Laser-Induced Breakdown Spectroscopy (LIBS) for the Rapid Identification and Classification of Pathogenic Bacteria

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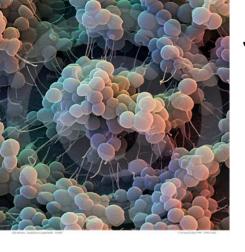


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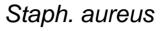
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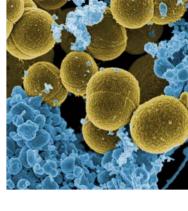






Staph. epidermidis



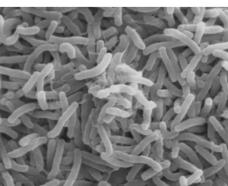


there is an urgent need right now in the military, civilian (hospital, food processing, environmental), and first responder communities for a "...rapid point-of-care (multiplex?) diagnostic for disease-causing pathogens."

E. coli



V. cholerae



How do we identify bacteria?

4 ways

- genetic
- serological (antigenic)
- microbiological
- compositional
 - LIBS
 - Raman
 - MALDI-TOF-MS



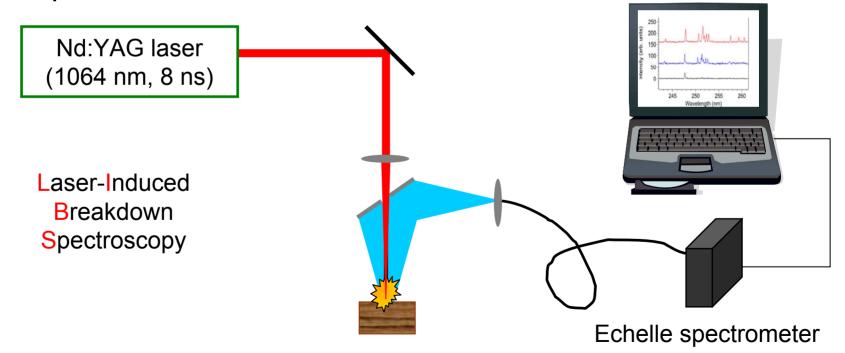
things that make a LIBS-based technology powerful

- 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



EMMA: Elemental Multivariate Microbiological Analysis

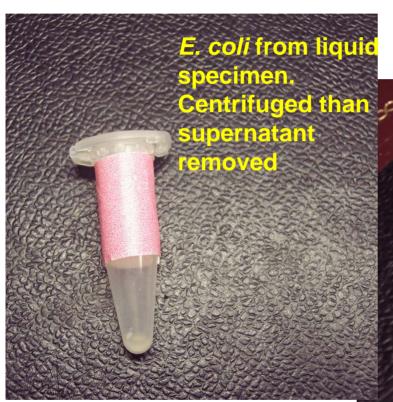
 utilizes laser-induced breakdown spectroscopy (LIBS) to measure the unique atomic or <u>elemental</u> composition of bacteria



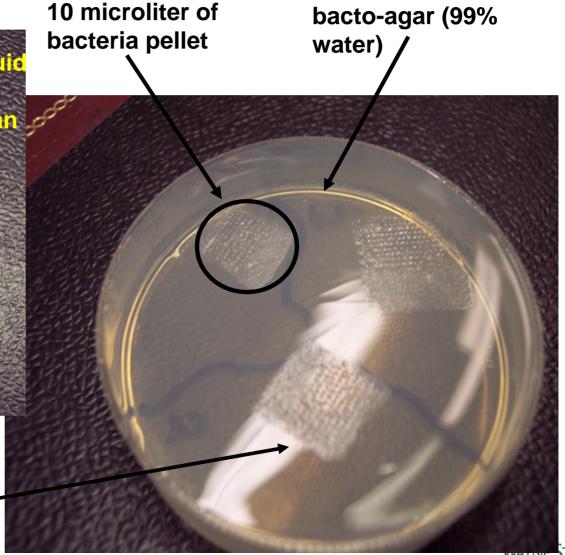
LIBS Spectrum is like a Spectral Fingerprint: Unique for Each Sample

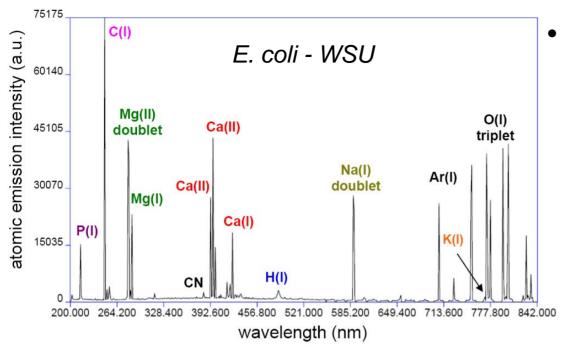


how we did it...

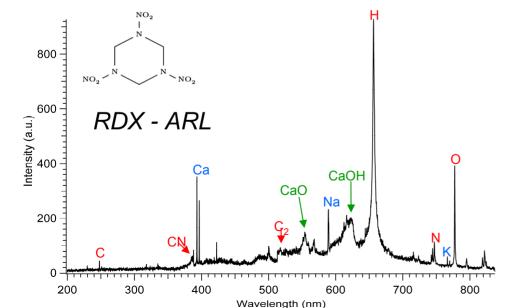


about 500-1500 bacteria per — sampling location





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

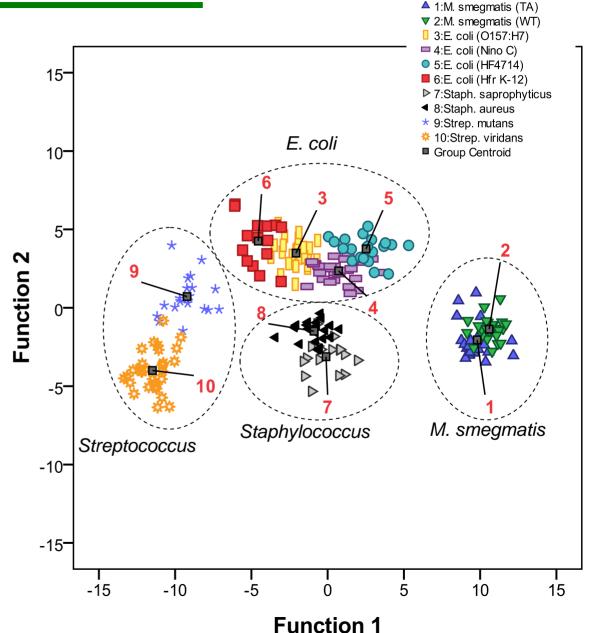


concentrations of elements (or ratios of concentrations) become independent variables in a chemometric multivariate analysis (e.g. PCA, DFA, LDA, PLS-DA)



Does it work? YES!

- "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)



Category

Group	Predicted Group Membership (%)										
	1	2	3	4	5	6	7	8	9	10	
1:M. smegmatis (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.</i> coli (HF4714)	0	0	0	0	100.0	0	0	0	0	0	
6:E. coli (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	



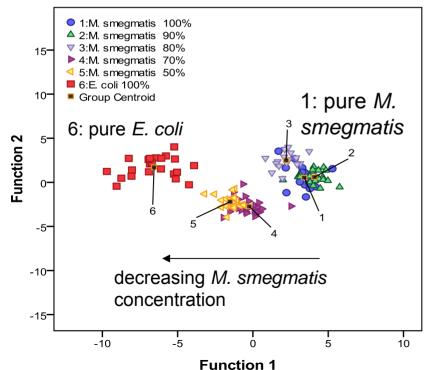
The Wayne State Team has already demonstrated...

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

"Mixed" Samples

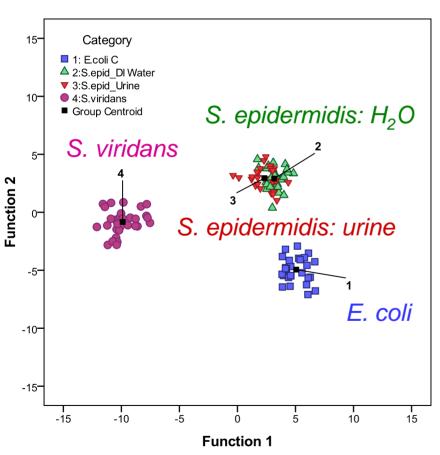
Category	# of Spectra	Classification Results					
Category	н от Бресиа	M. smegmatis	E. coli	S. viridans			
100% M. smegmatis, 0% E. coli	21	100%	0%	0%			
90% M. smegmatis, 10% E. coli	20	100%	0%	0%			
80% M. smegmatis, 20% E. coli	16	100%	0%	0%			
70% M. smegmatis, 40% E. coli	21	76%	24%	0%			
50% M.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



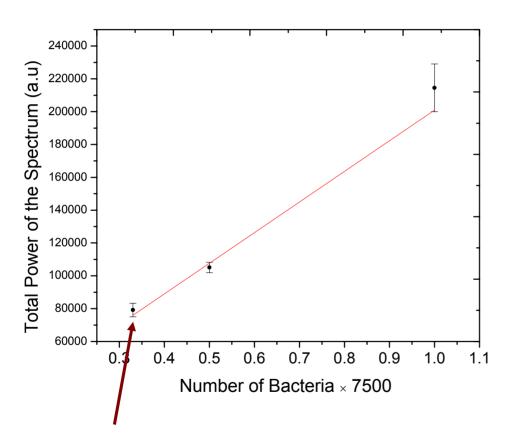
"Dirty" samples



- 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?

Strain discrimination confirmed by others...

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

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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

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- 100% accuracy exhibited in blind trials of 4 MRSA strains and one *E. coli* strain
- Iyophilized ("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?

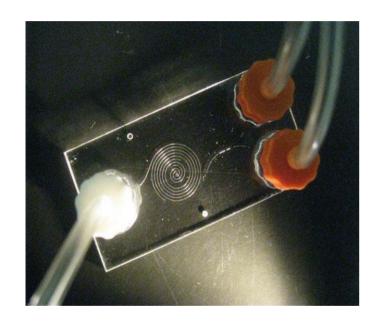
Develop protocols for clinical sample preparation (blood, urine, sputum)

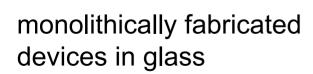
- isolation
- concentration under the laser focus

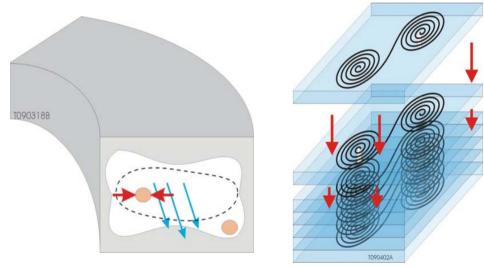
solutions

- 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)







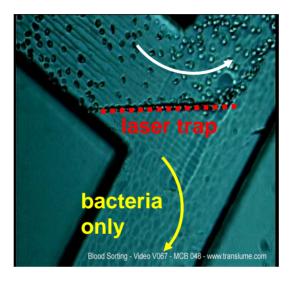
hydrodynamic (microfluidic) separation of heavier cells from lighter cells



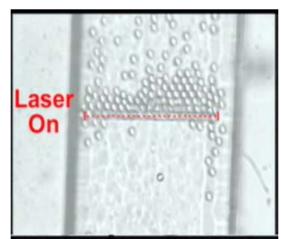
Microfluidic separation/concentration

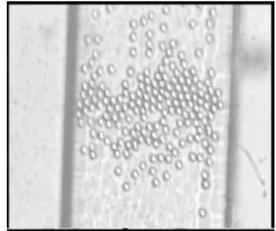
(Translume, Inc. Ann Arbor, MI)





optical trap-based separation of heavier cells from lighter cells



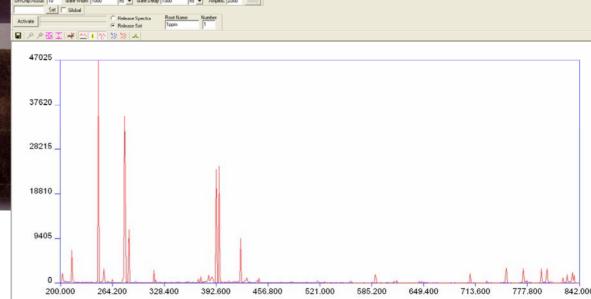




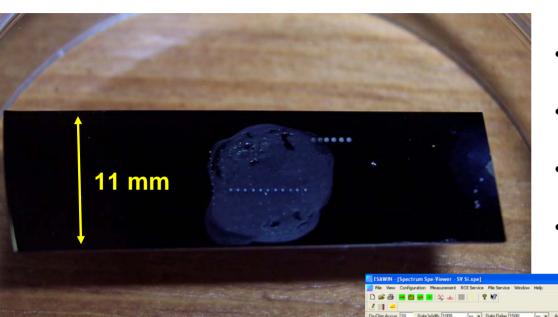
Novel substrates 1



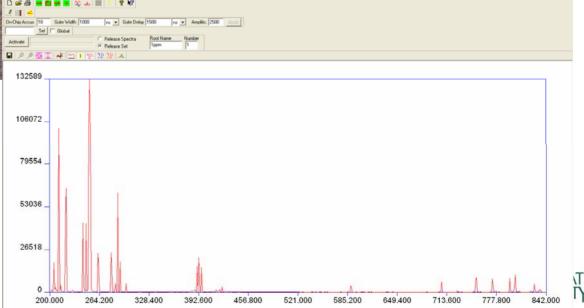
- 10 mL of a suspended bacterial culture pushed through a 0.22 or 0.44 µm cellulose (carbon) Millipore filter
- alternately, bacteria just deposited on filter (wicking)
- C line does "contaminate" spectrum, but only at 7% level (same as agar!)



Novel substrates 2



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



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.



My students

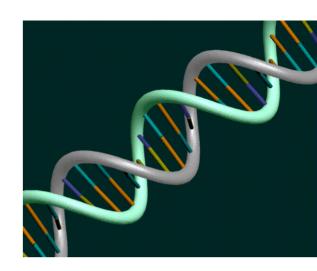


Thank you!



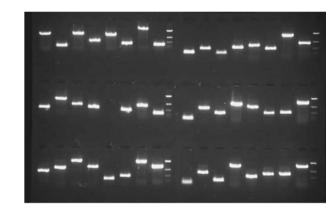
genetic

- PCR (polymerase chain reaction)
- (random primed) RAPID-PCR
- FISH (fluorescence in situly hybridization)



requires

 a priori knowledge of genetic sequence (16s RNA gene is conserved in most)



drawbacks

- amplification time (multiple generations needed)
- nonspecific reactivity
- still need to do gel electrophoresis
- very contamination sensitive



<u>serological</u>

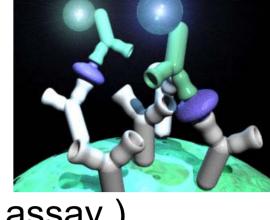
- immunoassays
- microwell devices
- ELISA (enzyme-linked immunosorbent assay)
- fluorescently labeled antibody techniques
- MEMS

requires

 a priori knowledge of serology (surface antigens)

drawbacks

- any mutation (common) undetectable
- antibodies are not stable (shelf-life)
- consumables
- binding affinities may be low





<u>microbiological</u>

- culturing and colony counting
- phenotyping
- sensitivity to immunochemicals
- Gram staining

requires

- time
- expertise
- LOTS of supplies
- a priori clinical knowledge (case-history)

drawbacks

- slow/labor intensive
- requires experts



