

***Elemental Multi-variate Microbiological Analysis
(EMMA):
a Stand-alone Technology and a Sensor-fusion
Component for the Rapid Field Identification and
Classification of Biomaterials and Biological
Agents***

Steven J. Rehse Wayne State University, Detroit, MI

Andrzej W. Miziolek US Army Research Laboratory, APG, MD

Leslie M. Collins Duke University, Durham, NC

Peter A. Torrione Duke University, Durham, NC

WAYNE STATE
UNIVERSITY

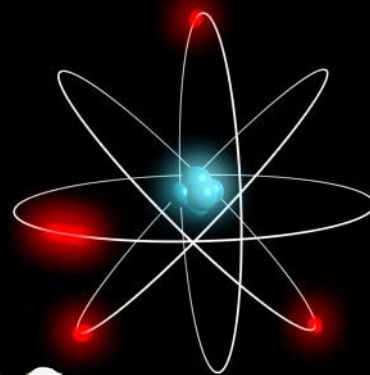
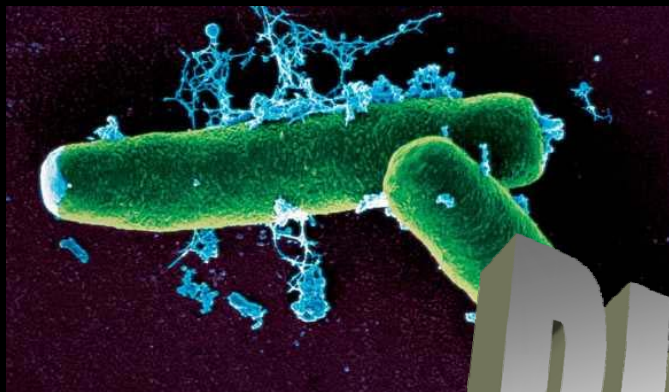


Duke UNIVERSITY

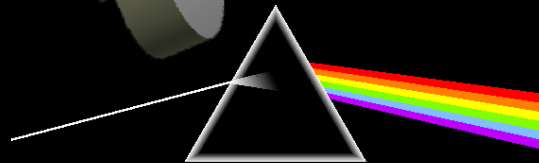
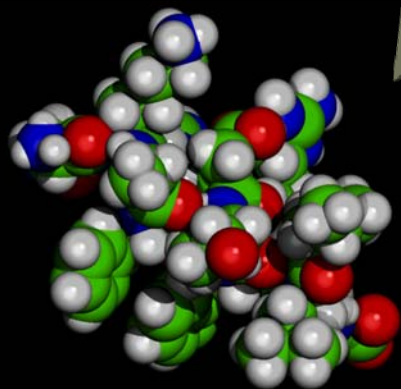


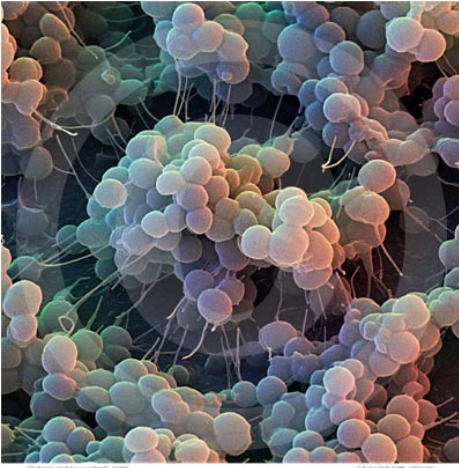
CBD S&T Conference, November 2009, Dallas, TX

The BIOMAS Project: Bacteria Identification by Optical, Molecular, and Atomic Spectroscopy

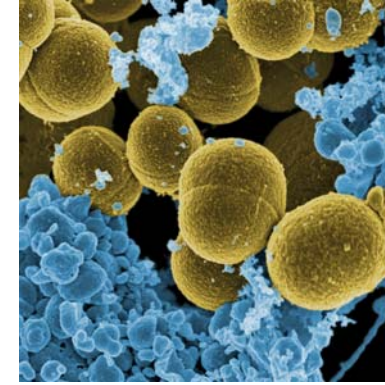


BIOMAS





Staph. epidermidis



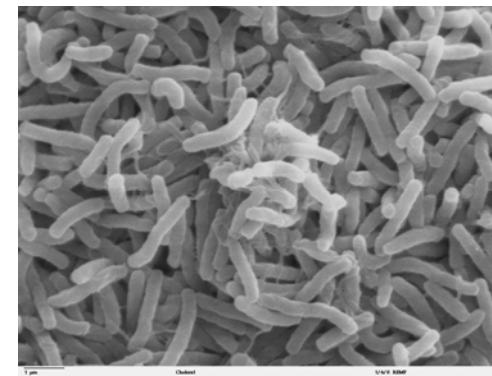
Staph. aureus

there is an urgent need right now in the military, civilian (hospital, food processing, environmental), and first responder communities for a “...rapid point-of-care (multiplex?) diagnostic for disease-causing pathogens.”

E. coli

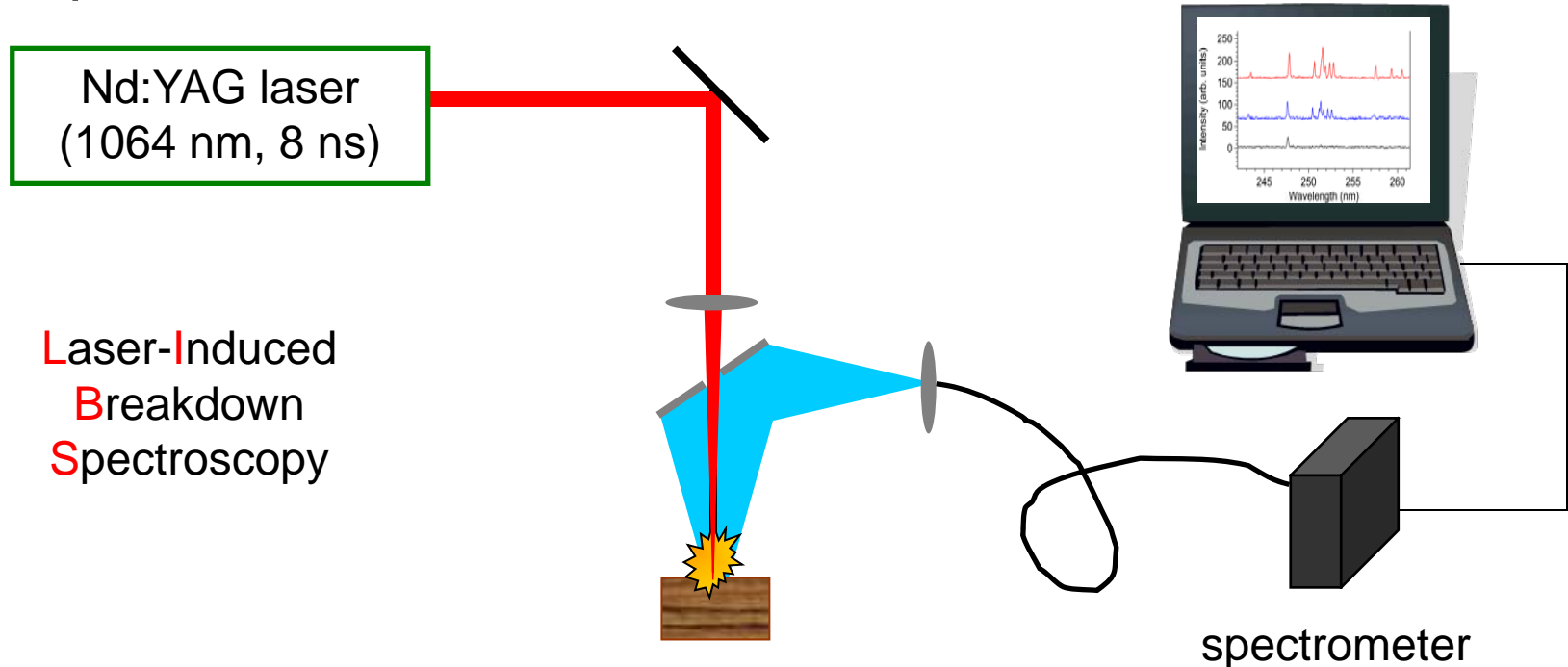


V. cholerae

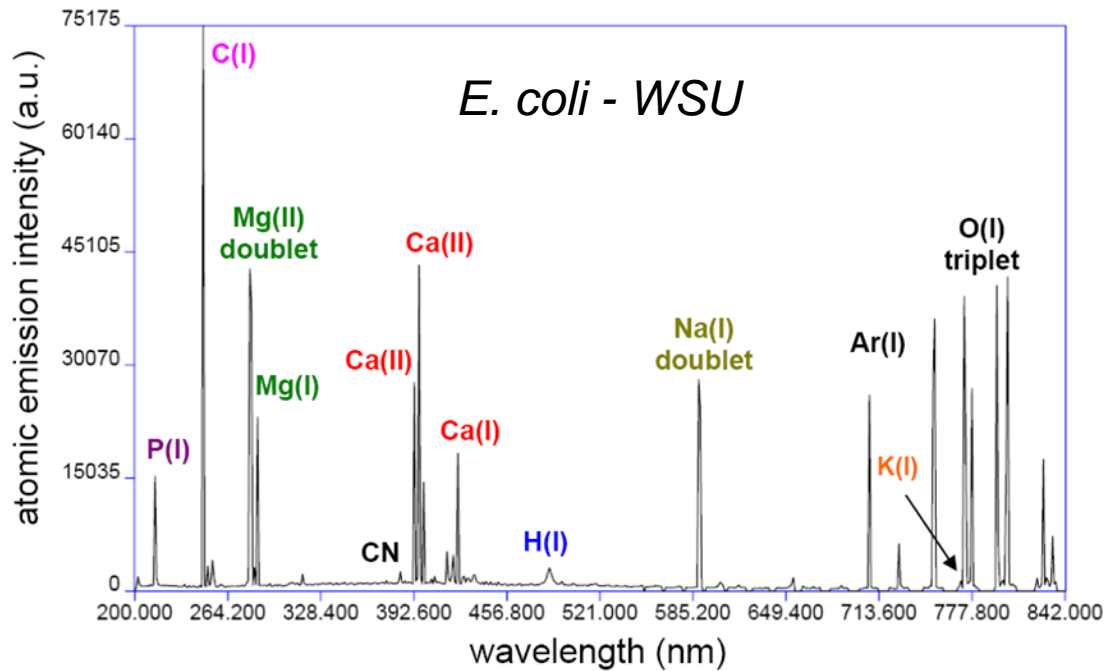


EMMA: Elemental Multivariate Microbiological Analysis

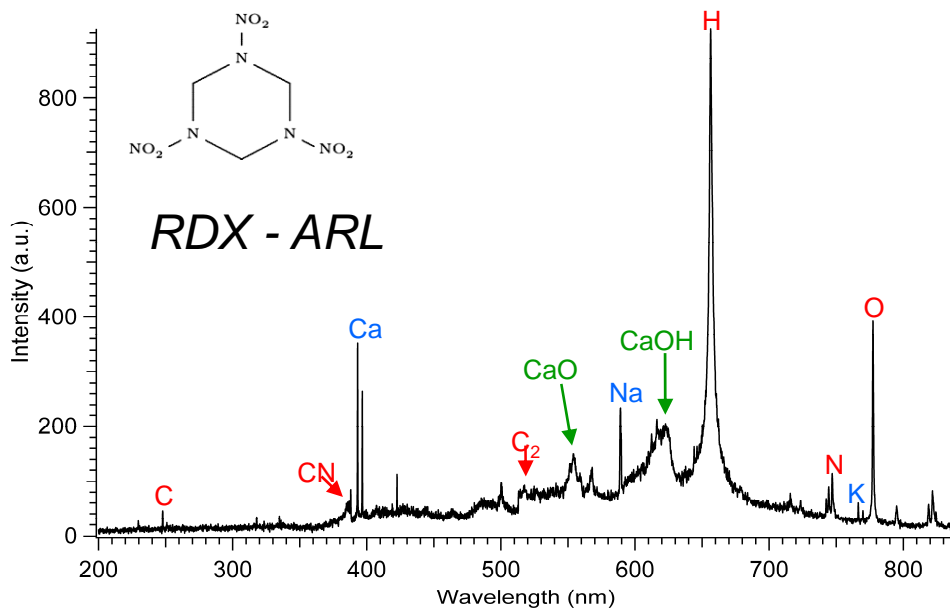
- utilizes laser-induced breakdown spectroscopy (LIBS) to measure the unique atomic or elemental composition of bacteria



LIBS Spectrum is like a Bar Code- Unique for Each Sample



- advanced signal-processing statistical techniques (“chemometrics”) classify/identify the unknown target on the basis of its unique atomic signature



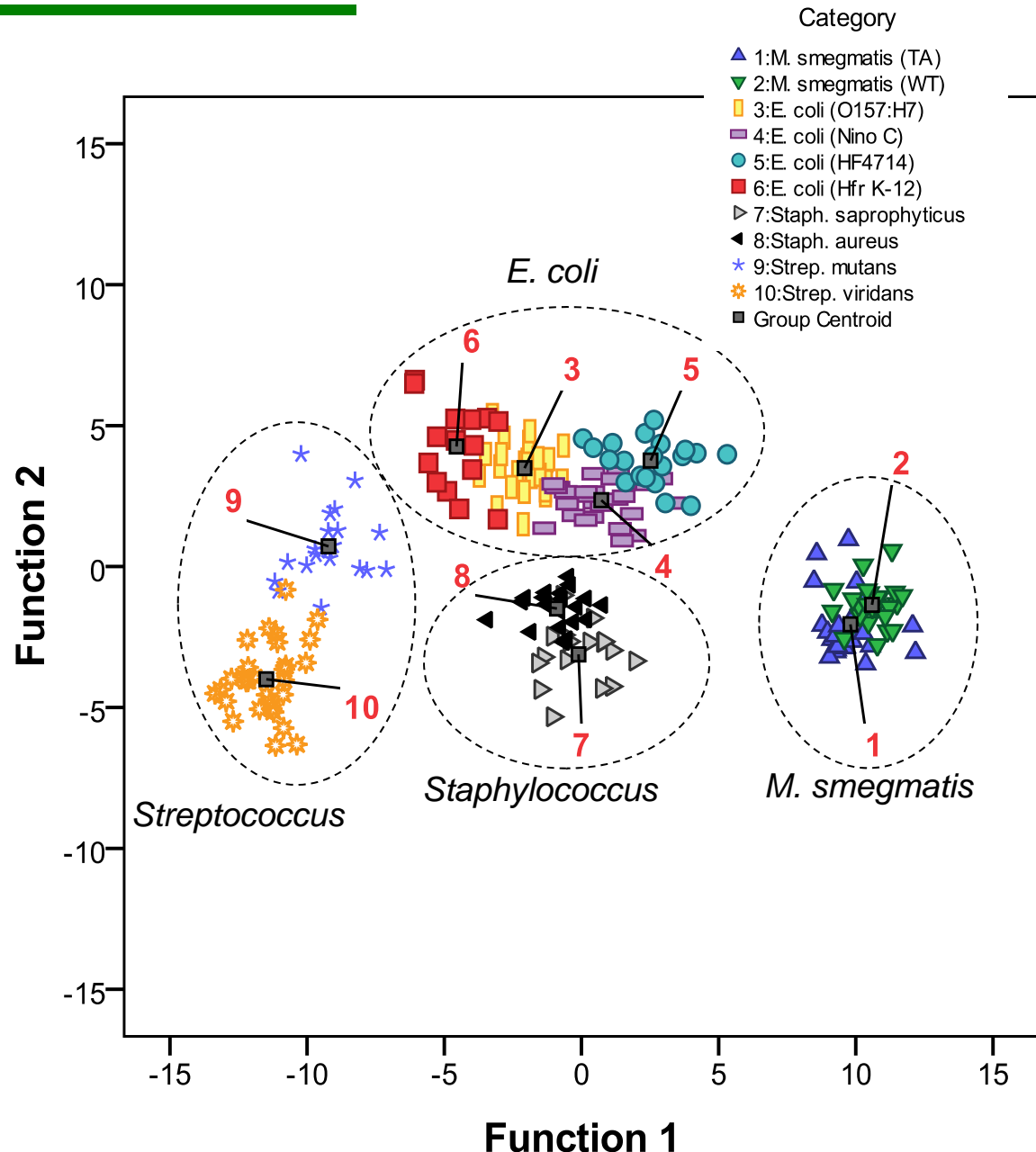
- concentrations of elements (or ratios of concentrations) become independent variables in a chemometric multivariate analysis

things that make EMMA *technology unique*

- speed / portability / durability (ruggedness)
 - “rapid point-of-care diagnostic...”
- lack of complicated sample preparation
- no expertise required
- no genetic or antigenic precursors (consumables) necessary
- same technology / hardware useful for explosives, chemical, other threats (CBRNE capable)
- capability of sensor fusion

Does it work? YES!

- Intensity of lines, ratios of intensities used in a statistical multi-variate analysis
- Discriminant function analysis (DFA)
 - Principal component analysis (PCA)
 - Partial least squares – discriminant analysis (PLS-DA)
 - Linear Discriminant Analysis (LDA)



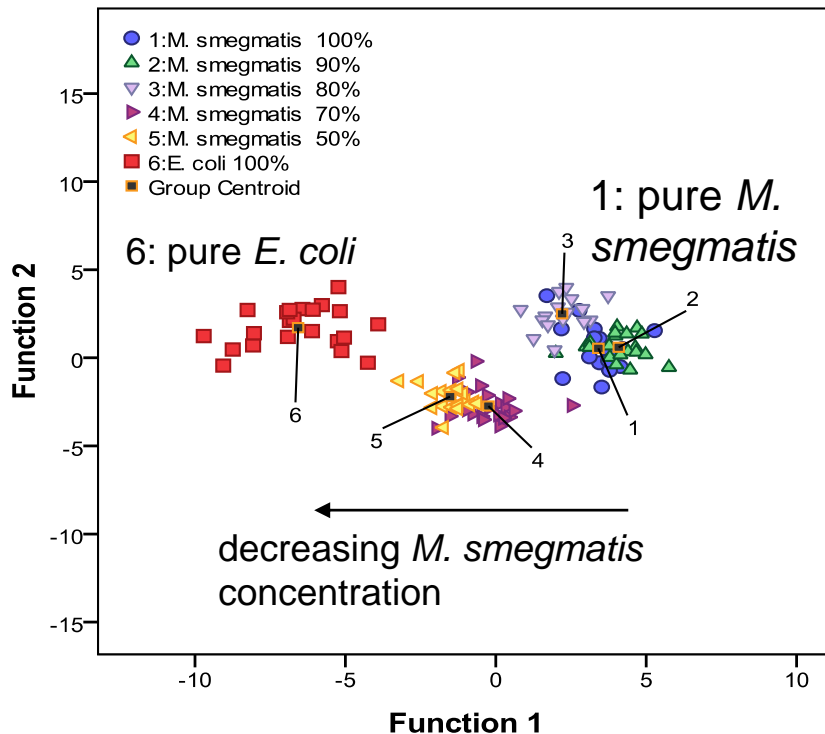
The Wayne State Team has already proven...

EMMA spectral fingerprint is:

- growth-medium independent
- independent of state of growth (how “old” the bacteria are)
- independent of whether the bacteria are live or dead
- obtainable even when other types of bacteria or contaminants are present (mixed samples)
- capable of strain discrimination
- obtainable from about 500 bacteria

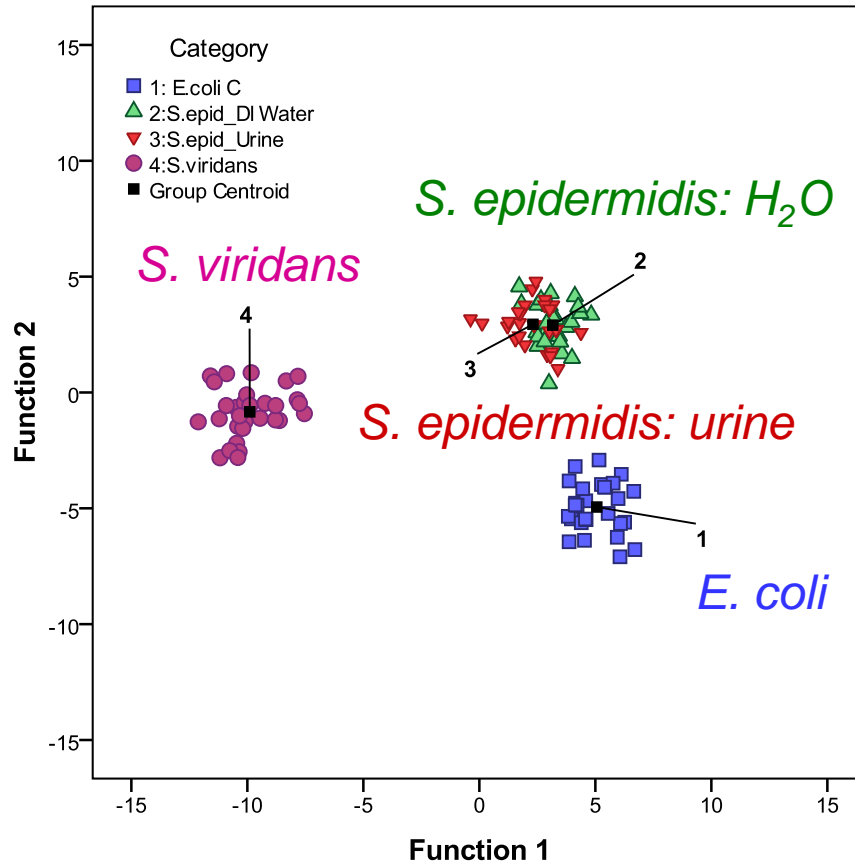
“Mixed” Samples

Category	# of Spectra	Classification Results		
		<i>M. smegmatis</i>	<i>E. coli</i>	<i>S. viridans</i>
100% <i>M. smegmatis</i> , 0% <i>E. coli</i>	21	100%	0%	0%
90% <i>M. smegmatis</i> , 10% <i>E. coli</i>	20	100%	0%	0%
80% <i>M. smegmatis</i> , 20% <i>E. coli</i>	16	100%	0%	0%
70% <i>M. smegmatis</i> , 40% <i>E. coli</i>	21	76%	34%	0%
50% <i>M. smegmatis</i> , 50% <i>E. coli</i>	19	47%	53%	0%
0% <i>M. smegmatis</i> , 100% <i>E. coli</i>	25	0%	100%	0%



- Mixtures of known mixing fraction were prepared from suspensions *M. smegmatis* and *E. coli* C.
- six separate mixtures were prepared with a ratio *M. smegmatis* to *E. coli* C given by $M_{1-x}:C_x$ with $x = 0.0, 0.1, 0.2, 0.3, 0.5, 1.0$.
- Multiple 1.5 mL tubes of these mixtures were prepared, thoroughly agitated via vortex mixing, then centrifuged for 3 minutes at 5000 rev/min.

“Dirty” samples



- Samples of *Staph. epidermidis* were prepared in DI water and sterile urine.
- Samples were collected and tested via EMMA with NO WASHING.
- EMMA spectral fingerprint from urine-originating bacteria were identical to water-originating bacteria.
- EMMA correctly classified 100% of the urine-originating bacteria as being consistent with *S. epidermidis*

Team

Wayne State University / Detroit Medical Center (micro samples, spectra)

Rehse, Palchaudhuri, Salimnia

Duke University (pattern recognition)

Collins, Torrione

ARL (enhancement, integration)

LIBS Group

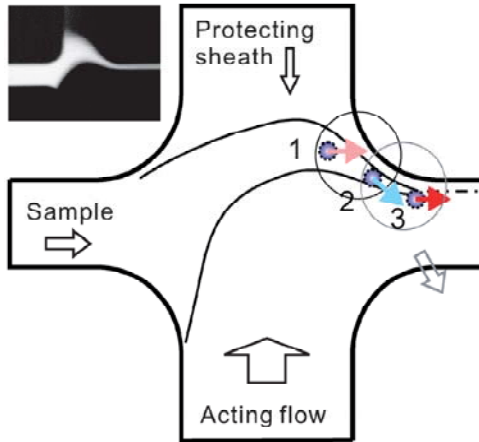
Translume, Inc. (sample prep. front end)

Haddock

Firm (device design and construction)

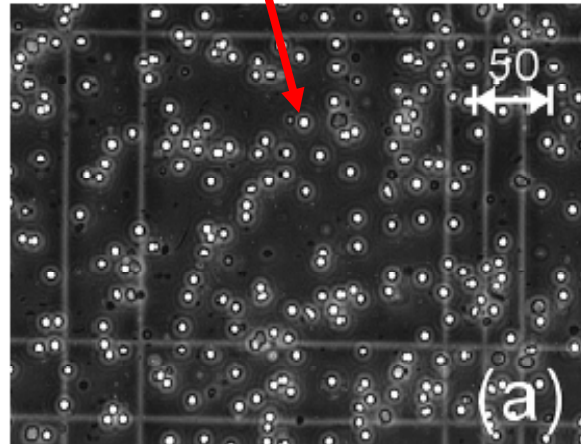
to be determined

Microfluidic separation/concentration



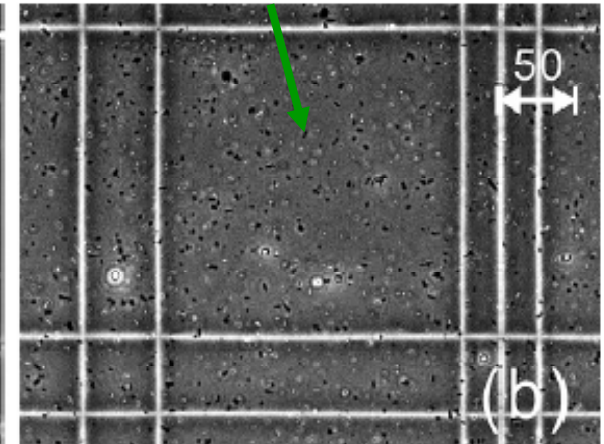
A "lab on a chip" microfluidic design by Wu et al. that has been shown to separate *E. coli* from red blood cells in human blood flow resulting in bacterial concentration

red blood cells



sample pre-separation

E. coli



sample after separation /
concentration

Soft inertial microfluidics for high throughput separation of bacteria from human blood cells

Zhigang Wu,^{*a} Ben Willing,^b Joakim Bjerketorp,^b Janet K. Jansson^{bc} and Klas Hjort^a

Lab Chip, 2009, **9**, 1193–1199

© The Royal Society of Chemistry 2009

Field-Portable Hardware

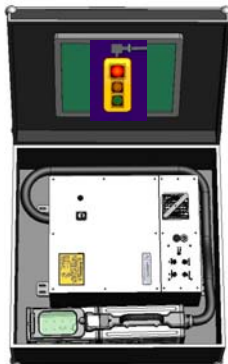
An example of current commercial LIBS hardware that operates on wallplug power



Applied Photonics Limited

Under development: battery operated for rapid field analyses

- eye-safe
- choice of sample chambers (smaller/larger)
- real-time chemometrics

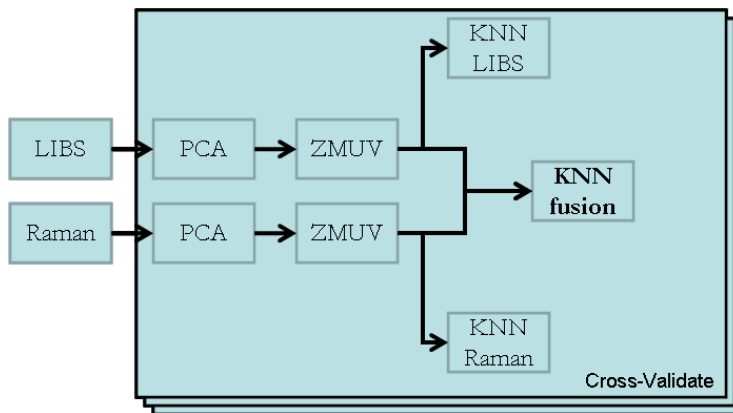


Sensor Fusion (with Raman)

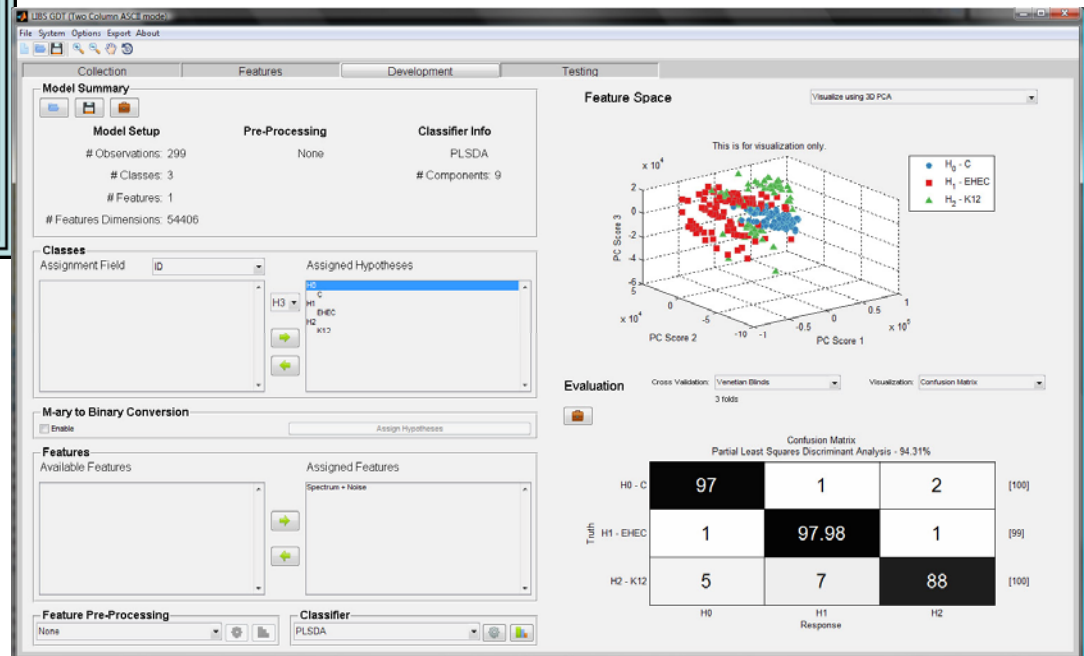
- Team at Duke are experts in real-time chemometric analysis
 - under US Army contract, delivered the very first truly “real-time” software for analysis of LIBS spectra (identified biological warfare simulant in one second!)
- Also working with us on LIBS/Raman data fusion

“Data Level” Sensor Fusion

- Flow chart below shows example of Data-Level-Fusion, explored for PCA, LDA, PLS on EMMA/Raman data from three *E. coli* strains.



New Folder Consulting



LIBS/Raman Data fusion with a PLS

- EMMA: 86%
(LIBS)

Libs PLS Confusion Matrix 86% Correct

	E Coli C	EHEC	Hfr K12
Truth			
E Coli C	98	0	2
EHEC	6	80	14
Hfr K12	8	11	81
	E Coli C	EHEC	Hfr K12
	Response		

Libs & Raman PLS Confusion Matrix 94% Correct

	E Coli C	EHEC	Hfr K12
Truth			
E Coli C	96	0	4
EHEC	3	92	5
Hfr K12	6	1	93
	E Coli C	EHEC	Hfr K12
	Response		

- Fusion: 94% correct
E. Coli identification

LIBS/Raman Data fusion with a LDA

- EMMA:91%
(LIBS)

Libs LDA Confusion Matrix 91% Correct

	E Coli C	EHEC	Hfr K12
Truth			
E Coli C	94	0	6
EHEC	4	94	2
Hfr K12	6	10	84
	E Coli C	EHEC Response	Hfr K12

- **Fusion: 100% correct *E. Coli* identification**

Libs & Raman LDA Confusion Matrix 100% Correct

	E Coli C	EHEC	Hfr K12
Truth			
E Coli C	100	0	0
EHEC	0	100	0
Hfr K12	0	0	100
	E Coli C	EHEC Response	Hfr K12

Conclusions

- All EMMA experiments to date have successfully shown the utility of LIBS to identify bacterial samples in a variety of growth conditions, in mixed samples, in dirty samples, etc.
- We are ready to move to testing real “clinical” type samples through our in-place organizational structure, which combines expertise in hardware development, software development, microbiological handling, and LIBS development
- Early result show LIBS can be combined with Raman for improved accuracy of identification: “sensor fusion.”