

## **Environment and Natural Resources Trust Fund**

2024 Request for Proposal

#### **General Information**

**Proposal ID: 2024-073** 

Proposal Title: Enhancing Wastewater Treatment while Bioprospecting for Novel Pharmaceuticals

## **Project Manager Information**

Name: Timothy LaPara

Organization: U of MN - College of Science and Engineering

**Office Telephone:** (612) 624-6028

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### **Project Basic Information**

**Project Summary:** We will generate genome sequences of bacteria growing in wastewater treatment bioreactors, allowing us to improve phosphorus and nitrogen removal from wastewater in Minnesota and to discover novel pharmaceutical compounds.

Funds Requested: \$690,000

Proposed Project Completion: June 30, 2027

LCCMR Funding Category: Water Resources (B)

## **Project Location**

What is the best scale for describing where your work will take place?

Statewide

What is the best scale to describe the area impacted by your work?

Statewide

When will the work impact occur?

In the Future

#### **Narrative**

#### Describe the opportunity or problem your proposal seeks to address. Include any relevant background information.

Municipal wastewater treatment is essential for protecting the quality of Minnesota's lakes and rivers. Historically, municipal wastewater treatment focused on the removal of biodegradable organic carbon (measured as "biochemical oxygen demand"), but modern wastewater treatment also focuses on removing phosphorus and nitrogen because of their ability to stimulate the growth of nuisance photosynthetic organisms (plants, algae, and cyanobacteria) in inland and coastal waters, respectively. Many municipal wastewater treatment facilities are already intentionally removing phosphorus and the Minnesota Pollution Control Agency is considering opportunities to reduce Minnesota's contribution of nitrogenous pollution to the hypoxic zone in the Gulf of Mexico from both point and non-point sources.

While there are numerous bioreactors designs that can successfully remove phosphorus and nitrogen from municipal wastewater, we have a relatively poor understanding of the microorganisms responsible for removing these contaminants. For example, our knowledge of ammonia-oxidizing bacteria (necessary for nitrogen removal) is based on a few model organisms that were isolated many years ago; even worse, the organism(s) necessary for phosphorus removal have never been isolated. Because of this limited knowledge, phosphorus and nitrogen removal from municipal wastewater can be unreliable, particularly in the cold climate that we enjoy in Minnesota.

What is your proposed solution to the problem or opportunity discussed above? Introduce us to the work you are seeking funding to do. You will be asked to expand on this proposed solution in Activities & Milestones.

The ideas for this proposed research come from two past/ongoing projects funded by LCCMR/ENRTF (M.L. 2016, Chp. 186, Sec. 2, Subd. 04k and M.L. 2021, Chp. 6, Art. 6, Sec. 2, Subd. 04j).

We propose to use proximity-ligation DNA sequencing to produce genome sequences of numerous bacteria from 30 municipal wastewater treatment bioreactor samples. These samples will be collected from 10 wastewater treatment facilities that perform enhanced biological phosphorus removal or that support novel/important nitrogen-removing microorganisms. We expect to obtain dozens of bacterial genome sequences from each of these samples; these genome sequences will then inform us on the optimal wastewater treatment conditions that are required to support these critically important organisms.

In addition, our prior work suggests that many of the organisms growing in wastewater treatment bioreactors can produce novel secondary metabolites that could have significant value to the pharmaceutical industry. We therefore also propose to "bioprospect" these genomic sequences for the potential to produce novel pharmaceuticals and then to use these same genomic sequences to guide the culturing of these organisms in the laboratory. These cultured organisms could then be used to produce these novel molecules on an industrially relevant scale.

What are the specific project outcomes as they relate to the public purpose of protection, conservation, preservation, and enhancement of the state's natural resources?

This project will generate genome sequences from bacteria growing in municipal wastewater treatment bioreactors. This data will help us identify the optimal bioreactor conditions for removing phosphorus and nitrogen from wastewater, which are current (phosphorus) and future (nitrogen) priorities for the State of Minnesota. In addition, we will also "bioprospect" these genome sequences for the ability to produce biomolecules of potential value to the pharmaceutical industry. Simultaneously, these genome sequences will suggest methods to cultivate these organisms in the laboratory.

#### **Activities and Milestones**

# Activity 1: Collect samples from municipal wastewater treatment bioreactors for proximity-ligation DNA sequence analysis

**Activity Budget:** \$100,000

#### **Activity Description:**

Bioreactor samples will be collected from 10 different wastewater treatment facilities within the State of Minnesota on two separate occasions (i.e., a "summer" sample and a "winter" sample). These wastewater treatment facilities will be selected based on prior research (M.L. 2016, Chp. 186, Sec. 2, Subd. 04k) in which microbiome profiles identified the prominent bacteria involved in nitrogen removal and phosphorus removal. In addition, monthly samples will be collected from a single wastewater treatment bioreactor so that community dynamics can be tracked for an entire calendar year. These samples will be sent to Phase Genomics, who will perform their proprietary proximity-ligation DNA sequencing and subsequent analysis. That is, Phase Genomics will sequence DNA from our samples, and in return, they will send us 20-50 genome sequences from each sample.

#### **Activity Milestones:**

Description	Approximate Completion Date
Collect wastewater samples and send them to Phase Genomics for DNA sequence analysis	August 31, 2025
Obtain all genome sequence data from Phase Genomics	December 31, 2025

## Activity 2: Analyze genomes sequences for the genetic ability for phosphorus and nitrogen removal

Activity Budget: \$206,722

#### **Activity Description:**

The first step of this activity will be to compare the genome sizes and sequences of specific populations of bacteria that are responsible for nitrogen (Nitrosomonas spp., Nitrosospira spp., Nitrotoga spp., and Nitrospira spp.) and for phosphorus removal (Tetrasphaera-like and Rhodocyclus-like); these genome sequences will also be compared to previously sequenced genomes that are already published in public databases. In the second step of this activity, we analyze the metabolic pathways that are found in these genome sequences, which should suggest the ideal bioreactor conditions (e.g., dissolved oxygen concentrations, pH, temperature, etc.) for each of these organisms to proliferate during municipal wastewater treatment. Finally, we will analyze the time-series data from a single wastewater treatment facility to determine how these organisms wax and wane in population density throughout the calendar year.

#### **Activity Milestones:**

Description	Approximate Completion Date
Perform comparative genomics between wastewater genomes and previously described bacterial genomes	August 31, 2026
Analyze metabolic pathways within bacterial genomes to determine optimal conditions for wastewater treatment	June 30, 2027

#### Activity 3: Bioprospecting for genes associated with novel pharmaceuticals

Activity Budget: \$191,639

#### **Activity Description:**

Microorganisms live in complex and competitive communities and they naturally produce molecules to inhibit the

growth of their neighbors. These are some of the antibiotics and antifungals we rely on in both medicine and agriculture, however, these therapeutic molecules are losing their effectiveness as microorganisms are becoming resistant to their effects. Antibiotic and antifungal discovery has long depended on bioprospecting, with much success derived from soil microorganisms. Wastewater treatment bioreactors undoubtedly harbor numerous bioactive compounds that await discovery. Our previous analysis of two representative wastewater samples has yielded multiple potential molecules. In this activity, therefore, we will use the genome sequence data to identify novel pharmaceuticals. As a result of the work done in Activity 1, we will have DNA sequence data equivalent thousands of bacterial genomes, many of which will be novel. We will use the antiSMASH software package, which has the capability of identifying 71 different types of DNA sequence signatures, each associated with a different type of potential pharmaceutical molecule. We will catalog the genes according to the novelty of the predicted molecules they produce to prioritize those that are most likely to yield potentially novel pharmaceuticals.

#### **Activity Milestones:**

Description	Approximate Completion Date
	Completion Date
Identify potential biosynthetic gene clusters	July 31, 2026

## Activity 4: Growth and selection for bacteria with antibiotic and antifungal activity

Activity Budget: \$191,639

#### **Activity Description:**

While computer analysis of genetic information can make predictions about the potential for the synthesis of novel pharmaceuticals, a critical next step is to isolate the bioactive compounds from the actual organisms. In this activity, we will grow organisms from the wastewater bioreactor samples from which sequence data was obtained. Culturing conditions that select for the organisms of interest will be chosen, varying nutrient levels and environmental conditions such as pH, temperature, and oxygen availability. Once bacteria are isolated, genetic markers from organisms that grow will be compared to those observed through antiSMASH analysis of genomic sequencing data (Activity 3) to focus our study on those organisms predicted to produce novel compounds. In the laboratory, these bacterial candidates will be tested for the production of antibiotic or antifungal activity using competition assays. In the simplest assay, newly isolated bacteria will be grown alongside tester strains that mimic some of the most recalcitrant bacterial superbugs. Bacteria that are found to inhibit their competition will be further studied by chemically isolating and identifying the inhibitor molecule to determine if it is indeed a novel potential pharmaceutical.

#### **Activity Milestones:**

Description	Approximate Completion Date
Grow bacteria isolated from bioreactors using culture conditions informed by genomic data	December 31, 2026
Screen isolated bacteria for antibiotic and antifungal activity	June 30, 2027

## **Project Partners and Collaborators**

Name	Organization	Role	Receiving Funds
Justin Donato	University of	Co-Investigator/Project Manager	Yes
	St. Thomas		
Joanna Klein	University of	Co-Investigator/Project Manager	Yes
	St. Thomas		

### Long-Term Implementation and Funding

Describe how the results will be implemented and how any ongoing effort will be funded. If not already addressed as part of the project, how will findings, results, and products developed be implemented after project completion? If additional work is needed, how will this work be funded?

This project should result in several high visibility research publications that will attract the attention of other scientists and engineers. This will, in turn, make receiving funding for additional research much easier from federal sources such as the Environmental Protection Agency, which is keenly interested in improving nutrient removal from municipal wastewater. The bioprospecting portion of this research would lead to array of new research to better understand the activity of and production of any discovered biomolecules.

### Other ENRTF Appropriations Awarded in the Last Six Years

Name	Appropriation	Amount
5 1 1: 0 1: 4 10:1 44: 1: 1		Awarded
Evaluating Coronavirus And Other Microbiological	-	-
Contamination Of Drinking Water Sources From		
Wastewater		
Evaluate Emerging Pathogens in Lakes, Rivers, and Tap	M.L. 2018, Chp. 214, Art. 4, Sec. 2, Subd. 04f	\$325,000
Water to Keep Drinking Water Safe		
Mapping Antibiotic Resistance in Minnesota to Help	M.L. 2018, Chp. 214, Art. 4, Sec. 2, Subd. 04h	\$750,000
Protect Environmental, Animal, and Human Health		
Improving Nitrogen Removal in Greater Minnesota	M.L. 2019, First Special Session, Chp. 4, Art. 2, Sec. 2,	\$325,000
Wastewater Treatment Ponds	Subd. 04e	
Monitoring Emerging Viruses in Minnesota's Urban	M.L. 2021, First Special Session, Chp. 6, Art. 6, Sec. 2,	\$416,000
Water Cycles	Subd. 04c	
Evaluating Coronavirus And Other Microbiological	M.L. 2021, First Special Session, Chp. 6, Art. 6, Sec. 2,	\$594,000
Contamination Of Drinking Water Sources From	Subd. 04g	
Wastewater		
Antibiotic Resistance And Wastewater Treatment:	M.L. 2021, First Special Session, Chp. 6, Art. 6, Sec. 2,	\$432,000
Problems And Solutions	Subd. 04j	
High Temperature Anaerobic Digestion of Sewage	M.L. 2022, , Chp. 94, Art. , Sec. 2, Subd. 04b	\$208,000
Sludge	,	

## Project Manager and Organization Qualifications

Project Manager Name: Timothy LaPara

Job Title: Professor

#### Provide description of the project manager's qualifications to manage the proposed project.

Professor LaPara has worked at the University of Minnesota in the Department of Civil, Environmental, and Geo-Engineering for more than 20 years. His research primarily focuses on the microbiology and microbial ecology of municipal wastewater and its treatment as well as public water supplies and its treatment. He has directed or codirected more than 50 research projects funded by the State of Minnesota, the National Science Foundation, the United States Department of Agriculture, local water/wastewater utilities, and other funding sources. He has published more than 90 papers in the peer-reviewed literature.

Organization: U of MN - College of Science and Engineering

#### **Organization Description:**

The University of Minnesota is one of the largest, most comprehensive, and most prestigious public universities in the United States (http://www1.umn.edu/twincities/about/index.html). The Department of Civil, Environmental, and Geo-Engineering provides the project manager sufficient laboratory, office space, and ancillary facilities that are needed for the proposed study.

## **Budget Summary**

Category / Name	Subcategory or Type	Description	Purpose	Gen. Ineli gible	% Bene fits	# FTE	Class ified Staff?	\$ Amount
Personnel								
Timothy LaPara		Project Manager			26.9%	0.36		\$89,295
Post- doctoral researcher		Researcher			20.4%	3		\$267,922
							Sub Total	\$357,217
Contracts and Services								
University of St. Thomas	Sub award	Collaborate throughout the entire project and lead the bioprospecting work (Activity 3) and bacterial isolation (Activity 4). Personnel (\$219,135; comanagers and undergrads; 7.4% fringe), equipment/freezer (\$15,000), expendable supplies (\$5,000; Petri dishes, growth media, etc), and DNA sequencing services from Phase Genomics (\$65,000).		X		3.75		\$314,137
							Sub Total	\$314,137
Equipment, Tools, and Supplies								
	Tools and Supplies	General laboratory supplies	We will need to collect and process 30 wastewater samples. We will also need to perform miscellaneous laboratory analyses to support the primary data that will be generated.					\$3,000
							Sub Total	\$3,000
Capital Expenditures								
							Sub Total	-
Acquisitions and Stewardship								

				Sub	-
Travel In Minnesota				Total	
	Miles/ Meals/ Lodging	30 trips to wastewater treatment facilities	Collect samples		\$1,500
	Conference Registration Miles/ Meals/ Lodging	1 conference	Attend a conference within the State of Minnesota to disseminate research results		\$1,500
				Sub Total	\$3,000
Travel Outside Minnesota					
				Sub Total	-
Printing and Publication					
	Publication	Open access fees	This will allow us to retain the copyright of all publications; this will help us disseminate our results widely		\$7,646
				Sub Total	\$7,646
Other Expenses					
		Hiring an internal (UMGC) and/or an external laboratory for DNA sequence analysis	We envision needing to perform a limited amount of DNA sequencing to validate and improve our results. Some of the sequencing technology we will likely need (Sanger sequencing) is no longer performed by the University of Minnesota Genomics Center		\$5,000
				Sub Total	\$5,000
				Grand Total	\$690,000

## Classified Staff or Generally Ineligible Expenses

Category/Name	Subcategory or	Description	Justification Ineligible Expense or Classified Staff Request
	Туре		
Contracts and Services - University of St. Thomas	Sub award	Collaborate throughout the entire project and lead the bioprospecting work (Activity 3) and bacterial isolation (Activity 4). Personnel (\$219,135; co-managers and undergrads; 7.4% fringe), equipment/freezer (\$15,000), expendable supplies (\$5,000; Petri dishes, growth media, etc), and DNA sequencing services from Phase Genomics (\$65,000).	A laboratory-grade deep freezer (-80 degrees celsius) is required for Activity 4 of this project (cost = \$15000). This will allow us to cryopreserve bacterial isolates indefinitely so that they can be isolated, preserved, and then studied in a reasonable time frame (i.e., otherwise, the bacteria would die). The University of St. Thomas does not currently have any freezer that fulfills this need. This freezer would be dedicated solely to this project. We acknowledge that we understand ENTRF requirements for repayment if the use of this capital equipment changes.

## Non ENRTF Funds

Category	Specific Source	Use	Status	Amount
State				
			State Sub	-
			Total	
Non-State				
In-Kind	University of Minnesota	Indirect costs that are not allowed to be charged to LCCMR/ENTRF	Secured	\$220,375
In-Kind	University of St. Thomas	Indirect costs that are not allowed to be charged to LCCMR/ENTRF	Secured	\$123,843
			Non State	\$344,218
			Sub Total	
			Funds	\$344,218
			Total	

#### **Attachments**

#### **Required Attachments**

Visual Component

File: 9218902b-374.pdf

#### Alternate Text for Visual Component

A flow diagram that shows wastewater bioreactors leading to bacterial genomes that lead to better surface water quality. Another pathway shows wastewater bioreactors leading to bacterial genomes to the discovery of novel genes that lead to novel pharmaceutical compounds....

#### **Optional Attachments**

Support Letter, Photos, Media, Other

Title	File
Letter of Commitment	<u>aeb2f574-f55.pdf</u>

#### Administrative Use

Does your project include restoration or acquisition of land rights?

Nο

Does your project have potential for royalties, copyrights, patents, or sale of products and assets?

Yes

Do you understand and acknowledge IP and revenue-return and sharing requirements in 116P.10?

Yes

Do you wish to request reinvestment of any revenues into your project instead of returning revenue to the ENRTF?

No

Does your project include original, hypothesis-driven research?

Yes

Does the organization have a fiscal agent for this project?

Yes, Sponsored Projects Administration

Does your project include the design, construction, or renovation of a building, trail, campground, or other capital asset costing \$10,000 or more?

No

Do you propose using an appropriation from the Environment and Natural Resources Trust Fund to conduct a project that provides children's services, as defined in Minnesota Statutes section 299C.61 Subd.7?

No