Presence of Antibiotic Resistant Bacteria and Antibiotic Res <u>Collin Toups (UNDE)¹, Christopher Oubre¹, Ramaraj Dr</u> ¹Department of Biological Sciences, Nicholls Stat ²Barataria-Terrebonne National Estuary Program, Nicho 3Department of Sciences, Thibodaux High

Abstract

Since the dawn of the commercialization of antibiotics, antibiotic resistant bacteria (ARB's) and antibiotic resistance genes (ARG's) have been a rapidly growing problem. As international travel is popularized, these ARB's and ARG's are able to move to new places using humans as a vector in a somewhat anthropogenic way. However, these potentially harmful bacteria and genes could be spreading in another way less reliant on human involvement. The Nicholls State University Biotechnology lab has been studying the increasing presence of ARBS' and ARG's in local waterways for a number of years now. These ARB's and ARG's have been correlated to the consumer, commercial, and hospital related improper disposal of antibiotics and their presence in the waters of Southeast Louisiana is well studied. Also present in these waterways are several species of migratory birds which use Louisiana either as a stopover point or endpoint for migration along the Mississippi Flyaway route. It is possible that these migratory birds could serve as a vector for the aforementioned ARB's and ARG's, allowing these antibiotic resistant bacteria and genes to travel all along the migratory route, from the southern coast of the United states to central Canada and everywhere in between. In this study, fecal and cloacal bacterial samples were taken from Louisiana migratory birds. These samples were enriched in TSB and streaked onto TSA to collect isolates. These isolates were then tested for the presence of ARB's and ARG's using the Kirby Bauer Assay and PCR techniques respectively. Identification of bacteria was achieved using 16srRNA sequencing techniques.

Introduction

Antibiotic resistance in bacteria is a natural phenomenon arising from the selective pressure of exposure to clinical antibiotics and is also caused by misuse and overuse of antibiotics. This can cause considerable public health problems. Within the past decade alone, the Center for Disease Control has identified various bacteria as posing serious, urgent, and concerning threats to the US healthcare system due to its resistance to antibiotics (CDC, 2013). Improper usage and disposal of antibiotics by consumers, hospitals, industries, and sewage treatment plants has furthered the emergence of antibiotic resistance in the waterways of Louisiana. Furthermore, antibiotic resistant bacteria (ARB) has been found in raw sewage and treated sewage which is released into the environment in the waterways. Ultimately, the ARB and antibiotic resistance genes (ARGs) in the treatment plants affect wildlife, where the treated water is being released (Naquin et al., 2015). These water sources are home to much of Louisiana's wildlife including migratory birds. Because Louisiana is part of the Mississippi Flyway migratory route, It is possible that birds may be serving as a vector for transportation of antibiotic resistant and antibiotic resistant bacterial genes from our waterways all throughout North America leading to an overall increase in antibiotic resistance on the continent.

Objectives:

Isolate and identify various antibiotic resistant bacteria (ARB) and antibiotic resistant genes (ARG) in the fecal matter and cloacae of migratory birds in Louisiana.

Study the ecological relevance of the ARB and ARGs in our waterways. Create public awareness on the presence of ARB and ARGs in the local water samples and seafood.

Methods:

1. Pure cultures of bacteria were isolated and identified from various migratory bird samples

Colonies were isolated as pure cultures in Tryptic Soy Agar (TSA) using quadrant streak method from various local water and seafood samples listed in the result section.

2. Antibiotic resistance was observed using Kirby-Bauer Assay:

Pure Cultures were streaked as a bacterial lawn onto Mueller Hinton (MH) Agar and antibiotic discs were stamped onto the agar using an antibiotic disc

dispenser.



sistant Genes in the Migratory Birds of Louisiana		
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te University, Thibo	daux, LA 70310	Nicholls Biotech
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	Results cont.:	
n (CC), Vancomycin (VA),		AMCBVAPCCSCommon Name: Northern Cardinal Species Code: NOCAFConsortium23.87000014.23Isolate a21.45*000015.39
and Penicillin (P).		Age: AHY C Consortium 12.3 11.62 12.7 5.56 18.88 16.24 Sex: M Isolate b 12.82 10.32 8 0 16.26 20.27 Location: HOME Isolate b 11.76 10.95 2 0 19.84 21.08
no color), Intermediate		Table 10. Kirby Bauer results for Northern Cardinal #1
	Image 12. Northern Cardinal (<i>Cardinalis cardinalis</i>). Photograph obtained by Delaina LeBlanc	AMC B VA P CC S Common Name: Northern Cardinal Species Code: NOCA F Consortium 21.14 0 0 0 13.54 Band Number: 2451-36346 Isolate b 19.09 0 0 0 19.17 Age: AHY C Consortium 16.28 8.56 0 0 0 10.9
ResistantIntermediateSusceptiblehycin (CC) ≤ 14 $> 14 \leq 20$ > 20 hycin (VA) ≤ 9 $> 9 \leq 11$ > 11 mycin (S) ≤ 11 $> 11 \leq 14$ > 14	a contraction of the contraction	Sex: M Isolate a 13 11.61 18.12 0 10.55 12.04 Date: 1-19-20 Isolate b 0 0 0 0 0 14.16 Location: HOME B Consortium Isolate h Isolateh Isolateh Isola
Implified $S = 11$ $> 11 \le 14$ > 14 Illin (AMC) ≤ 13 $> 13 \le 17$ > 17 sin (B) ≤ 8 $> 8 \le 12$ > 12 n (P) ≤ 19 $> 19 \le 27$ > 27 e 1. Antibiotic susceptibility standards		AMCBVAPCCSCommon Name: Northern Cardinal Species Code: NOCAFConsortium9.79000013.68Isolate a20.140000000
I for assessing Kirby Bauer assay		Band Number: 2451-53033 Isolate b 21.24 0 0 0 0 14.52 Age: SY C Consortium 6.1 0 4.16 22.2 26.7 10.2 Sex: F Isolate b 12.7 13.3* 14.16 9.9 21.95 15.23 Date: 1-19-20 Isolate b 16.5 12.97 17.96 10.09 24.01 16.53 Location: HOME B Consortium Isolate b Isolate b </td
	Year-round	Table 12. Kirby Bauer results for Northern Cardinal #3
AMC B VA P CC S ned Kinglet F Consortium 11.37 0 0 0 0 11.76 Isolate a 0 0 0 0 0 11.21* Isolate b 10.74* 0 0 0 0 17.16*	Image 13. Ruby-crowned Kinglet range map ta from the Cornell Lab's website "All About Birds	ken s"
C Consortium 17.09* 0 0 0 0 19.61 Isolate a 0 0 0 0 0 19.61 Isolate b 0 0 0 0 0 14.11* Isolate b 0 0 0 0 0 13.43 1B Consortium Ier results for Ruby-crowned Kinglet #1	Discussion and Conclusions	
Image: American series American series P CC S ned F Consortium 0 0 0 0 0 14.2* Isolate a 0 0 0 0 0 11.06 Isolate b 0 0 0 0 12.78	Bird fecal and cloacae samples were collected from several birds at a few	
B Consortium	different sites in Louisiana and both the consortiums and isolates taken from the consortiums were tested for the presence of antibiotic resistant bacteria.	
AMC B VA P CC S vned Kinglet F Consortium 8.39* 0 0 0 0 12.14 Isolate a 0 0 0 0 0 11.76	• All the bird fecal samples and most of the bird cloacae samples tested showed the presence of various antibiotic resistant bacteria.	
B Consortium	• The presence of antibiotics in our waterways will exert selection pressure on bacteria to develop antibiotic resistance which will inevitably lead to migratory	
	 birds spreading these antibiotic resistant bacteria throughout North America. Horizontal transfer of antibiotic resistance genes will allow non-native 	
AMC B VA P CC S d Vireo F Consortium 26.67 0 0 0 14.07*	bacteria introduced by birds to spread their antibiotic resistance to native bacteria.	
Isolate a 19.65 0 0 0 0 15.24 Isolate b 18.49 0 0 0 0 11.8 C Consortium 15.0 10.34 28.24 27.13 18.84 Isolate a 0 0 0 0 0 10.57 Isolate b 0 0 0 0 0 13.57	a Bacterial transformation Release of DNA	
iuer results for White-eyed Vireo	Donor cell b Bacterial transduction b Bacterial transduction b Bacterial transduction	
	and the second s	
Image: Second state AMC B VA P CC S Image: Second state Consortium 0 0 0 0 0 0 Isolate a 0 0 0 0 0 15.45 Isolate b 0 0 0 0 14.58	Phage-infected donor cell phage Recipient cell C Bacterial conjugation	
C Consortium 37.58 14 16.16 18.72 28.66 23.74 Isolate a 15. 15. 19.36 17.66 Isolate b 25. 19.03 10.14 22.68 12.25 B Consortium 1 1 1 1 1	Transposon Donor cell Recipient cell Image 14. Horizontal transfer of antibiotic resistance genes	
nuer results for Orange-crowned Warbler #1AMCBVAPCCSFConsortium14.57000014.53Isolate a11.76*00000		
Isolate b11.72000014.71BConsortiumImage: state sta	Future Work:	
uer results for Orange-crowned Warbler #2	Future work will focus on identifying th sample consortiums and identifying the	e isolates obtained from the bird e presence of antibiotic resistant genes
	within these isolates. Future work will a from the sites where the bird samples w	also focus on obtaining water samples were collected, and testing these sites
AMC B VA P CC S arrow F Consortium 0 0 0 0 14.69 Isolate a 0 0 0 0 17.7 Isolate b 0 0 0 0 11.55	for the presence of antibiotics, antibiot resistant genes as well as performing w	ic resistant bacteria, and antibiotic ater quality tests on the areas.
uer results for Swamp Sparrow	Citat	tions:
	Najdenski H, Dimova T, Zaharieva MM, Nikolov B, Pe Nikolova I, Zehtindjiev P, Peev S, Trifonov	trova-Dinkova G, Dalakchieva S, Popov K, Hristova- va-Hristova A, Carniel E, Panferova YA, and Tokarevich
	pathogens." Canadian Journal of Microbiology.: 91 Naquin, A., Shrestha, A., Sherpa, M., Nathaniel, R., ar	nd Boopathy, R. (2015). Presence of antibiotic
row F Consortium 32.5 24.5 14.03* 9.04 34 23.09	resistance genes in a sewage treatment p technology, 188, 79-83. Ramey AM, Hernandez J, Tyrlöv V, Uher-Koch BD. Sc	biant in Thibodaux, Louisiana, USA. Bioresource hmutz JA, Atterby C, Järhult JD, Bonnedahl J. 2018.
Isolate a 24.01 0 0 0 0 24.12 Isolate b 20.36 10.76 15.24 0 4.04 19.06 C Consortium 12.12* 0 0 0 0 19.16 Isolate a 0 0 14.7* 0 0 0 Isolate b 12.24 0 15.65 0 17.51 12.35 B Consortium 1 1 1 1 1 1	"Antibiotic-Resistant Escherichia coli in N 72-81	Migratory Birds Inhabiting Remote Alaska." Ecohealth:
er results for House Sparrow	Resistance Mediated by Wild Birds." Frontier www.cdc.gov/drugresistance/biggest_threats.html?	s in Microbiology. 9: 745 CDC_AA_refVal=https%3A%2F%2Fwww.cdc.gov%2Fdr
	ugresistance%2Fthreat-report- 2013%2 https://www.allaboutbirds.org/guide/Ruby-crowned	Findex.html. I_Kinglet/maps-range
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