

**Microbial Resources:
Vital Infrastructure for
Research and Extension Endeavors**

Maria Auxilia T. Sirigan
auxie.nsri@gmail.com

**Microbiological Research and Services Laboratory
Natural Sciences Research Institute
University of the Philippines, Diliman, Quezon City**



**Microbiological Research
and Services Laboratory
Natural Sciences Research Institute
University of the Philippines-Diliman,
Quezon City, Philippines**

Vision

**A nationally and internationally recognized
vital microbial resource center**

Mission

**Providing quality technical expertise, services
and microbial resources to a wide range of clients**

Outline

**Microorganisms: Resources for Research
Metabolites for the industry**

**Microorganisms:
Resources for Extension/Technical Services
As tools for assays and efficacy testing**

**Microorganisms:
Resources for Research**

Metabolites for the industry

Research endeavors

Microbial diversity studies

- Exploring metabolic diversity for industrial applications
- Screening for useful metabolites

As sources of:

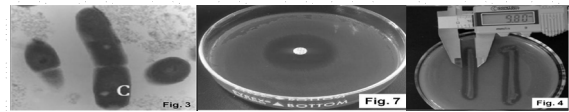
Enzymes

Biosurfactant

Bioactive compounds

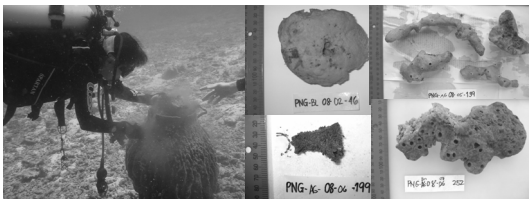
Microorganisms as sources of enzymes

- Halophilic bacterium from hyperalkaline spring: *Exiguobacterium* sp.
Exhibited amylase, pectinase and protease



Microorganisms as sources of biosurfactant

- Isolates were derived from marine sponges from different regions in the Philippines



Microorganisms as sources of biosurfactant

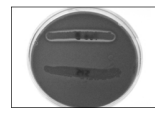
- Screening for biosurfactant production



Drop
Collapse Test



Oil Displacement
Test



Hemolysis
Test

Microorganisms as sources of biosurfactant

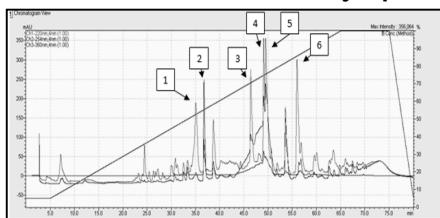
- Extracts of 6 microbial isolates
Determination of TLC profile
TLC bands – Test for emulsification activity
- *P. aeruginosa* R2A 766B
Best hemolytic activity
Determination of HPLC profile

Sponge-derived MMOs with Biosurfactant Activity

Identification	Isolate Code
<i>Bacillus subtilis</i>	R2A 515A
<i>Gordonia</i> sp.	EM 767A
<i>Pseudomonas aeruginosa</i>	ISP3 768A
<i>Pseudomonas aeruginosa</i>	R2A-766B
<i>Pseudomonas stutzeri</i>	R2A 709A
<i>Pseudomonas stutzeri</i>	R2A 685A

Microorganisms as sources of biosurfactant

- *P. aeruginosa* R2A 766B
HPLC Profile of extract – 6 major peaks

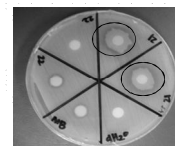


Bradford assay of R2A 766B biosurfactant:
possibly a glycopeptide.

Microorganisms from Marine Sponges As sources of bioactive compounds

Exploring anti-bacterial properties

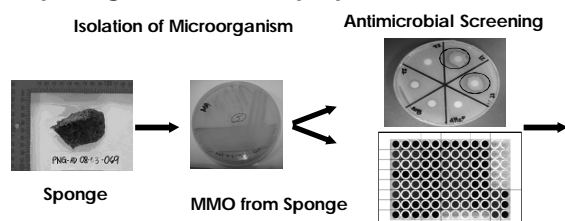
- Preliminary screening
Disk diffusion assay



- Test organisms:
Pseudomonas aeruginosa ATCC 12600
Staphylococcus aureus ATCC 10145

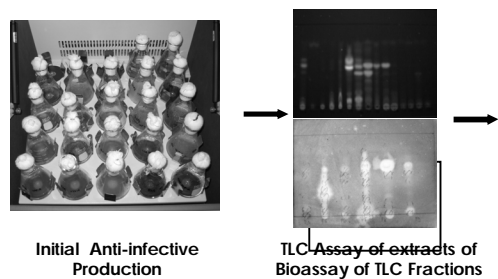
Microorganisms from Marine Sponges As sources of bioactive compounds

Exploring anti-bacterial properties



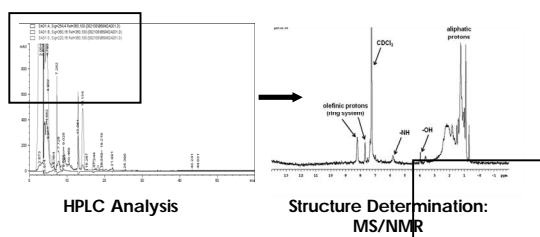
Microorganisms from Marine Sponges As sources of bioactive compounds

Exploring anti-bacterial properties

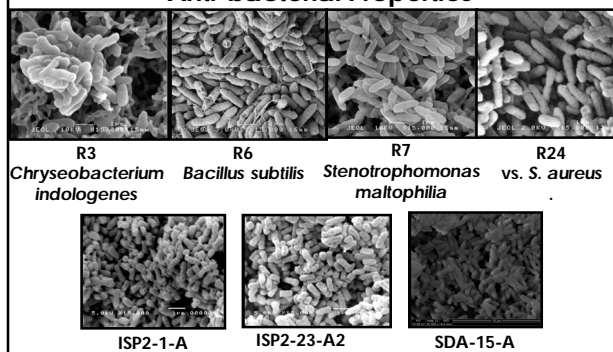


Microorganisms from Marine Sponges As sources of bioactive compounds

Exploring anti-bacterial properties

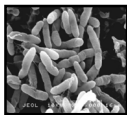


Microorganisms from Sponges: Anti-bacterial Properties

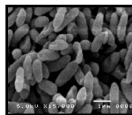


Identification of all isolates: *Pseudomonas aeruginosa*

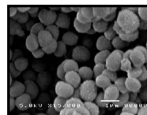
Microorganisms from Sponges: Anti-bacterial Properties



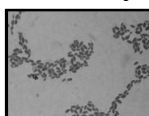
P142
Achromobacter
xylosoxidans



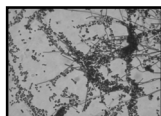
SDA-430-A
Pseudomonas
oryzae



ISP2-602-A
Micrococcus luteus

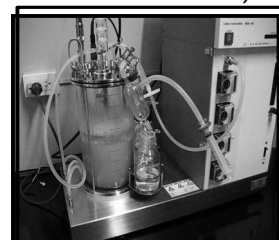


R2A-512-
Micrococcus luteus



ISP2-569-A
Streptomyces sp.

Antimicrobial Production by ISP2-569-A *Streptomyces* sp. (add flasks in NSRI video)

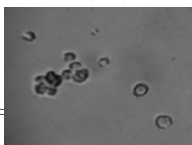


Optimization studies: Determination of C and N sources for enhanced yield of antimicrobial compound from ISP2-569-A

Microorganisms from Marine Sponges As sources of bioactive compounds

Exploring anti-protozoan properties

- Determination of % reduction in the viable *Trichomonas vaginalis* cell count



Trichomonas vaginalis

Microorganisms from Sponges: Potential anti-protozoan compounds vs. *Trichomonas vaginalis*

Sponge-derived MMOs	Isolate Code	Methanolic Extracts: Percent Kill of <i>T. vaginalis</i> (%)	Acetone Extracts: Percent Kill of <i>T. vaginalis</i> (%)
<i>Achromobacter xylosoxidans</i>	P-142A-I-1-A	96.7	99.5
<i>Chryseobacterium indologenes</i>	P-046-I-1-A	97.2	95.7
<i>Micrococcus luteus</i>	ISP2-602-A	98.1	99.1
<i>Penicillium griseofulvum</i>	ISP2-576-A	99.3	99.8
<i>Pseudomonas aeruginosa</i>	ISP2-1-A	100.0	98.0
<i>Pseudomonas aeruginosa</i>	ISP2-23-A2	92.6	99.4
<i>Pseudomonas oryzae</i>	SDA-430-A	97.8	99.2
<i>Rhodotulula mucilaginosa</i>	R2A-512-B	99.6	97.2
<i>Stenotrophomonas maltophilia</i>	P-203-I-1-A	98.0	99.1
<i>Streptomyces rubiginosus</i>	ISP2-569-A	99.8	99.9

Microorganisms: Resources for Extension and Technical Services

As tools for assays and efficacy testing

Extension and Technical Services

Test organisms – Tools for Efficacy Testing

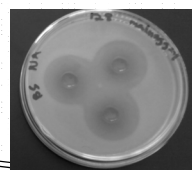
- For the evaluation of materials and products with biocidal agents
- For the evaluation of sterilization technology

- Test organisms – most are ATCC cultures

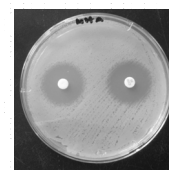
Test Organisms	Accession Numbers
Bacteria	
<i>Bacillus subtilis</i>	ATCC 6633
<i>Escherichia coli</i>	ATCC 25922
<i>Pseudomonas aeruginosa</i>	ATCC 15442; ATCC 27853
<i>Salmonella typhimurium</i>	ATCC 14028
<i>Staphylococcus aureus</i>	ATCC 6638; ATCC 29213
Fungi	
<i>Aspergillus niger</i>	
<i>Candida albicans</i>	ATCC 10231
<i>Trichophyton mentagrophytes</i>	UPCC 9533

- Test organisms: Applications

Antimicrobial Assay
Qualitative screening of natural and synthetic compounds

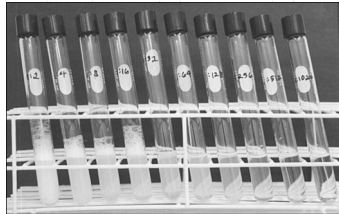


Antimicrobial assay
of Moringa extract



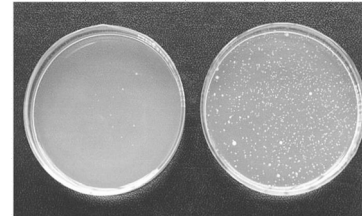
Antimicrobial assay
of an antibiotic formulation

▪ Test organisms: Applications



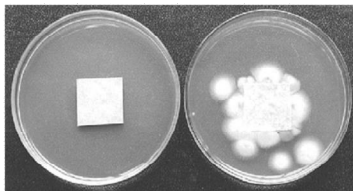
Determination of Minimum Inhibitory Concentration (MIC) of Biocidal Agents

▪ Test organisms: Applications



Percent kill Test
Efficacy testing of Disinfectants and other biocidal-containing materials and formulations

▪ Test organisms: Applications



Mold Resistance Test
American Society for Testing and Materials
ASTM Designation: D-4576-01

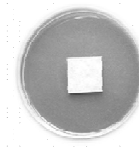
Determination of efficacy of antifungal agent in the material/matrix

▪ Test organisms: Applications

Materials tested include:



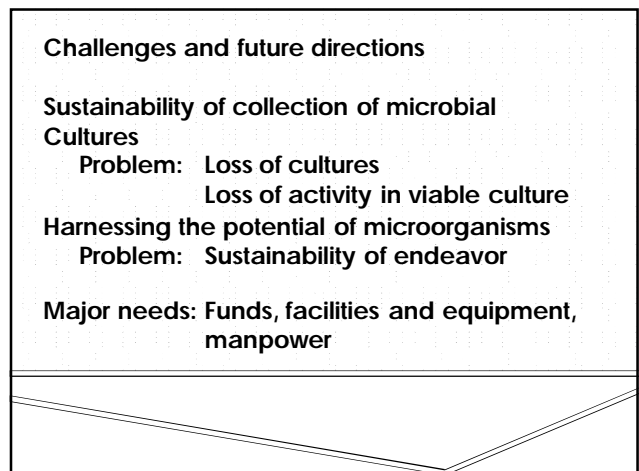
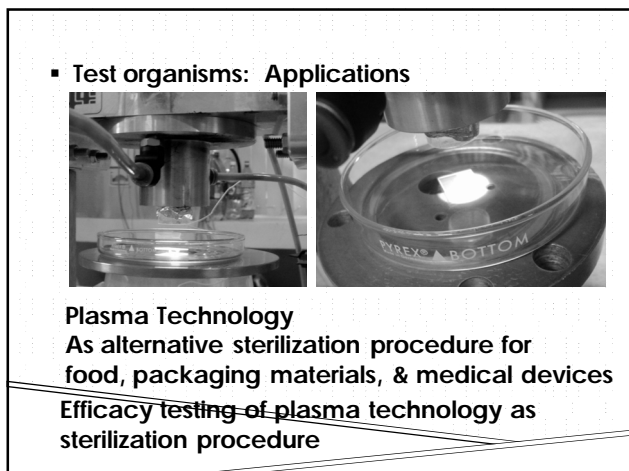
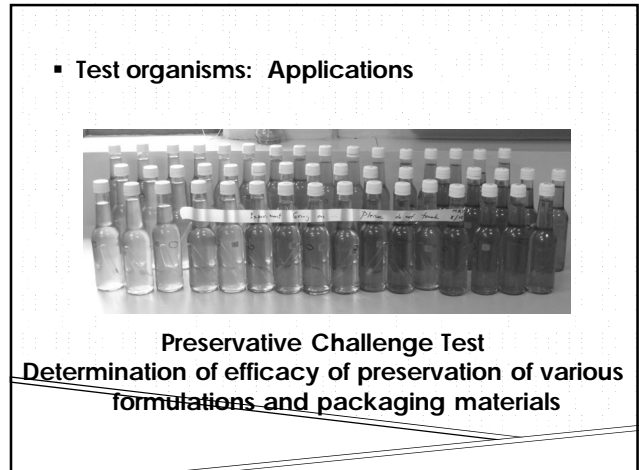
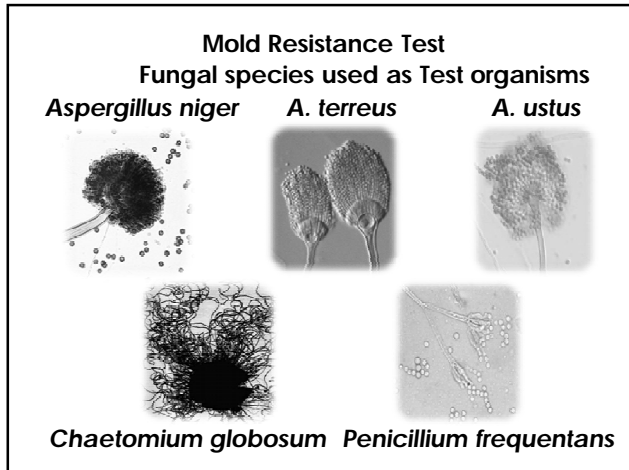
Baseball glove



Packaging material
(e.g. soap box)



Military boots



Future directions

- Upgrading microbial holdings
- Sustainable long-term preservation
- Upgrading of facilities to support culture collection
- Enhancing database
 - Data generation:
 - Phenotypic traits
 - Genotypic traits
 - Useful genes through genome sequencing
- Capacity building
 - Training in key techniques used in microbial systematics
 - Training in metagenomics relevant to metabolite production