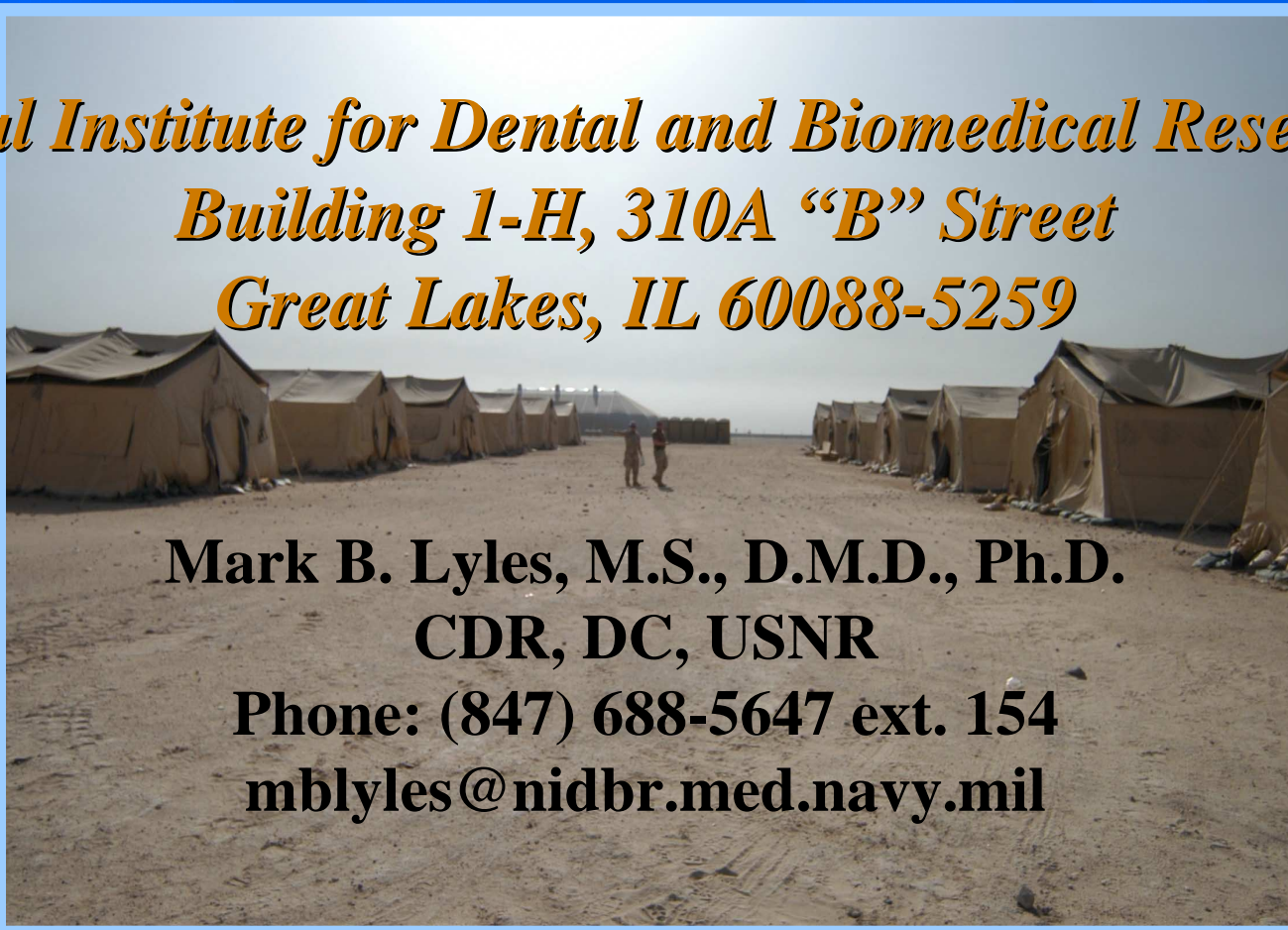


# **The Physical, Biological and Chemical Characterization of Micro-particulates from the Middle East**

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# Acknowledgements

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DN Joe Noble

HM1 Steve Kaufman

DT1 Brad Noble

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## **Corp of Engineers**

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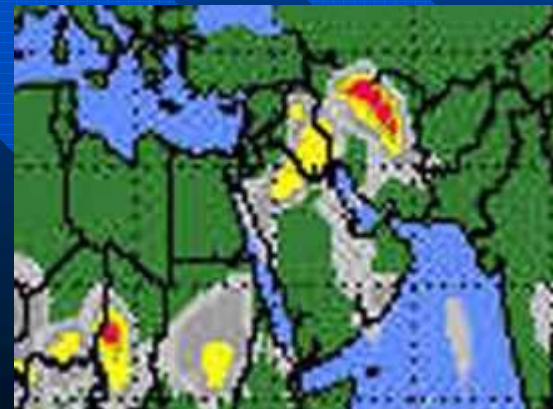
Dr. Herb Frederickson

Dr. Tony Bednar

Dr. Lillian Wakeley

Mr. John Furey

Ms. M. Richmond





# Background

- Dust sample arrives from Iraq October 2003 for ADAL project.
- Scanning Electron Micrograph and Fluorescent Microscopy analysis of dust sample indicates: 1) significant micro-particulate concentration (<20  $\mu\text{m}$ ), 2) micro-particulates are not crystalline or are coated, and 3) particles exhibit unique physical properties.
- Pneumonia and other respiratory health problems among US military personnel in the US Central Command Area of Responsibility (USCENTCOM AOR). Significant post-operative infection rate among U.S. and coalition wounded.
- Exposure of airborne micro-particulates, especially during duststorms, is ubiquitous for all coalition forces.
- Information passed to ONR which funds a pilot project.

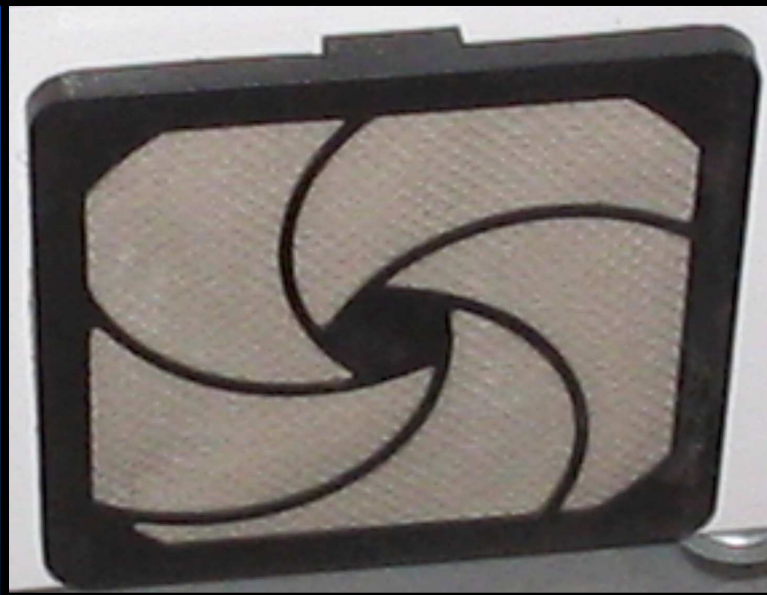
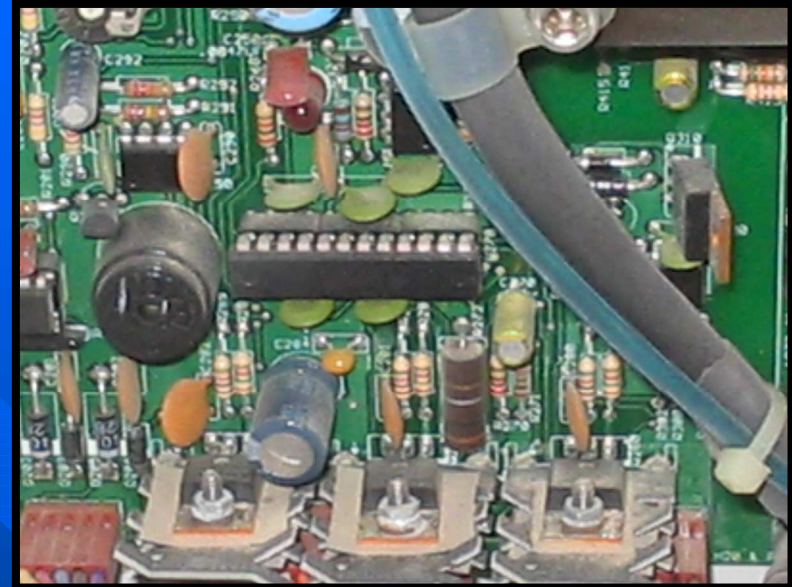
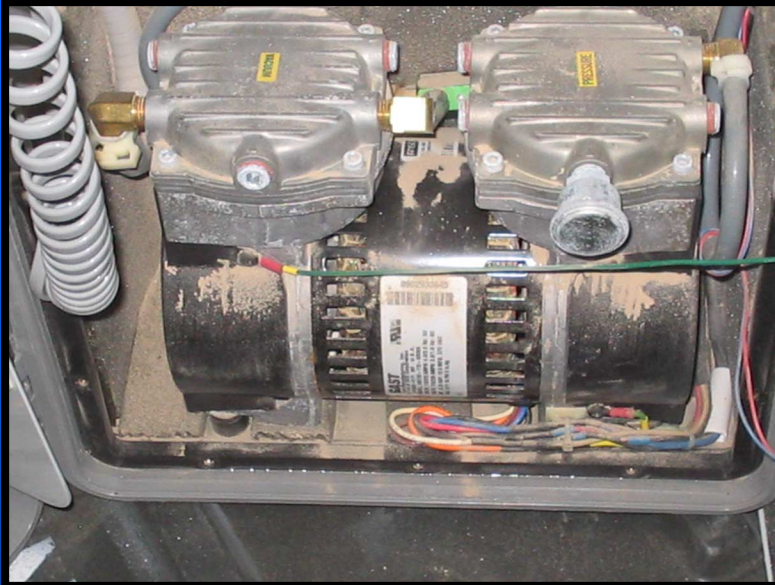


# “The Battlespace Environment” to the ‘Warfighter’ this means:



- ❖ The “terrain”, the battlefield *landscape*
- ❖ Living *on* the “dirt”
- ❖ Traveling *across* the “dirt”
- ❖ Fighting *in* the “dirt”
- ❖ In Iraq this also means *breathing* the “dirt”

# Micro-particulate Exposure

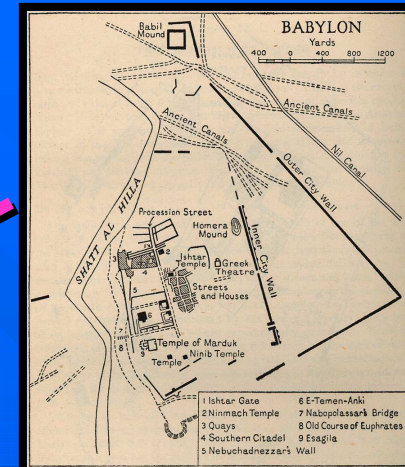


# Specific Aims

- ✓ Scientifically collect and process sand/dust samples for transport back to CONUS for further testing.
- ✓ Characterize the sand/dust environment.
- ✓ Physically characterize samples as to particle size distribution and concentration, and other physical qualities.
- ✓ Chemically analyze samples based on size fraction as to elemental content and concentration as well as poly-anion and cation identification.
- ✓ Biologically characterize samples based on size fraction as to anaerobic/ aerobic bacterial populations, yeasts, fungi, and viruses. Special identification of known pathogens.



# Sampling Locations



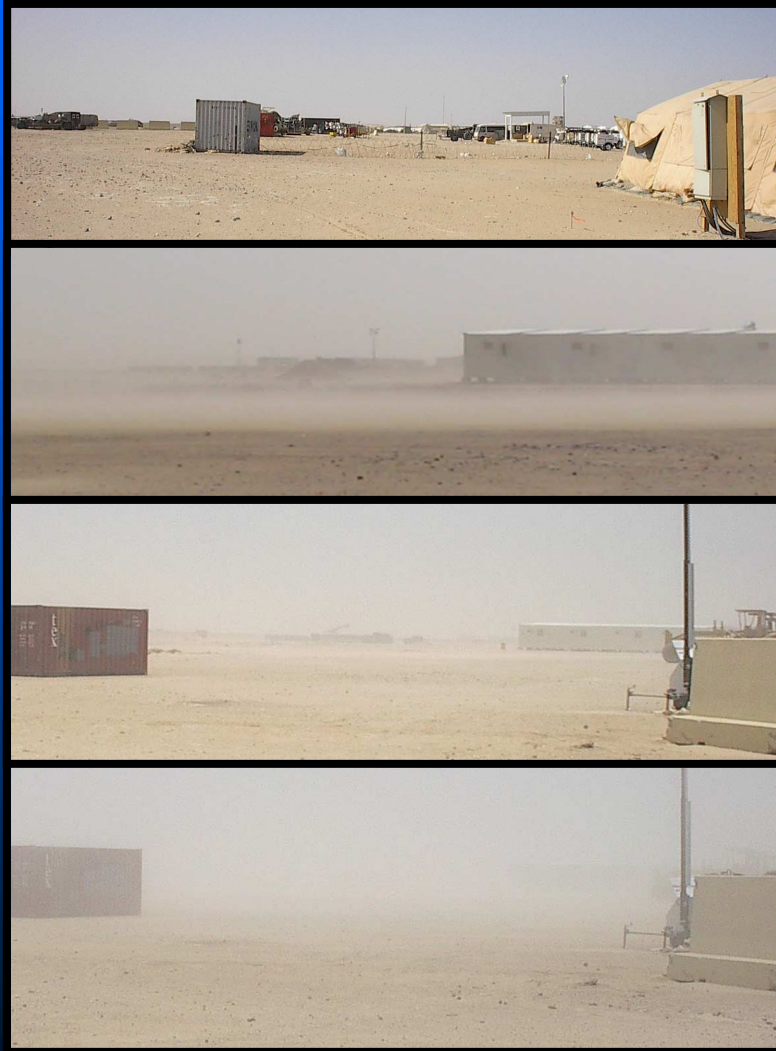


# Sample Processing

<u>Sieve #</u>	<u>Pore size (<math>\mu\text{m}</math>)</u>
40	420
100	149
120	125
230	63
325	44
635	20



# Physical Characterization of the Desert Environment



- Particle size distribution & conc.
- Average Wind Velocity
- Humidity
- Temperature





# Equipment Used to Characterize the Desert Environment

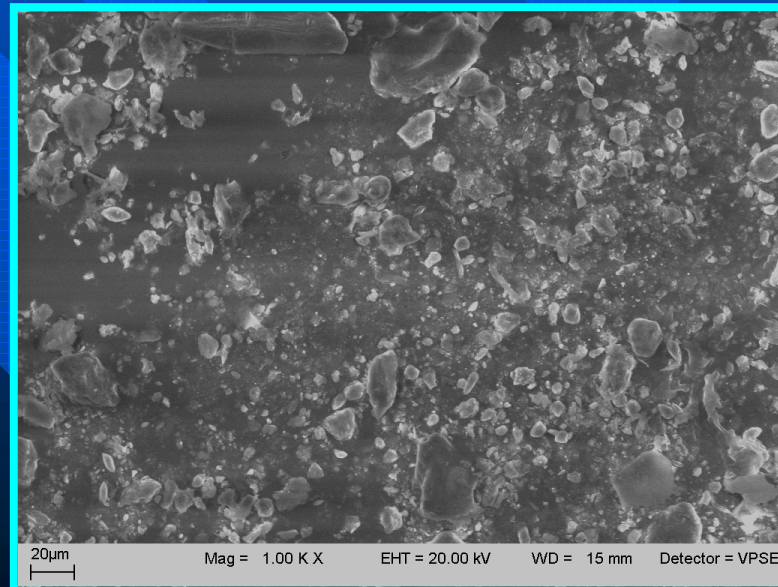
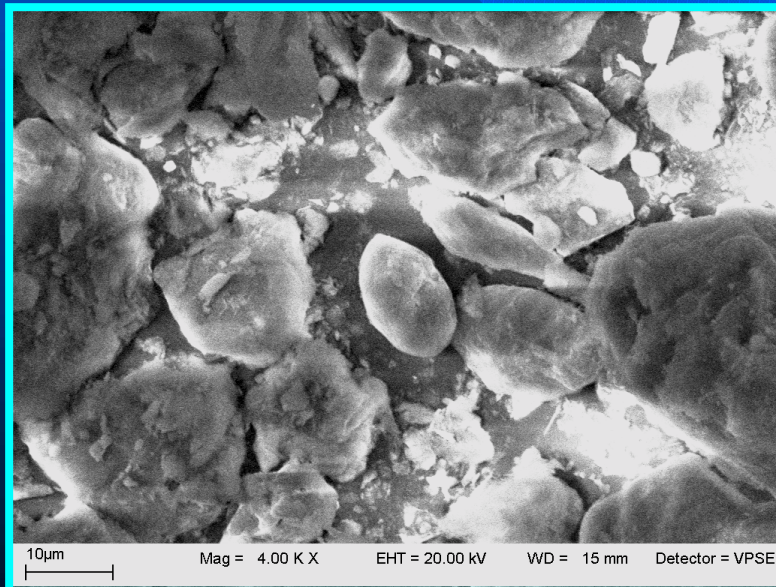
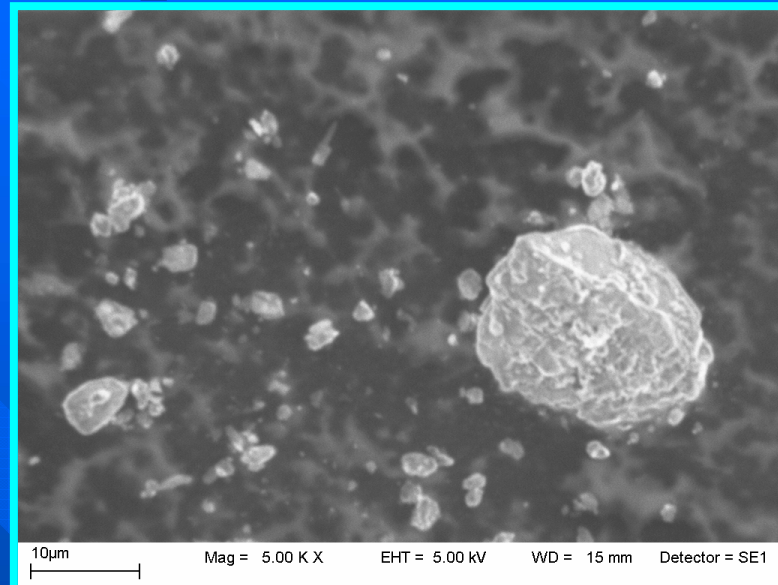
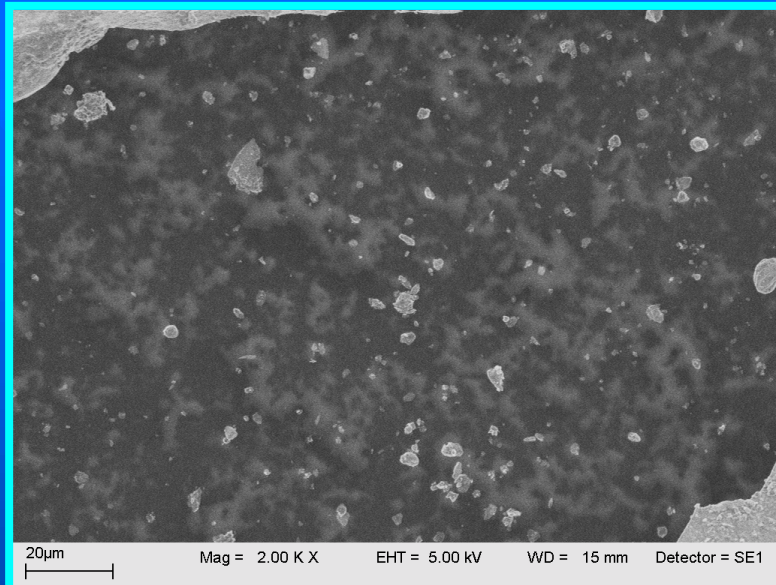


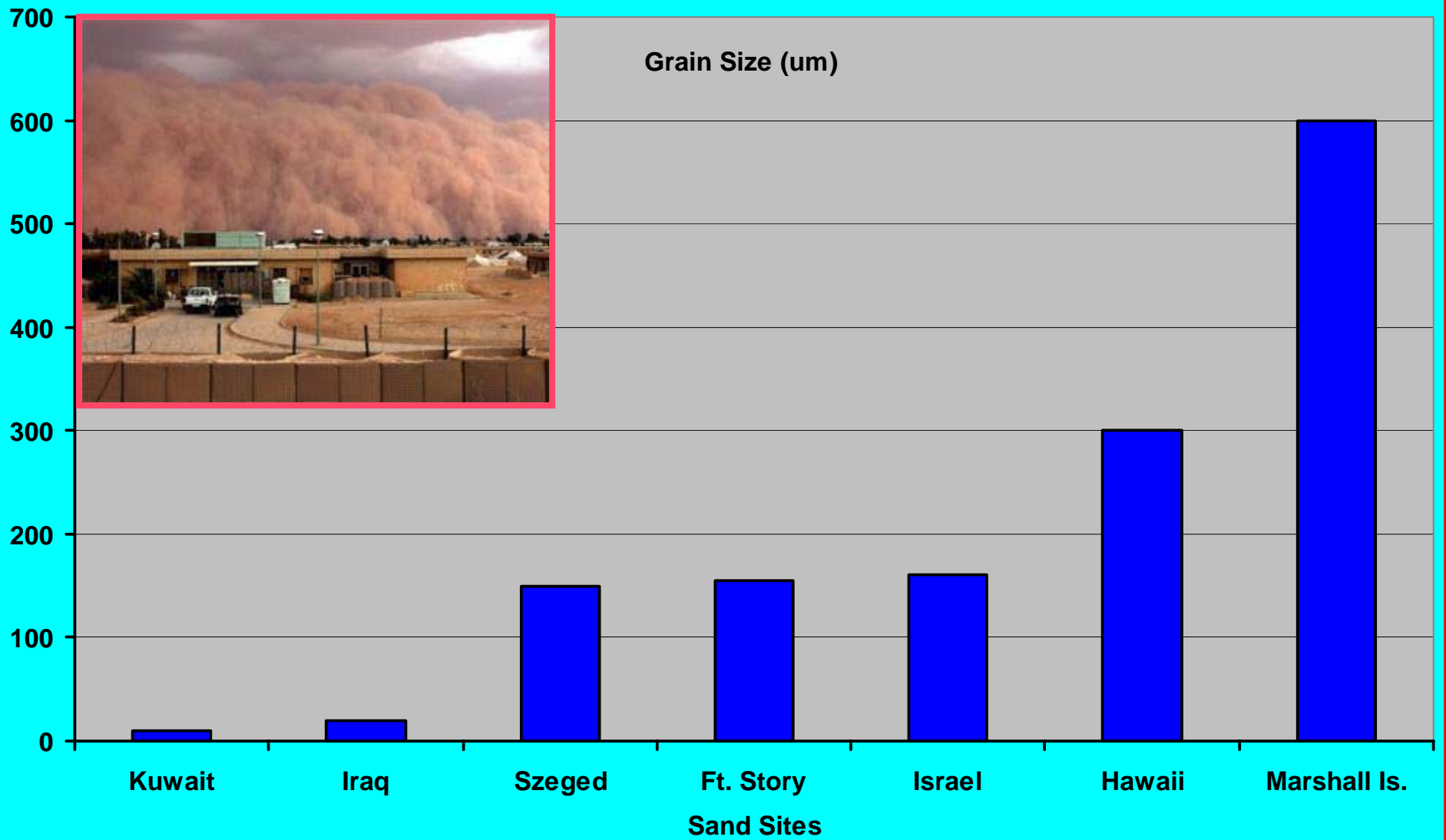
Particle size and distribution, temperature and rel. humidity recorded with a Aerocet 531 hand-held monitoring device (Met One Instruments, Inc.).



Wind speed monitor (CFM Master Thermo-Anemometer Model DCFM 8901; Mannix Testing and Measurement, Inc.).

# Sand / Dust Micro-particulates









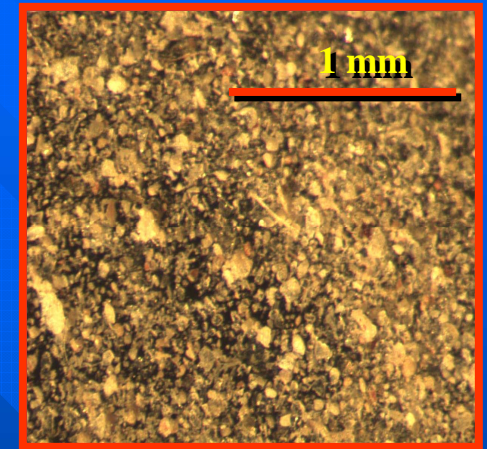
# Soils of Lower Mesopotamia



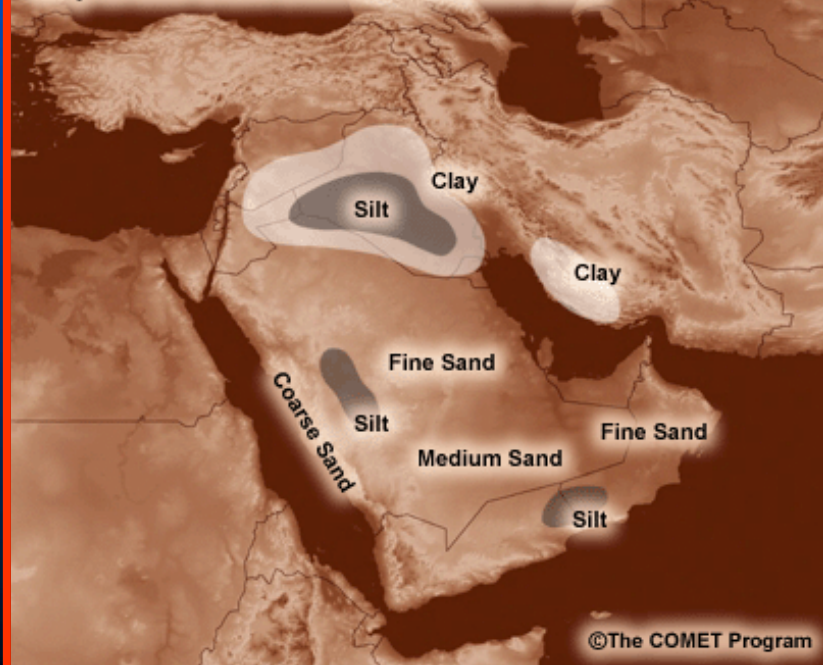
20-60% of surface soil mass is comprised of particles  $< 2.5 \mu\text{m}$  in size ( $\text{PM}_{2.5}$ )

As, Cr, Pb, Os are elevated

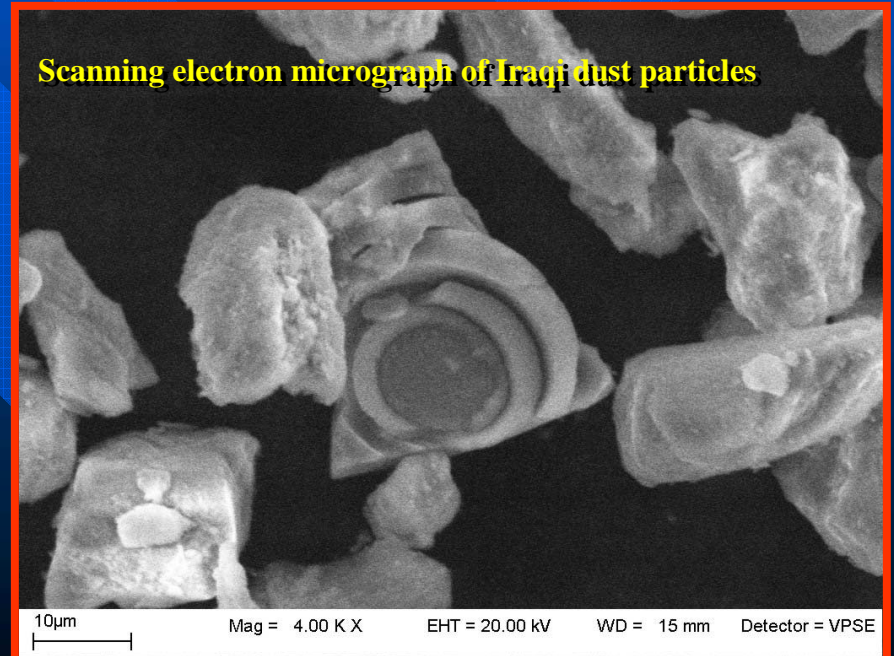
Silicate mineral component mixed with significant proportions of calcium carbonate



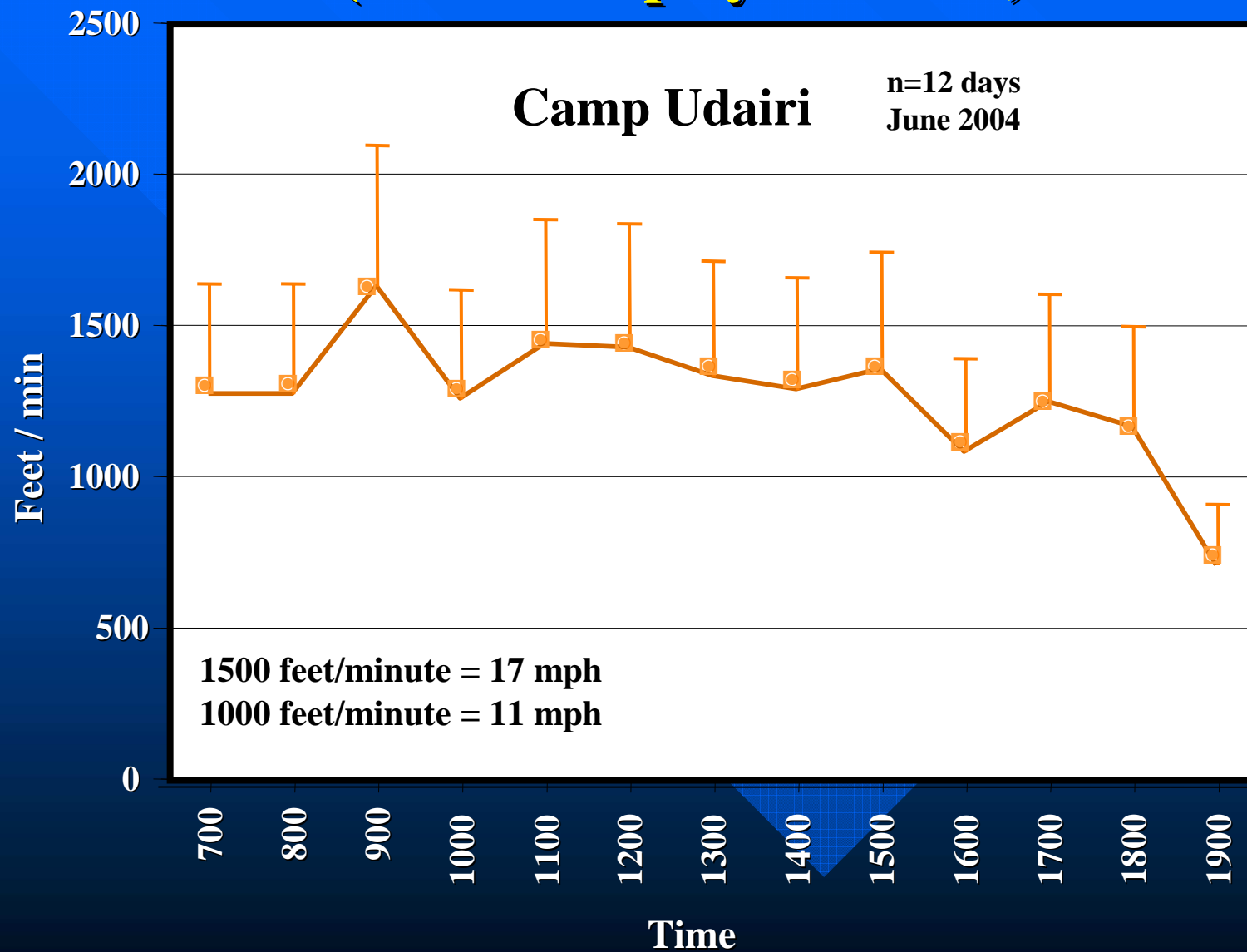
Map of Soil Grain Sizes in the Middle East



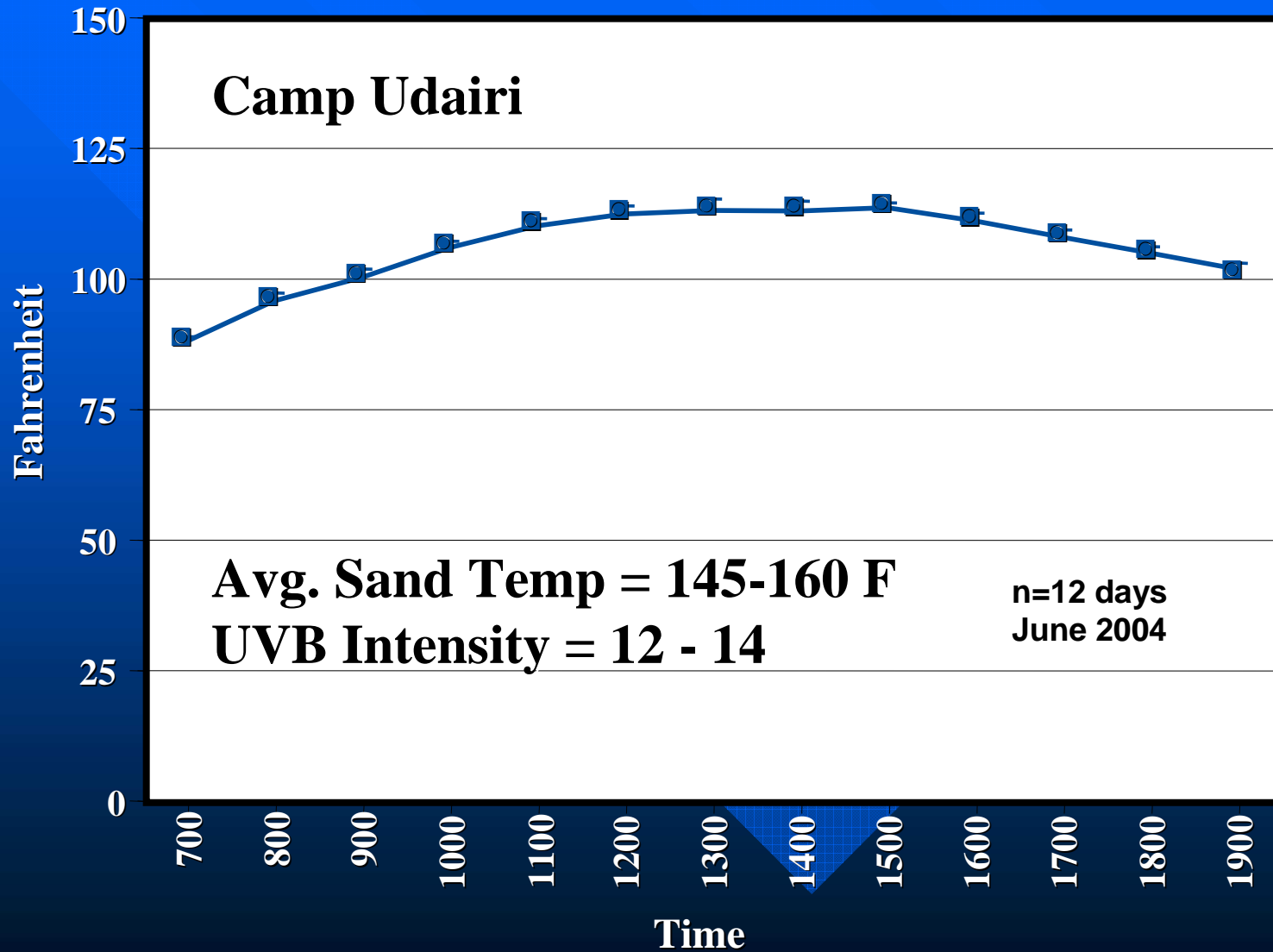
Scanning electron micrograph of Iraqi dust particles



# Average Hourly Wind Velocity (Kuwait Deployment Site)

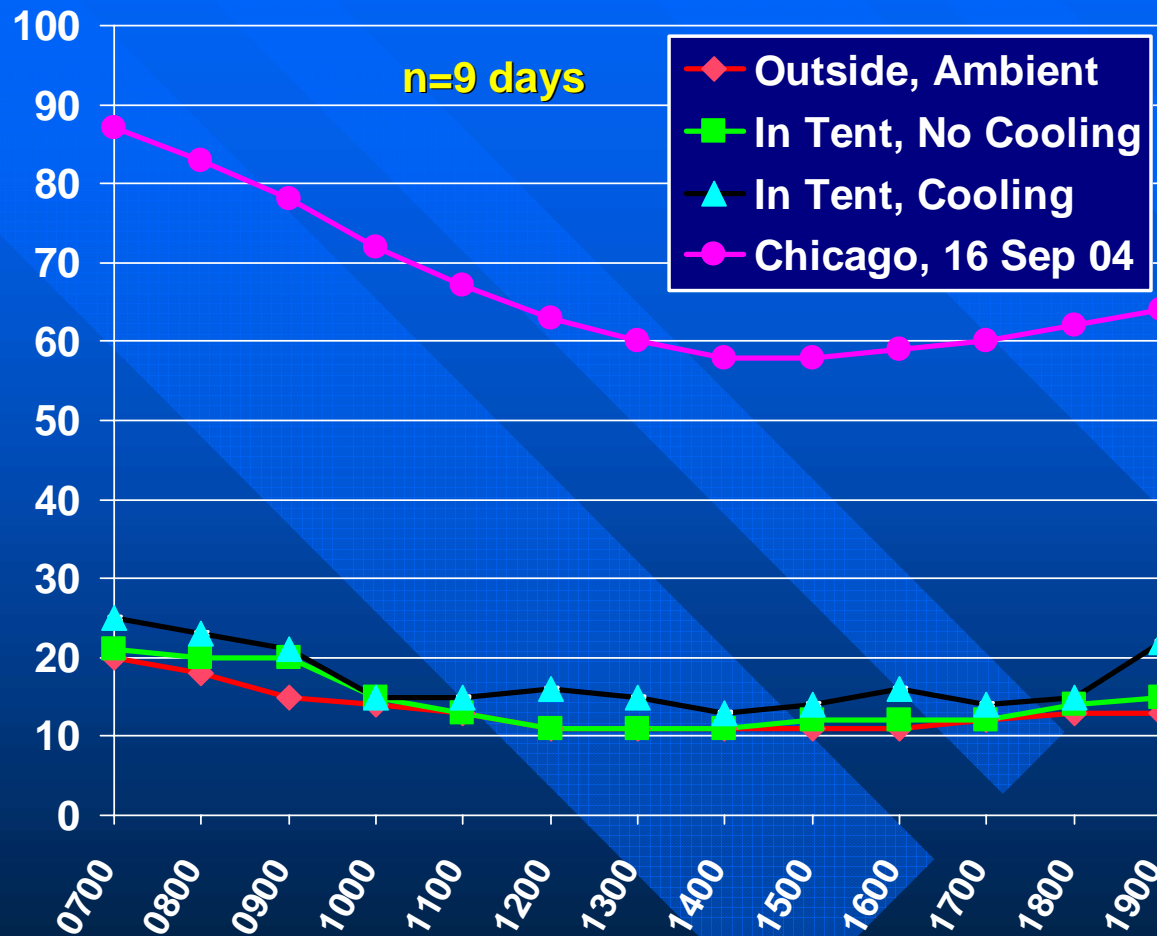


# Average Hourly Temperature (Kuwait Deployment Site)

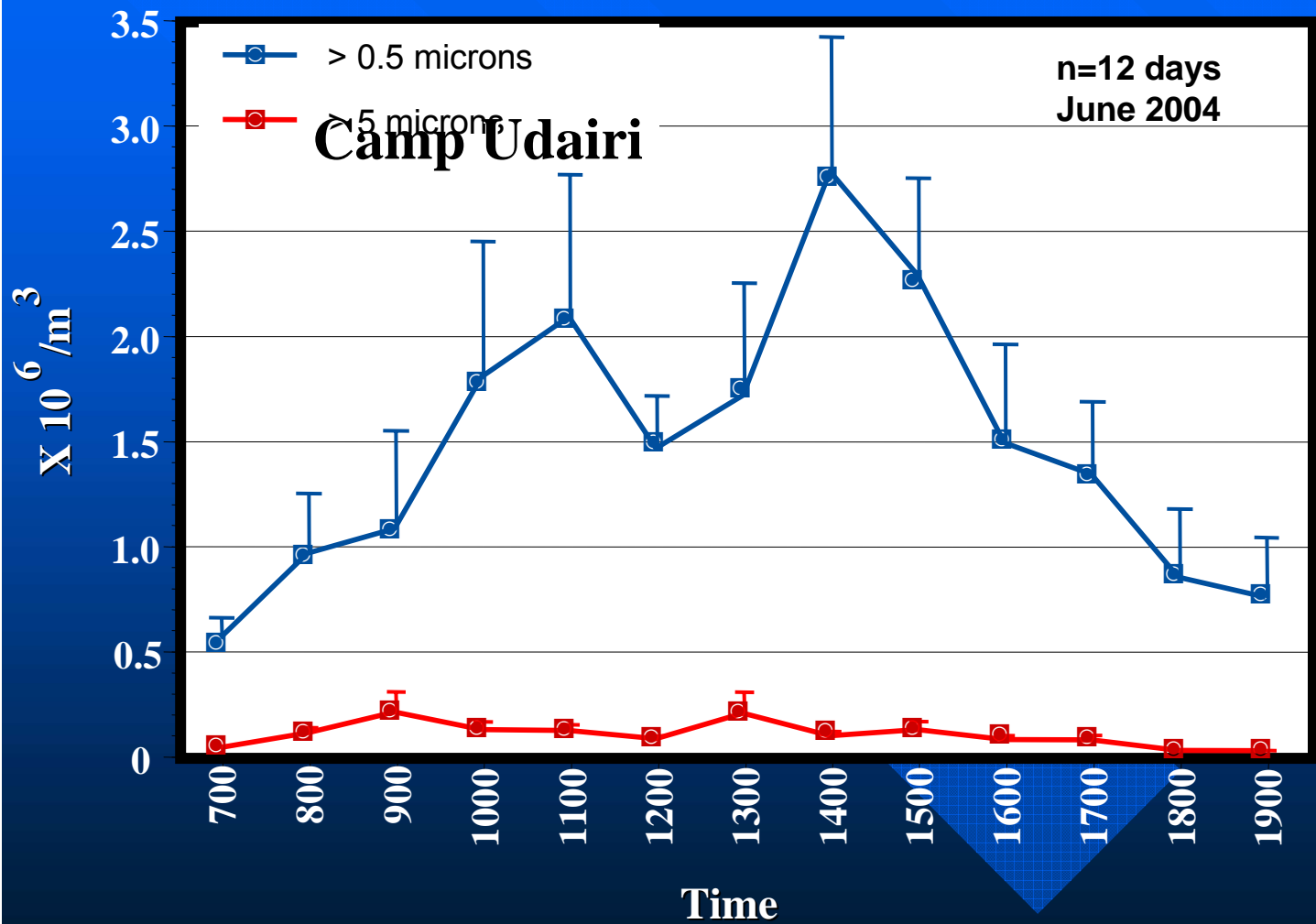




# Average Hourly Relative Humidity



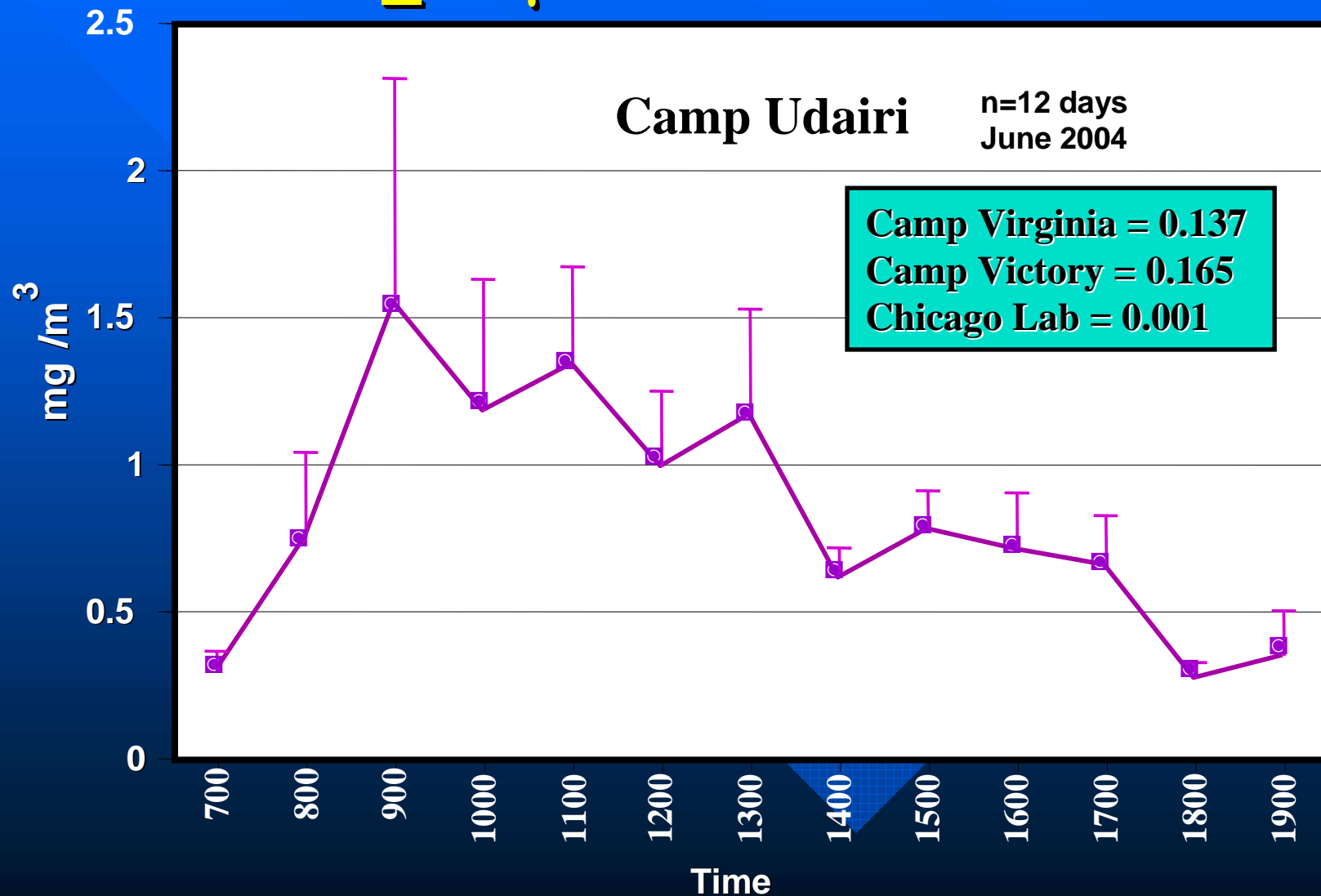
# Mean Ambient Airborne Concentrations



**>0.5um**  
 Camp Virginia = 348,040  
 Camp Victory = 201,510  
 Chicago Lab = 37,234

**>5um**  
 Camp Virginia = 14,380  
 Camp Victory = 13,150  
 Chicago Lab = 1,034

# Mean Total Suspended Particle Mass $\leq 10 \mu\text{m}$ in Ambient Air



# Chemical Analysis: Heavy Metals

Acid Extractables  
Tent 1

Particles less than 250  $\mu\text{m}$   
are airborne at 15 MPH.

20-40  $\mu\text{m}$  deposited in nose,  
throat, trachea.

20-10  $\mu\text{m}$  deposited in all  
upper airways of the lung.

$\leq 10 \mu\text{m}$  deposited in lowest  
airways of lungs.

98% of Tent Dust is less  
than 149  $\mu\text{m}$ .

~ 60% of Sand Samples  
are less than 149  $\mu\text{m}$ .

Sample	>120um	>90um	> 63um	>44um	>20um	<20um
Mass	0.2627	0.2596	0.2488	0.2626	0.2441	0.2504
Element	% dry wt	%dry wt	%dry wt	%dry wt	%dry wt	%dry wt
<b>Sr</b>	0.0697	0.0642	0.0995	0.1978	<b>0.2718</b>	0.2436
<b>Ba</b>	0.0068	0.0072	0.0081	0.0192	0.0308	<b>0.0463</b>
<b>P</b>	0.0160	0.0170	0.0234	0.0433	0.0549	<b>0.0649</b>
<b>S</b>	2.4413	2.4230	3.0444	<b>4.0062</b>	3.6646	3.0458
<b>Mg</b>	0.6844	0.8718	1.2672	1.5505	1.7234	<b>1.7784</b>
<b>V</b>	0.0022	0.0026	0.0032	0.0041	0.0046	<b>0.0049</b>
<b>Na</b>	0.1759	0.1963	0.1672	0.2056	0.2123	<b>0.2225</b>
<b>Al</b>	0.2969	0.3832	0.4948	0.6351	0.7164	<b>0.7521</b>
<b>Ca</b>	9.0134	10.3057	11.7495	13.9148	15.3535	<b>16.7133</b>
<b>Zn</b>	0.0053	0.0039	0.0042	0.0070	0.0112	<b>0.0206</b>
<b>Cu</b>	0.0060	0.0050	0.0036	0.0054	0.0077	<b>0.0268</b>
<b>Ni</b>	0.0089	0.0094	0.0169	0.0197	0.0305	<b>0.0564</b>
<b>Y</b>	0.0009	0.0006	0.0006	0.0007	0.0009	<b>0.0010</b>
<b>K</b>	0.0502	0.0653	0.0612	0.0942	0.1186	<b>0.1422</b>
<b>Mn</b>	0.0174	0.0222	0.0268	0.0305	0.0331	<b>0.0352</b>
<b>Fe</b>	0.3506	0.4844	0.6889	0.8419	0.9601	<b>0.9736</b>
<b>Cr</b>	0.0027	0.0032	0.0039	0.0049	0.0052	<b>0.0052</b>
<b>Pb</b>	0.0111	0.0038	0.0049	0.0056	0.0076	<b>0.0138</b>



# Summary

## TSP (Total Suspended Particle Mass) (mg/m<sup>3</sup>) PM10 (10 µm) and below

- ≡ 0.001 mg/m<sup>3</sup> (NIDBR Lab, Great Lakes, IL)
- ≡ 0.137 mg/m<sup>3</sup> (Camp Virginia Clinic, Kuwait - indoors)
- ≡ 2.469 mg/m<sup>3</sup> (Highest hourly average - 8 AM)
- ≡ 9.114 mg/m<sup>3</sup> (Highest TSP reading)
- ≡ 2.051 mg/m<sup>3</sup> (Highest daily maximum - 18 June at 1300)
- \* *NOTE: >9.999 mg/m<sup>3</sup> readings recorded during peak dust storms*

## Count (Total Number of Suspended Particles) (/ft<sup>3</sup>)

### Size Range = 0.5 µm to 10 µm

- ≡ 37,234 /ft<sup>3</sup> (NIDBR Lab, Great Lakes, IL)
- ≡ 348,040 /ft<sup>3</sup> (Camp Virginia Clinic, Kuwait - indoors)
- ≡ 3,037,298 /ft<sup>3</sup> (Highest average hourly maximum at 1300) (SD = 1,556,266/ft<sup>3</sup>)
- ≡ 16,668,250 /ft<sup>3</sup> (Highest daily maximum - 18 June at 1300)
- ≡ 3,614,455 /ft<sup>3</sup> (Highest average daily (0700-1900) maximum 13 June) (SD = 971,589/ft<sup>3</sup>)
- \* *NOTE: >20,000,000 counts /ft<sup>3</sup> readings recorded during peak dust storms*

### Size Range = 5.0 µm to 10 µm

- ≡ 1,034 /ft<sup>3</sup> (NIDBR Lab, Great Lakes, IL)
- ≡ 14,380 /ft<sup>3</sup> (Camp Virginia Clinic, Kuwait - indoors)
- ≡ 194,945 /ft<sup>3</sup> (Highest average hourly maximum at 1300) (SD = 117,305/ft<sup>3</sup>)
- ≡ 1,262,120 /ft<sup>3</sup> (Highest daily maximum - 18 June at 1300)
- ≡ 148,512 /ft<sup>3</sup> (Highest average daily maximum - 13 June) (SD = 102,861/ft<sup>3</sup>)
- \* *NOTE: >9,999,999 counts/ft<sup>3</sup> readings recorded during peak dust storms*

**NOTE: 1 cu ft = 28.31685 Liters = 0.02831685 m<sup>3</sup>**

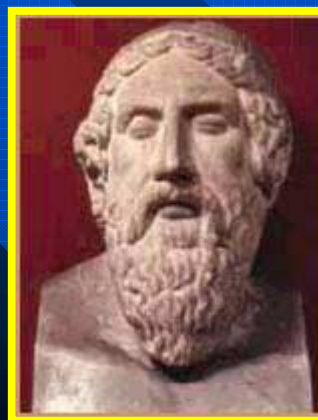
# Microbiological Study of Micro-particulates

**Fifth Plague of Egypt** – *“Murrain of beasts” Exodus 9:6*

**Sixth Plague of Egypt** – *“And it shall become small dust in all the land of Egypt, and shall be a boil breaking forth [with] blains upon man, and upon beast, throughout all the land of Egypt”*. Exodus 9:10

**25 BC: Poet Virgil**

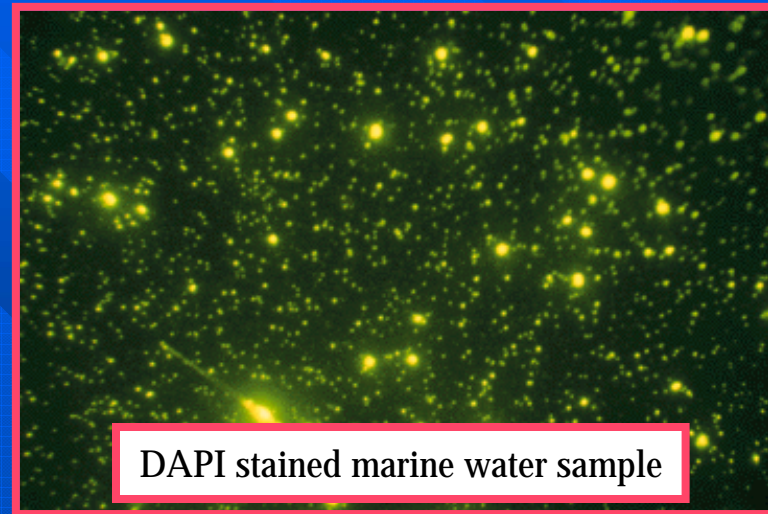
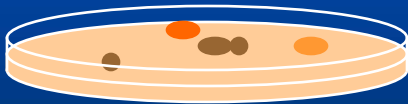
- *The Iliad (Homer) “the burning wing of plague...”*
- *Middle Ages: European pandemic “Black Bane” killed 60,000 cattle.*



# The great plate count anomaly

microbial community

↓  
plating

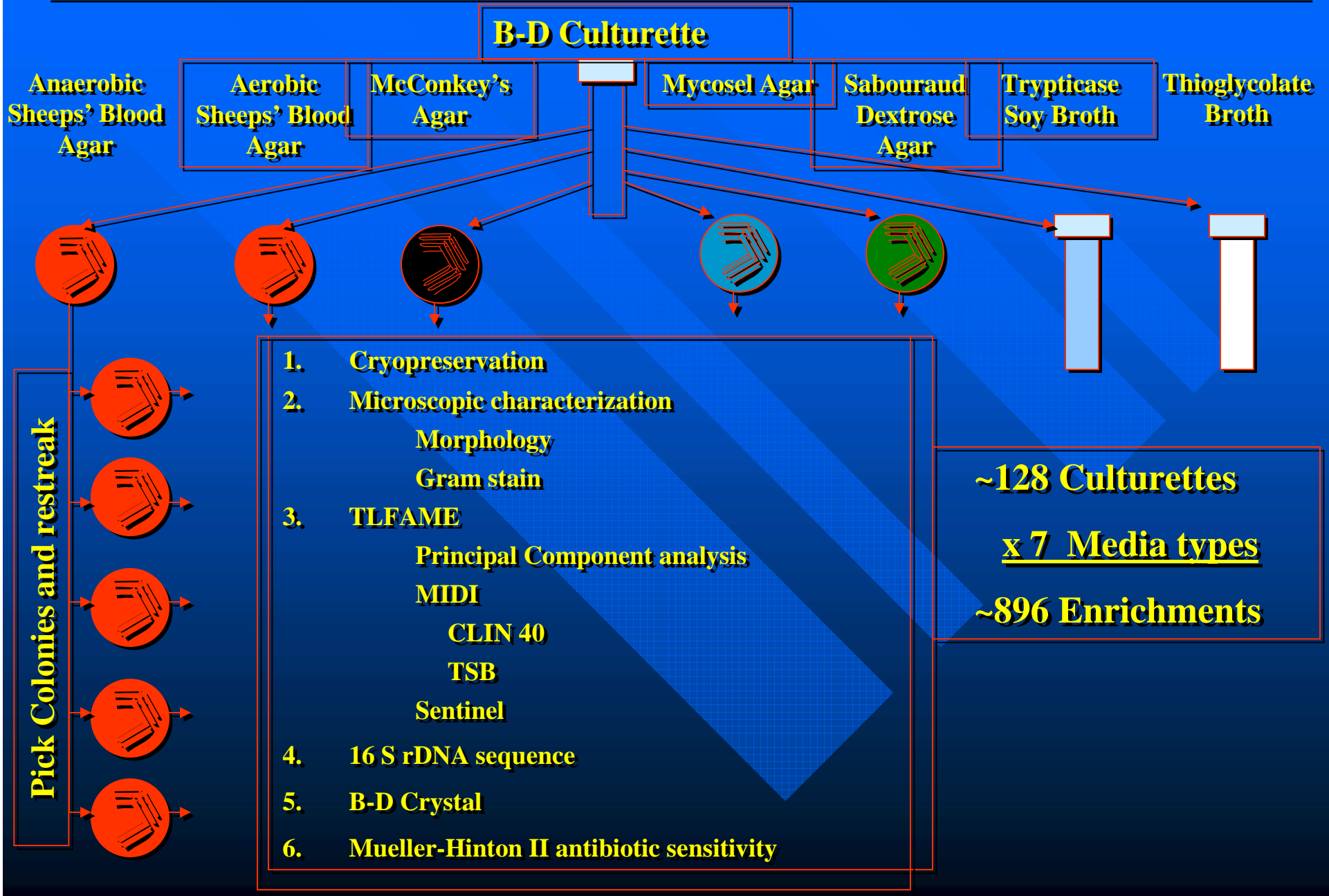


DAPI stained marine water sample

< 1% of observable bacteria grow on standard culture media



# Microbial Isolation and Characterizations



# Microbiology Summary

NO.	Site	Hemolysis on Blood agar	MIDI @ DE	Similarity	MIDI @ MS	Similarity	MIDI @ MS	Similarity	MIDI 500 bp rDNA sequence analysis			Best ID thus Far
			Environmental Database ID	Index	CLIN 40 Database ID	Index	Sentinel Database ID	Index	% Difference			
2	Babylon	No	Pseudomonas stutzeri	0.597	Pseudomonas stutzeri	0.503	Neisseria meningitis	0.357				Neisseria meningitis
							Neisseria meningitis	0.29				
							Neisseria cinerea	0.29				
5	Babylon	No	Staphylococcus epidermidis	0.827	Staphylococcus aureus	0.676	Staphylococcus aureus	0.609				Staphylococcus aureus
			Staphylococcus epidermidis	0.78	Staphylococcus warneri	0.596	Staphylococcus epidermidis	0.576				
			Staphylococcus capitis	0.753	Staphylococcus aureus	0.569	Staphylococcus hominis	0.497				
8	Babylon	No	Bacillus circulans	0.61	No match/Too dilute	N/A	No match					Bacillus circulans
9	Udairi	Alpha hemoly	Not sent to MIDI	N/A	Not Extracted	N/A						None
10	Udairi	Alpha hemoly	Ewingella americana	0.778	Neisseria cinera	0.204	Providencia rettgeri	0.023	Pantoea agglomerans	0.95%	Species	Pantoea agglomerans
			Salmonella typhimurium	0.592	Aeromonas veronii	0.175	Arcobacter skirrowii	0.018				
			Pantoea agglomerans	0.568	Neisseria cinera	0.169	Erwinia amylovora	0.017				
11	Udairi	Alpha hemoly	Pseudomonas stutzeri	0.896	Not Extracted	N/A			Pseudomonas agrici	1.34%	Genus	Pseudomonas agrici
			Pseudomonas balearica	0.659								
			Pseudomonas resinovorans	0.584								
12	Udairi	No	Not growing when others	N/A	Vibrio alginolyticus	0.366	Ralstonia paucula	0.127				Ralstonia paucula
					Aeromonas hydrophilia	0.366	Erwinia mallotivora	0.103				
					Neisseria mucosa	0.335	Ralstonia basilensis	0.1				
14	Udairi	No	Not growing when others	N/A	Staphylococcus epidermidis	0.419	Staphylococcus pasteurii	0.207				Staphylococcus pasteurii
							Staphylococcus caprae	0.185				
							Staphylococcus warneri	0.135				
15	Udairi	No	Virgibacillus pantothenicus	0.677	Bacillus coagulans	0.432	Arthrobacter atrocyaneus	0.414	Arthrobacter crystallopoietes	0.00%	Species	Arthrobacter crystallopoietes
			Micrococcus luteus	0.499	Dermobacter hominis	0.327	Agromyces ramosus	0.283				
			Bacillus atropheus	0.477	Kocuria-variens(Micrococcus)	0.316						
16	Udairi	No	Staphylococcus warneri	0.881	Pseudomonas stutzeri	0.44	Pseudomonas balearica	0.097				Pseudomonas balearica
			Staphylococcus epidermidis	0.754	N/A	N/A						
			Staphylococcus epidermidis	0.61	N/A	N/A						
17	Udairi	Beta/Alpha	Paenibacillus thiaminolyticus	0.534	Not Extracted	N/A			Paenibacillus thiaminolyticus	2.97%	Genus	Paenibacillus thiaminolyticus
			Bacillus atropheus	0.464								
18	Udairi	Beta hemolyti	Bacillus subtilis	0.901	Bacillus subtilis	0.52	Bacillus vedderi	0.656				Bacillus vedderi
			Bacillus atropheus	0.697	N/A	N/A	Bacillus mojavensis	0.642				

# Summary of Soil Isolates

<b>Best ID thus Far</b>	<b>Comment</b>
<i>Neisseria meningitidis</i>	meningitis
<i>Staphylococcus aureus</i>	cystic fibrosis
<i>Bacillus circulans</i>	gastro-enteritis
<b>NONE</b>	Unidentified
<i>Pantoea agglomerans</i>	septic arthritis
<i>Pseudomonas agrici</i>	
<i>Ralstonia paucula</i>	opportunism-septicemia, peritonitis, abscesses
<i>Staphylococcus pasteurii</i>	various infections
<i>Arthrobacter crystallopoietes</i>	
<i>Pseudomonas balearica</i>	cystic fibrosis
<i>Paenibacillus thiaminolyticus</i>	bacteremia
<i>Bacillus vedderi</i>	obligate alkaliphile
<i>Bacillus subtilis</i>	
<i>Pantoea agglomerans</i>	epiphyte
<i>Pseudomonas pseudobalcaligenes</i>	
<i>Cryptococcus albidus</i>	septicemia and meningitis
<i>Bacillus clausii</i>	Oral bacteriotherapy
<i>Kurthia gibsonii</i>	Diarthorea
<i>Bacillus firmus</i>	alkaliphile; bread spoilage
<i>Staphylococcus kloosii</i>	various infections
<i>Bacillus mojaviensis</i>	biosurfactant
<i>Bacillus licheniformis</i>	food poisoning
<i>Pseudomonas oryzae</i>	Hickman catheter biofilm







# B-D Crystal Biochemical Characterization of Hemolytic Isolates



Sample	Gram + Organism	Confidence	Associated Disease
#25	Kytococcus sedentarius	0.467	Pitting of human epidermis in keratolysis
#36	Kocuria rosea	0.9927	Catheter related bacteremia
#72	No Match	*	
#56	Gardnerella vaginalis	0.9982	Human bacterial vaginosis
#69	Leifsonia aquaticum	0.9999	Rare bacteremia
#22	Leifsonia aquaticum	0.9941	Rare bacteremia
#16	Corynebacterium pseudotuberculosis	0.6877	Ventral lymphadenitis, abscesses, and ulcerative dermatitis in cattle
#70	Leifsonia aquaticum	0.9951	Rare bacteremia



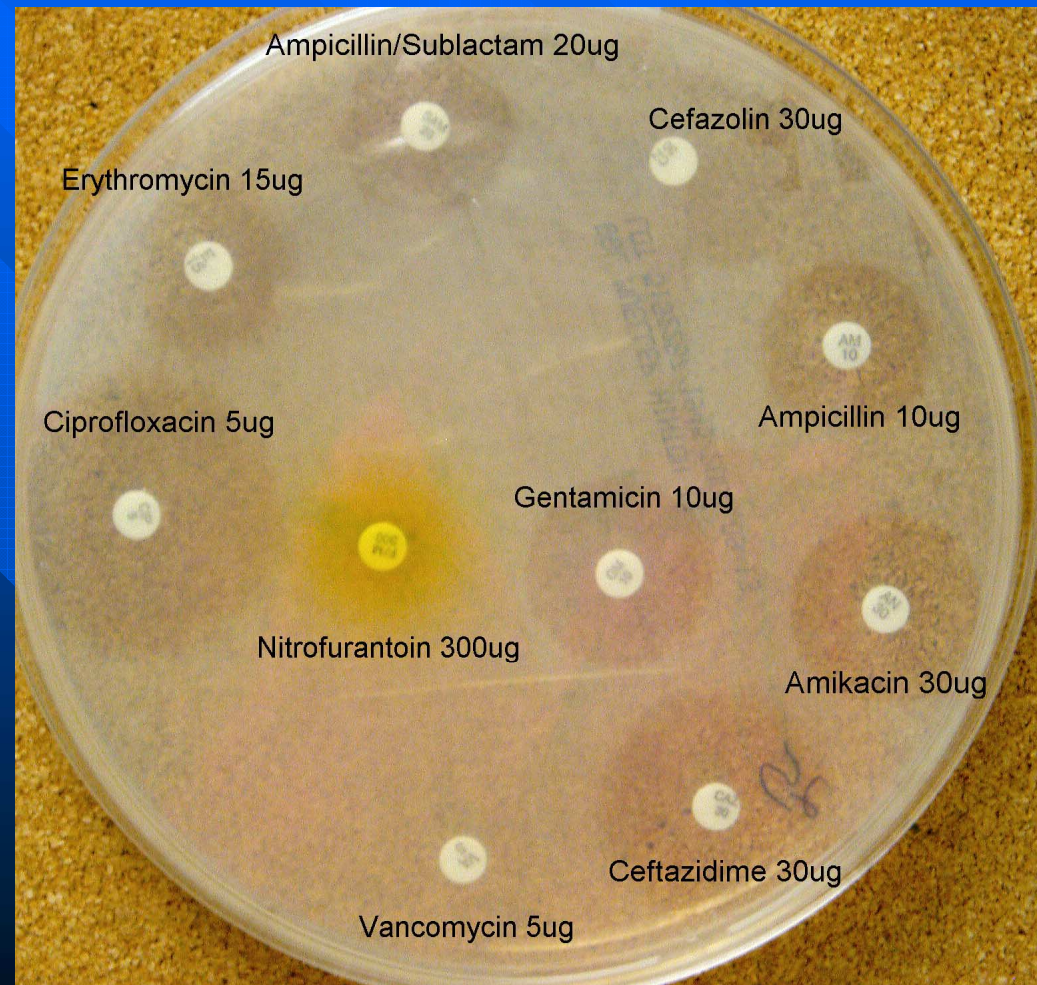
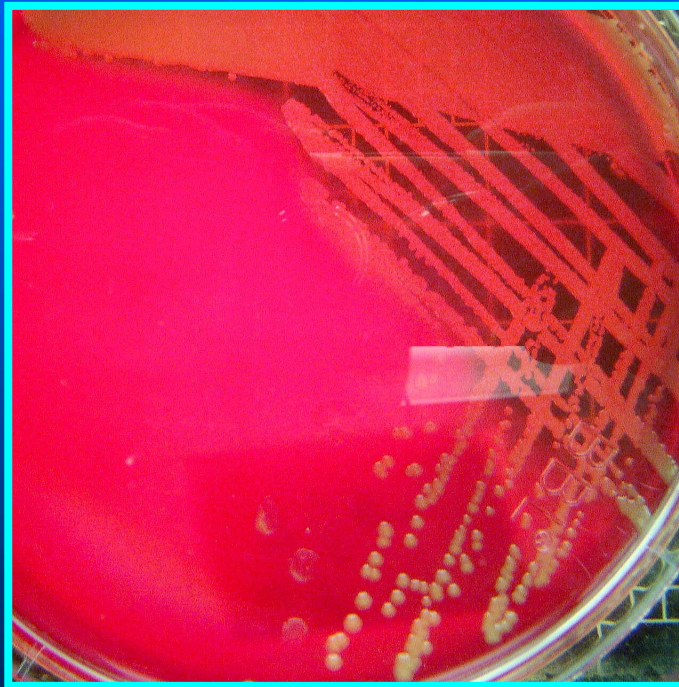
Sample	Gram neg. Organism	Confidence	Comments
#22	No Match	*	Additional testing required to establish ID
#16	No Match	*	Additional testing required to establish ID
#36	<i>Yersinia pestis</i>	0.9384	*
#25	<i>Pseudomonas stutzeri</i>	0.8598	Supplemental testing recommended
#69	No Match	*	Additional testing required to establish ID
#72	No Match	*	Additional testing required to establish ID
#56	No Match	*	Additional testing required to establish ID
#70	No Match	*	Additional testing required to establish ID



# Antibiotic Sensitivity



## Beta Hemolytic Isolate Number 69





# Bacteria Isolated from Kuwait and Iraq that have Shown Antibiotic Resistance.

Culture#	Description	Location	Culturette	Hemolysis	Colony Morphology	MIDI @ DE Environmental	Similarity Index	Comment
8	BSSI	Babylon	Green	No	Dry Fungal type colonies; White spreader on TSA & Blood	Bacillus circulans	0.61	N/A
12	>20<44 $\mu$ m	Udairi	Green	No	Cream colored mucoid colonies on Blood and TSA;	<b>Not growing when others sent off</b>	N/A	N/A
16	>44<63 $\mu$ m	Udairi	Green	No	Small mucoid colonies on Blood; Spreading mucoid on TSA;	Staphylococcus wameri	0.881	N/A
20	>44<90 $\mu$ m	Udairi	Green	alpha	Small dry cream colored colonies	Pantoea agglomerans Pantoea agglomerans	0.82 0.711	GC subgroupB GC subgroupC
24	<20 $\mu$ m	Udairi	Green	No	Shiny yellowish/cream spreading colonies on TSA; Purple spreader on Blood;			
28	TAB II Sand A	Tallil	Green	No	Large shiny mucoid colonies	Not sent to MIDI	N/A	N/A
32	TAB II Sand B	Tallil	Green	beta	Clear white cauliflower colony on TSA; Shiny clear runny colony on Blood;	Not sent to MIDI	N/A	N/A
Culture#	MIDI @ MS CLIN 40	Similarity Index	Comment	MIDI @ MS Sentenial	Similarity Index	MIDI 500 bp rDNA sequece analysis % Diff	Comments	
8	No match/Too dilute	N/A	N/A	No match				
12	Vibrio alginolyticus Aeromonas hydrophilia	0.366 0.366	N/A N/A	Ralstonia paucula Erwinia mallotivora	0.127 0.103			
16	Pseudomonas stutzeri	0.44	N/A	Pseudomonas balearica	0.097			
20	Pantoe agglomerans Aeromonas hydrophilia	0.623 0.386	GC subgroup N/A	Ralstonia paucula Buttiauxella gaviniae	0.274 0.175	Pantoea agglomerans	0.85%	Plant/Human Pathogen
24								
28	Not Extracted	N/A	N/A					
32	Not Extracted	N/A	N/A			Flavimonas oryzihabitans	0.10%	Hickman Cath. Pathogen





# Fungal Isolates

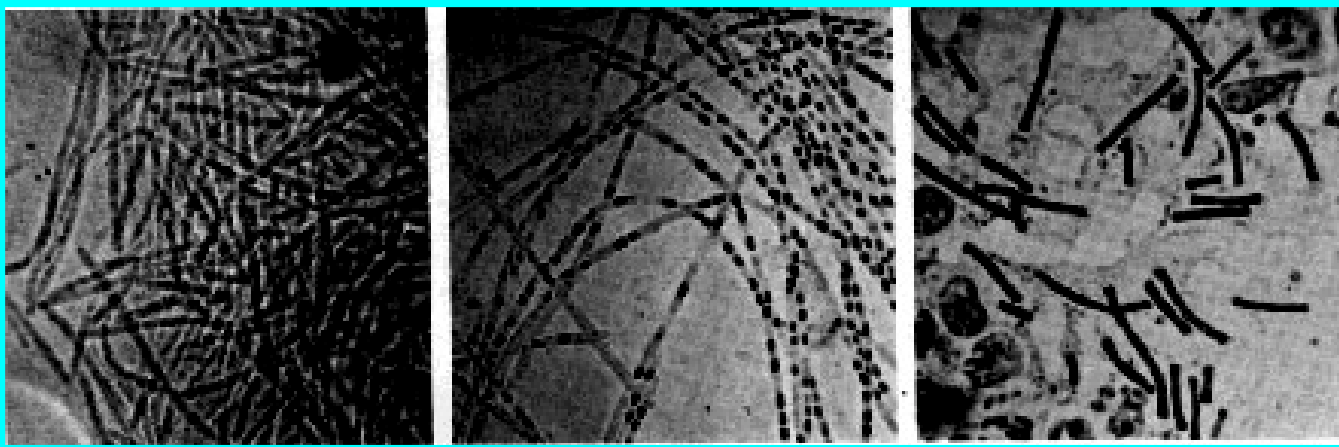


~300 bp of D2 region of LSU rDNA

Microseq Library database							
Midi D2(300 bp)LSU rRNA	% diff	LSU D2 Genbank Database	% ID	Associated Disease			
Allewia eureka	0.31	Ulocladium sp.	99				
Allewia eureka	4.64	Cryptococcus uzbekistanensis	100				
Allewia eureka	0.31	Ulocladium sp.	99	None			
Alternaria alternata	0	Alternaria sp	100	Plant pathogen			
Rhodotorula minuata	5.73	Rhodotorula minuata	99	Eye infections			
Cryptococcus albidus	2.44	Cryptococcus sp.	100	Cryptococcus neoformans - meningoencephalitis			
Ulocladium chartarum	0.31	Stemphylium sp	99	Fungal biocontrol agent			
Filobasidium uniguttulatum	4.64	Cryptococcus uzbekistanensis	100	Teleomorph of Cryptococcus, non pathogenic yeast			
Ulocladium consortiale	0	Stemphylium sp	100	cutaneous mycoses			
Ulocladium chartarum	0.31	Stemphylium sp	99				
Mortierella polycephala	7.1	Mortierella polycephala	92	Pulmonary mycosis in cattle			
Embellisia chlamydospora	0	Ulocladium sp.	99	none			
Filobasidium uniguttulatum	4.64	Cryptococcus uzbekistanensis	100				
Penicillium camembertii	0	Penicillium sp.	100				
Cryptococcus albidus	0	Cryptococcus albidus	100				
Allewia eureka	0.31	Ulocladium sp.	99				
Embellisia chlamydospora	0	Ulocladium sp.	99				
Filobasidium uniguttulatum	4.64	Cryptococcus uzbekistanensis	100				
Embellisia chlamydospora	0	Ulocladium sp.	99				
Filobasidium uniguttulatum	4.64	Cryptococcus uzbekistanensis	100				
Penicillium camembertii	0	Penicillium sp.	100				
Allewia eureka	0.31	Ulocladium sp.	99	Plant pathogen			
Phoma glomerata	0	Phoma herbarum	99				

# Summary of the Biological Analysis

- *Hemolytic microorganisms*
- *Gram positive spore-formers*
- *Gram negative opportunists*
- *Fungi*





# Conclusions Thus Far



- **Relatively abundant microbial community in dust**
  - **Including PM<sub>2.5</sub>**
- **Hemolytic strains isolated/characterized**
- **Microbial identification systems cause uncertainty**
  - **B-D Crystal biochemical characterization**
  - **TLFAME – MIDI CLIN 40, Sentinel TS, MIDI Env**
- **Methods bias perspective of microbial community**
- **Need a defensible trigger for concern**



