### "B is for Biological." Using a LIBS-based Elemental Microbiological Multivariate Analysis (EMMA) to detect bacteriological threats.

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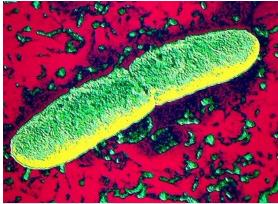


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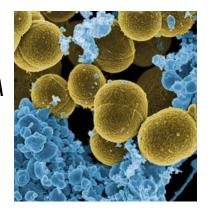




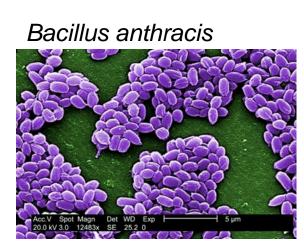
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Yersinia pestis

**MRSA** 



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."



<u>C</u>hemical <u>B</u>iological <u>R</u>adiological <u>N</u>uclear <u>E</u>xplosive Acinetobacter baumannii



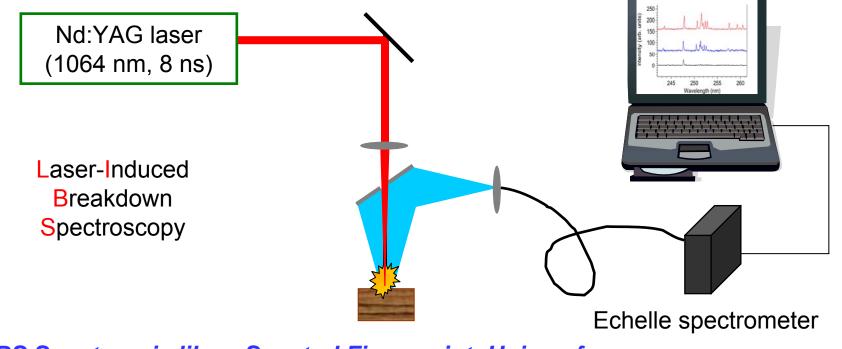
Due to certain well-recognized advantages, laserinduced breakdown spectroscopy (LIBS) is an attractive diagnostic candidate technology

- speed / portability / durability (ruggedness)
- 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



### <u>EMMA: Elemental Multivariate</u> <u>Microbiological Analysis</u>

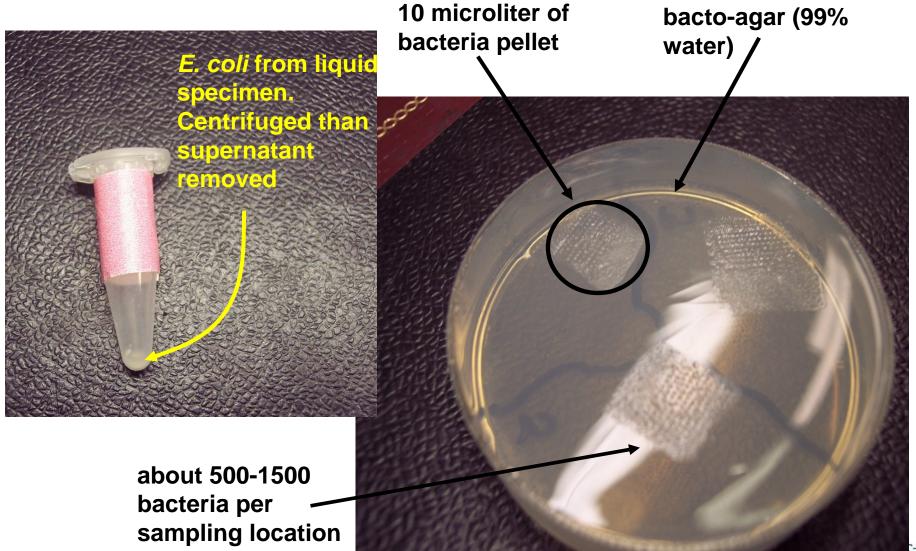
utilizing laser-induced breakdown spectroscopy (LIBS) to measure the unique atomic or <u>elemental</u> composition of bacteria



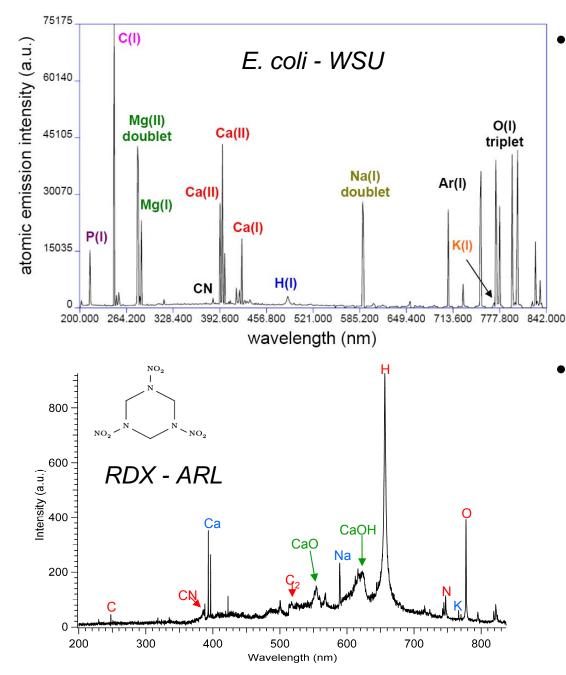
LIBS Spectrum is like a Spectral Fingerprint: Unique for Each Sample (courtesy of A. Miziolek)



### How we've been doing it...







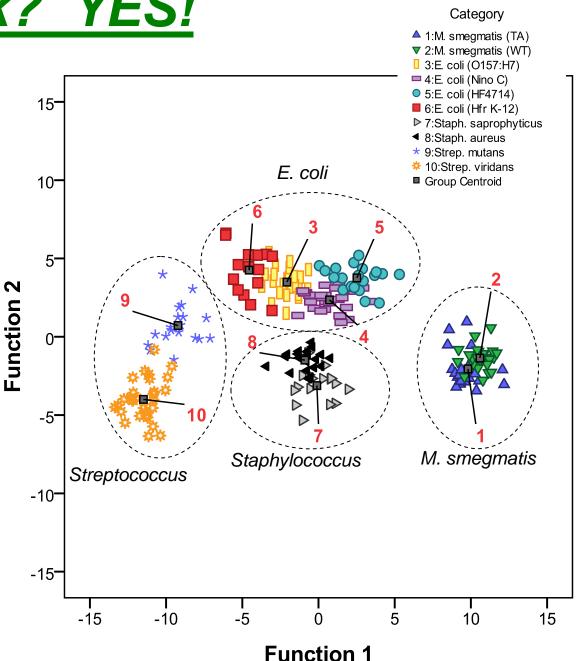
 high signal-to-noise atomic emission lines from inorganic elements allow a classification of the unknown target on the basis of its unique atomic spectrum

 concentrations of elements (or ratios of concentrations) become independent variables in a chemometric multivariate analysis (e.g. PCA, DFA, LDA, PLS-DA)



# • "Area under the curve" of 13 emission lines from 6 inorganic elements input as independent variables into a DFA.

 This test shows only the first two discriminant function scores for 10 different bacterial types (multiple genera, species, strains)



Crown	Predicted Group Membership (%)										
Group	1	2	3	4	5	6	7	8	9	10	
1: <i>M. smegmatis</i> (TA)	82.4	17.6	0	0	0	0	0	0	0	0	
2:M. smegmatis (WT)	28.0	72.0	0	0	0	0	0	0	0	0	
3: <i>E. coli</i> (O157:H7)	0	0	96.0	4.0	0	0	0	0	0	0	
4: <i>E. coli</i> (C)	0	0	3.6	96.4	0	0	0	0	0	0	
5: <i>E. coli</i> (HF4714)	0	0	0	0	100.0	0	0	0	0	0	
6 <i>:E. coli</i> (HfrK-12)	0	0	6.7	0	0	93.3	0	0	0	0	
7:Staph. saprophyticus	0	0	0	0	0	0	94.1	5.9	0	0	
8:Staph. aureus	0	0	0	0	0	0	0	100.0	0	0	
9:Strep. mutans	0	0	0	0	0	0	0	0	95.0	5.0	
10:Strep. viridans	0	0	0	0	0	0	0	0	0	100.0	

<u>The Wayne State Team has</u> <u>already demonstrated...</u>

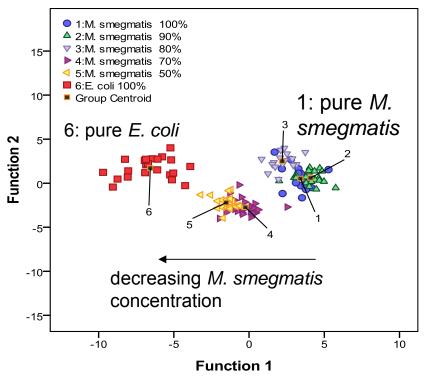
EMMA spectral fingerprint is:

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

6 publications in Applied Physics Letters, Journal of Applied Physics, Applied Optics, and Spectrochimica Acta B

### "Mixed" Samples

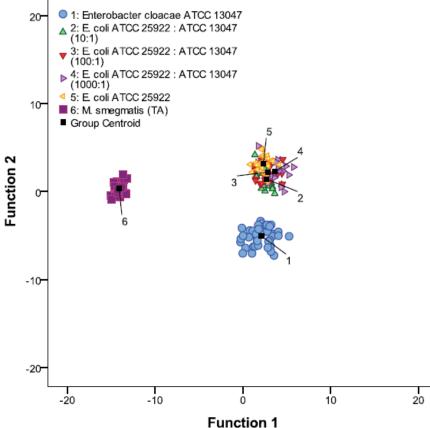
Category	# of Spectra	Classification Results					
		M. smegmatis	E. coli	S. viridans			
100% M. smegmatis, 0% E. coli	21	100%	0%	0%			
90% M. smegmatis, 10% E. coli	20	100%	0%	0%			
80% M. smegmatis, 20% E. coli	16	100%	0%	0%			
70% M. smegmatis, 40% E. coli	21	76%	24%	0%			
50% M. smegmatis, 50% E. coli	19	47%	53%	0%			
0% M. smegmatis, 100% E. coli	25	0%	100%	0%			



- Six separate mixtures of known mixing fraction were prepared from suspensions *M. smegmatis* and *E. coli* C.
- As long as the majority bacterium comprised 80% of the mixture, we saw 100% identification.



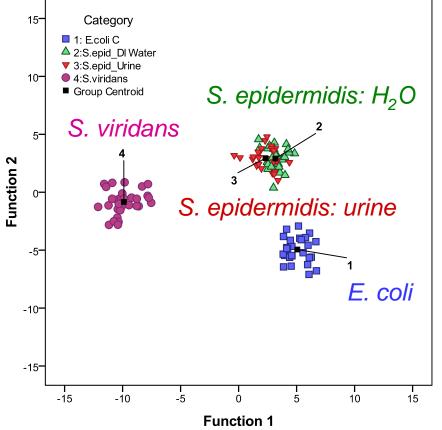
### "Mixed" Samples



- Mixtures of known mixing fraction were prepared from suspensions *E. coli* C and *E. cloacae*.
- Mixing represent "clinical" contaminations and/or mixtures (i.e. 10:1, 100:1, 1000:1).



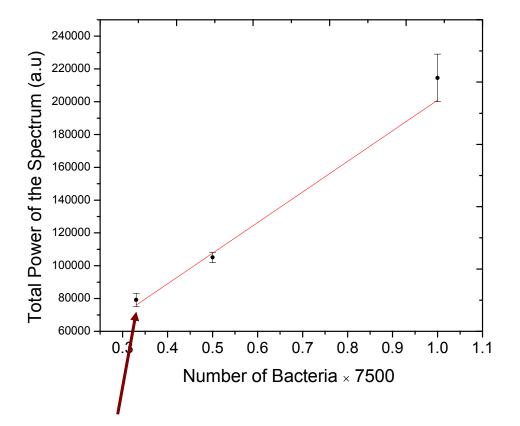
### "Dirty" clinical samples



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



### LIBS intensity linearly dependent on number of bacteria

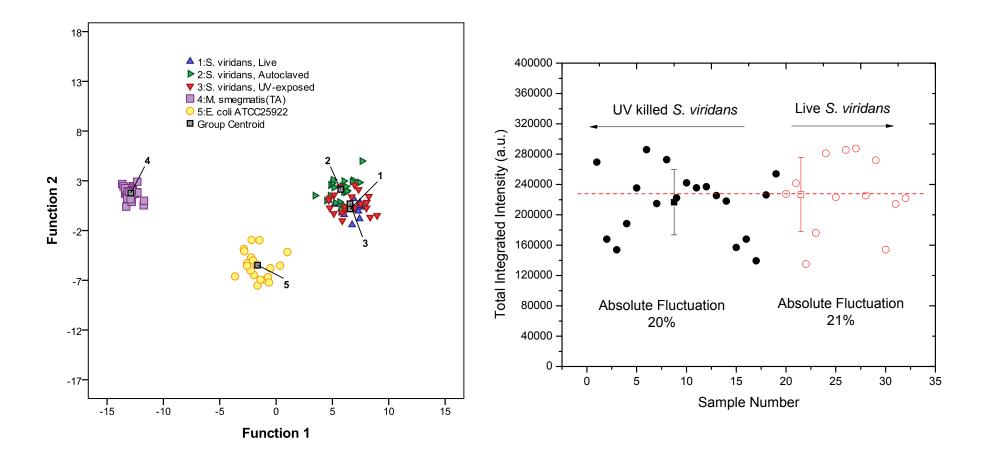


5 laser sampling locations~500 bacteria per locations

- Samples of *E. coli* with different titer tested on agar.
- Each data point is the average of 5 sampling locations.
- As expected, spectra demonstrate a linear dependence with cell number.
- All spectra were 100% correctly identified (specificity <u>not</u> dependent on number of cells).
- Suggests an antibiotic resistance test?

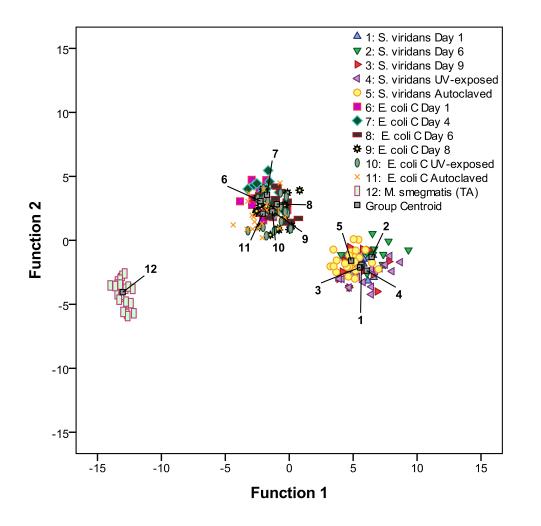


### LIBS specificity and sensitivity not dependent on bio-activity of the bacteria



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### LIBS specificity and sensitivity not dependent on bio-activity of the bacteria



- Two species of bacteria tested
- All specimens prepared separately and left to sit on a nutrient-free medium for up to 9 days at room temperature
- This graph also includes the UV-irradiated and the autoclaved specimens
- All species 100% accurately identified



## Strain discrimination confirmed by others...

#### The Use of Laser-Induced Breakdown Spectroscopy for Distinguishing Between Bacterial Pathogen Species and Strains

ROSALIE A. MULTARI,\* DAVID A. CREMERS, JOANNE M. DUPRE, and JOHN E. GUSTAFSON

Applied Research Associates, Inc., 4300 San Mateo Blvd NE Suite A-220, Albuquerque, New Mexico 87110 (R.A.M., D.A.C.); and Department of Biology, New Mexico State University, P.O. Box 30001, Las Cruces, New Mexico, 88003-8001 (J.M.D., J.G.)

APPLIED SPECTROSCOPY Volume 64, Number 7, 2010

- 100% accuracy exhibited in blind trials of 4 MRSA strains and one *E. coli* strain
- lyophilized ("freeze-dried") specimens used



### We Must Proceed, and Faster...

LIBS research must proceed along two equally important avenues:

- fundamental research to explore the microbiological diversity that can occur in specimens
- specimen preparation and handling protocols and techniques to isolate pathogens from contaminants of biological origin

NOTE: we do NOT need to fingerprint hundreds and hundreds of "new" bacteria



# what must we do to make LIBS a clinical tool?

Develop hardware and protocols for clinical sample testing (blood, urine, sputum)

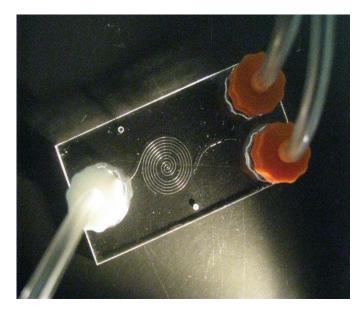
- isolation
- concentration under the laser focus



### <u>solutions</u>

- 1. differential centrifugation
- 2. filtration (sequential?)
- 3. optical trapping / separation
- 4. microfluidic separation
- 5. antibody isolation/phage display technology (consumables!)

### Microfluidic separation/concentration (Translume, Inc. Ann Arbor, MI)



hydrodynamic (microfluidic) separation of heavier cells

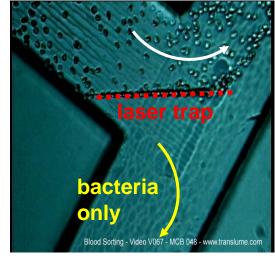
from lighter cells

monolithically fabricated devices in glass

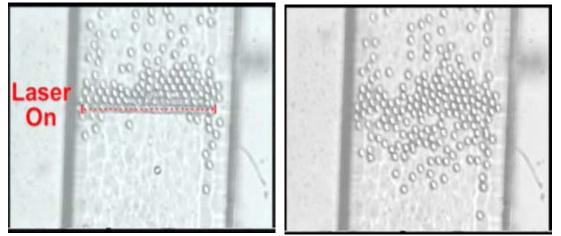


### Microfluidic separation/concentration (Translume, Inc. Ann Arbor, MI)





optical trap-based separation of heavier cells from lighter cells







- 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.



#### My students



Thank you!

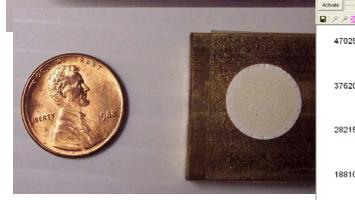


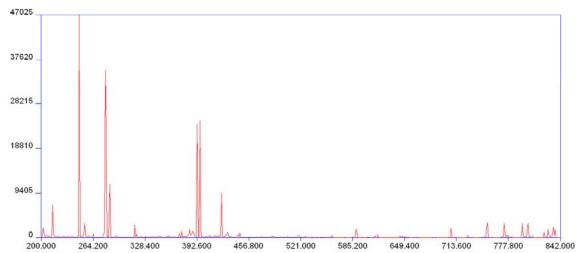


### **Novel substrates 1**



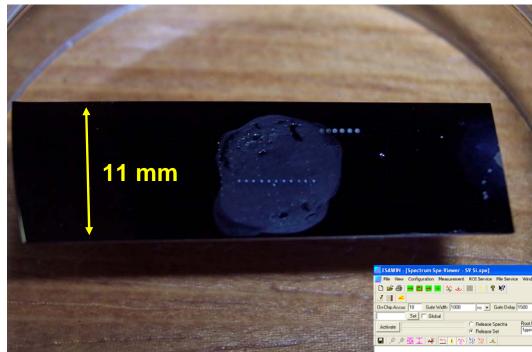
- 10 mL of a suspended bacterial culture pushed through a 0.22 or 0.44 µm cellulose (carbon) Millipore filter
- alternately, bacteria just deposited on filter (wicking)
- C line does "contaminate" spectrum, but only at 7% level (same as agar!)





ns 
Amplific. 2500 Apply

### **Novel substrates 2**



- Acid etched "porous" silicon
- Bacteria fixed with polyacrimide
- High SNR LIBS spectrum
- Si lies do not contaminate spectrum

