

Fluorescence In Situ Hybridization

EMD Team Fact Sheet—November 2011

This fact sheet, developed by the ITRC Environmental Molecular Diagnostics (EMD) Team, is one of 10 designed to provide introductory information about and promote awareness of EMDs. Please review the Introduction to EMDs Fact Sheet along with this one. A glossary is included at the end of this fact sheet.

Why is fluorescence in situ hybridization relevant?

Fluorescence in situ hybridization (FISH) is a molecular biology technique that can be used to detect microorganisms known to biodegrade contaminants. When combined with traditional measuring of changes in contaminant concentration over time, FISH provides project managers valuable information for site management, including site conceptual model development, remedy selection, and optimization and determination of contaminant attenuation rates.

What does FISH do?

To implement FISH, environmental samples (soil, water, sediment) are taken to a laboratory and in a series of steps a fluorescent dye is attached to a particular gene of interest in microorganisms or families of microorganisms. These targeted microorganisms can then be observed and their abundance and spatial distribution determined under a microscope by detecting the fluorescent light emitted. For the purposes of environmental investigation, the targeted microorganisms are typically ones capable of degrading specific contaminants.

Originally developed in the 1990s, FISH is routinely used in medical fields such as genetic counselling, disease identification, and microorganism species identification. More recently, it has been applied to the study of environmental processes, mostly in the wastewater treatment field, but also for contaminant biodegradation at sites with coal tar, herbicides, or chlorinated solvents, as well as at wetlands used to treat acid mine drainage. The FISH technique also has potential applicability in other environmental monitoring efforts, such as tracking of fecal microorganisms, so-called "microbial source tracking," and identifying and tracking microorganisms potentially related to chemical compounds originating from industrial or energy-related activities.

How are the data used?

FISH can provide valuable insights for environmental remediation alone and in combination with other EMDs, like polymerase chain reaction (PCR). For example, this method can reveal whether key organism(s) needed for biodegradation are present in the sample material and allow estimation of their abundance, similar to other EMDs. However, this method can also allow investigators to explore their structure, form, and spatial distribution and association with other microorganisms. FISH signals can provide some information about activity of the target organisms although no precise rate information can be obtained.

Example Environmental Remediation Questions FISH Can Help Answer

• Site Characterization

- Are the right microorganisms and/or genes present that are capable of degrading the contaminant? If so, how many and where?
- What other microorganisms are present in the environment, and what impact do they have on the microorganisms or processes of interest?

Monitoring

Does the microbial community change in response to an amendment?

How does it work?

Short sequences of single-stranded nucleic acids (such as DNA), called "gene probes," are designed to match a portion of a gene or metabolic product of the organism or population of interest. A fluorescent dye is attached to the probe so that when the probe binds to target sequences within a cell, it emits fluorescent light that can be observed through a microscope (i.e., using an epifluorescent microscope) or sorted with flow cytometry. Cells emitting a fluorescent light are called "hybridized cells." In flow cytometry, labeled cells are diluted or concentrated (depending on the initial cell concentration in the sample) so that individual cells pass through a laser beam that detects and counts fluorescently labeled cells. Flow cytometry can be significantly more efficient than counting cells using a microscope. Cells in environmental samples are handled in such a way that the cell structure remains intact while still allowing the comparatively large gene probe to enter the cell and bind to the target gene within the microorganism of interest (Figure 1). Under ideal conditions, only cells that contain the target gene are recognized by the probe and become fluorescently labeled. Various cell staining procedures are sometimes combined with FISH probes to allow quantification of various parameters such as the total number of microorganisms or the presence of specific compounds. Table 1 presents selected FISH probes and cellular stains.

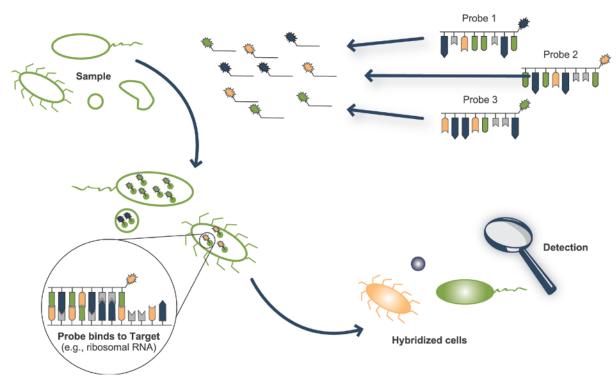


Figure 1. Fluorescence in situ hybridization method.

How are the data reported?

Depending on the method, FISH results can be presented in two different ways:

- If FISH is evaluated using a microscope and manual counting of labeled cells, the results are presented as cells per unit (liter of liquid or gram of solid) analyzed.
- If FISH is evaluated with advanced microscopy techniques and digital image processing, the results
 are usually presented on a relative volume or area basis, which can be converted by the laboratory to
 cells per unit of liquid or solid.

FISH probes or **Contaminants** Target microorganism(s) Reference cellular stains DAPI NA DNA of all microorganisms (live This is a very common laboratory cellular stain. Not unique to and dead) environmental contaminants. DNA of all microorganisms (live Acridine orange NA This is a very common laboratory and dead) cellular stain. Not unique to environmental contaminants. Dhe1259t Chlorinated Some Dehalococcoides spp. 16S Yang and Zeyer 2003 solvents rRNA Dhe1259c Some Dehalococcoides spp. 16S Chlorinated solvents rRNA KT1phe Trichloroethene Ralstonia eutropha KT1 16S rRNA Tani et al. 2002 Ac627BR Naphthalene Naphthalene dioxygenase (nahAc) Bakermans and Madsen 2002 mRNA s-Triazine RhLu Rhodococcus wratislaviensis 16S Grenni et al. 2009 herbicides rRNA

Table 1. Selected FISH probes or cellular stains

Advantages

- FISH does not require cultivation of the organisms or any technology-based gene amplification (see PCR Fact Sheet), which can lead to false negatives and positives.
- In contrast to some other EMDs, FISH allows visualization of whole cells that are important to
 environmental remediation activities. FISH can thus provide complementary information to other
 EMDs, such as morphology of the cells or association of groups of microorganisms with relationship
 to one another.
- FISH can target several different genes simultaneously, for example, genes associated with specific degrading species of interest (e.g., *Dehalococcoides*) and broader microbial groups, such as methane-producing organisms.
- Depending on the species, and in combination with other appropriately validated activity-targeted approaches, FISH can provide general information about the activity of the organisms or populations of interest.
- FISH enables single-cell microbial studies and allows for subsequent studies, such as gene sequencing (see the Microbial Fingerprinting Methods Fact Sheet).

Limitations

- The detection limit of FISH is high (~10⁶ cells/mL). However, in some cases high detection limits can be corrected by sample concentration or cell extraction methods which lower the detection limits to a few hundred cells per concentrated sample.
- Validated probes and FISH procedures are not available for a wide range of organisms within the bioremediation field. Additionally, standard protocols for sample collection and storage prior to FISH analysis have not yet been developed.
- FISH can also be used to target not only ribosomal genes (which indicate the type of organism) but also functional genes (via mRNA) relevant in bioremediation. These other genes indicate what the microorganisms can do with regards to contaminant biodegradation, for example, naphthalene dioxygenase or reductive dehalogenase. However, laboratory protocols are often time-consuming and complicated and not yet validated for field applications.
- The FISH method is not widely commercially available. Currently, mainly specialized research laboratories are performing these analyses to explore and optimize the potential of FISH for validated and cost-effective applied studies.
- The FISH method is currently expensive because of the expertise and labor needed for development of validated FISH protocols and direct microscopic counting. Once validated protocols have been

developed, FISH can be automated to some extent by using flow cytometry to count target cells more efficiently, reducing the analytical costs. However, when using flow cytometry for cell counting, all information regarding spatial relationships (among and between the cells) is lost.

Sampling Protocols

Sample matrices that can be analyzed by FISH include most kinds of environmental samples, such as wastewater, groundwater, filters, soil, and sediments. However, depending on the sample type, different types of sample preparations and FISH protocols may have to be employed. Basic sampling for microbiological samples can be easily incorporated into routine environmental monitoring programs. The following items are typical requirements for microbiological sampling: (a) use of aseptic sample collection techniques and sterile containers, (b) shipment of the samples to the laboratory within 24 hours of sample collection, and (c) maintenance of the samples at an appropriate temperature (e.g., 4°C during handling and transport to the laboratory). Sample collection techniques and containers may vary depending on the matrix sampled and the laboratory analyzing the samples. Users should work with the analytical laboratory to ensure sampling protocols for collecting, handling, and transporting the samples are in place and understood.

Quality Assurance/Quality Control

To date, most EMDs do not have standardized protocols accepted by the U.S. Environmental Protection Agency or other government agencies. However, most laboratories work under standard operating procedures (SOPs) and good laboratory practices, which can be provided to the user (e.g., consultant, state regulator) on request.

Currently, users can best ensure data quality by detailing the laboratory requirements in a site-specific quality assurance project plan (QAPP). This plan should include identification of the EMDs being used; the field sampling procedures, including preservation requirements; the SOPs of the laboratory performing the analysis; and any internal quality assurance/quality control information available (such as results for positive and negative controls). Sample collection, preservation, and laboratory protocols for FISH have been standardized for only certain types of organisms and ecosystems.

Additional Information

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Glossary

- **biodegradation**—A process by which microorganisms transform or alter (through metabolic or enzymatic action) the structure of chemicals introduced into the environment (EPA 2011).
- **Dehalococcoides**—A specific group (genus) of bacteria. *Dehalococcoides* species are the only microorganisms that have been isolated to date which are capable of complete reductive dechlorination of PCE and TCE to ethene. Some *Dehalococcoides* species are also capable of reductive dechlorination of chlorinated benzenes, chlorinated phenols, and polychlorinated biphenyls.
- **DNA (deoxyribonucleic acid)**—A nucleic acid that carries the genetic information of an organism. DNA is capable of self-replication and is used as a template for the synthesis of RNA. DNA consists of two long chains of nucleotides twisted into a double helix (EPA 2004).
- epifluorescent microscope—A type of microscope that uses a high-energy light source (e.g., ultraviolet light) and specialized filters to visualize fluorescently stained specimens. Epifluorescent microscopy procedures can be used to determine both the total number of cells and total number of viable or active cells in a sample.
- **FISH probes**—Short sequences of single-stranded DNA carrying a fluorescent label. When the probe binds to the target DNA/RNA sequence of the microorganism(s) of interest in an environmental sample, the target cell fluoresces and can be visualized and counted using a specialized microscope or a flow cytometer.
- **flow cytometry**—A method whereby cells or particles move in a liquid stream past a laser or light beam and a sensor detects the relative light scattering and fluorescence of the particles.
- **functional gene**—A segment of DNA that encodes an enzyme or other protein that performs a known biochemical reaction. For example, the functional gene *tceA* encodes the reductive dehalogenase enzyme that initiates reductive dechlorination of TCE. Other genes can code for RNA entities which can regulate the activity of other DNA target sequences.
- **gene**—A segment of DNA containing the code for a protein, transfer RNA, or ribosomal RNA molecule (based on Madigan et al. 2010).
- **labeled cell**—A microorganism in which a gene probe has bound to a matching sequencing within the microorganism and released a fluorescent dye, resulting in a cell that is emitting fluorescent light.
- **metabolic product**—Products generated by a microorganism whose structure and function are defined by DNA sequences also called genes. Example metabolic products include RNA and proteins or enzymes.
- **probes**—(1) short DNA strands (see *microarray probes*, Microarray Fact Sheet; *FISH probes*; *qPCR probes*, qPCR Fact Sheet); (2) surrogate compounds (see *enzyme activity probes*, EAP Fact Sheet).
- **RNA** (ribonucleic acid)—Single-stranded nucleic acid that is transcribed from DNA and thus contains the complementary genetic information.

whole cell—The entirety of a microbial cell, without extraction of DNA, RNA, etc. A whole-cell preparation does not modify the cell but evaluates it as unit.

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