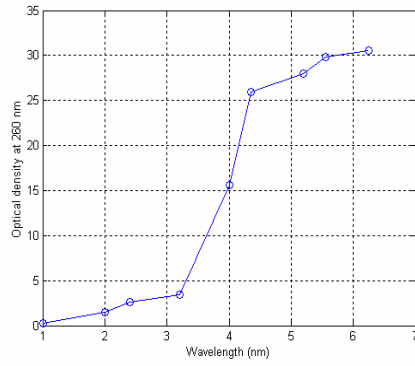
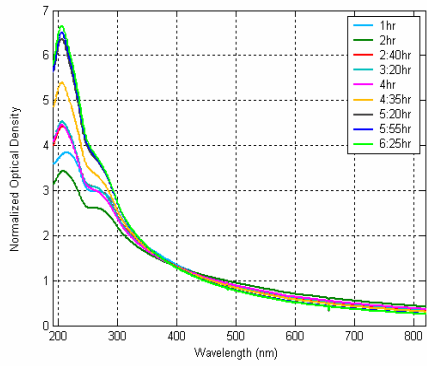


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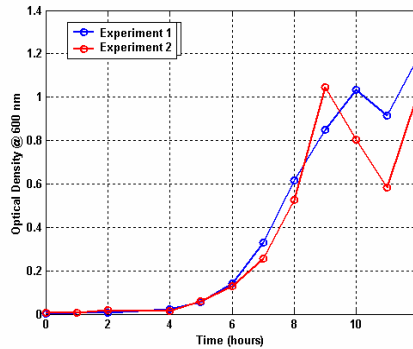
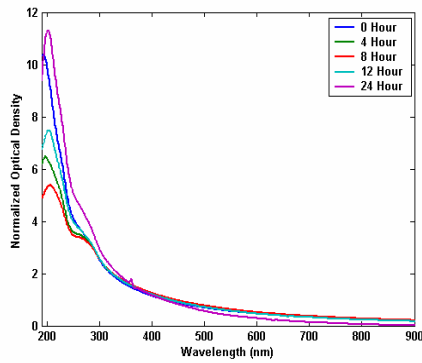
Growth behavior: *E. Coli*



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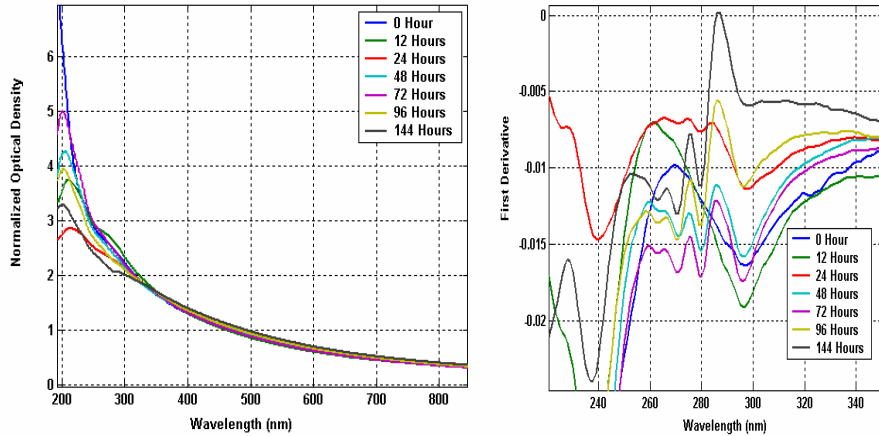
Growth Behavior: *Pantoea agglomerans* (Yersinia Pestis)



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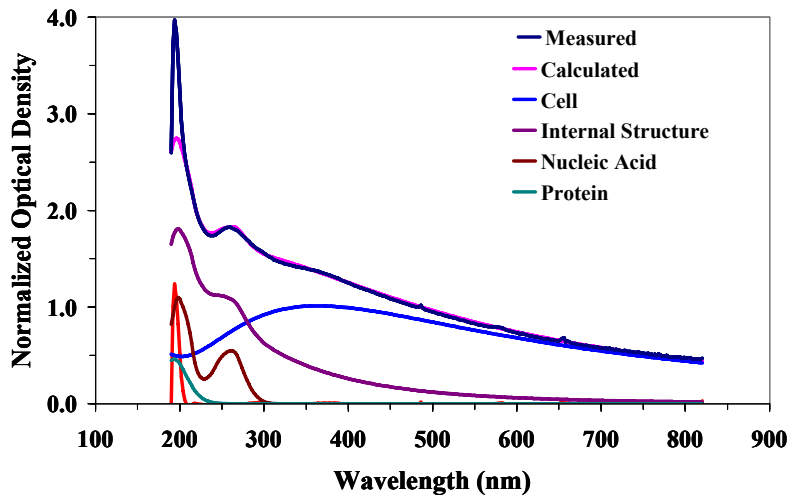
Sporulation Media: *B. Subtilis*



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Deconvolution of the *Cryptosporidium* Transmission Spectrum



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Quantitative Interpretation of the Transmission Spectrum for *Cryptosporidium*

Microorganism

Average Size = 4.07 μm
 Particle No = 341,000 per mL

Corpuscles/Organelles (internal structure)

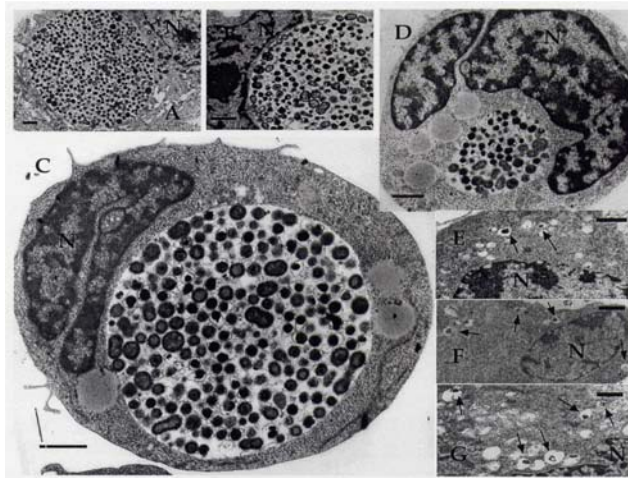
Hypochromicity = 0
 Average Size = 114 nm
 Volume Fraction = 0.4
 Fraction DNA+RNA+Nucleotides = 0.06
 Corpuscle Concentration = 5.21 $\mu\text{g/mL}$ suspension
 Corpuscle Enumeration = 6.46 billion/mL suspension

Max. No. of Corpuscles/cell = 46,481
 Estimated No corpuscles/cell = 18,950
 DNA+RNA+Nucleotides = 0.94 pg/cell

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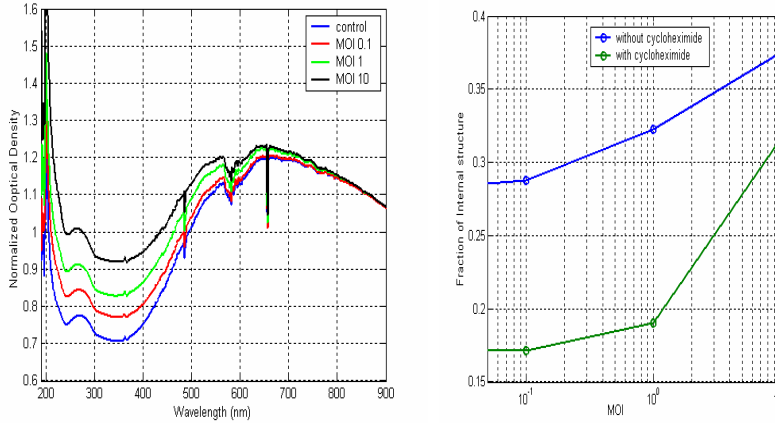
Chlamydia pneumoniae infected cells: HEp-2 (A); monocyte THP-1 (B, E); lymphoid Molt (C, F; lymphoid P3HR1 (D,G). Infected with 10 bacteria/cell after 72 Hrs of incubation. From Y. Yamamoto *et al. private Comm. October 2002.*



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Representative data from the normalized optical density spectra of *THP-1 Cells (human monocyte cell)* infected with *C. pneumoniae*



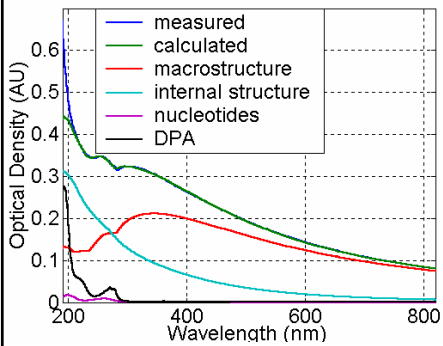
MOI = *Chlamydia cells/Host cells*, Efficiency of infection = 1% (Yamamoto)

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Quantitative Deconvolution

Bacillus globigii (spores)

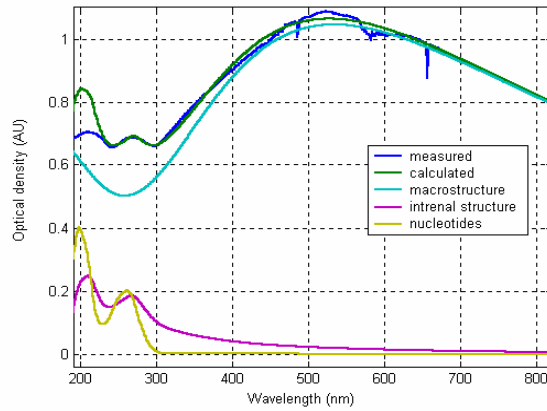


Volume, μm^3	0.782
Literature values	<1
Equivalent diameter, μm	1.14
DNA+RNA, g/cell	1.52×10^{-14}
DPA, %	6.22
(% dry cell (Murrell, 1969))	(5.6-13.55)
Fraction of internal structure, %	41.47
Average size internal structure, nm	132

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Quantitative Deconvolution of the Spectral Fingerprint of *Saccharomyces Cerevisiae* (Baker's Yeast)

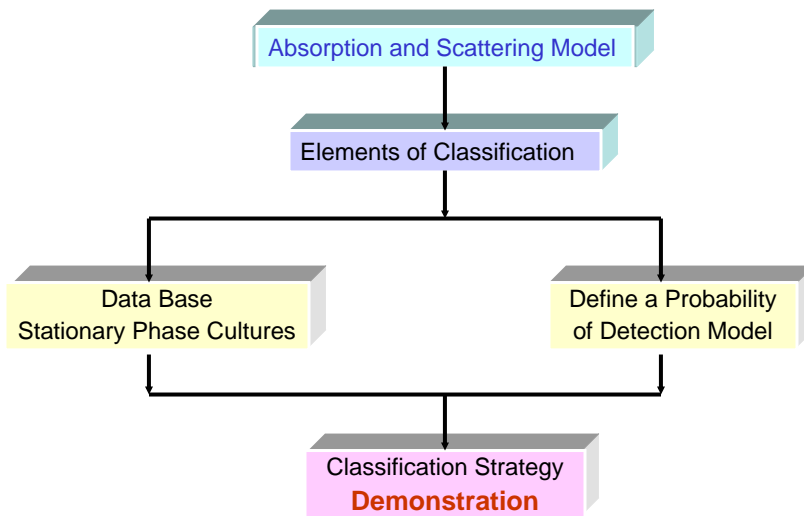


	Literature		Quantitative Deconvolution
	Diploid	Haploid	
Size (um)	4	5 x 6	5.011
DNA+RNA Content (pg/cell)	1.217	1.934	1.85

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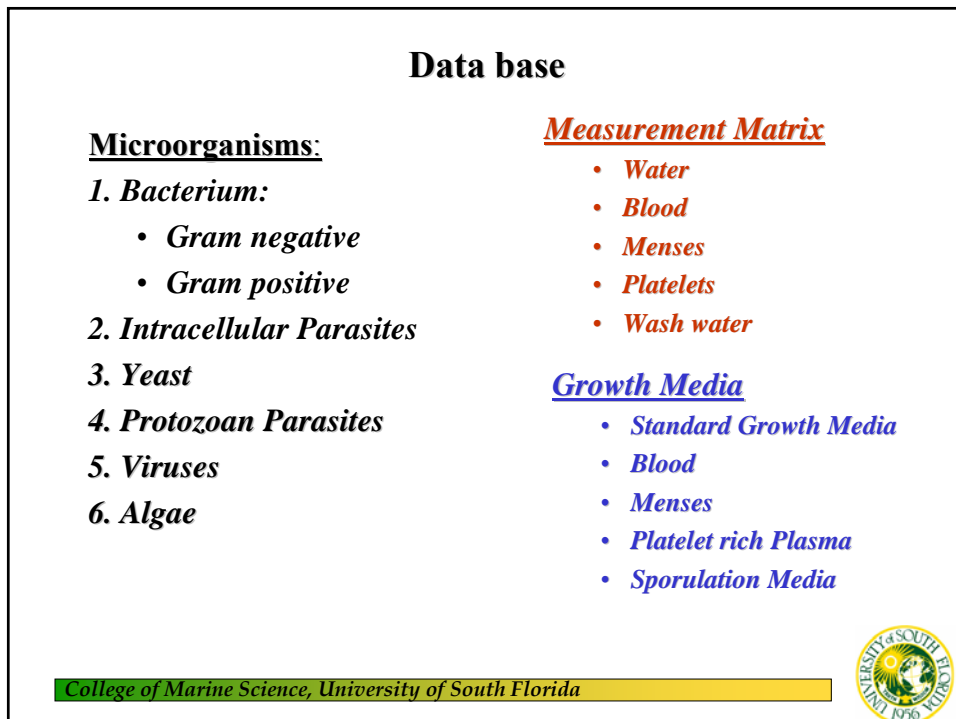
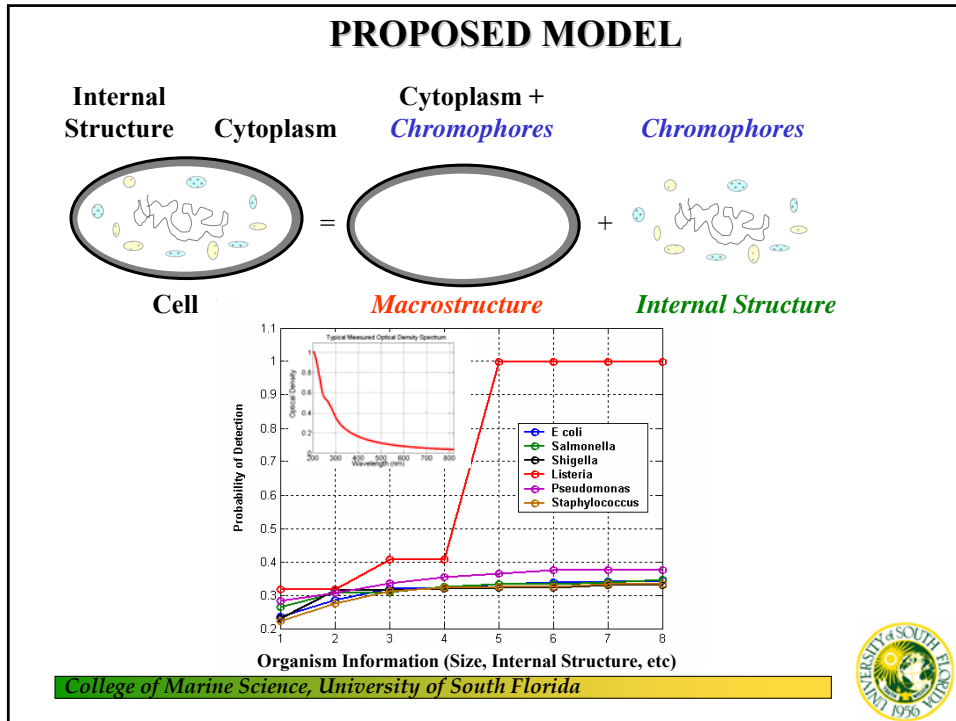


Schematic Representation for Classification Strategy



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Pathogens in data base

	Genus Name	Species Name	Strains	Comments
Gram Negative Bacteria	<i>Escherichia</i> +2	<i>coll</i>	10	+2
	<i>Pantoea</i> +	<i>agglomerans</i>	1	+
	<i>Salmonella</i>	<i>choleraesuis</i>	7	
	<i>Shigella</i>	<i>sonnei</i>	2	
	<i>Shigella</i>	<i>boydii</i>	1	
	<i>Shigella</i>	<i>flexneri</i>	1	
	<i>Pseudomonas</i> +	<i>aeruginosa</i>	1	+
	<i>Pseudomonas</i>	<i>fluorescens</i>	1	
Gram Positive Bacteria	<i>Enterococcus</i>	<i>faecalis</i>	1	
	<i>Staphylococcus</i>	<i>aureus</i>	6	
	<i>Staphylococcus</i>	<i>hominis</i>	1	
	<i>Staphylococcus</i> +2	<i>epidermidis</i>	1	+2
	<i>Listeria</i>	<i>monocytogenes</i>	3	
	<i>Listeria</i>	<i>seeligeri</i>	1	
	<i>Bacillus</i> +*	<i>subtilis</i>	1	+*
<i>Bacillus</i> +*	<i>anthracis Sterne</i>	1	+*	
Intercellular Parasites	<i>Chlamydia</i> 1	<i>pneumoniae</i>	1	1
Yeast	<i>Candida</i> 1,3	<i>albicans</i>	1	1,3
Protozoan Parasites	<i>Cryptosporidium</i>	<i>parvum</i>	1	
	<i>Giardia</i>	<i>muris</i>	1	
	<i>Giardia</i>	<i>lamblia</i>	1	
	<i>Plasmodium</i> 1	<i>vivax</i>	1	1
	<i>Plasmodium</i> 1	<i>falciparum</i>	1	1
Viruses	<i>Enterovirus</i>	<i>poliovirus</i>	1	
	Group B Arbovirus	<i>dengue</i>	1	
Algae	<i>Synechococcus</i>	<i>sp.</i>	1	
	<i>Karenia</i>	<i>brevis</i>	1	

+ denotes strains with vegetative growth curves
 * denotes strains with spore formation over time
 1. denotes strains grown in blood
 2. denotes strains grown in platelets
 3. denotes strains grown in menses

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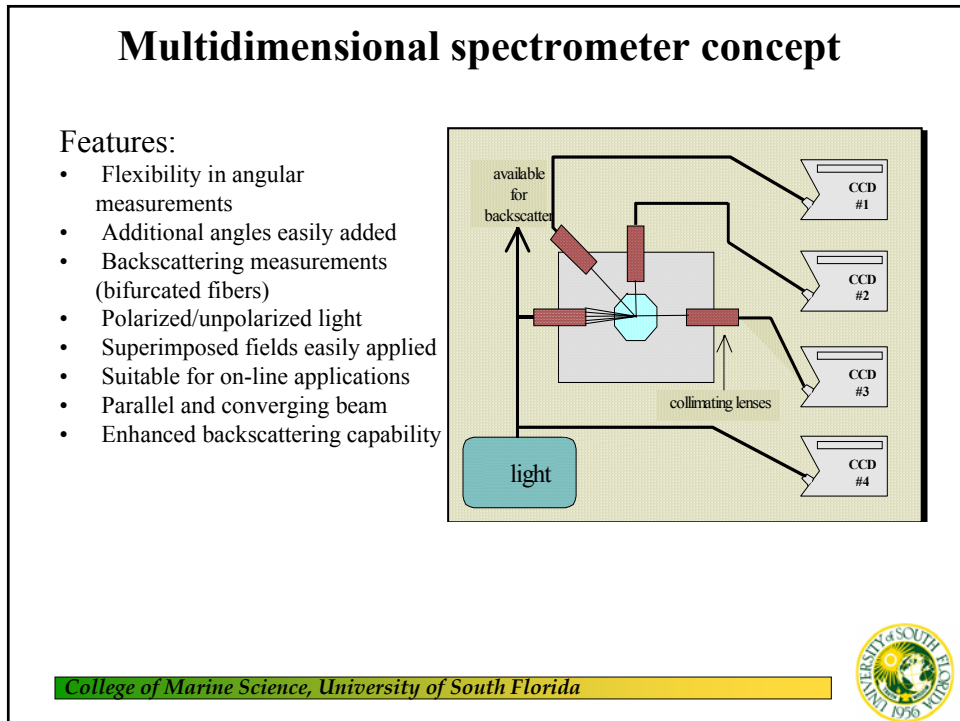
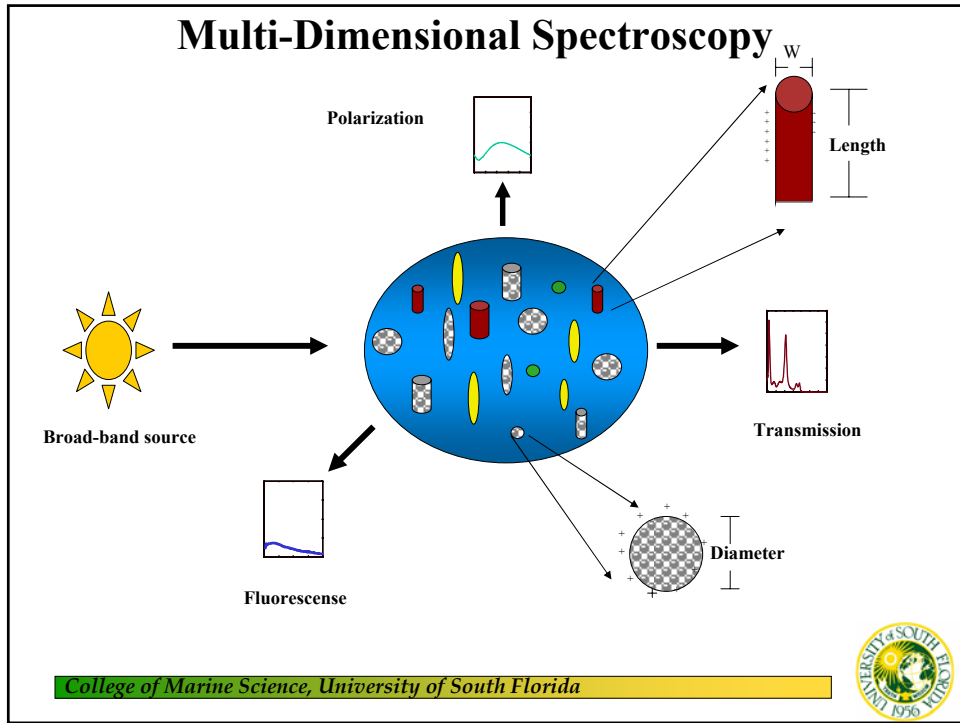


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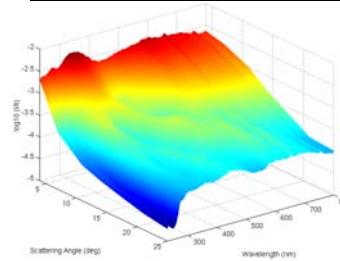
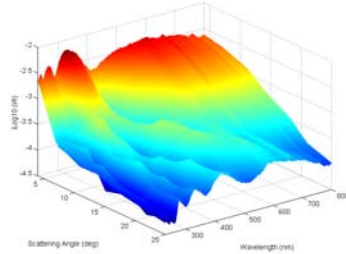
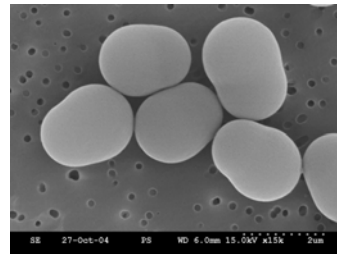
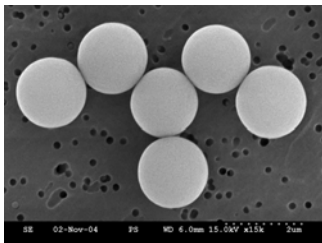
MAMW Prototype LANL



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UV-VIS MAMW RESPONSES: Particle Shape



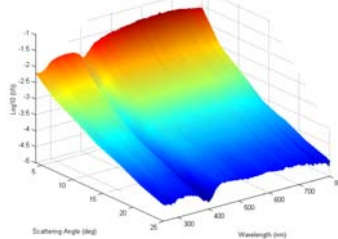
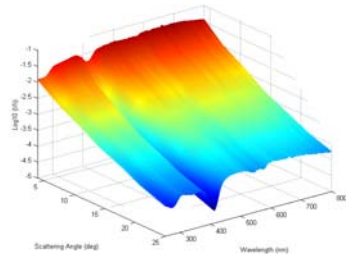
Spherical Particles: $D= 1.9 \mu\text{m}$

Peanut Shape Particles: $1.87 \mu\text{m}$

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UV-VIS MAMW RESPONSES: NORMAL & SICKLED RBC's



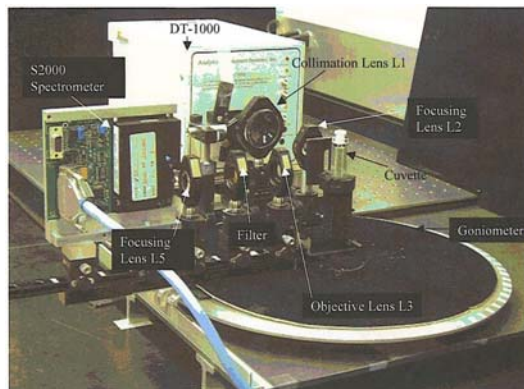
Normal whole blood sample

Sickled whole blood sample

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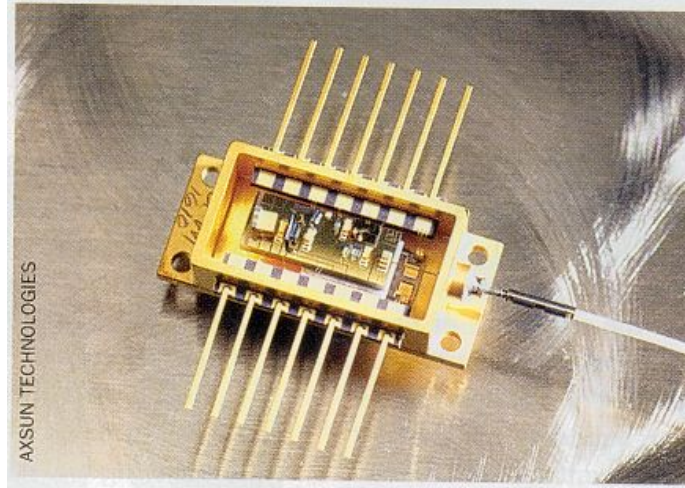
Photograph of the Fluorescence Measurement set up for the Multidimensional Spectrometer



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Miniaturized Spectrometers



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