

Technology Transition Workshop | Robert Driscoll, M.F.S.

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Introduction to Fluorescent In Situ Hybridization (FISH)

Fluorescent In Situ Hybridization (FISH)

- FISH is a cytogenetic technique used to detect the presence or absence of specific chromosomes and/or sequences
- Interphase FISH techniques incorporate probes which pass through cellular membranes and into the nucleus eliminating the need to lyse cells during processing
- Fluorescence compatible microscopes are typically employed to visualize the multicolor probes used in these hybridizations

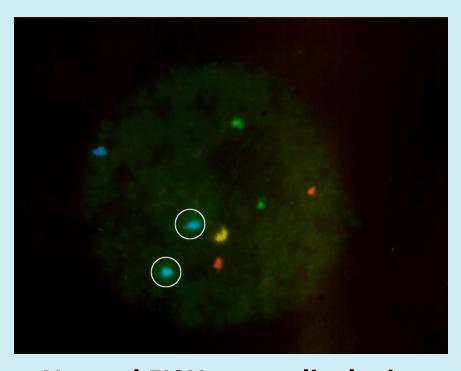
Traditional FISH Applications

- Used to detect a variety of diseases caused by chromosomal abnormalities:
 - Prader-Willi Syndrome
 - Angelman Syndrome
 - Cri-du-chat
 - Down Syndrome
 - Chronic myelogenous leukemia
 - Acute lymphoblastic leukemia
- Used to compare genomes of two biological species to deduce evolutionary relationships
- Gene mapping





Prenatal Screening for Down Syndrome

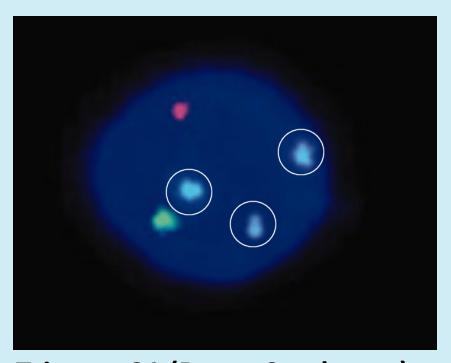


Normal FISH assay displaying 2 copies of chromosome 21

Image courtesy of: http://www.parkavefertility.com/pre-implantation_genetic_diagnosis.html



Prenatal Screening for Down Syndrome



Trisomy 21 (Down Syndrome) - 3 copies of chromosome 21

Image courtesy of: http://www.parkavefertility.com/pre-implantation_genetic_diagnosis.html

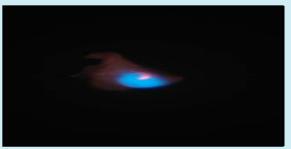
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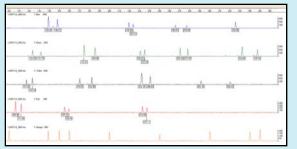


Methodology of LM-FISH Processing

- Isolate cells of evidence mixtures
- Identify contributor by gender
- Capture, process, and generate clean profile







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Images courtesy of Rob Driscoll

FISH Overview

- Preparation of slide
- Pre-treatment
 - Prepare sample for hybridization by reducing background fluorescence
- Hybridization
 - Apply probes and hybridization buffer to the sample
 - Denature and hybridize sample/probes
- Post-hybridization
 - Remove excess probe and salts
 - Apply DAPI counterstain to stain nucleus blue and increase fluorescence of probes

Preparation of Slide

- Elute cells from substrate.
- 2. Re-suspend cells in Carnoy's fixative.
 - 3:1 methanol/glacial acetic acid
 - Fixes cells to slides
 - Helps cells spread well
- 3. Steam slide for 5 seconds.
 - Allows for further membrane spreading
- 4. Dry on slide warmer.
- Bake in biological oven.
- 6. Store in desiccator overnight.



Pre-Treatment

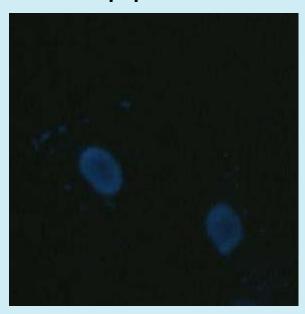
- SSC buffer wash
 - Isotonic buffer solution
 - Contains sodium chloride and sodium citrate
- Pepsin buffer wash
 - Protease digests the cell wall and cytoplasm, leaving the nucleus behind
 - Increases fluorescence of probes by reducing background fluorescence caused by cytoplasm



Pre-Treatment

Effect of pepsin buffer on cells

10 minute pepsin buffer wash



No pepsin buffer wash





Pre-Treatment Steps

- PBS/MgCl2 wash
 - Removes pepsin buffer from slide
- Formalin (or formaldehyde) buffer wash
 - Reduces background fluorescence
- PBS wash
- Serial ethanol washes
 - Dehydrates cells
- Slides are stored in dessicator until hybridization



Pre-Treatment

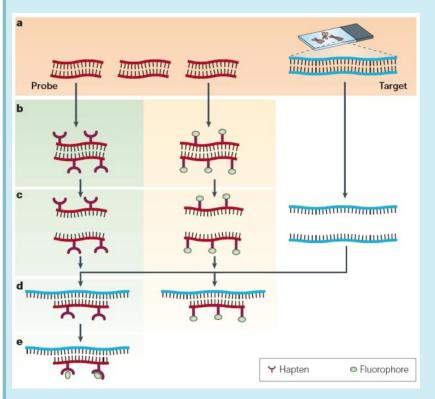


Serial ethanol washes

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Image courtesy of Rob Driscoll



- 1. Identify the DNA probe and a target sequence.
- 2. Probe is labeled.
- 3. Labeled probe and target DNA are denatured.
- 4. Denatured probe and DNA are combined and complimentary DNA sequences anneal.
- Indirectly labeled probe undergoes additional processing for visualization.

Image from reference #9 (See Handout, Relevant Scientific Literature)

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Hybridization Buffer

- **Contains:**
 - Formamide denaturing agent
 - SSC isotonic buffer, controls salt concentration
 - Dextran sulfate high molecular weight polymer
- Buffer components and hybridization temperature affect the stringency
 - Stringency adjustment of conditions to control hybridization of the probe binding to the target sequence
 - The higher the stringency, the lower the probability of hybridization
 - High stringency high temp, lower salt concentration
 - Low stringency low temp, higher salt concentration



Hybridization Probes

- Chromosome enumeration probes (CEP®) bind to highly conserved sequences of individual chromosomes
 - Highly repetitive human satellite DNA sequences
 - Strong, crisp signals
- CEP Y[®] Satellite III SpectrumOrange™
 - Y chromosome hybridized with TRITC-labeled CEP Y[®]
 (DYZ1) probe (orange)
- CEP X[®] Alpha Satellite SpectrumGreen™
 - X chromosome hybridized with FITC-labeled CEP X[®]
 (DXZ1) probe (green)

FISH - Male Epithelial Cells CEP Y® Satellite III SpectrumOrange™



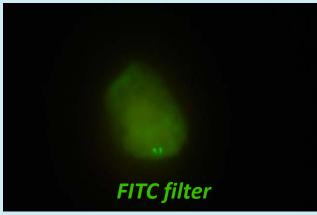
- CEP Y[®] (DYZ1)
 SpectrumOrange™
 probe
 - Chromosome positionYq12
 - Aquired via Abbott Molecular®/Vysis™
 - TRITC fluorophore
- Blue DAPI counterstain

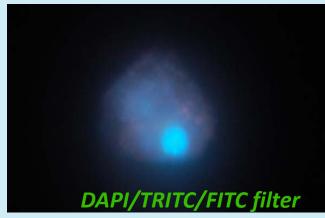
Image courtesy of Rob Driscoll

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FISH – Female Epithelial Cells CEP X® Alpha Satellite SpectrumGreen™

- CEP X[®] (DXZ1)
 SpectrumGreen[™] probe
 - Chromosome position Xp11.1
 - Aquired via Abbott
 Molecular®/Vysis™
 - FITC fluorophore
- Blue DAPI counterstain





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Images courtesy of Rob Driscoll

- Denaturation: Breaks the hydrogen bonds between double-stranded segments of DNA, allowing new bonds to form between the probes and target DNA during hybridization
- Denaturant solution contains formamide, SSC, and water
- **Process:**
 - Denature at 75 to 80°C in a water bath.
 - Complete serial ethanol washes.
 - Dehydrates the cells
 - Combine hybridization buffer, probes, and water.
 - Denature at 75 to 80°C in a water bath

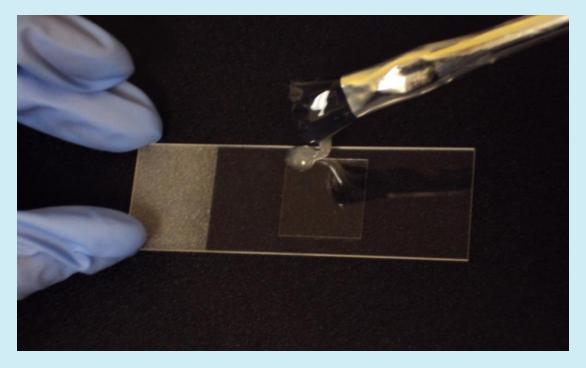


 Hybridization: Probes specifically hybridize to their complementary sequences on the target DNA

Process:

- Apply denatured probes to the denatured sample slide.
- 2. Cover with coverslip.
- 3. Seal with rubber cement.
 - Prevents evaporation of sample
- 4. Hybridize at 37 to 42°C in a humidified chamber.





Sealing coverslip with rubber cement to prevent evaporation of sample

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Images courtesy of Rob Driscoll

- Co-denaturation and hybridization on StatSpin®
 ThermoBrite™ Slide Hybridization/Denaturation
 System
 - 1. Combine hybridization buffer, probes, and water.
 - Apply to slide.
 - 3. Cover with coverslip.
 - 4. Seal with rubber cement.
 - 5. Denature at 73°C.
 - 6. Hybridize at 37 to 42°C.

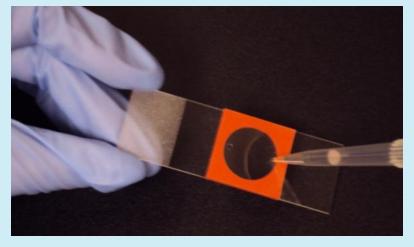


StatSpin® ThermoBrite™ Slide Hybridization/Denaturation System

- Microprocessor controlled bench top hot plate with lid
- Three operating modes: denaturation, hybridization, fixed temperature
- Stores 40 programs
- 12 slide capacity
- Humidity control to prevent evaporation of samples from slides
- Purchased to provide a more controlled environment for heating slides
 - Increased consistency and reproducibility of temperatures



- Use of re-useable perfusion chambers in combination with the ThermoBrite™ eliminates the need for water baths
 - 1. Place chamber on the slide.
 - 2. Fill chamber with buffer.
 - 3. Place slide on the ThermoBrite™ for incubation.



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Image courtesy of Rob Driscoll



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Image courtesy of: http://international.abbottmolecular.com/ThermoBrite 7633.aspx#

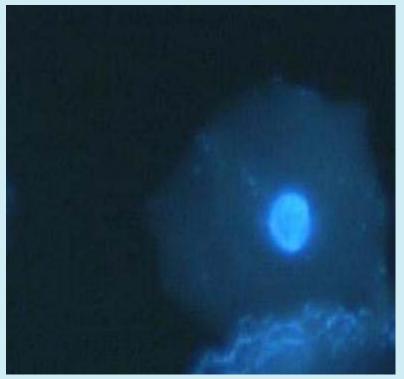
Post-Hybridization

- SSC/Tween® 20 wash buffer
 - High stringency wash
 - Removes excess unbound probe
- SSC wash
 - Low stringency wash
- Water rinse
 - Removes residual buffer salts to reduce nonspecific background fluorescence
- DAPI VECTASHIELD® counterstain
 - Fluorescent nuclear stain (blue)
 - Increases fluorescence of probes



- Utilize Vysis™ CEP X® Alpha Satellite and CEP Y®
 Satellite III probes to visually identify male and
 female cells.
- 2. Apply a DAPI counterstain to stain nuclei.
- Detect fluorescently labeled nuclei using DAPI/FITC/TRITC bandpass filters.
- 4. Capture cells using a laser microdissection system.





DAPI filter for detection of nuclei

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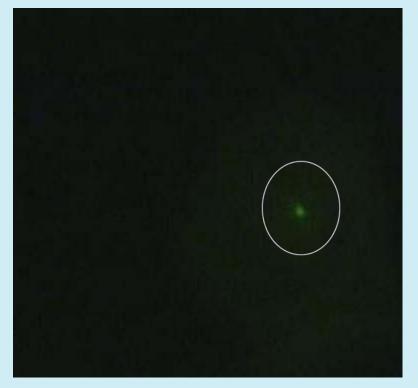




Rhodamine filter for detection of Y chromosome

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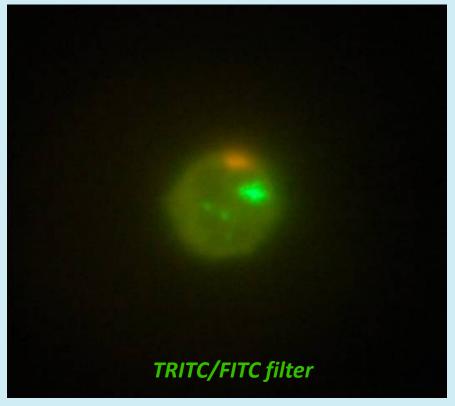


FITC filter for detection of X chromosome(s)

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Male White Blood Cells Stained with X and Y FISH Probes

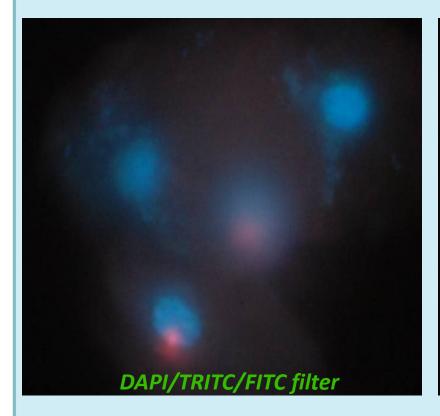


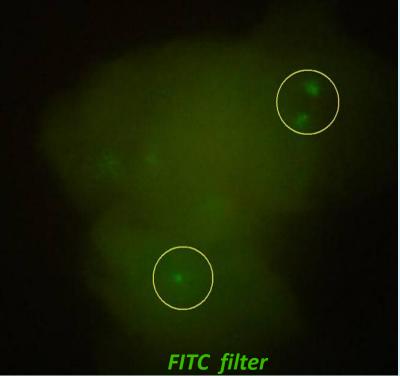
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Image courtesy of Rob Driscoll

Male/Female Epithelial Cell Mixture



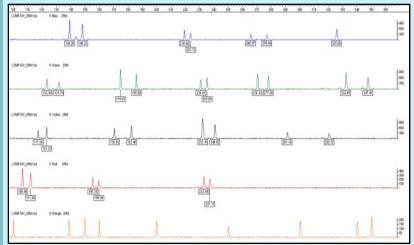


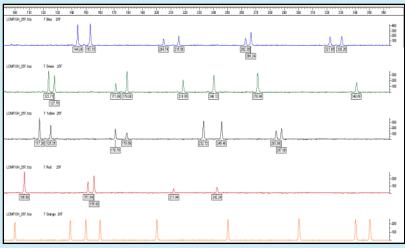
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Images courtesy of Rob Driscoll

Initial STR Profiling of FISH Samples





- Profiles generated from the separation of approximately 30 FISH-treated male and 30 FISH-treated female epithelial cells from a 1:1 mixture
- Extracts were amplified for 30 cycles in a 7 μl Identifiler® reaction
- Demonstrates a clean separation of male and female components

Images courtesy of Rob Driscoll

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Initial STR Profiling of FISH Samples

15 FISH-Stained Male White Blood Cells Separated From a Male/Female White Blood Cell Mixture



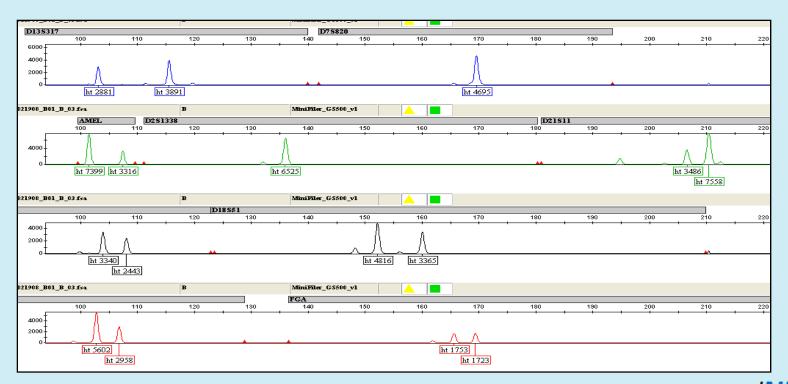
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Image courtesy of Rob Driscoll

Initial STR Profiling of FISH Samples

Profile From 15 FISH-Stained Male Epithelial Cells Separated From a Male Epithelial/Female White Blood Cell Mixture



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Ongoing Research

- The current FISH X/Y probing technique is limited
 - Only allows for separation of cells of different gender
- Research is underway regarding the use of FISH and LM to separate same gender mixtures
- Key objective is to separate cellular mixtures of similar morphology and same gender by using sequence-specific FISH probes based upon the genetic polymorphisms associated with SNPs and the ABO blood groups

Ongoing Research

- SNPs and the ABO blood groups were chosen as the basis for these assays because of the existence of their prevalent polymorphisms throughout multiple populations
- Both systems have multiple polymorphisms which can be targeted through multicolor probe assays
- Primer probes can be designed to target genetic loci associated with SNPs and the ABO blood groups to allow for visual identification of individuals in forensic mixtures

Advantages of FISH

- Allows for differentiation of male and female cells of similar morphological mixtures
- Allows for successful processing of previously unusable items of evidence
- Reduces need for complicated LCN mixture deconvolution



Limitations of FISH

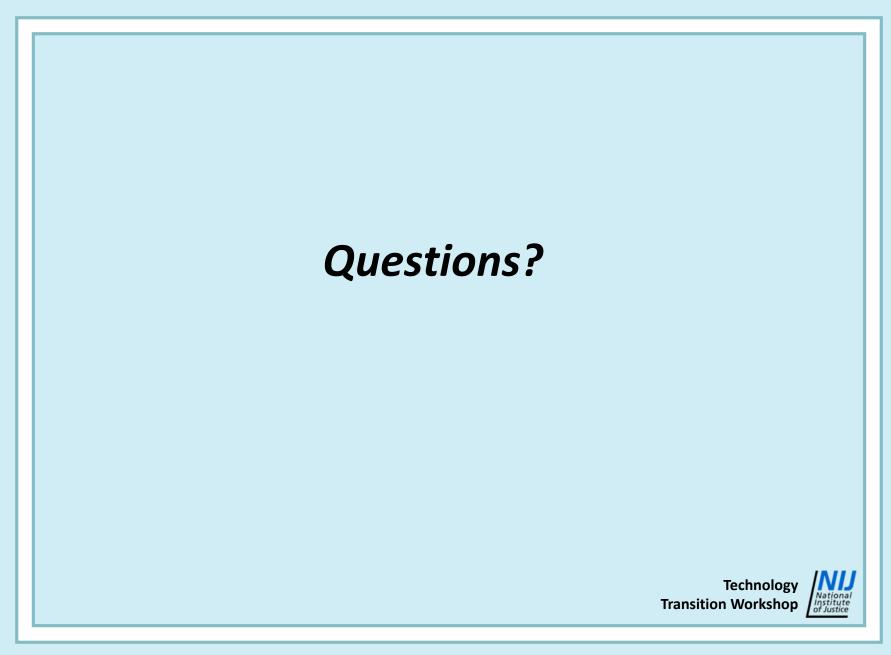
- 8 to 10 hours for FISH processing
- Probes are \$30 per μl
- Difficult to troubleshoot because multiple factors can affect successful hybridization
- Formalin and hybridization buffers contain toxic chemicals



Summary

- FISH processing utilizing X/Y probes is a novel method for differentiating male and female cells of similar morphological mixtures
- New systems are being explored that may allow for differentiation of cell mixtures of same morphology and gender





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