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The Role of Antibiotics in Modulating Secondary Metabolite Production of Streptomycesgrown in Co-Culture

Bryan Sierra-Rivera
California State University, Monterey Bay

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The role of antibiotics in modulating secondary metabolite production of *Streptomyces* grown in co-culture.

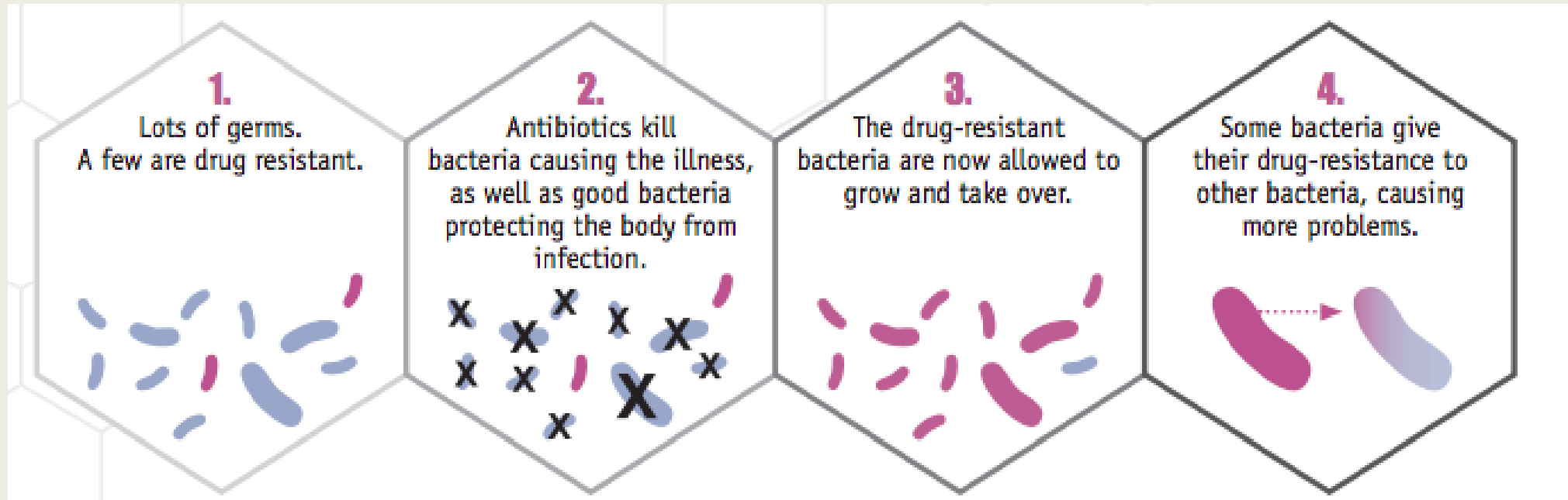
Bryan Sierra-Rivera ¹
Dr. Gregory C. Palmer ²



California State University ¹
MONTEREY BAY
Undergraduate Research Opportunities Center

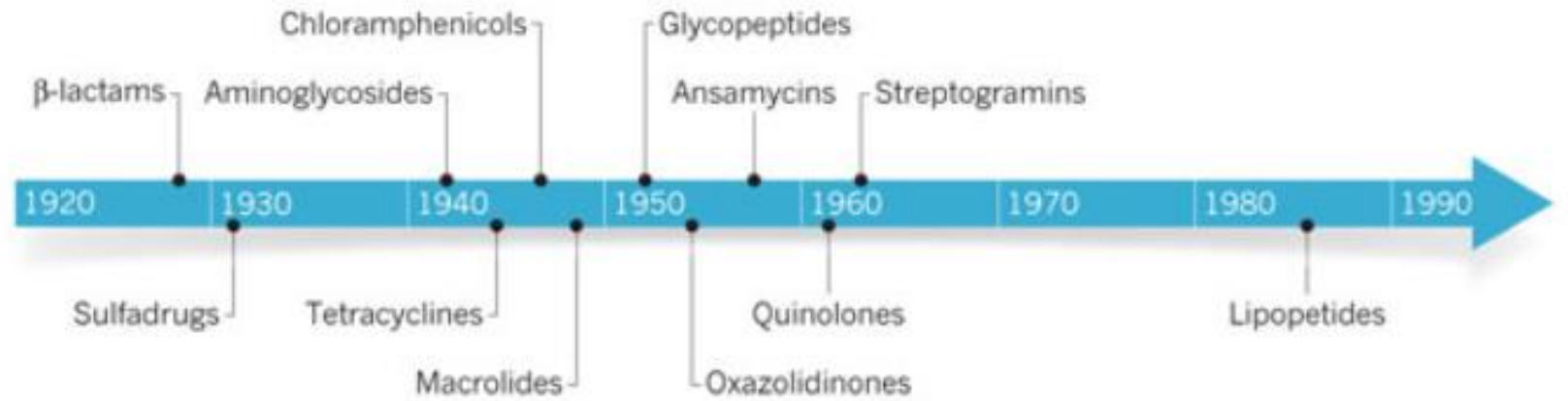
The University of Texas at Austin ²
College of Natural Sciences

How Antibiotic Resistance Happens



Examples of how it spreads:

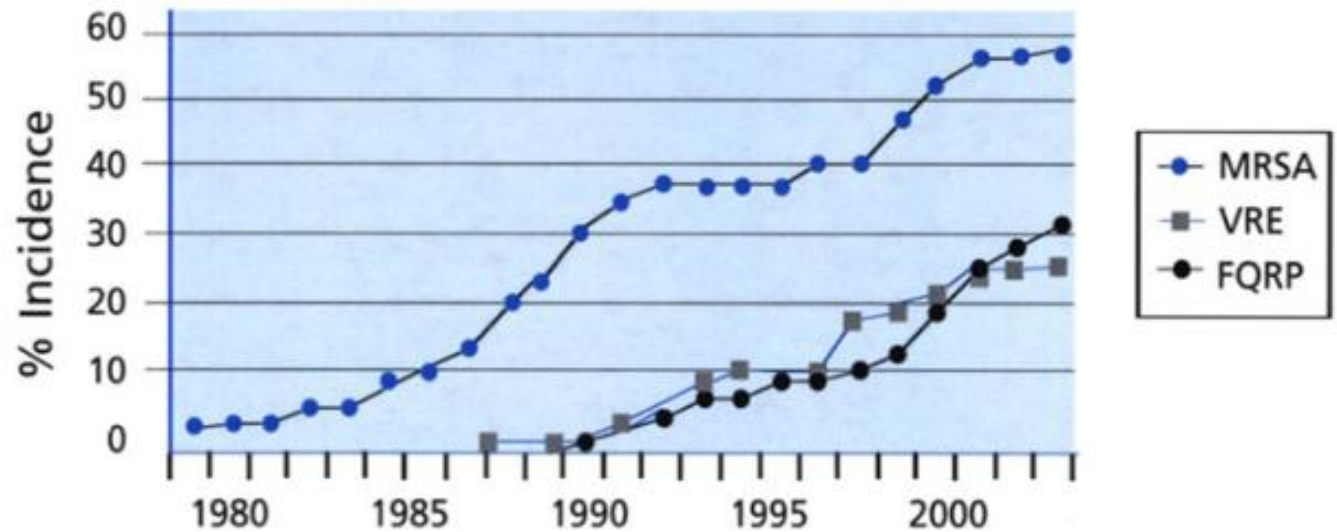
1. Livestock
2. Prescriptions



The Golden Era of Antibiotics

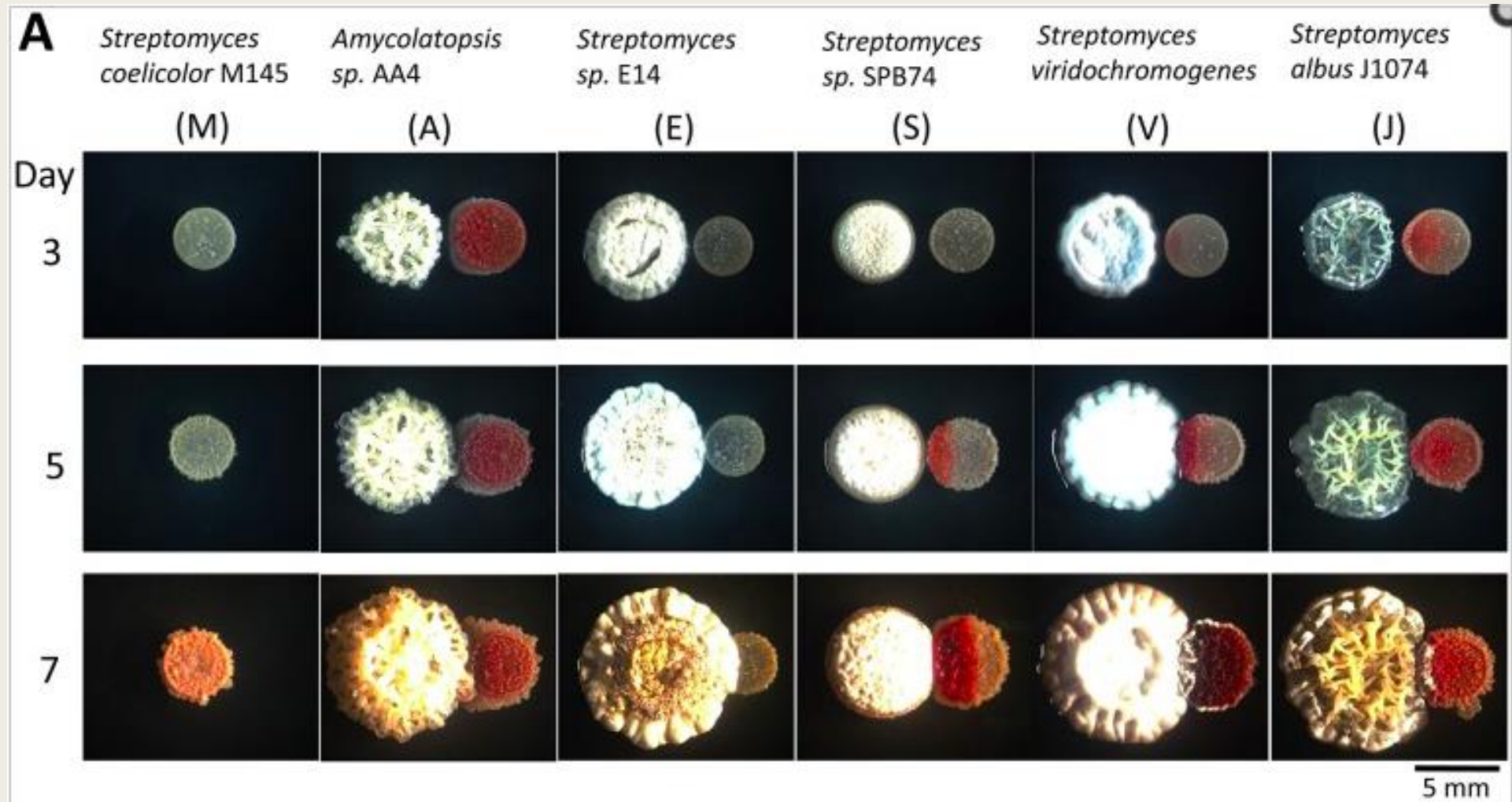
- Considered to be between the 20th century; specifically the 50's – 60's

Antibiotic resistance



MRSA = methicillin-resistant *Staphylococcus aureus*; VRE = Vancomycin-resistant *enterococci*
 FQRP = Fluoroquinolone-resistant *Pseudomonas aeruginosa*

Interspecies Interactions in co-cultures has led to new secondary metabolites.



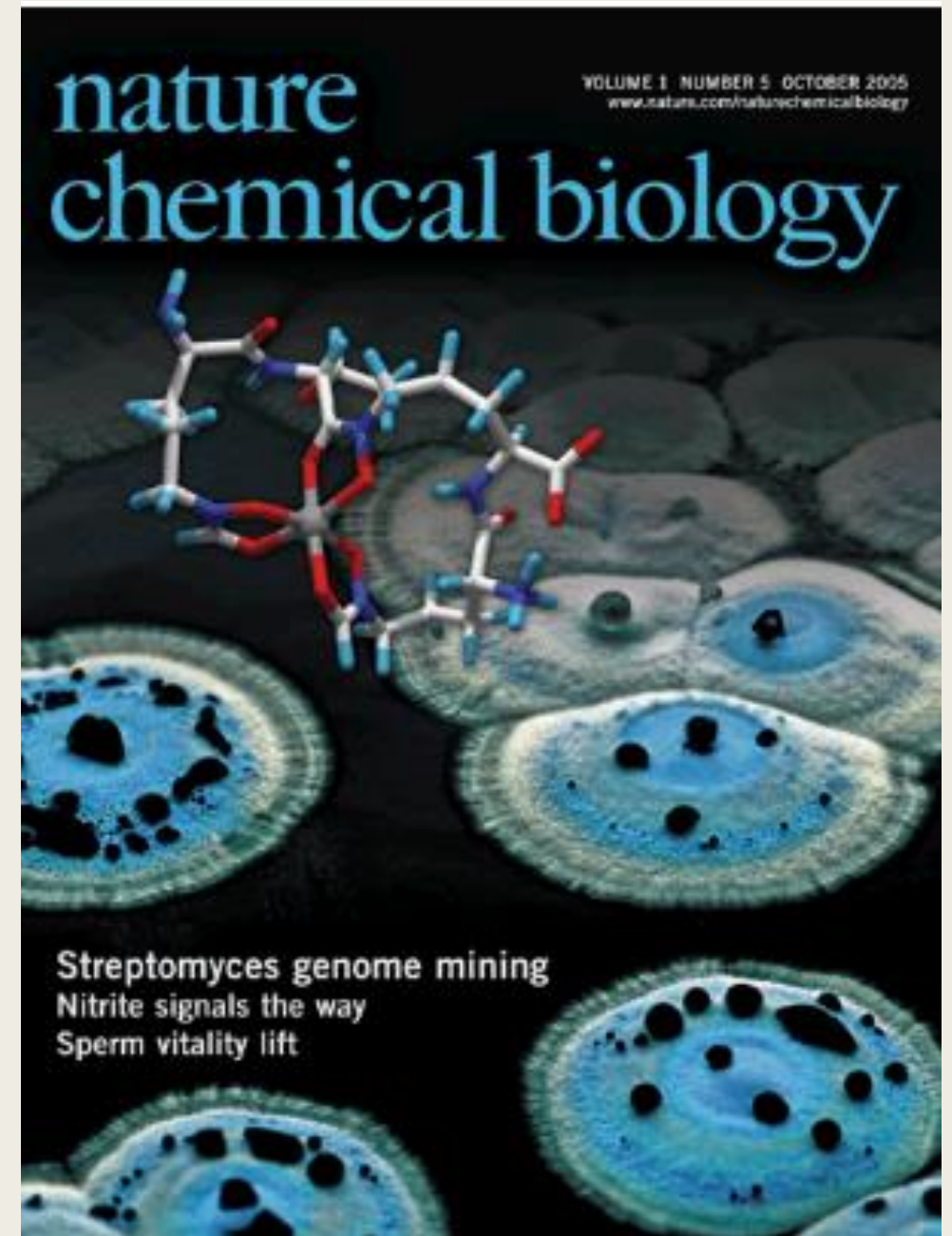
Traxler et al. 2013



PURPOSE:
**CAN ANTIBIOTICS BE USED IN CO-CULTURE
EXPERIMENTS TO PRODUCE NEW
SECONDARY METABOLITES**

An Introduction to *Streptomyces*

- *Streptomyces* is a genus of bacteria that contains many antibiotic producers
- *Streptomyces coelicolor* (wild type)
- *Streptomyces coelicolor* (M145)
- *Streptomyces osmaniensis*



Nature Chemical Biology cover from 2005

Isolating potential *Streptomyces*

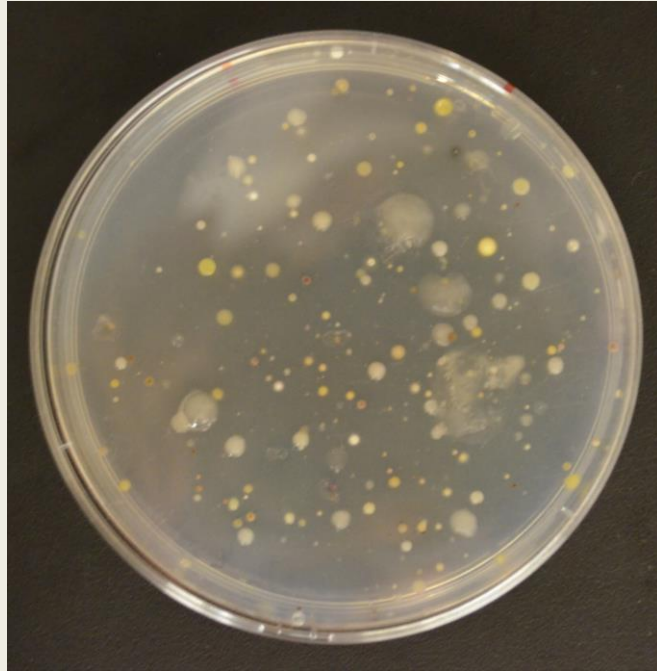


Plate by Matthew Co



Streptomyces osmaniensis gene for 16S ribosomal RNA, partial sequence, isolate: CDF2L1D13
Sequence ID: [LC136922.1](#) Length: 1434 Number of Matches: 1

Range 1: 163 to 1163		GenBank	Graphics	▼ Next Match	▲ Previous Match
Score	Expect	Identities	Gaps	Strand	
1805 bits(977)	0.0	989/1001(99%)	0/1001(0%)	Plus/Plus	
Query 1	CTGGANGGGGGTTNNAAGCTCCGGCGGTGAAGGANGAGCCCGCGCCTATCAGCTTGTG	60			
Sbjct 163	CTGGACGGGGGTTAAAAGCTCCGGCGGTGAAGGATGAGCCCGCGCCTATCAGCTTGTG	222			
Query 61	GTGAGGTAATGGCTCACCAAGGCGACGACGGGTAGCCGGCTGAGAGGGCGACCGGCCAC	120			
Sbjct 223	GTGAGGTAATGGCTCACCAAGGCGACGACGGGTAGCCGGCTGAGAGGGCGACCGGCCAC	282			
Query 121	ACTGGGACTGAGACACGGCCACAGACTCCCTACGGGAGGACAGTGGGGAATATTGCACAA	180			
Sbjct 283	ACTGGGACTGAGACACGGCCACAGACTCCCTACGGGAGGACAGTGGGGAATATTGCACAA	342			
Query 181	TGGGCGAAAGCCTGATGCAGCGACGCCCGGTGAGGGATGACGGCCTTCGGGTGTAAACC	240			
Sbjct 343	TGGGCGAAAGCCTGATGCAGCGACGCCCGGTGAGGGATGACGGCCTTCGGGTGTAAACC	402			
Query 241	TCTTTCAGCAGGGAAGAAGCGAAAAGTGACGGTACCTGCAGAAGAAGCGCGCTAACTAC	300			
Sbjct 403	TCTTTCAGCAGGGAAGAAGCGAAAAGTGACGGTACCTGCAGAAGAAGCGCGCTAACTAC	462			
Query 301	GTGCCAGCAGCCGCGGTAATACGTAGGGCGCAAGCGTTGTCCGGAATTATTGGGCGTAAA	360			
Sbjct 463	GTGCCAGCAGCCGCGGTAATACGTAGGGCGCAAGCGTTGTCCGGAATTATTGGGCGTAAA	522			
Query 361	GAGCTCGTAGGGCGGTTGTACAGTCGGGTGTGAAAGCCCGGGCTTAAACCCGGGTCTGC	420			
Sbjct 523	GAGCTCGTAGGGCGGTTGTACAGTCGGGTGTGAAAGCCCGGGCTTAAACCCGGGTCTGC	582			
Query 421	ATTCGATACGGGCTAGCTAGAGTGTGGTAGGGAGATCGGAATTCCTGGTGTAGCGGTGA	480			
Sbjct 583	ATTCGATACGGGCTAGCTAGAGTGTGGTAGGGAGATCGGAATTCCTGGTGTAGCGGTGA	642			
Query 481	AATGCGCAGATATCAGGAGGAACACCGGTGGCGAAGGCGGATCTCTGGGCCATTACTGAC	540			
Sbjct 643	AATGCGCAGATATCAGGAGGAACACCGGTGGCGAAGGCGGATCTCTGGGCCATTACTGAC	702			
Query 541	GCTGAGGAGCGAAAAGCGTGGGAGCGAACAGGATTAGATACCTCGGTAGTCCACGCCGTA	600			
Sbjct 703	GCTGAGGAGCGAAAAGCGTGGGAGCGAACAGGATTAGATACCTCGGTAGTCCACGCCGTA	762			
Query 601	AACGGTGGGAAC TAGGTGTTGGCGACATTCACGTCGTCGGTGCAGCAGTAAACGCATTA	660			
Sbjct 763	AACGGTGGGAAC TAGGTGTTGGCGACATTCACGTCGTCGGTGCAGCAGTAAACGCATTA	822			
Query 661	AGTTCCCCGCTGGGGAGTACGGCCGCAAGGCTAAAACTCAAAGGAATTGACGGGGGCC	720			
Sbjct 823	AGTTCCCCGCTGGGGAGTACGGCCGCAAGGCTAAAACTCAAAGGAATTGACGGGGGCC	882			
Query 721	GCACAAGCAGGGAGCATGTGGCTTAATTCGACGCAACCGCAAGAACCTTACCAAGGCTT	780			
Sbjct 883	GCACAAGCAGGGAGCATGTGGCTTAATTCGACGCAACCGCAAGAACCTTACCAAGGCTT	942			
Query 781	GACATACCCGAAACGCTCTGGAGACAGGCGCCCTTGTGGTCCGGTGTACAGGTGGTGC	840			
Sbjct 943	GACATACCCGAAACGCTCTGGAGACAGGCGCCCTTGTGGTCCGGTGTACAGGTGGTGC	1002			
Query 841	ATGGCTGTCGTCAGCTCGTGTGAGATGTTGGGTTAAGTCCCAGCAACGAGCGCAACCC	900			
Sbjct 1003	ATGGCTGTCGTCAGCTCGTGTGAGATGTTGGGTTAAGTCCCAGCAACGAGCGCAACCC	1062			
Query 901	TTGTTCTGTGTTGCCAGCATGCCCTTCGGGTGATGGGACTCACAGGAGACCGCNGGG	960			
Sbjct 1063	TTGTTCTGTGTTGCCAGCATGCCCTTCGGGTGATGGGACTCACAGGAGACCGCNGGG	1122			
Query 961	TCAACTCGGAGNAGNNGGGGACGACGTCAGTCNTCNTGC	1001			
Sbjct 1123	TCAACTCGGAGGAGGTGGGGACGACGTCAGTCATCATGC	1163			

Streptomyces osmaniensis was identified through sequencing.

Liquid cultures

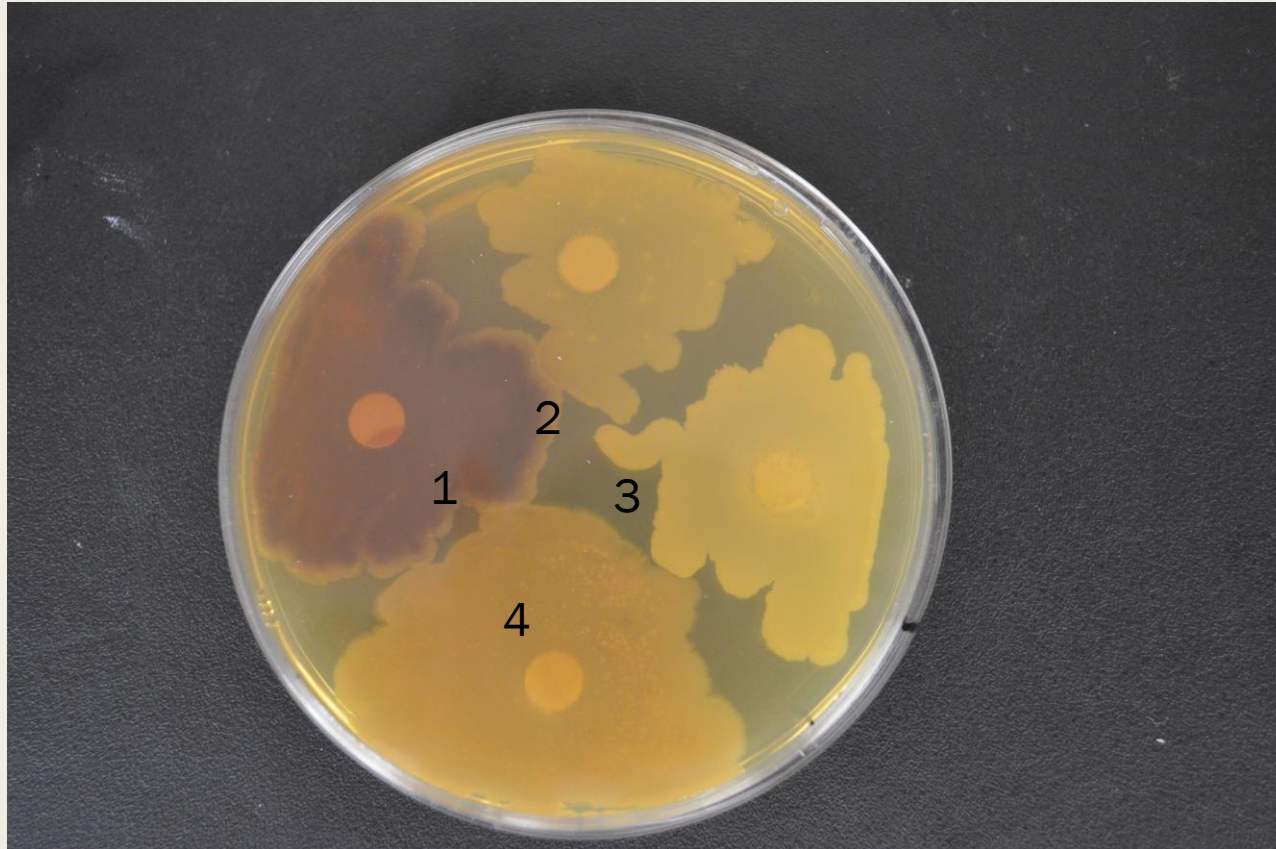


Image 1

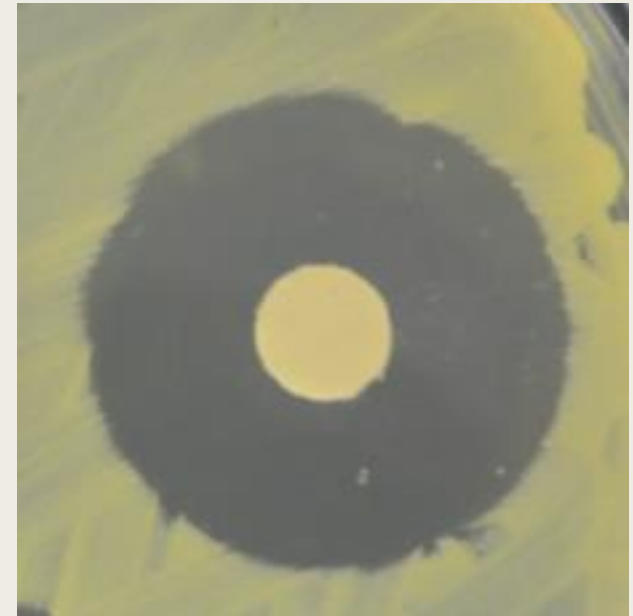
- TSB-YE¹
- ISP2²

Identifying what media is ideal for secondary metabolite production.

Cell Disk Assay

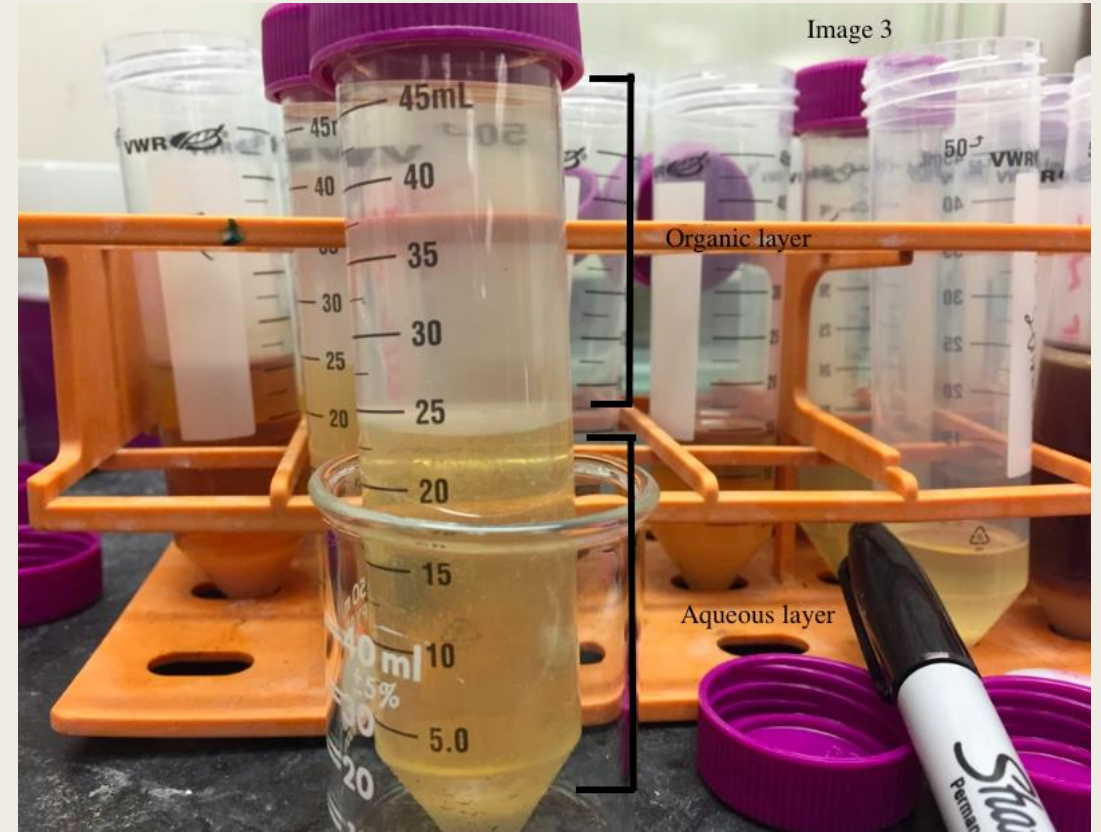
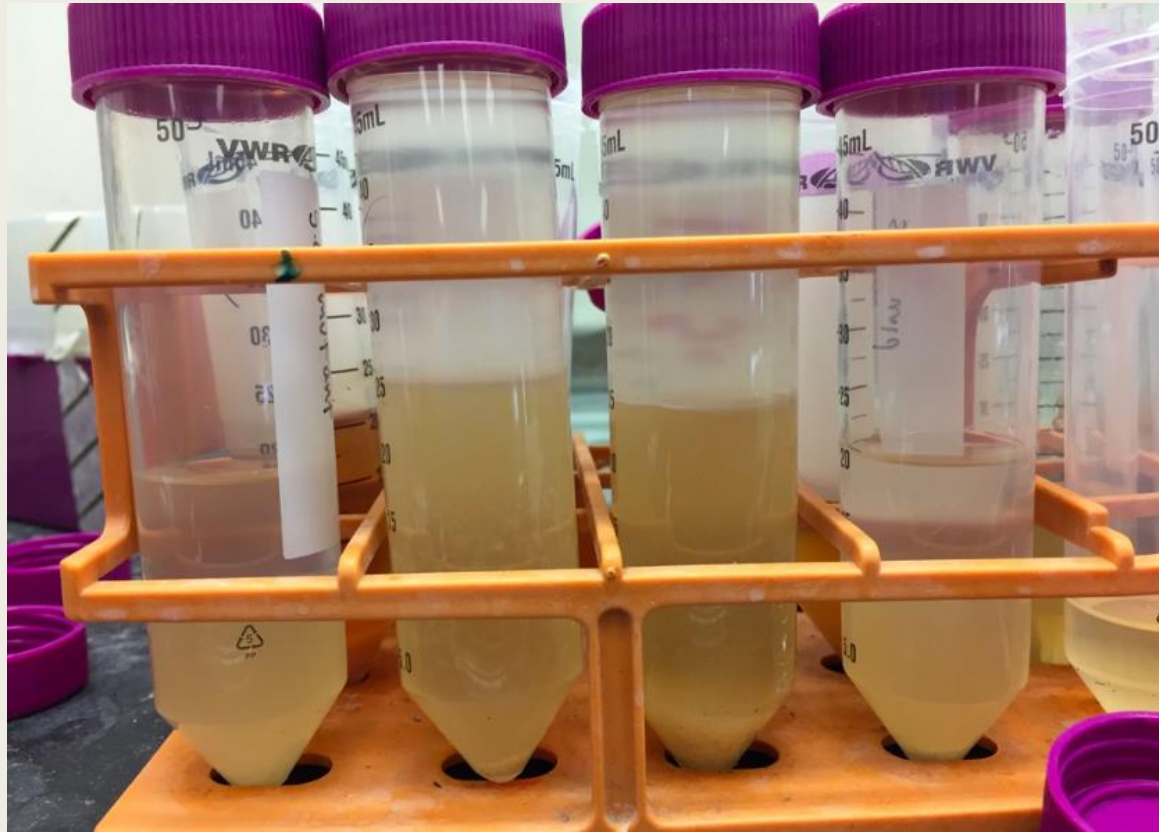


1. *Serratia marcescens*
2. *Pseudomonas sp.*
3. *Staphylococcus aureus*
4. *Bacillus sp.*



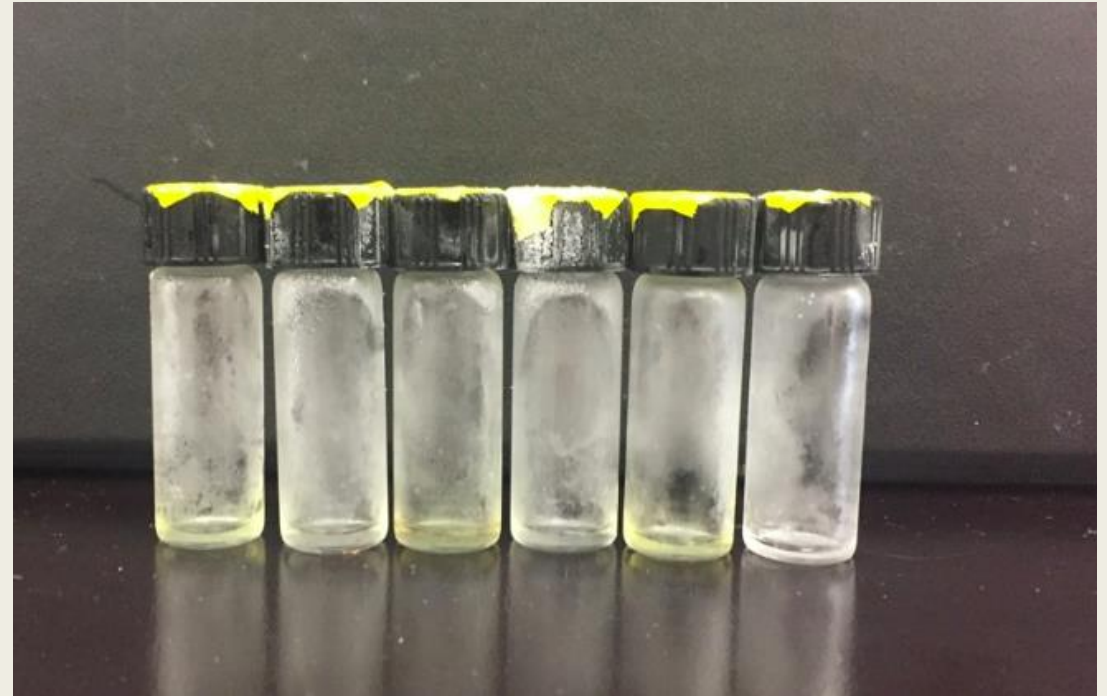
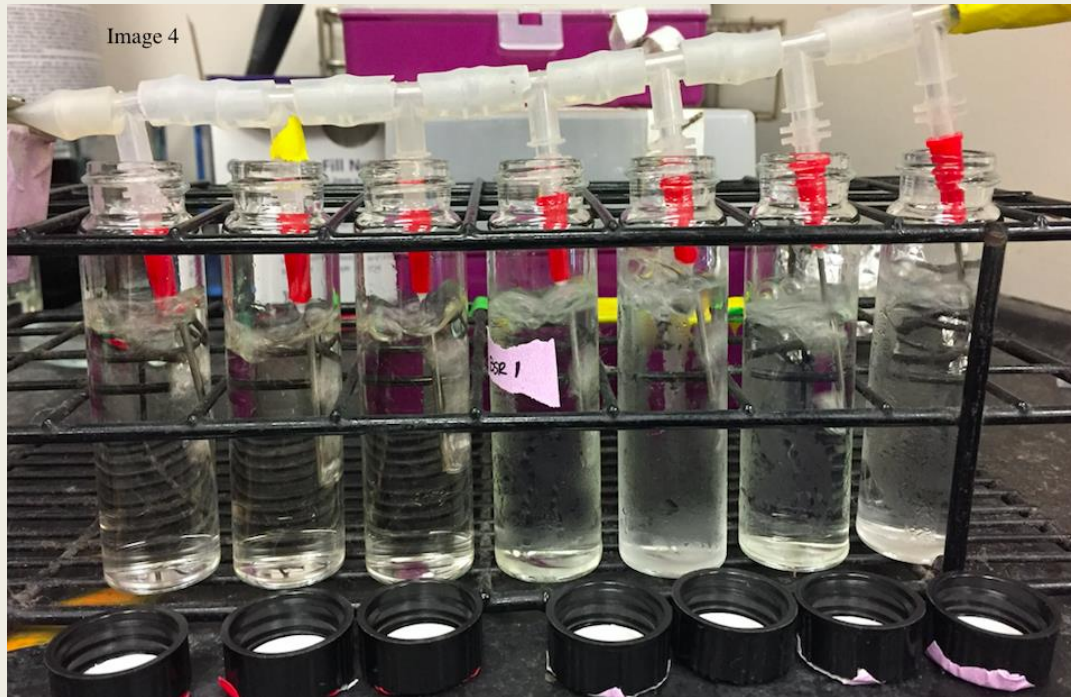
No inhibition when conducting "live" cell disk assays.

Ethyl Acetate extractions



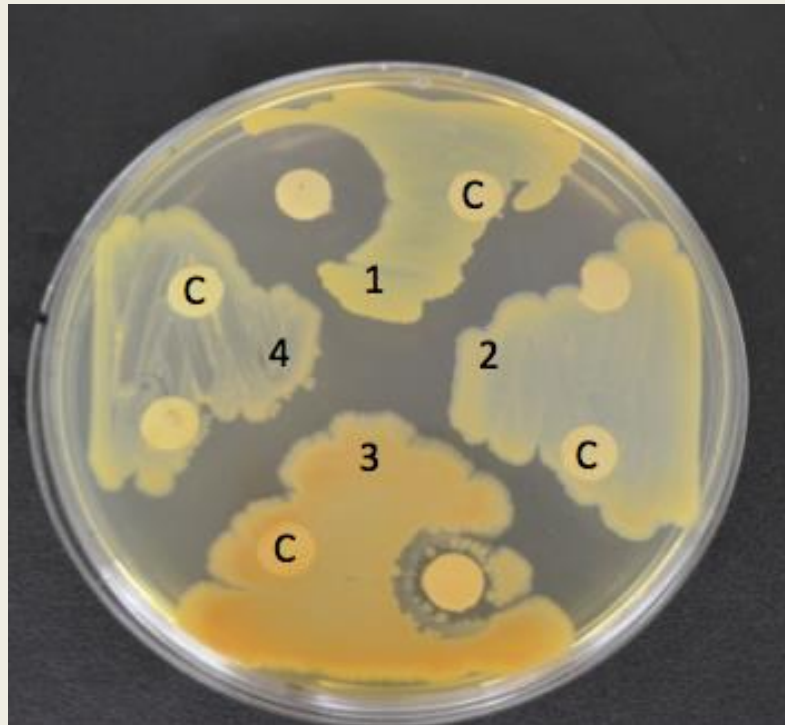
Ethyl Acetate extractions help isolate and test for antibiotics more efficiently.

Drying down



Organic layer dried down to collect potential antibiotics.

Extract disk assay

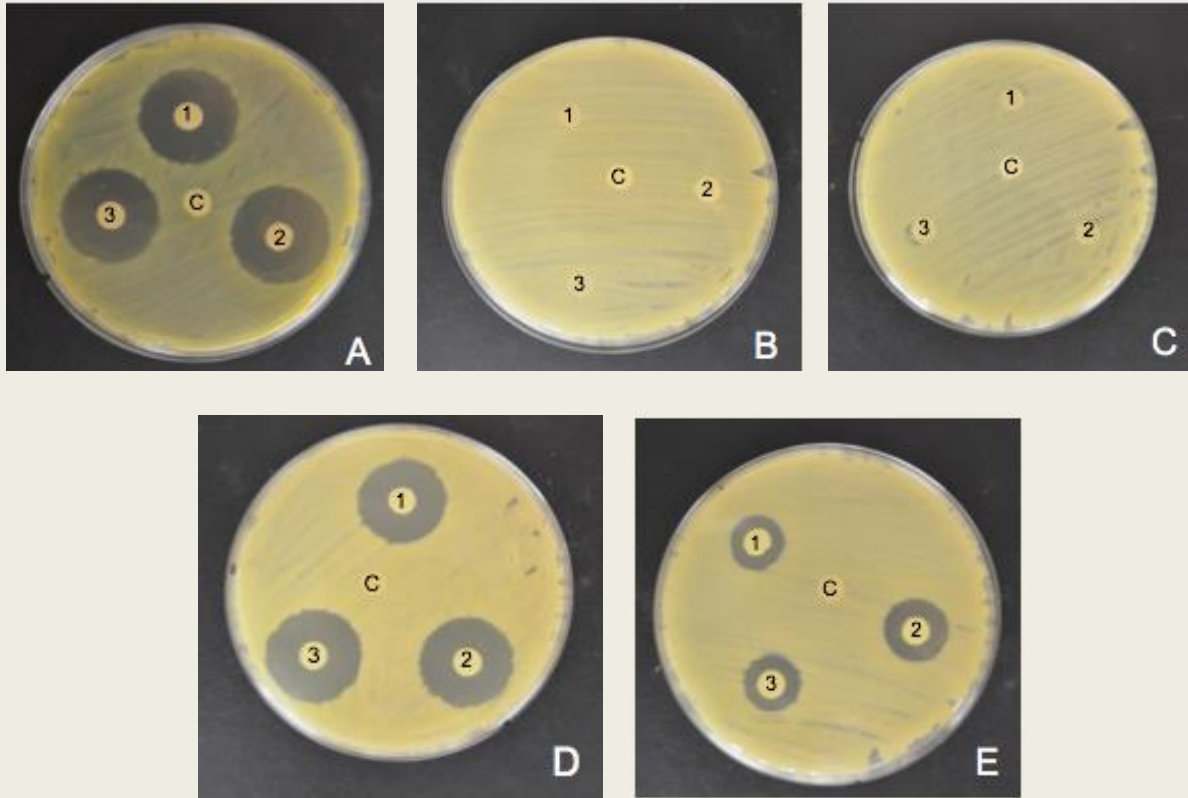


1. *Pseudomonas sp.*
2. *Rhodococcus sp.*
3. *Bacillus sp.*
4. *E. coli*

C : Control

Tests using Ethyl Acetate extractions led to inhibition on *Pseudomonas*.

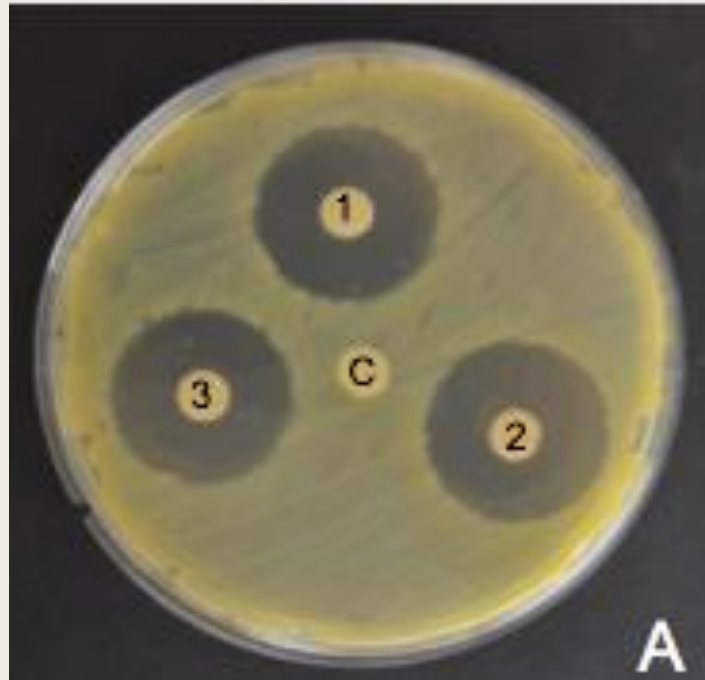
Disk Assays



- A. *S. osmaniensis*
- B. *S. coelicolor*
- C. *S. coelicolor* (M145)
- D. *S. osmaniensis* + *S. coelicolor* (M145)
- E. *S. osmaniensis* + *S. coelicolor*

Compared inhibition of mono-cultures vs. co-cultures to determine benefits of interspecies interactions.

Results



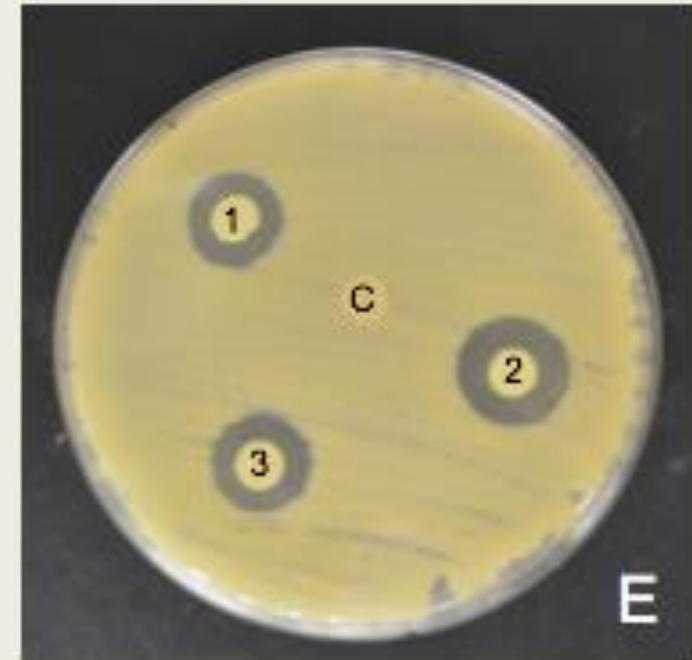
S. osmaniensis

24 mm



S. osmaniensis + *S. coelicolor*
(M145)

22 mm



S. osmaniensis + *S. coelicolor*

15 mm

Average diameter (mm) of mono-cultures of *S. osmaniensis* led to larger zones of inhibition compared to that of co-cultures with both wild and mutant type.

Future Directions

- Co-culturing of different antibiotic producing *Streptomyces spp.*
- Different nutrient media
- Antibiotics produced by the isolate should be characterized by using Liquid Chromatography Mass Spectrometry.



Antibiotic secretion of *Streptomyces spp.* colony

Acknowledgements

- I thank Dr. Gregory C. Palmer and the mentors in the Antibiotics Discovery and Function laboratory at the University of Texas at Austin for their guidance and support and Matthew Co for his soil sample plate image. I thank the Summer Undergraduate Program for Experiential Research (SUPER) 2017 program at the University of Texas at Austin and the Undergraduate Opportunities Center (UROC) at California State University, Monterey Bay.

Reference

- Traxler M., Watrous J., Alexandrov T., Dorrestein P., Kolter R. Interspecies Interactions Stimulate Diversification of the *Streptomyces coelicolor* Secreted Metabolome. *PubMed*. 2013. Available from: <https://www.ncbi.nlm.nih.gov/pmc/articles/PMC3747584/>
- Nature Chemical Biology, 2005
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- Davies J. Where have All the Antibiotics Gone? *The Canadian Journal of Infectious Diseases & Medical Microbiology*. 2006;17(5):287-290.

Thank you for your time.
Any Questions?