

Predicting Bacteria Concentrations On The Nation's Beaches

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*Year of Water:
Thirty Years of Progress
Through Partnering*

Environmental Issue

With the possible exception of those who value solitude above all else, no one wants beaches closed as a result of high bacterial water concentrations. Although microbiology is constantly making new breakthroughs in microbial detection and identification, beach closure decisions still frequently rely on time-intensive tests for indicator bacteria which do not differentiate between sewage plumes from distant sources reaching a beach and the visit of flocks of seagulls to the beach the previous day. Current procedure allows only for a 10-fold reduction in high bacterial counts, costing municipalities untold dollars in lost revenues and remediation actions.

Approach

ERD scientists are developing models that will in time be able to accurately track the movement of water and sources onto the nation's beaches. Models, like the ERD Visual Plumes model, will be modified to track both point sources (like sewage treatment plants) and non-point sources (like birds on beaches). Another aspect of this research seeks to develop and improve bacterial submodels that will be specific to actual pathogens, rather than indicators. To advance, pathogens must be positively identified and modeled. Microbial Source Tracking (MST) is a technique for quantifying pathogens. ERD researchers are developing a fast and reproducible alternative DNA-based MST methodology to estimate complex source mixtures. This work envisions a model, working title "Visual Beach", that will use real-time data and advanced biological inputs, for example, Microbial Source Tracking (MST), GIS data (bathymetry), forecast data, particularly weather data, and in coastal areas, astronomical tide factors.

Impact

Microbial contamination leads to beach closure and poses a threat to human health. The purpose of this work is to help protect aquatic ecological systems and recreational resources. If successful, the model will help devise effective source control strategies and help differentiate between dangerous episodes, requiring closure, and benign ones not requiring drastic measures. It would enable health officials to discriminate between dangerous and benign exposure levels and help treatment plant operators develop strategies to prevent most beach closures due to important point sources.

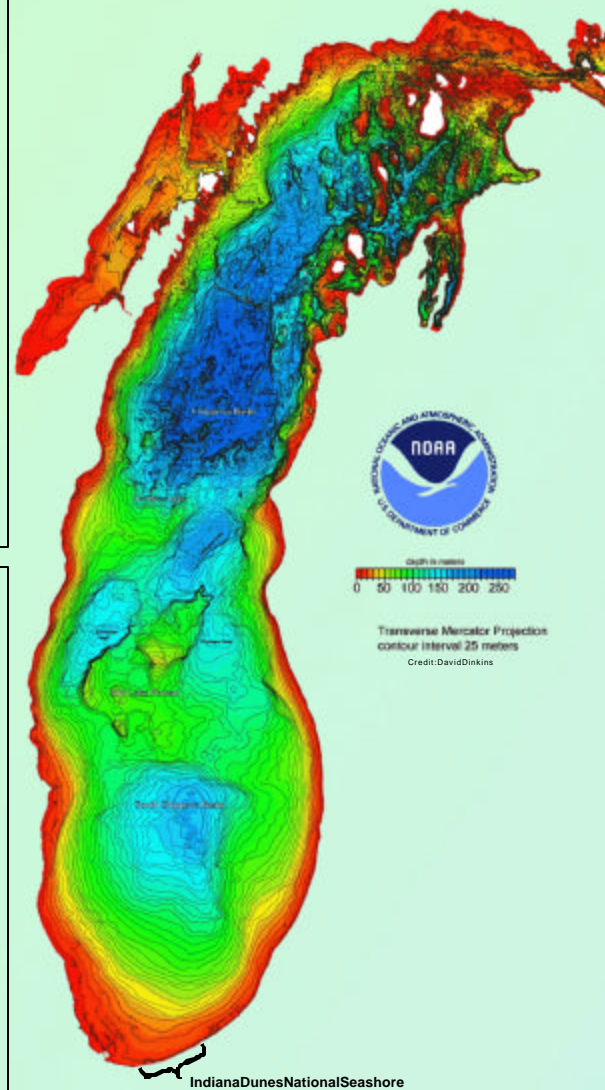
National Scope

An example of a national problem, Huntington Beach, California was closed in 1999 due to elevated bacterial counts. Visual Plumes has been successfully tested on this problem.

July-August 1999 Beach Closure Map locates the closed beach (green stripe) and the major Orange County 4-mile long sewage outfall discharging at 60m depth. Insets show beach scene and treatment plant.



Future Study Site Lake Michigan Bathymetry



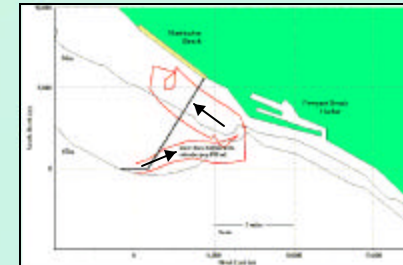
Indiana Dunes National Seashore The model described here will be tested on West Beach (Indiana Dunes National Seashore, Lake Michigan) data this summer.



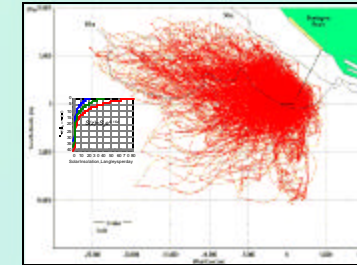
Values, Resources, and Problems

Modeling

A Visual Plume simulation. A release of sewage forming a plume (red outline) tracked for 36 hours (plume elements moved in 2-hr increments) and forming a plume that encroaches Huntington Beach in winter 2001, a rare event.

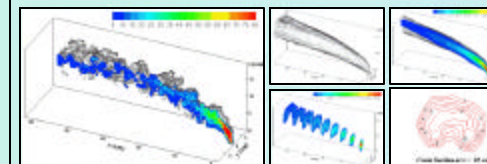


Not so paghetti, a month-long Visual Plume simulation. June-July 2000 simulation shows a reached by similarly contaminated plumes during that period. Note, the source did not impact the beach during that time. Inset diagram shows solar intensity (a killer of waterborne bacteria) factored into Visual Plumes) decreases rapidly with depth.

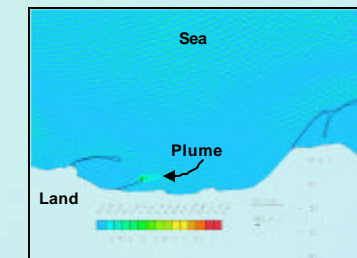


Hydrodynamics and PCR

The beach bacteria modeling spans two main disciplines, hydrodynamics and microbiology. Shown here is a laboratory plume image produced through laser-induced fluorescence imaging. Both surface features and false color cross-sectional features are shown.



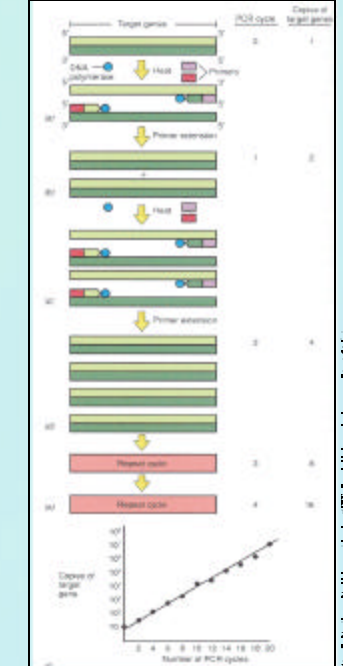
Plans for the Lake Michigan Model The next generation of beach bacteria models will include dense current information based on finite-element hydrodynamical model that will make forecasts possible. This example shows a current field (light green arrows) and an embedded plume representing a combined sewer overflow.



Polymerase Chain Reaction (PCR)

PCR is a method by which very low concentrations of DNA are amplified to improve detection of target organisms. However, conventional PCR methods cannot be used to determine bacterial numbers. Recently, researchers have developed an alternative PCR method commonly known as real-time PCR, where the number of amplicons (amplified DNA products) from each reaction cycle are measured. Product accumulation is quantified during the exponential phase of the amplification process. The procedure can be used with 16S-rDNA primers specific to a group or species of fecal indicator bacteria. This application will allow accurate monitoring and quantification of specific fecal indicator bacteria in recreational waters and may reduce detection time to 3 hours. The schematic diagram shows principles of the application of polymerase chain reaction technology.

Schematic of the polymerase chain reaction (PCR)



Conclusion

The Visual Beach modeling approach attempts to provide a holistic conceptual and numerical model of bacterial transport and fate. It attempts to realistically model the most important transport and fate mechanisms. It answers questions like: Where did the water come from? What were the concentrations of bacteria initially? To what stressors were the bacteria exposed during their transit from source to receptor regions? And so on. The basic physics that are increasingly better models will help better understand and model bacterial transport and fate and ultimately help reduce beach closures.



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