



Hybridization techniques and their practical applications

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Department of Pathology**

**Basic Molecular Pathology Course
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Hybridization techniques and their practical applications

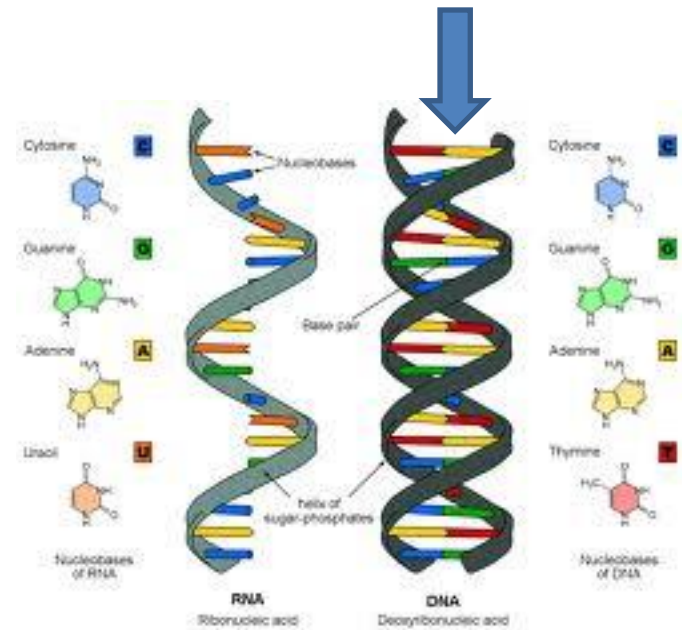
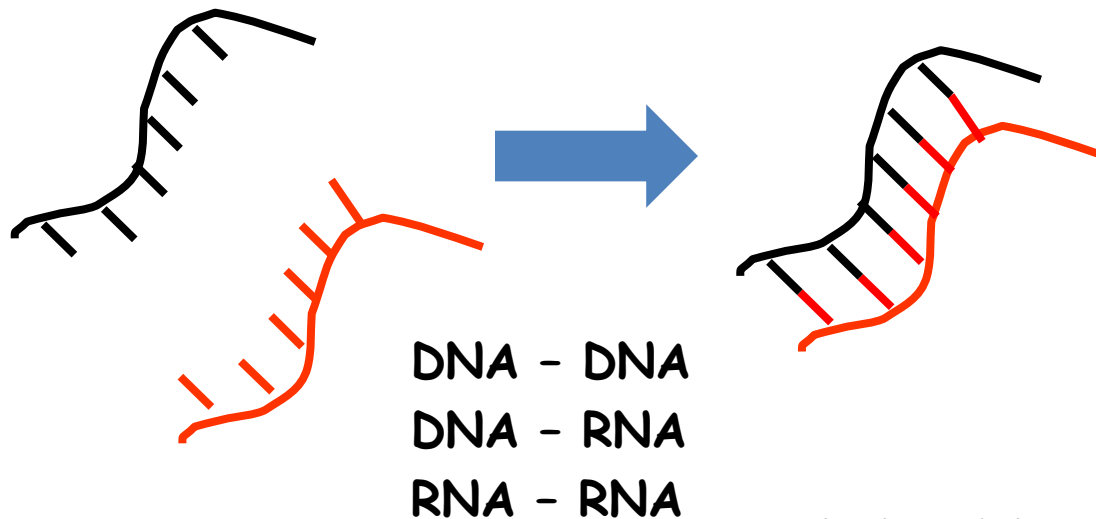


- Definition and terminology
- Explanation of the technique
- Interpretation
- Applications for diagnostic / research Pathology.

TERMINOLOGY

Hybridization

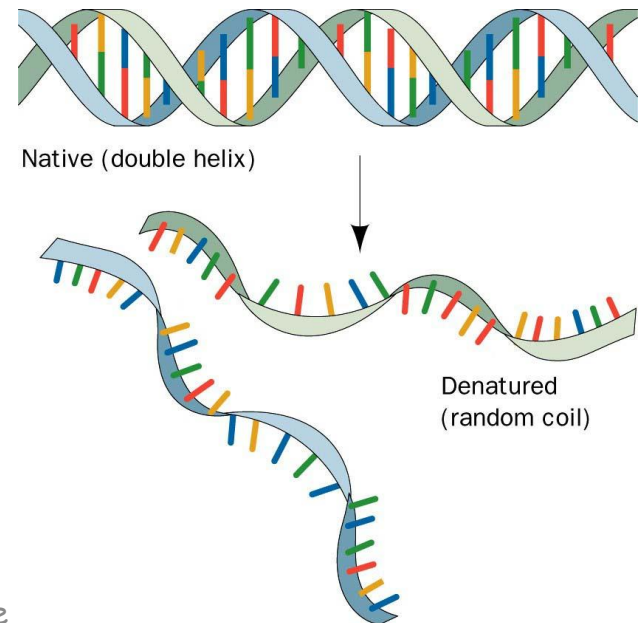
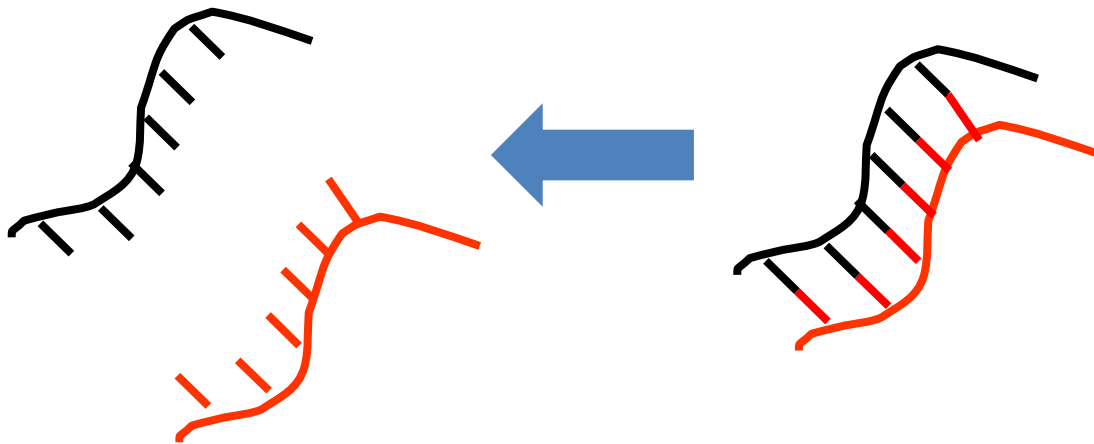
Process of establishing a non-covalent, sequence-specific interaction between two complementary strands of nucleic acids into a single complex



TERMINOLOGY

Denaturation

DNA melting, is the process by which double-stranded nucleic acid separates into single-stranded strands through the breaking of hydrophobic stacking attractions between the bases.

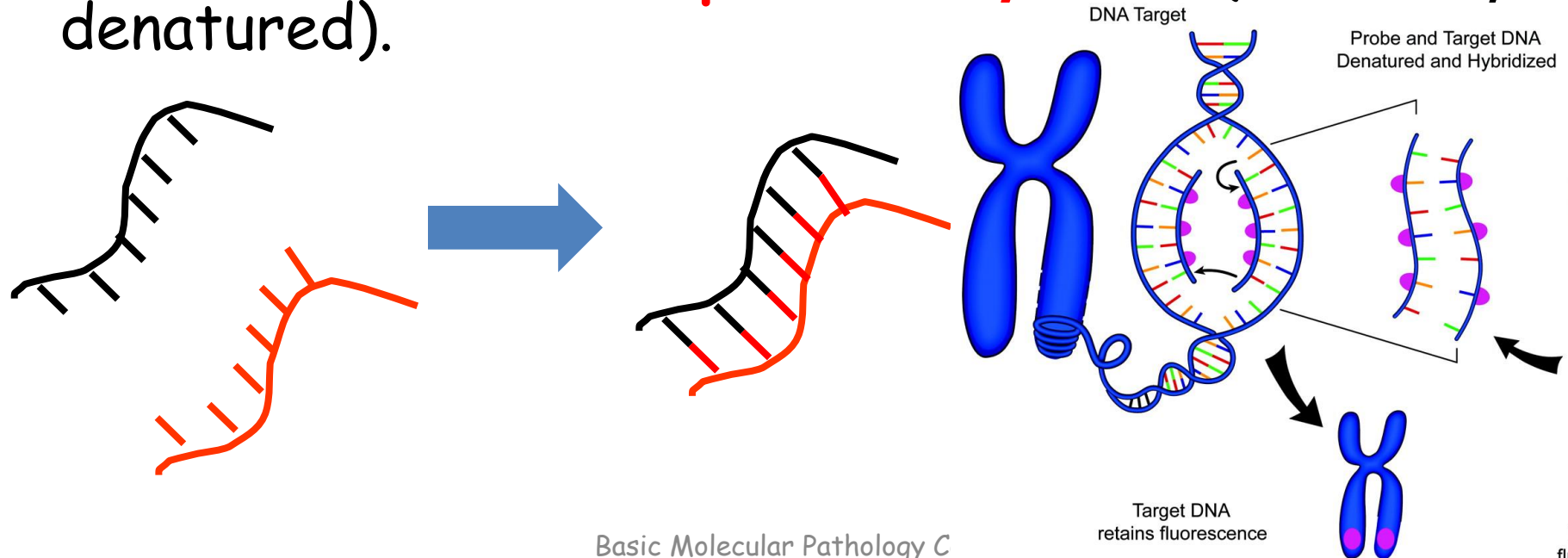


TERMINOLOGY

Annealing

For complementary sequences of single-stranded DNA or RNA to pair by hydrogen bonds to form a double-stranded nucleotide complex.

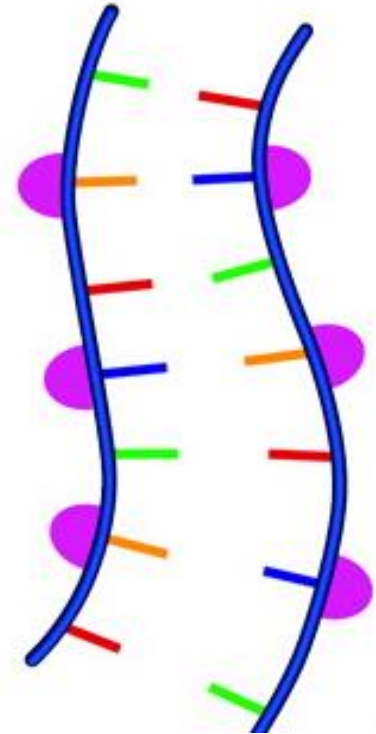
Binding of a probe. The term is also often used to describe the **renaturation of complementary strands that were separated by heat** (thermally denatured).



TERMINOLOGY

Probe

A fragment of **labelled** DNA or RNA of variable length (usually 100-1000 bases long) which is used in DNA or RNA samples to detect the presence of nucleotid sequences that are complementary to the target sequence.



HYBRIDISATION

DNA - DNA

DNA - RNA

RNA - RNA

THE MAIN PRINCIPLE OF
"HYBRIDIZATION TECHNIQUES"
ARE TO VISUALIZE
THIS REACTION

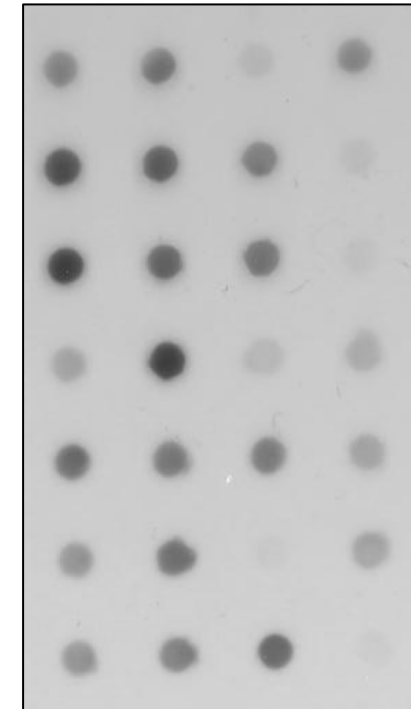
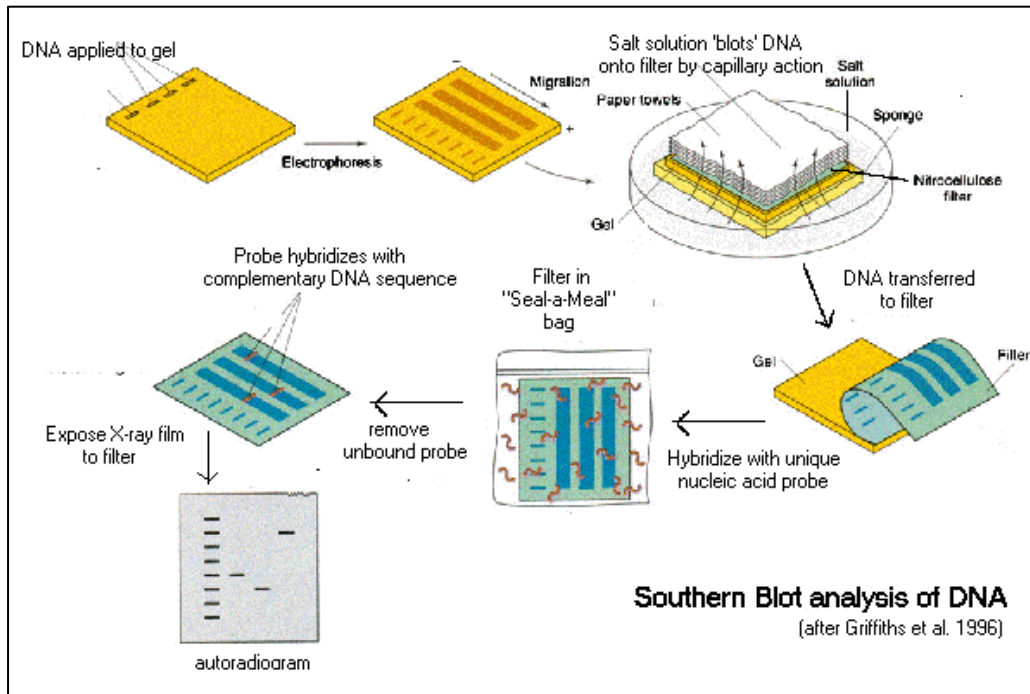


HYBRIDIZATION TECHNIQUES

- 1- Blotting
- 2- Array technologies
- 3- In situ hybridization
- 4- Comparative Genomic Hybridisation

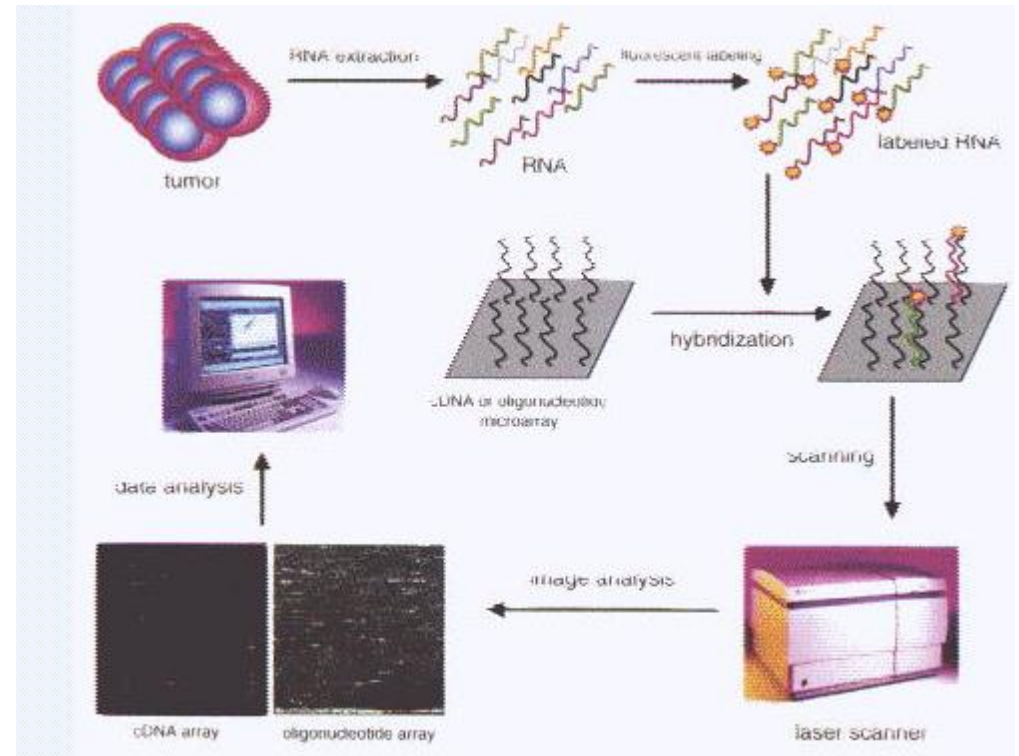
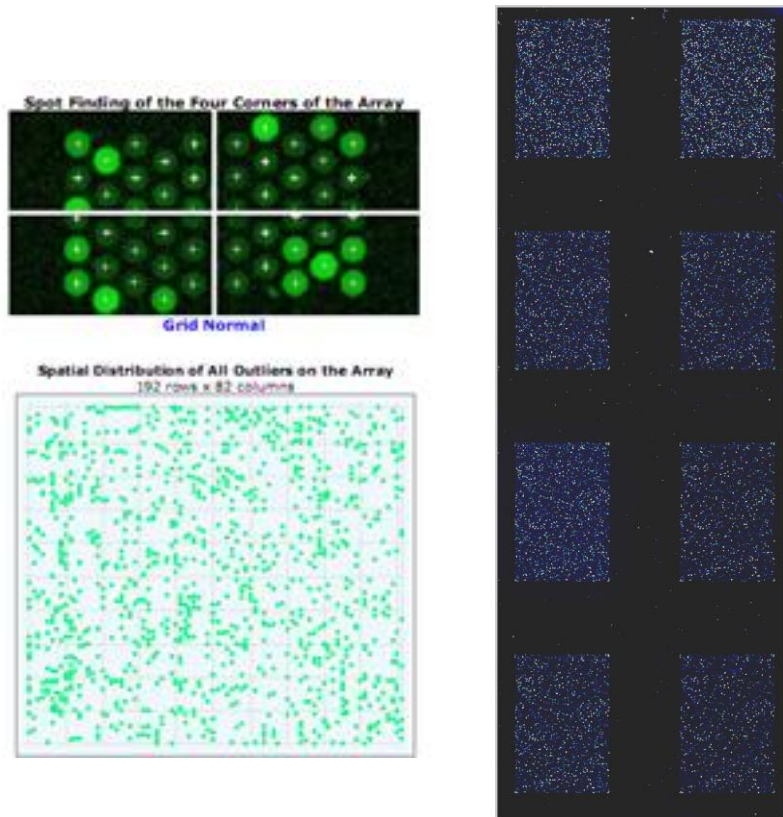
HYBRIDIZATION TECHNIQUES

1- Blotting : Southern Blot - Northern Blot, Dot blot



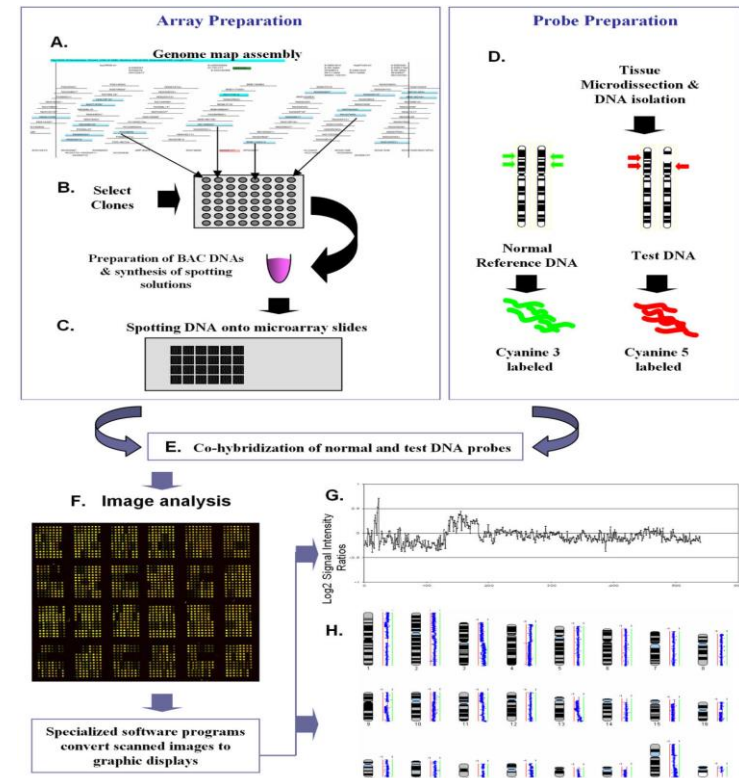
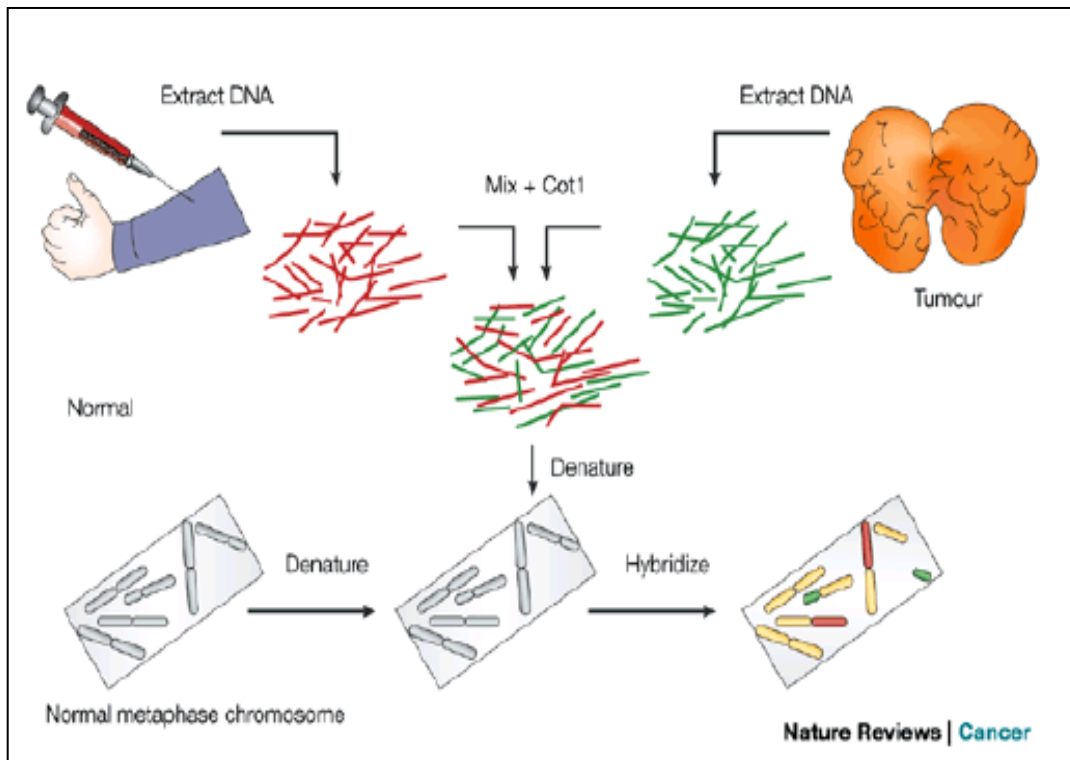
HYBRIDIZATION TECHNIQUES

2-Array technologies : (macroarray ve microarray, gene expression profile detection).



HYBRIDIZATION TECHNIQUES

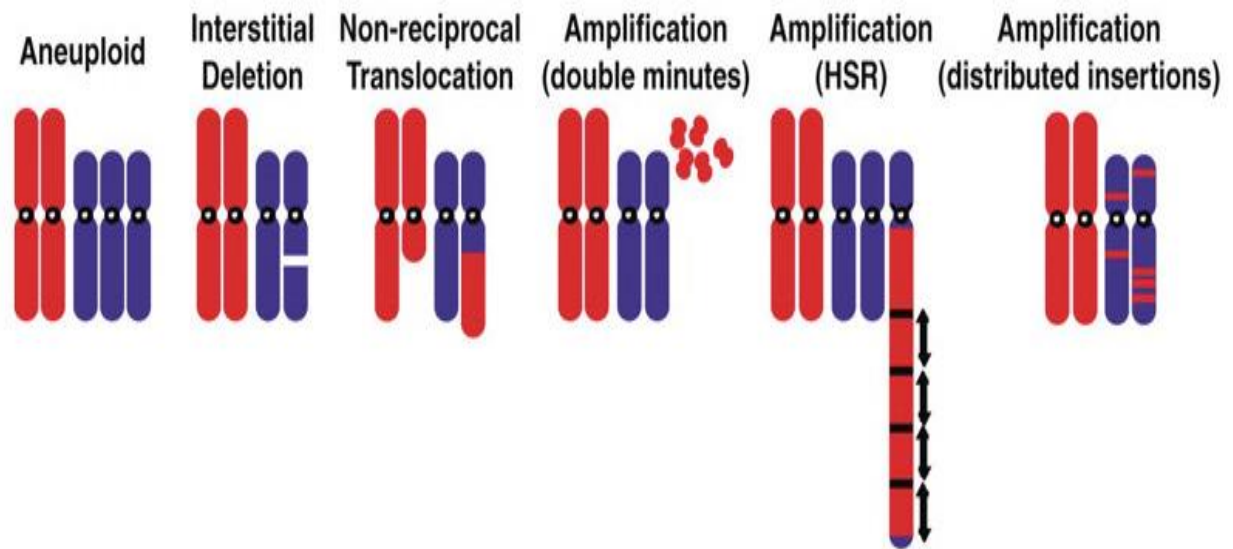
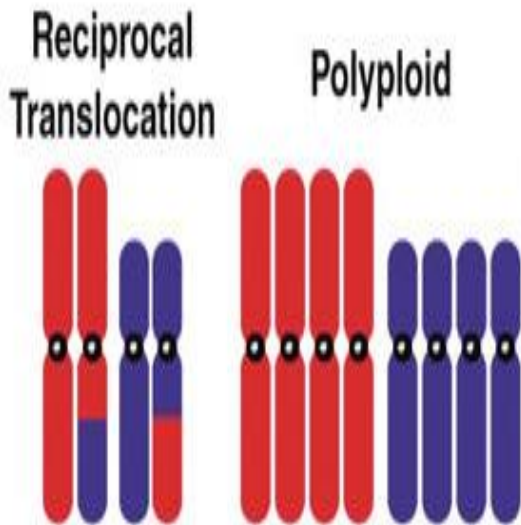
4- Comparative Genomic Hybridisation (CGH, ARRAY -CGH)



LIMITATIONS OF CGH

Can not be detectable
Balanced
chromosomal
abnormalities

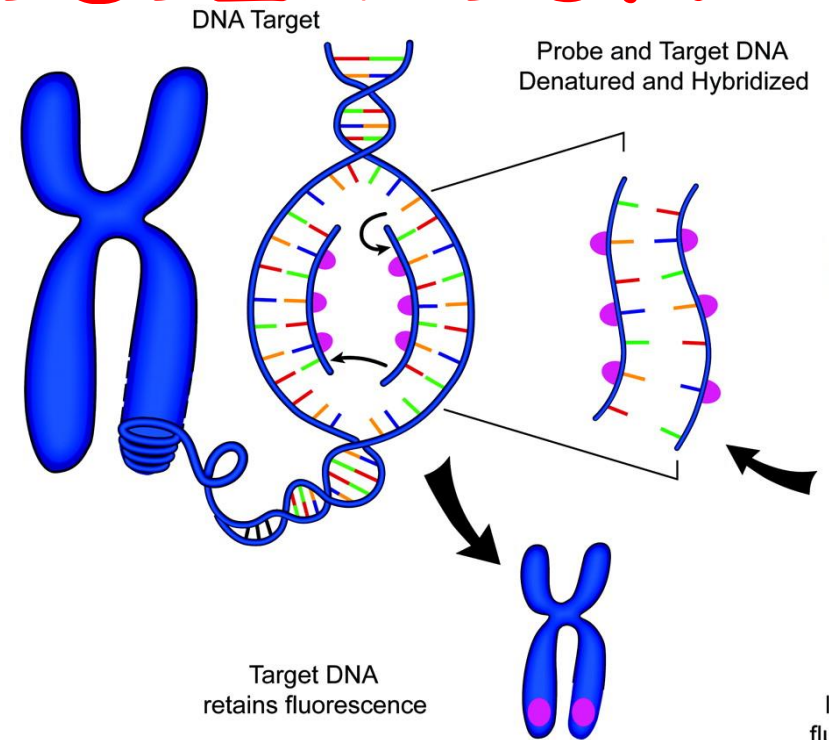
Can be detected
Unbalanced chromosomal
abnormalities



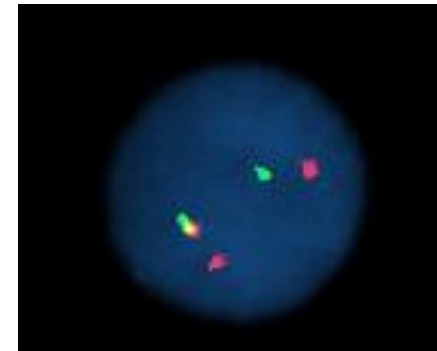
IN SITU HYBRIDIZATION

The principle of ISH

1- The **annealing of a labelled probe** to its complementary strand within the chromosomes of fixed cells or tissues

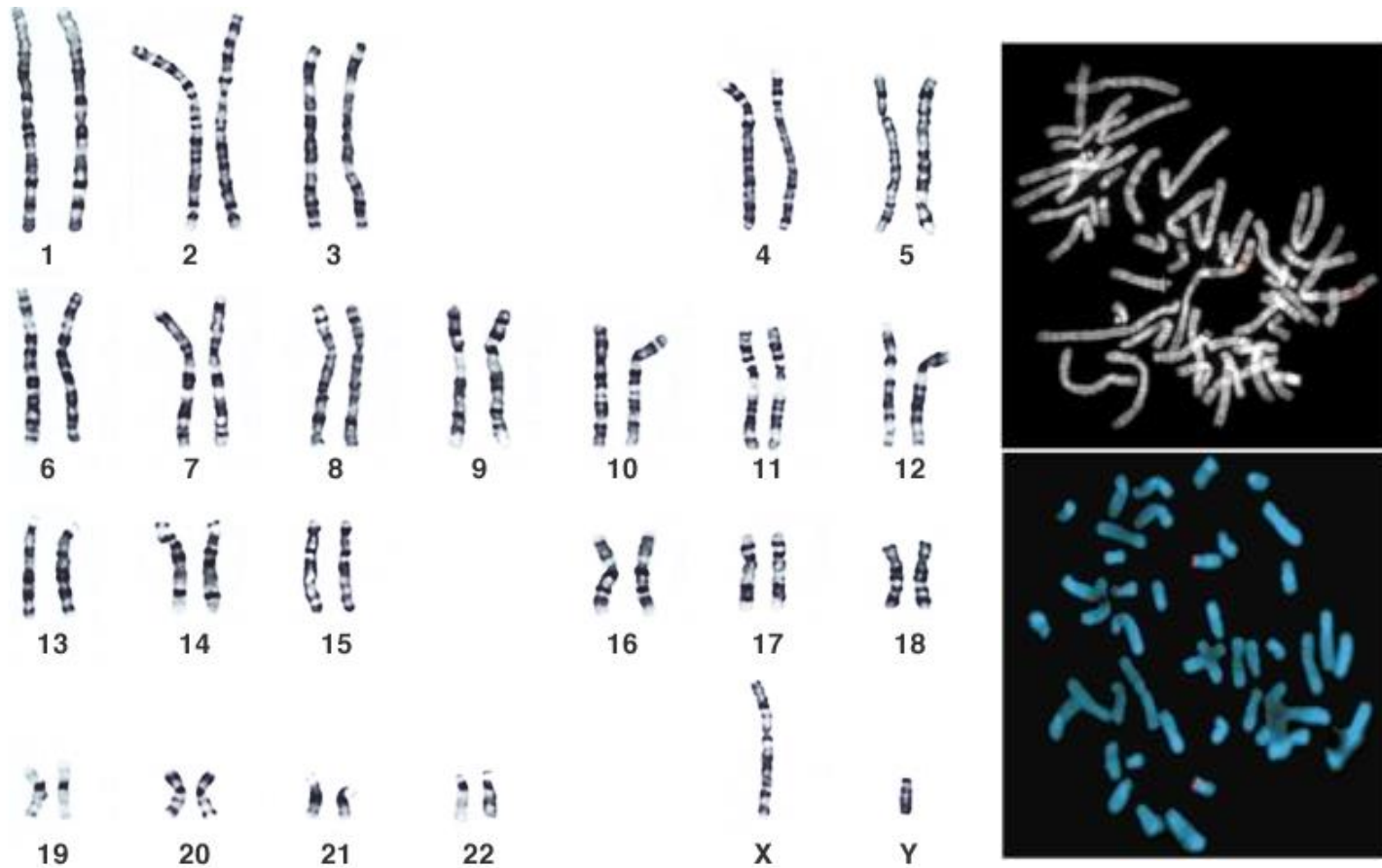


2- The **detection of the label.**



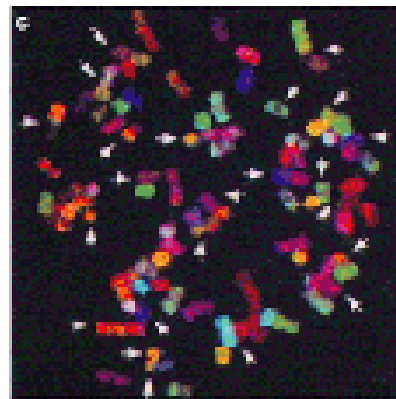
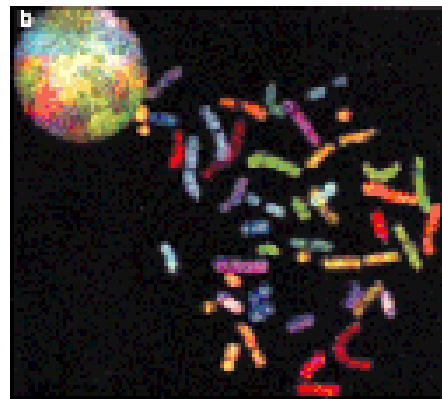
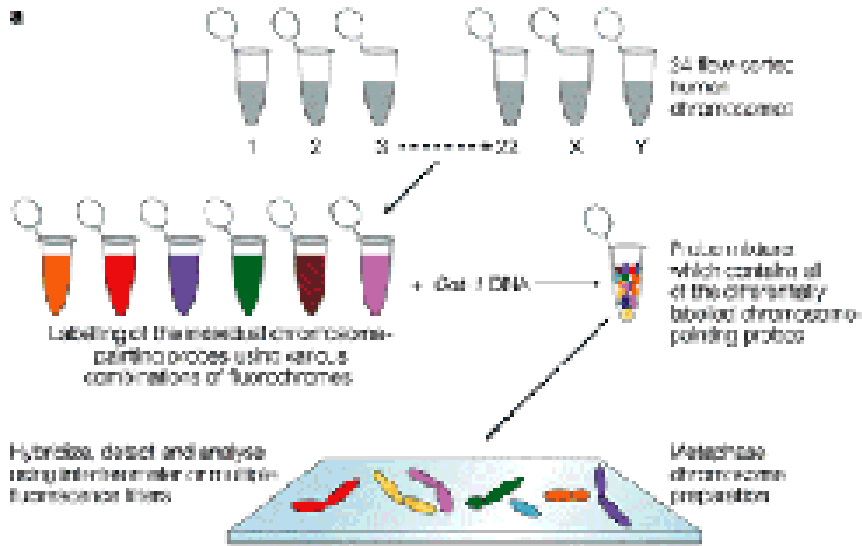
IN SITU HYBRIDIZATION

important tool for cytogenetic analysis

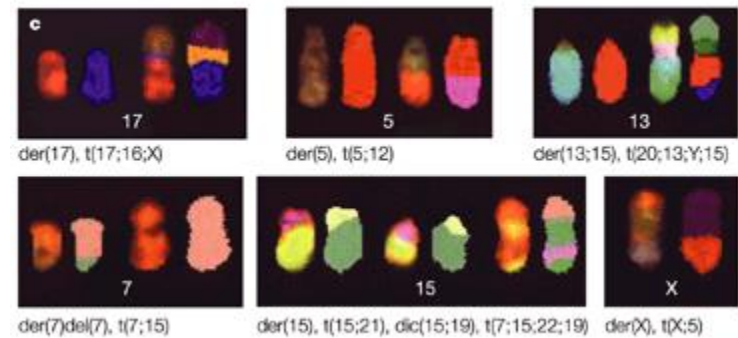
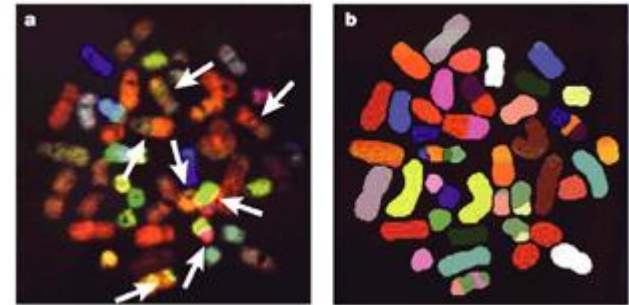


HUMAN CHROMOSOMES IN METAPHASE

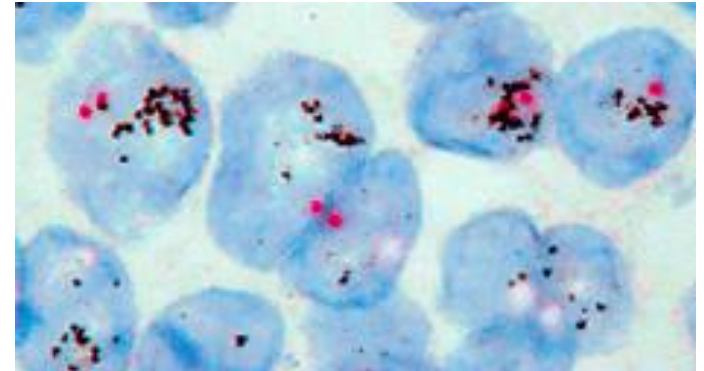
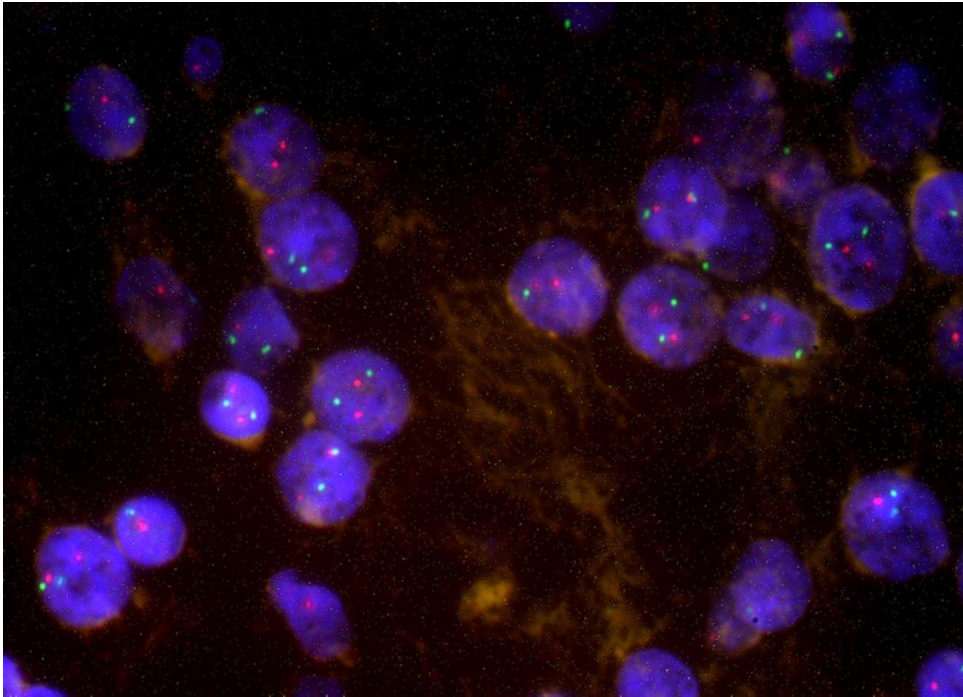
SPECTRAL CARYOTYPING



Nature Reviews | Genetics



ISH FOR PATHOLOGY



Chromosomes on interphase

WHAT COULD BE SEEN BY USING ISH?



GENE COPY NUMBER INCREASE (AMPLIFICATIONS)

GENE OR CHROMOSOME LOSS (DELETIONS)

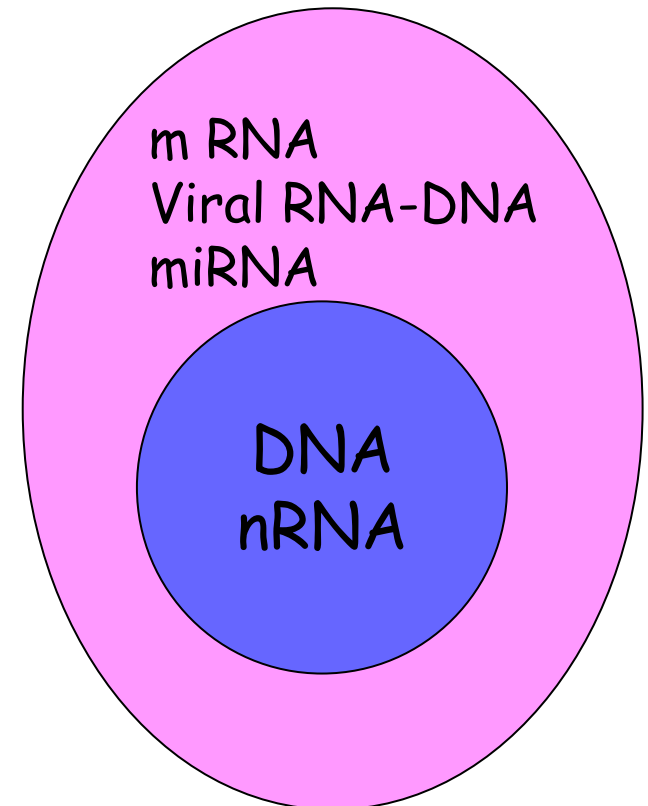
GENE REARRANGEMENTS (TRANSLOCATIONS)

INFECTIOUS AGENTS

PROTEIN EXPRESSION (mRNA)

CHIMERISM (X, Y)

miRNA's



PROBE TYPES

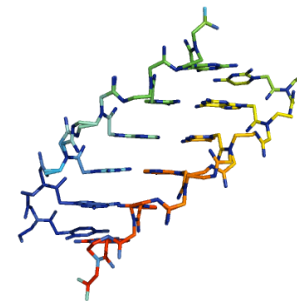
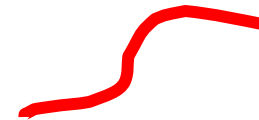
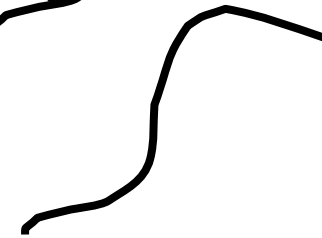
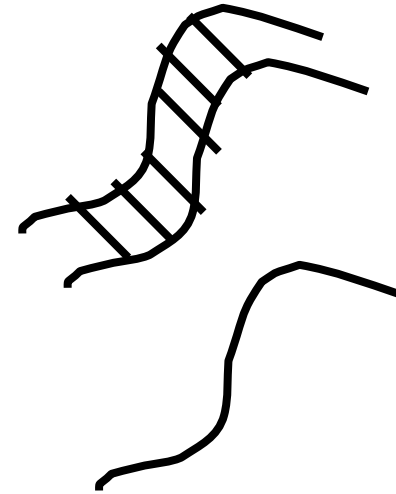
Double stranded DNA

Single stranded DNA

RNA

Oligonucleotide

PNA (peptide nucleic acid)



PROBE LABELS

- 1- Radioactive labeled probes
- 2- Non radioactive labeled probes

Biyotin-Avidin-Fluorophore conjugated

Biyotin-Avidin-Enzyme (Alkalin phophatase / HRP)

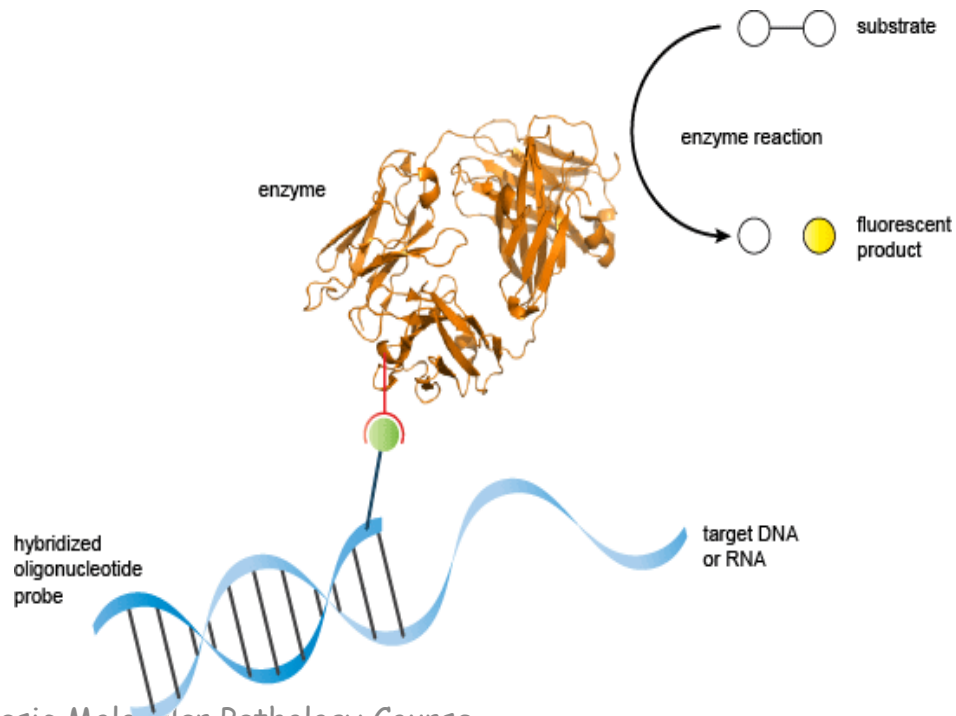
Hapten (Digoxigenin (DIG) , Dinitrophenyl (DNP))

Fluorophore

Enzyme
+
Substrate



Colourimetric,
Chemiluminescent
Fluorescent
product



PROBE SYNTHESIS

The probes (DNA or RNA) usually prepared by polymerase enzyme-based methods

Nick translation,
Random priming
PCR

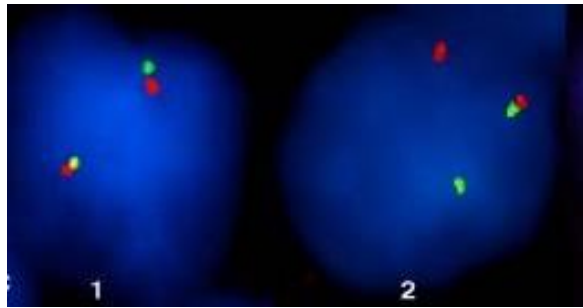
} incorporation of fluorescently-labelled deoxynucleotide triphosphates

The length of a DNA probe can be between 100 bp and 1000 bp.

Longer probes increase non-specific background fluorescence

Shorter probes difficult to detect - low levels of labelling.

Fluorescence in situ hybridization

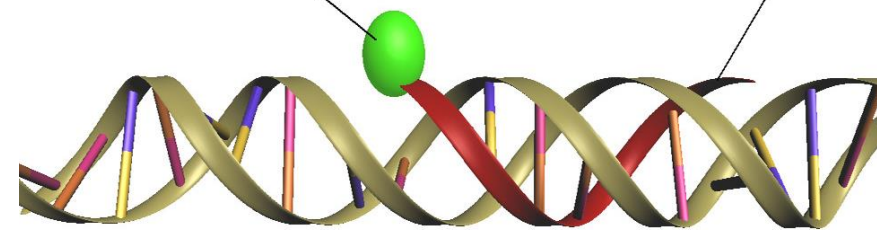


(FISH)

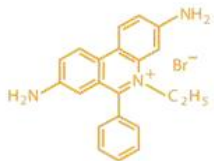


Fluorescent label

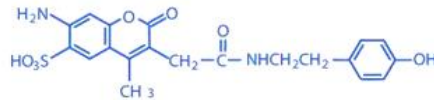
Labeled probe



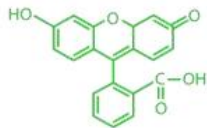
Definition of Fluorescence



Ethidium bromide

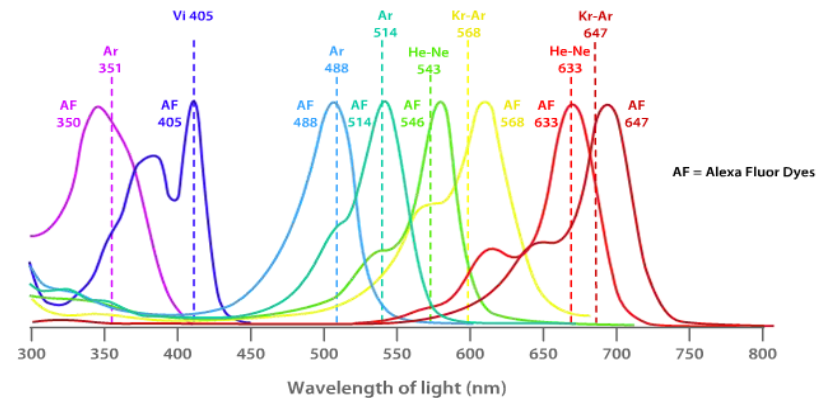


Alexa Fluor 350

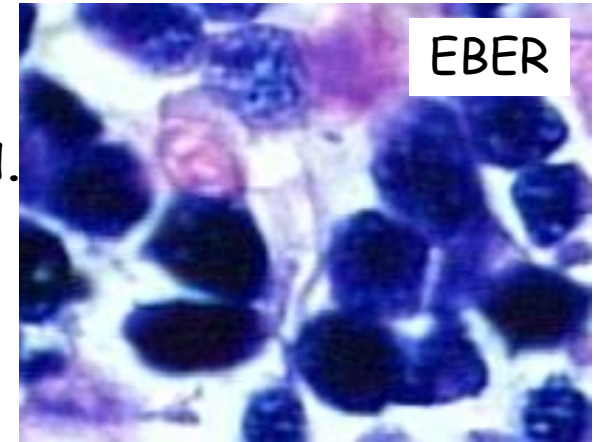
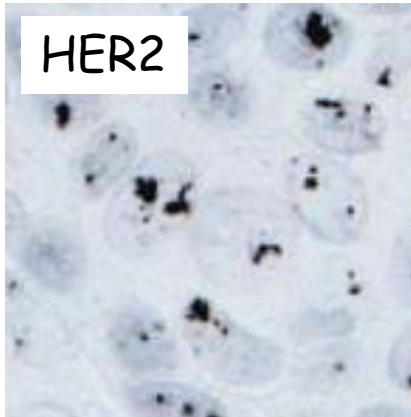


Fluorescein

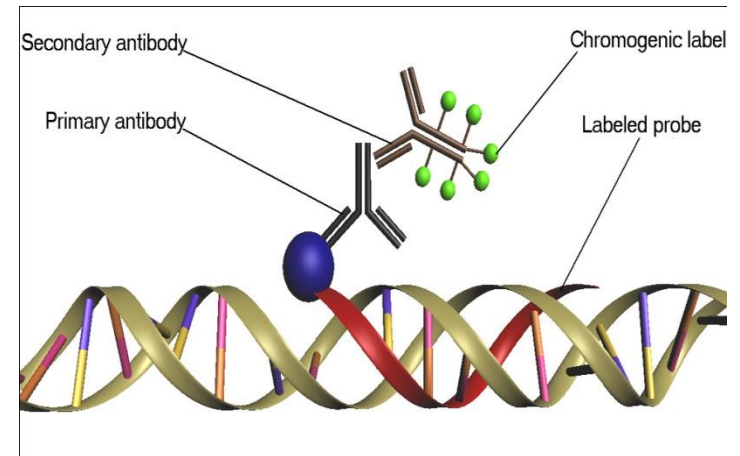
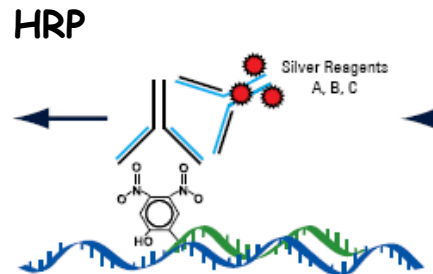
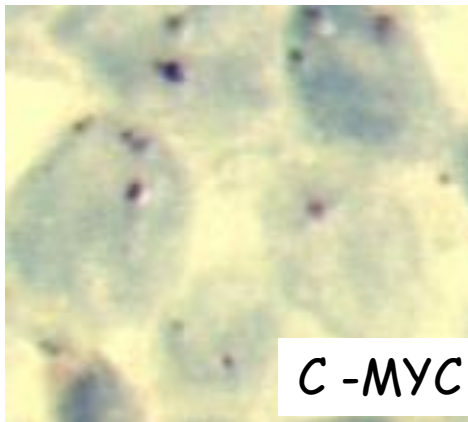
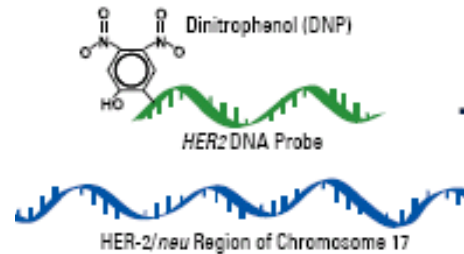
How and why do these
dyes and stains emit
different colors of light?



Chromogenic in situ hybridization (CISH)

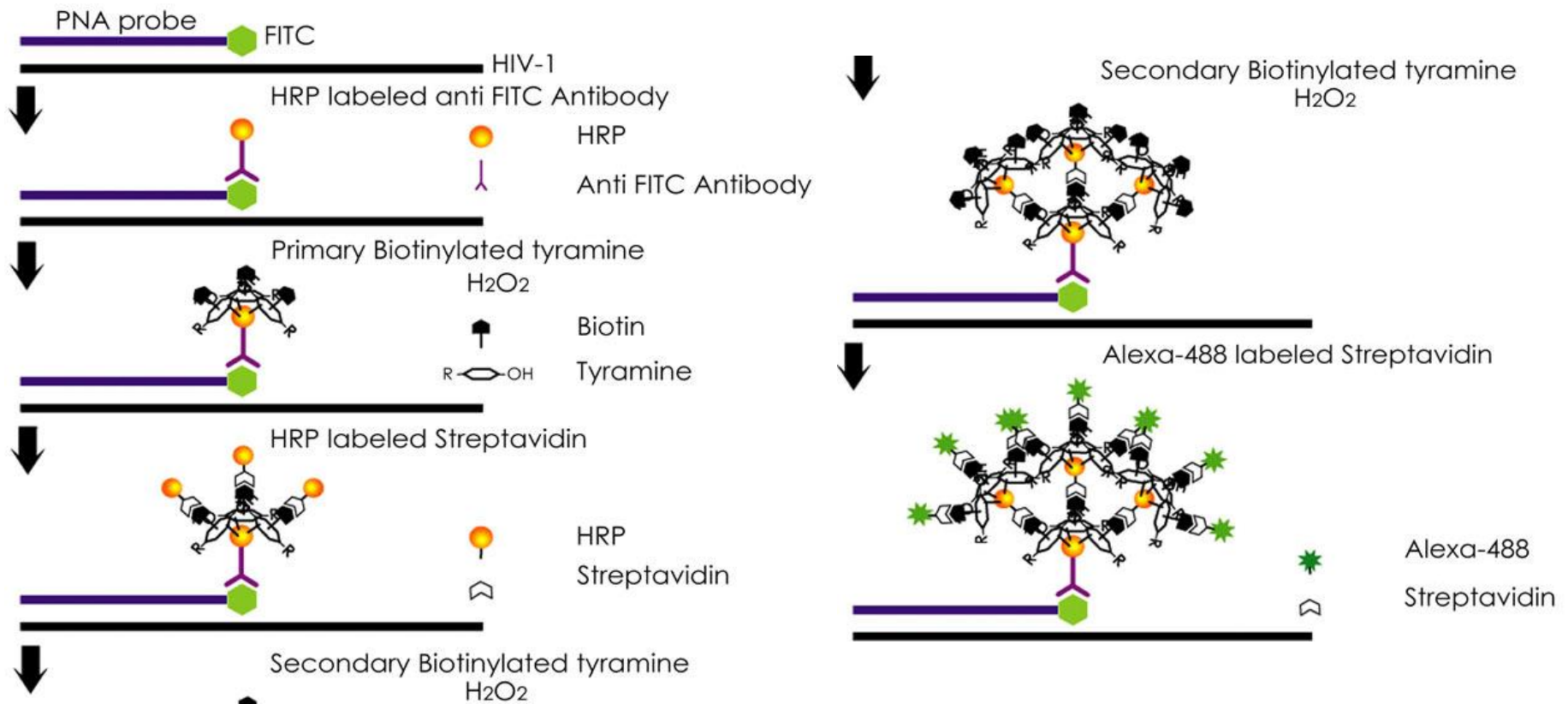


Less sensitive than FISH.
Better for seaching amplification signals



<http://www.histalim.com/pages/en/in-situ-hybridization.php>

PROBE SIGNAL AMPLIFICATION



Hagiwara T, 2006, Methods in Mol Biol 326 Ch 9, ED: Darby IJ

VISUALIZATION OF THE HYBRIDISATION



The target must be accessible by the probe

The target must retain in situ, not degraded by nuclease enzymes.

DNA is more stable than RNA



MOST FREQUENTLY USED MATERIALS FOR ISH APPLICATIONS

- **CULTURED CELLS**
- **ASPIRATION SMEARS**
- **FRESH FROZEN TISSUE SECTIONS**
- **PARAFFIN SECTIONS**



IN SITU HYBRIDISATION PROCEDURE ON PARAFFIN TISSUE

- TISSUE PRETREATMENT
- PROBE PREPERATION
- PROBE INCUBATION
- WASHING
- MOUNTING
- EXAMINATION

SPACE NEEDED FOR THE PROCEDURE



Full automated systems



TISSUE FIXATION

- Probe penetration
- RNA damage
- DNA damage
- Preservation of the tissue architecture

Most frequently used fixatives

Alcohol-acetic acid : allows probe penetration /
RNA damage

Gluteraldehyde : poor probe penetration /
RNA preservation

Paraformaldehyde (%4) Formalin (%10) 😊



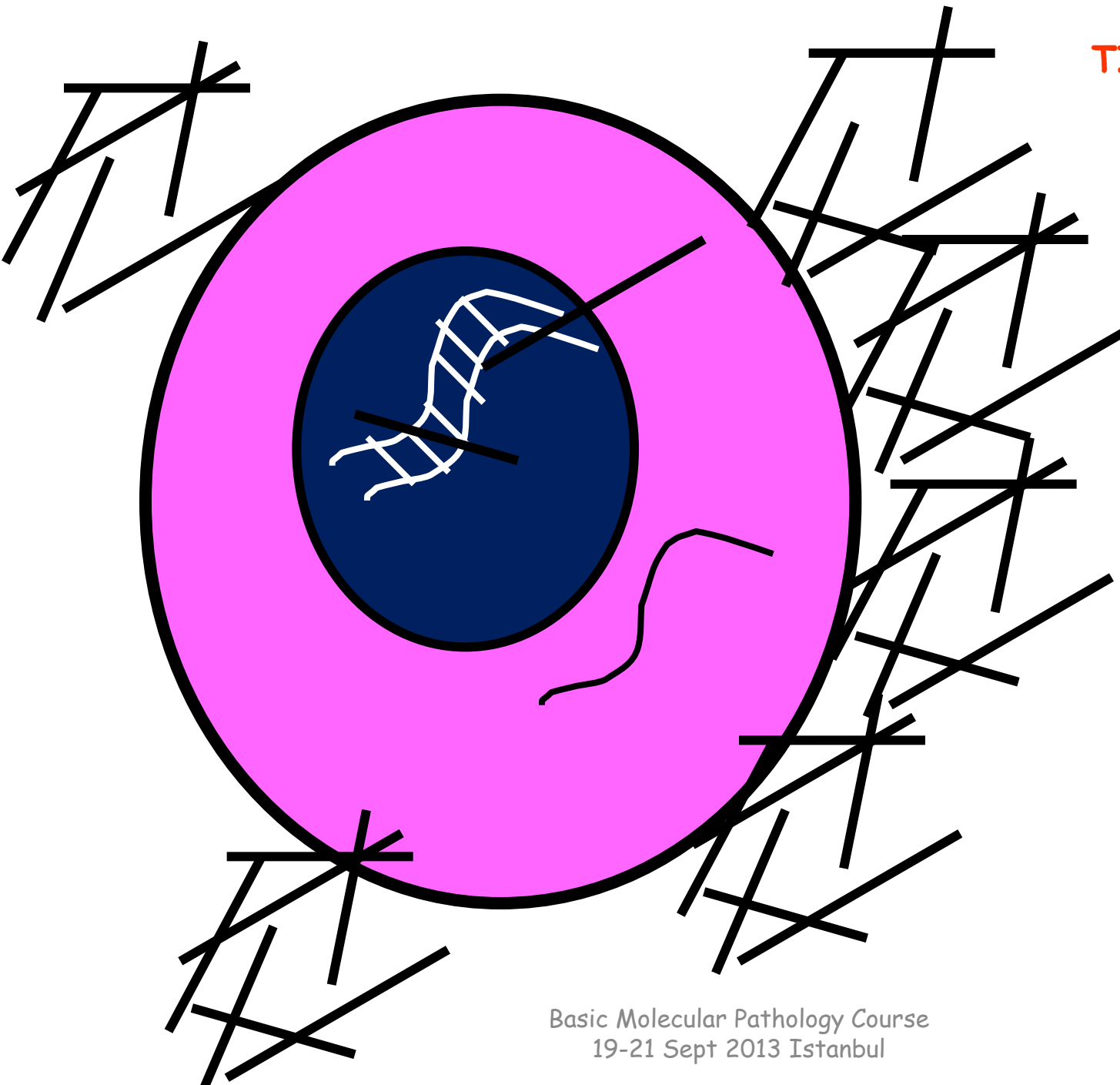
PRETREATMENT OF THE TISSUE BEFORE ISH

- DEPARAFFINIZATION
- INCREASING PROBE PENETRATION (HCl, enzyme, heat)
- DENATURATION
- STABILIZATION OF DENATURED DNA

DEPARAFFINIZATION



TISSUE



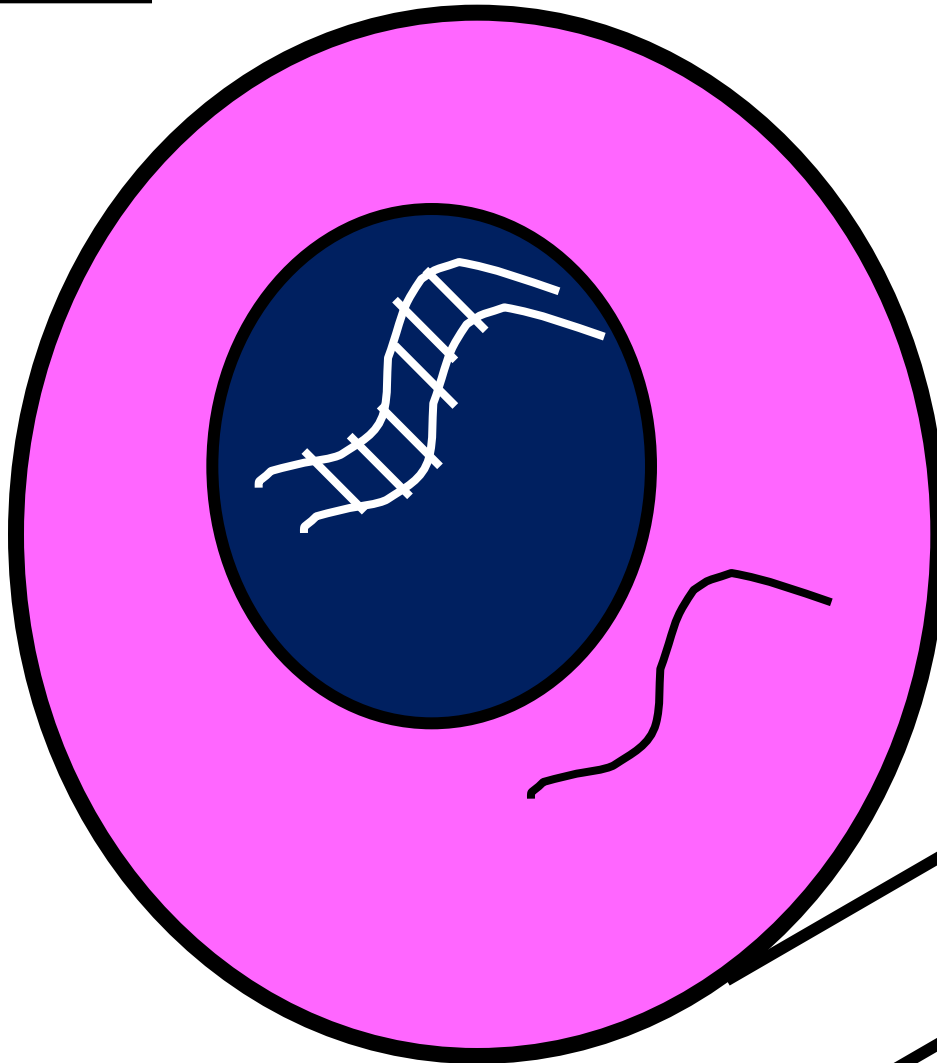
INCREASING PROBE PENETRATION



TISSUE PRETREATMENT

Removing lipids and some
other matrix matters

Detergents & HCl



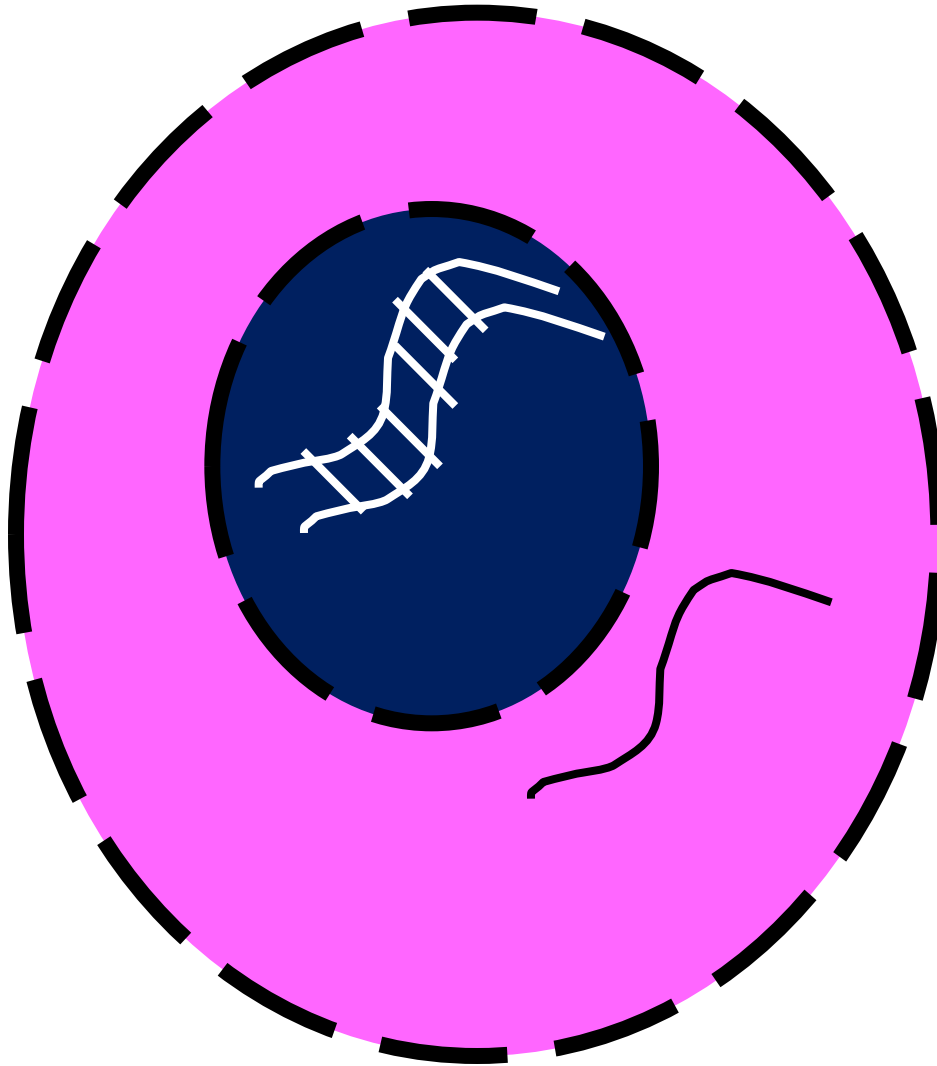
INCREASE PROBE PENETRATION



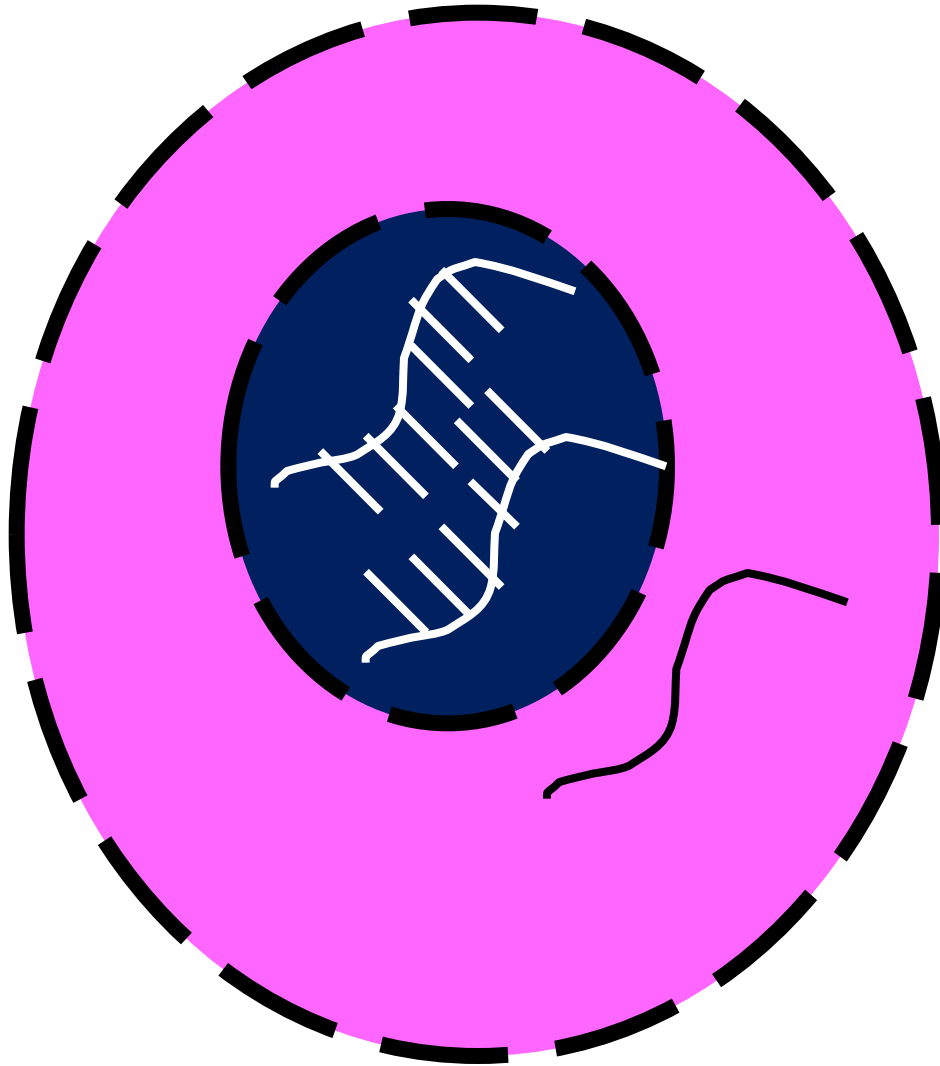
TISSUE PRETREATMENT

- *Removing matrix proteins
- *Making pores on cytoplasmic and nuclear membranes

Detergents & Enzyme



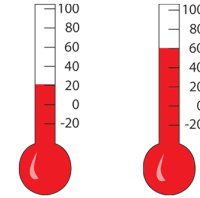
DENATURATION & STABILIZATION OF GENOMIC DNA



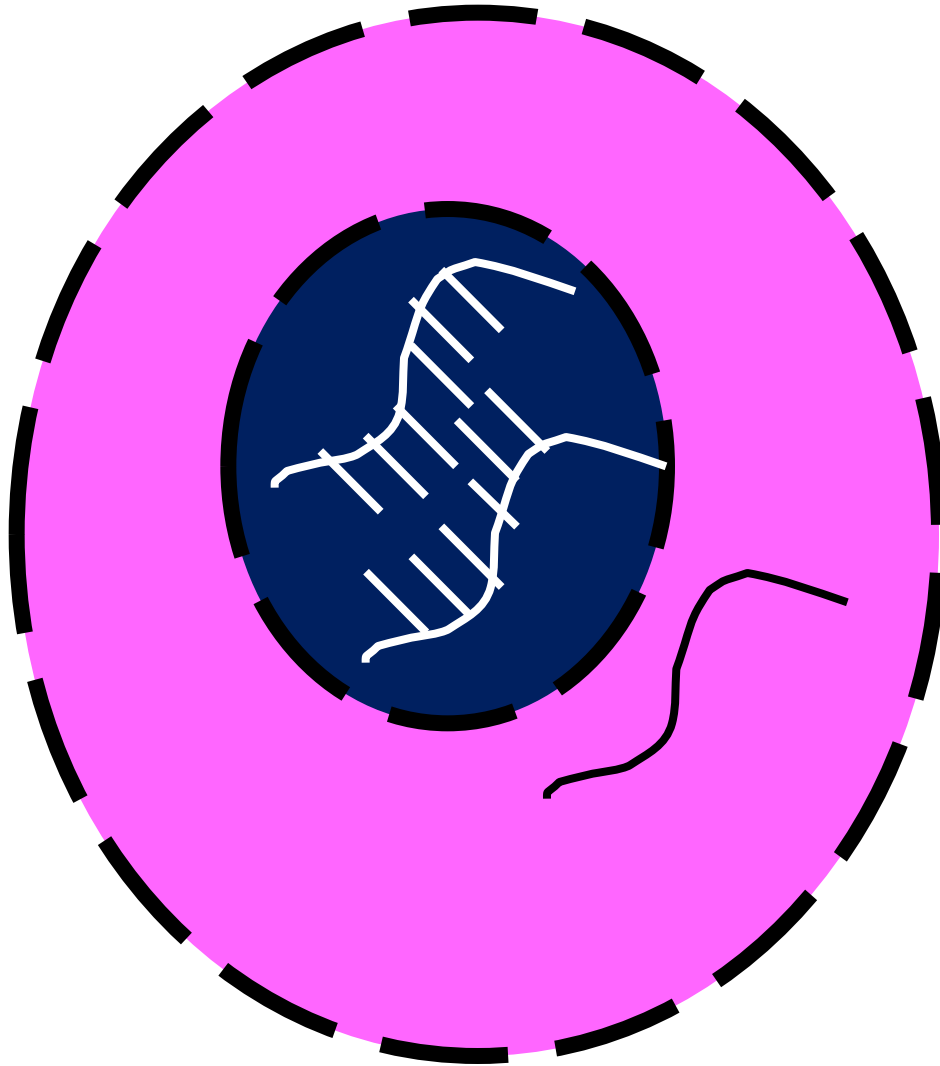
-HEAT

+

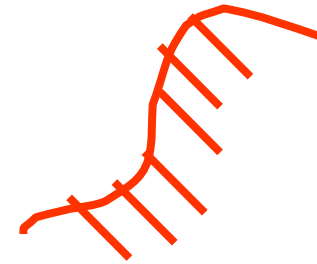
- Buffers containing
Formamide



PROBE INCUBATION



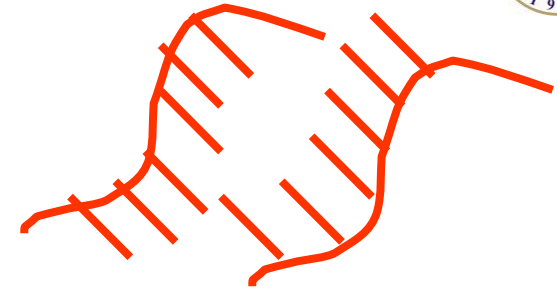
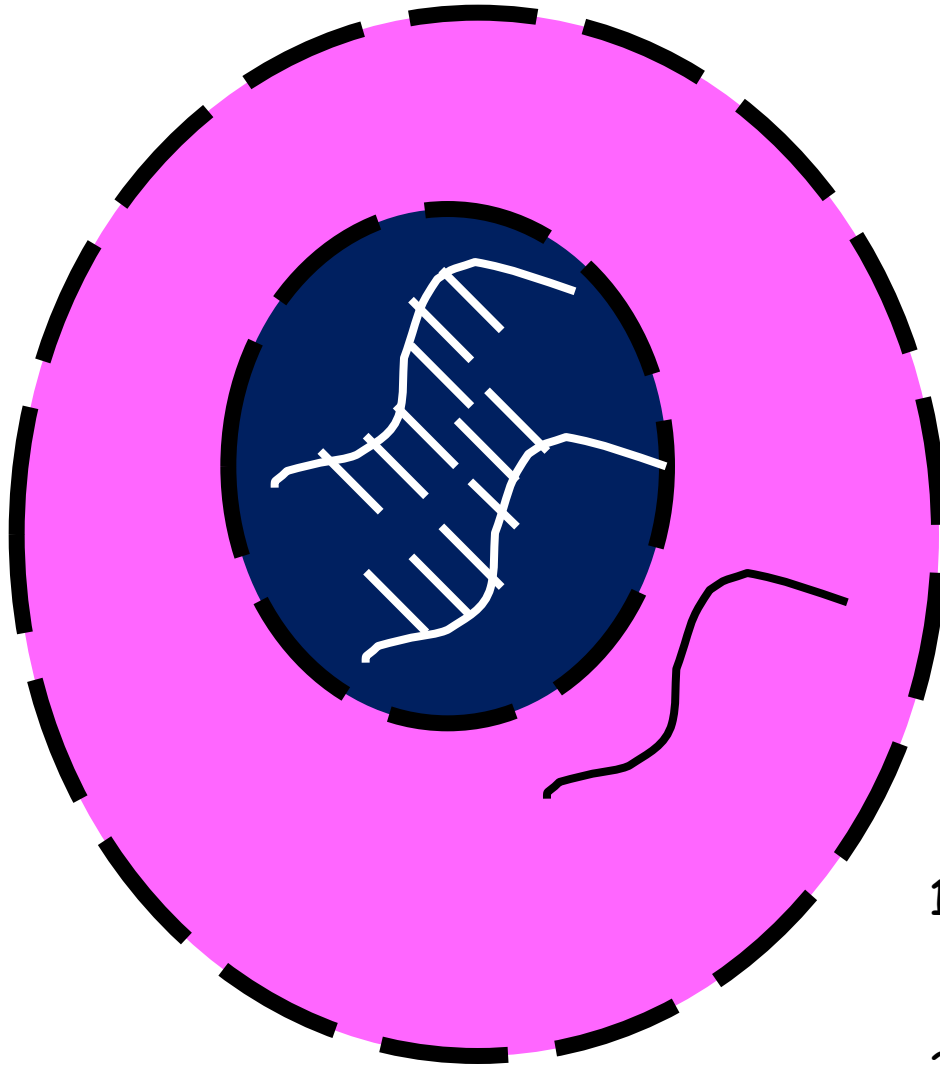
1. SINGLE STRANDED PROBE



2. DOUBLE STRANDED PROBE

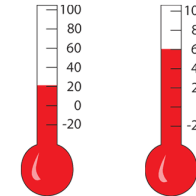


DENATURATION & STABILIZATION OF THE DOUBLE STRANDED PROBES



-HEAT

+



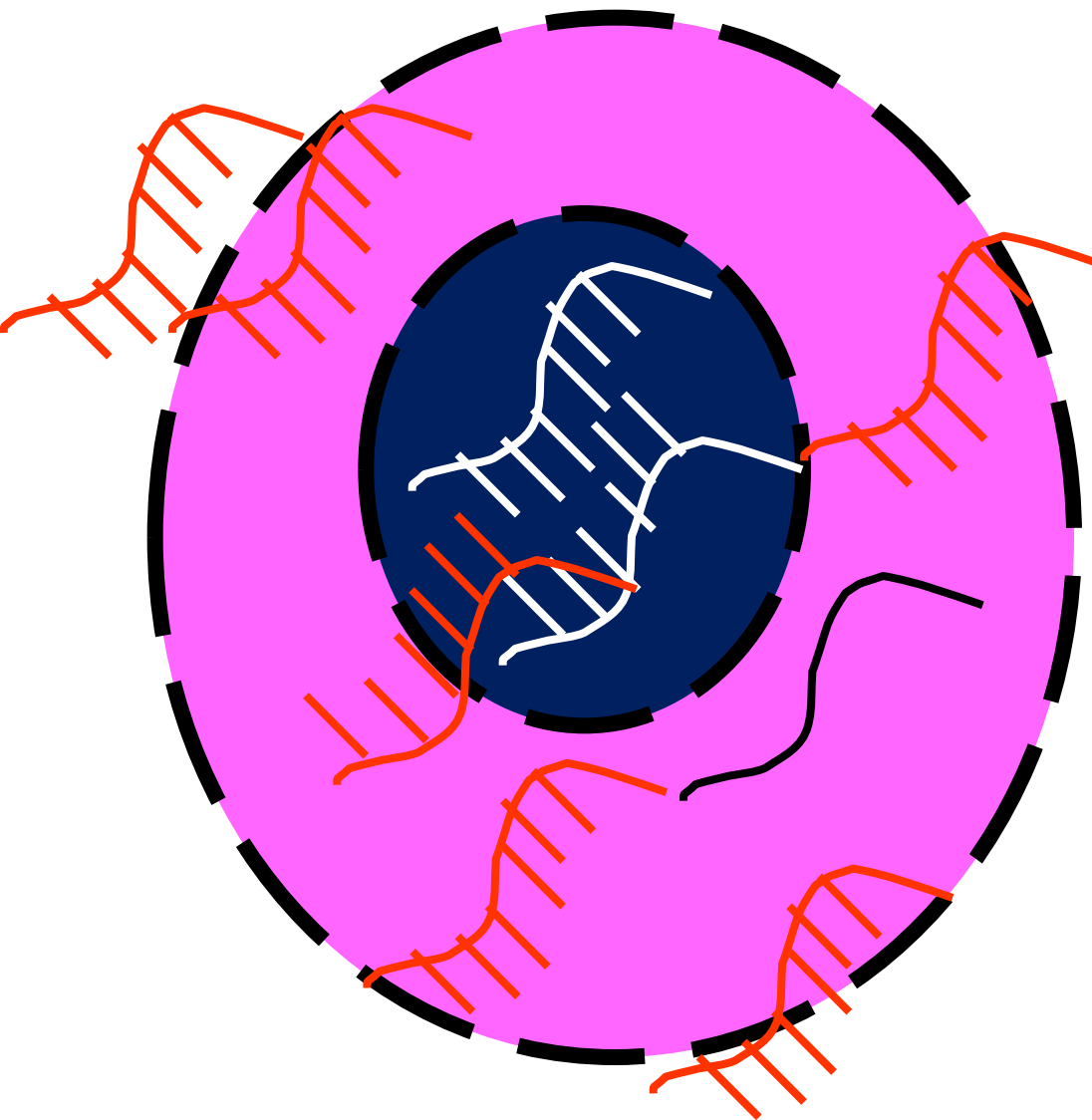
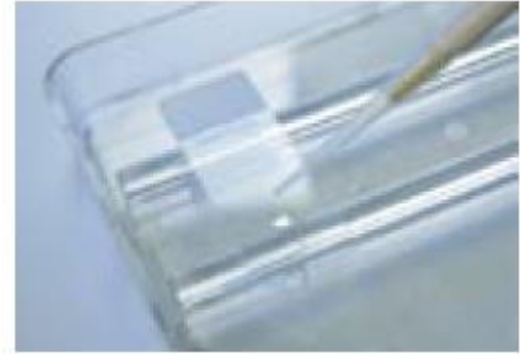
**- Buffers containing
Formamide**

1-SEPERATE



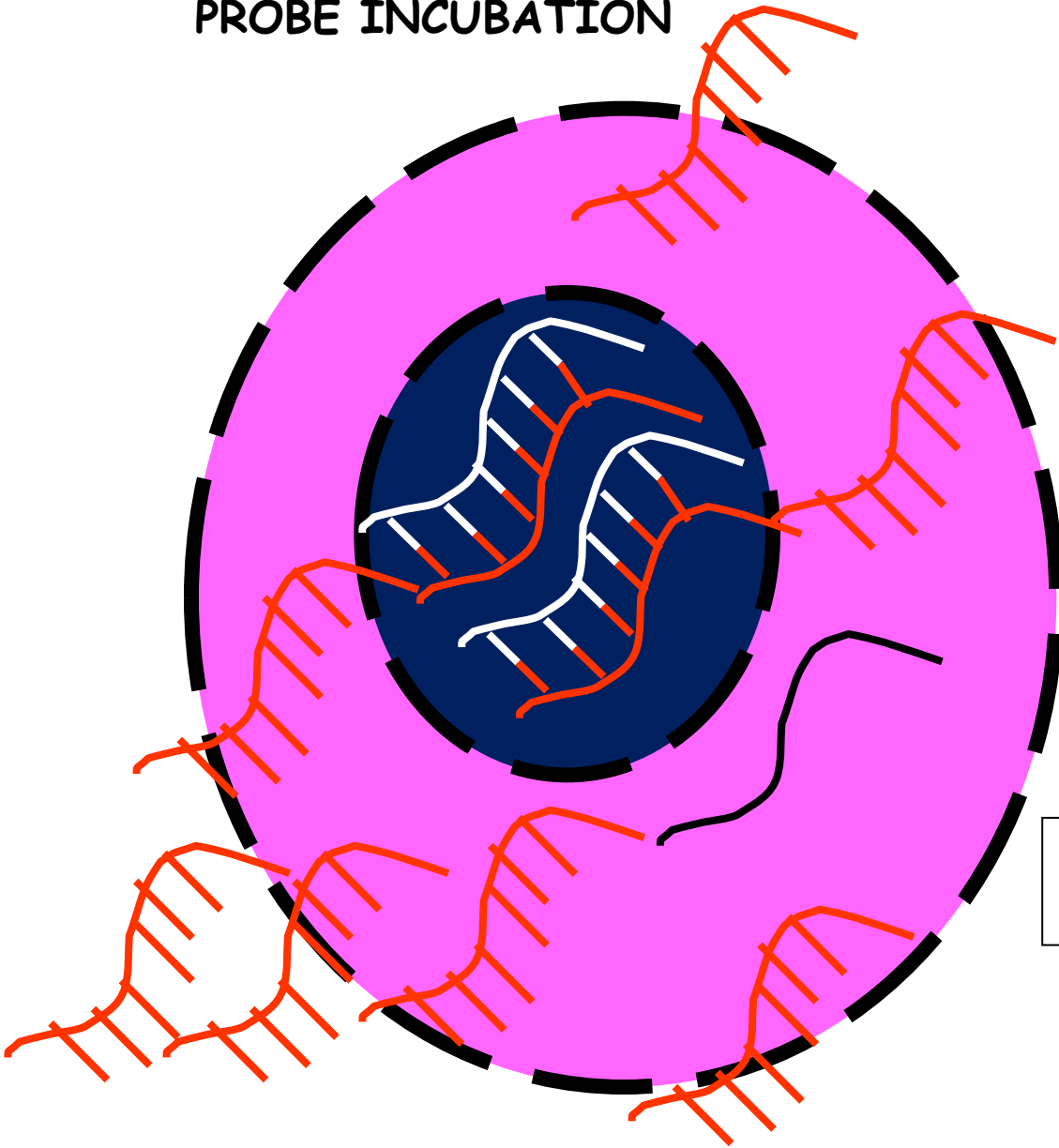
2- IN SITU ON THE TISSUE

PROBE INCUBATION

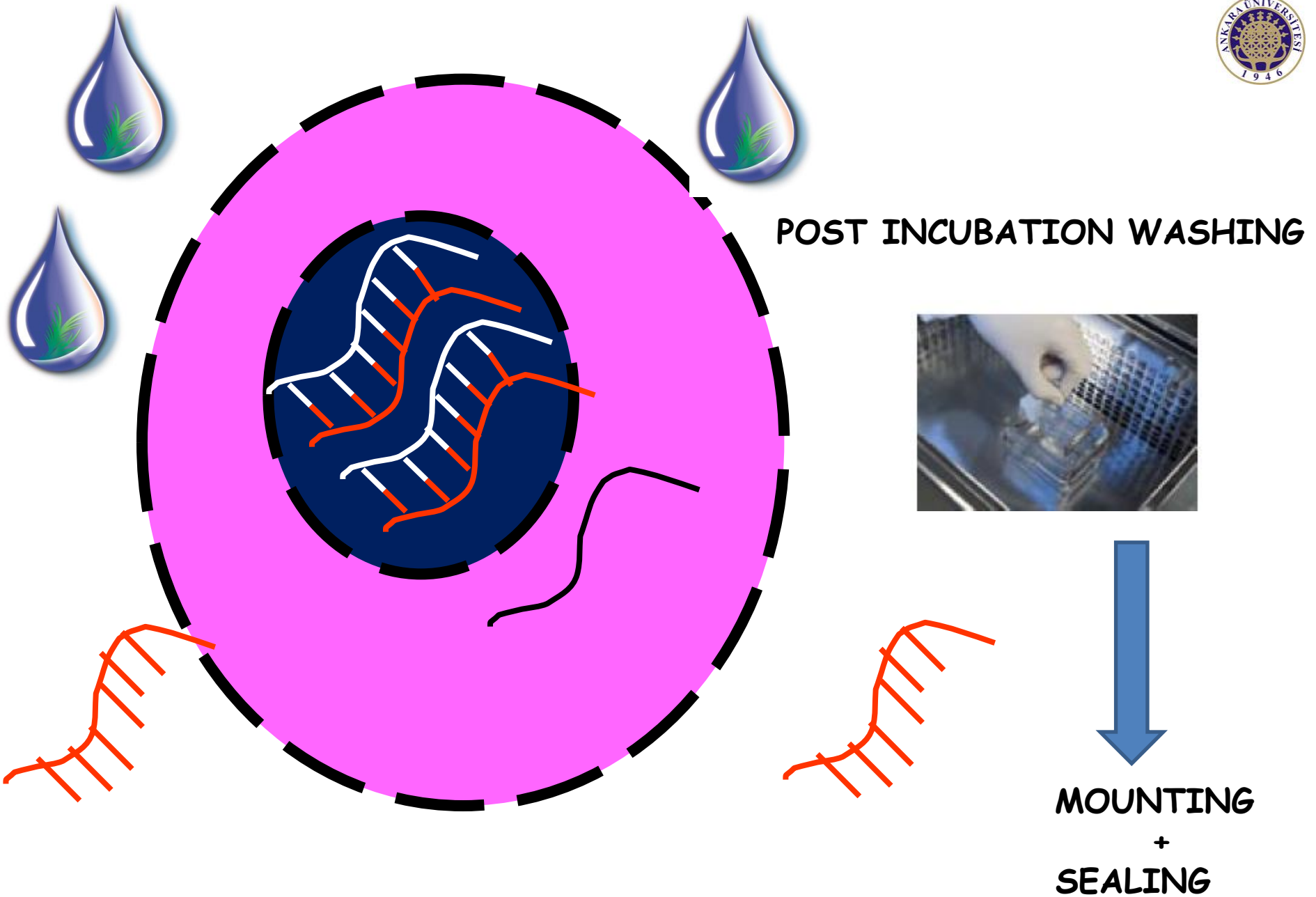


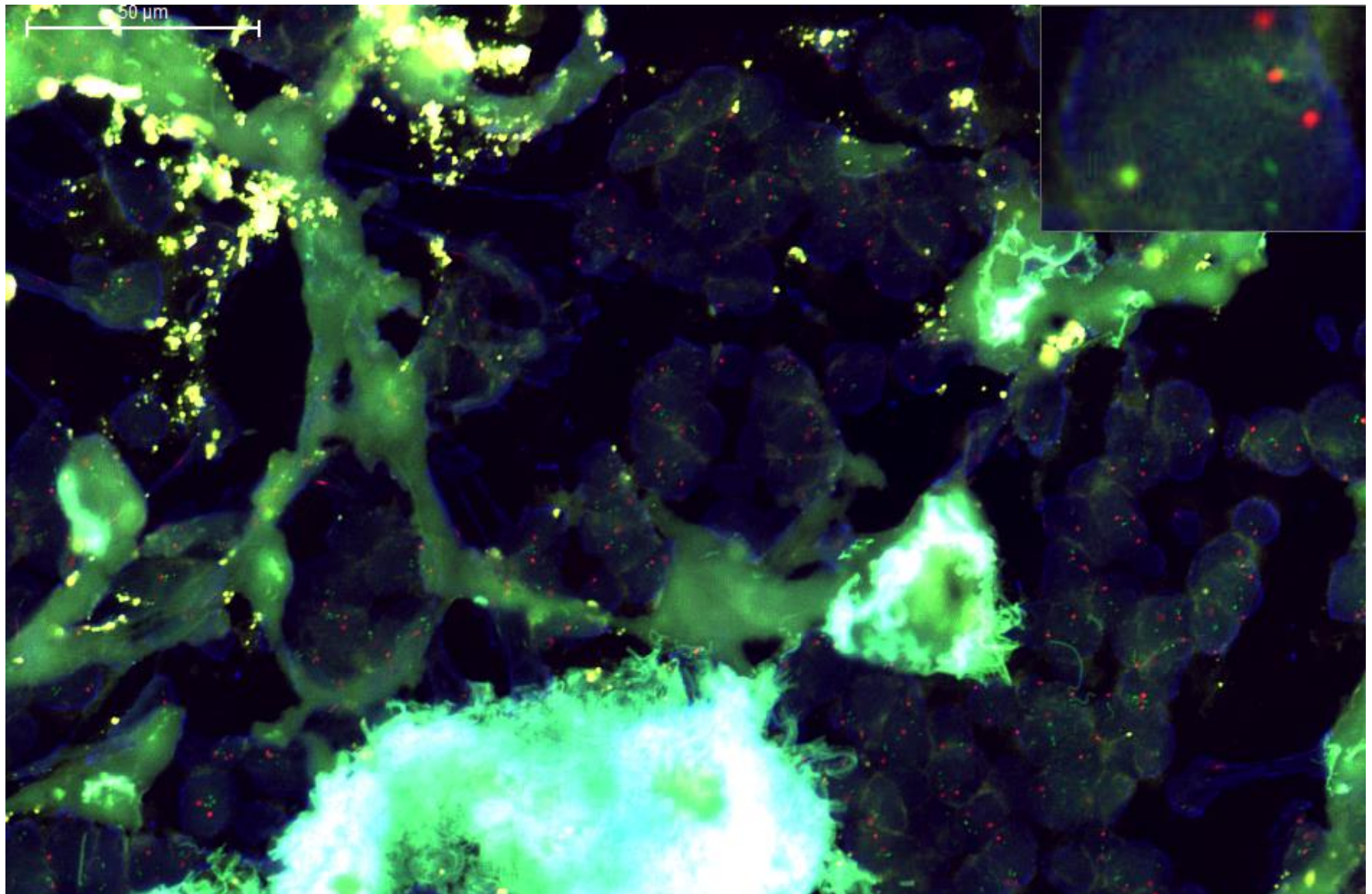
TIME (4-24 hr)
TEMPERATURE
HUMIDITY

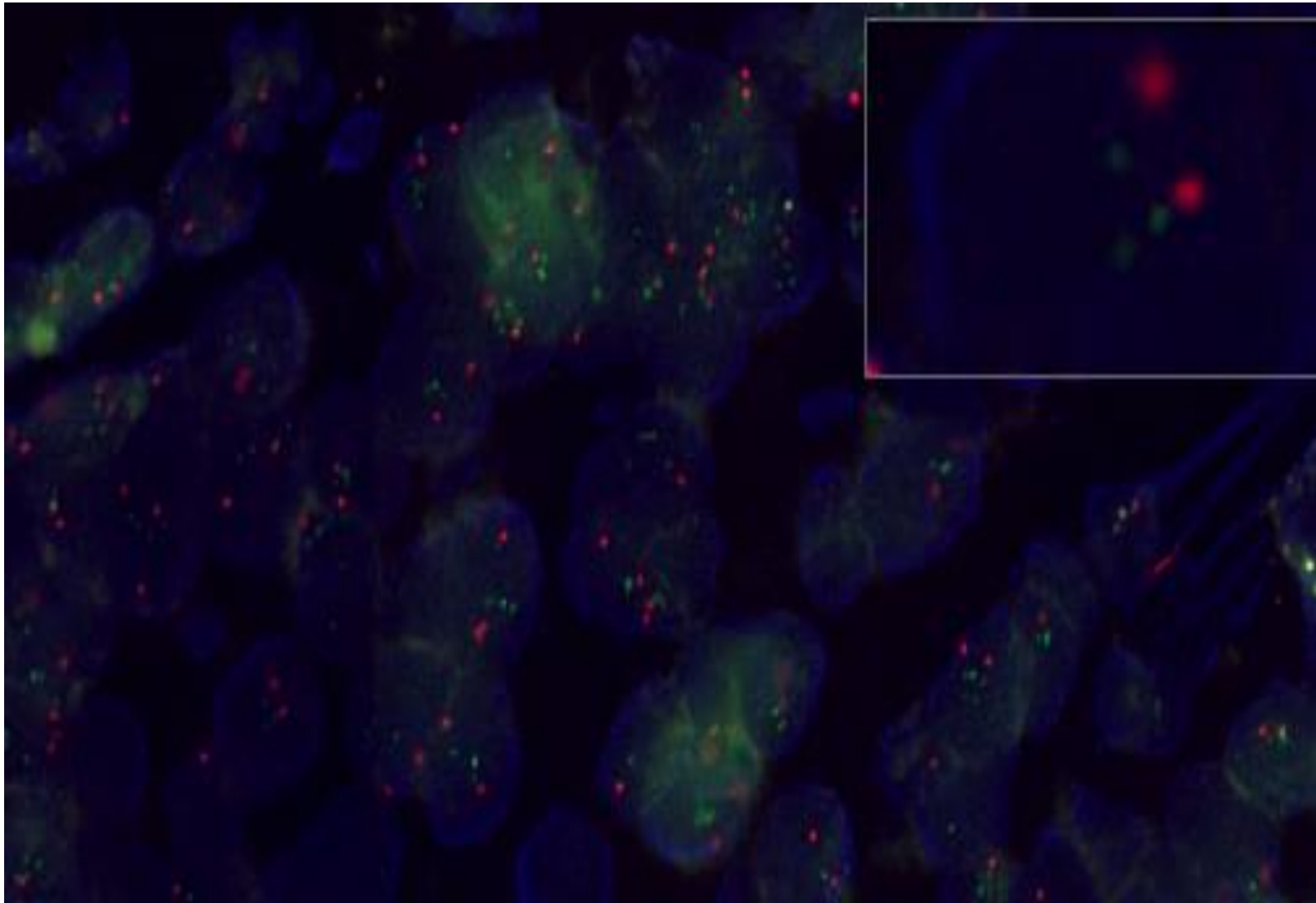
PROBE INCUBATION



**INCUBATION TEMPERATURE
(37-42°C)**



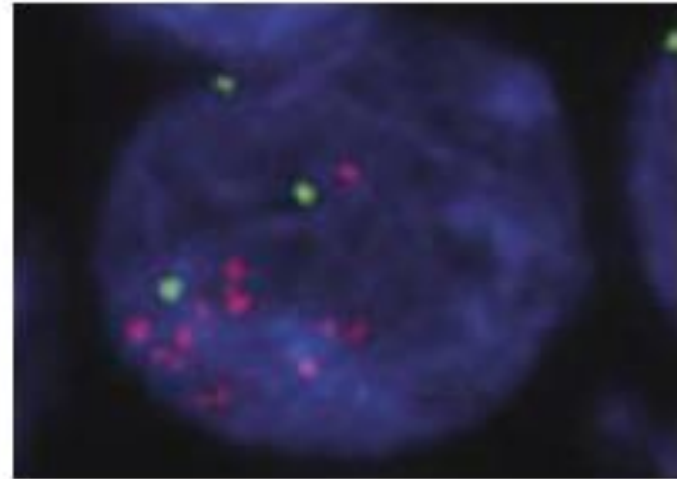




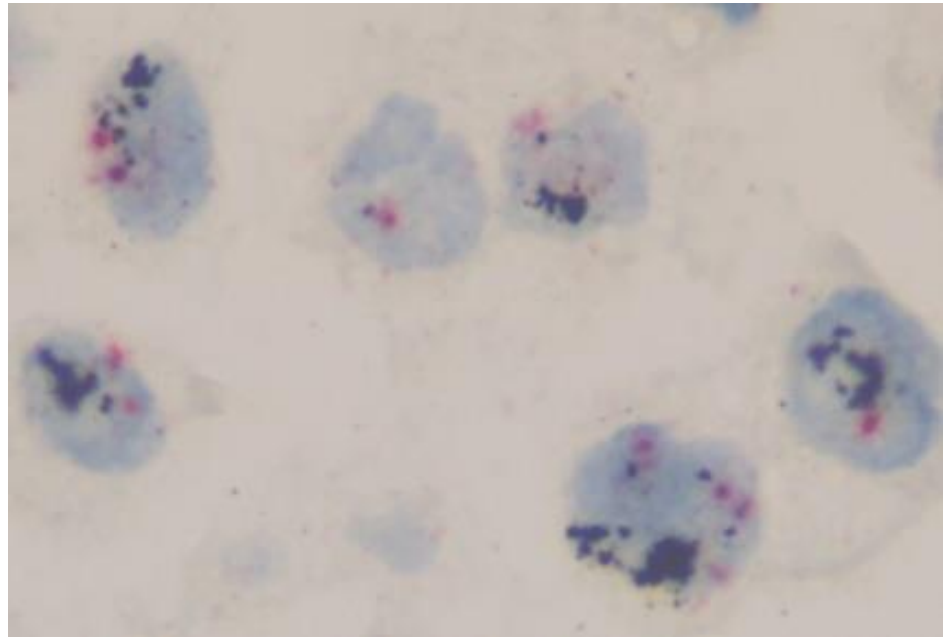
EXAMINATION



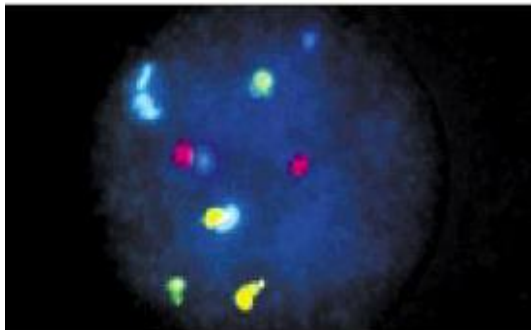
**DARK
FIELD**



**BRIGHT
FIELD**



MICROSCOPE FILTERS FOR FISH EXAMINATION

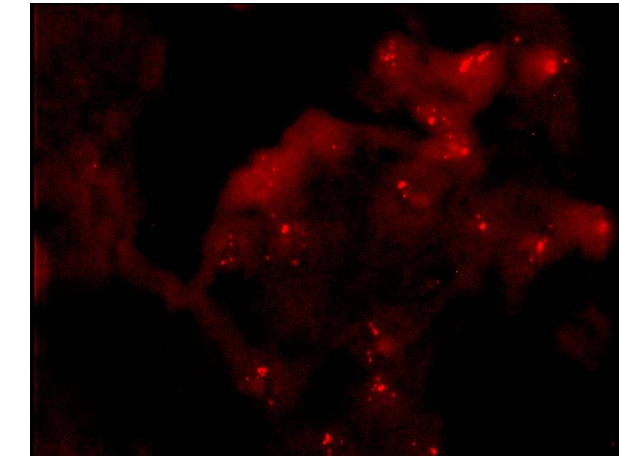
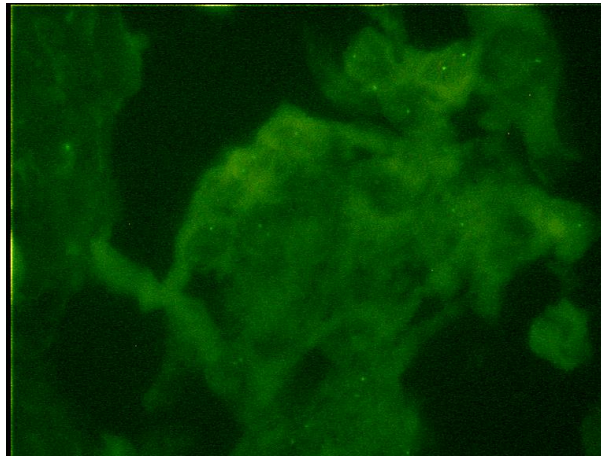
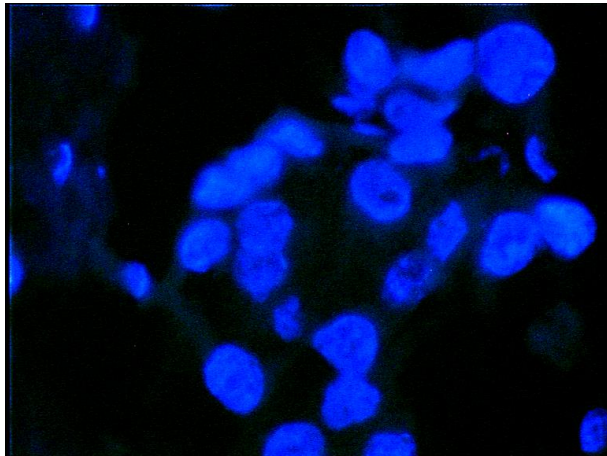
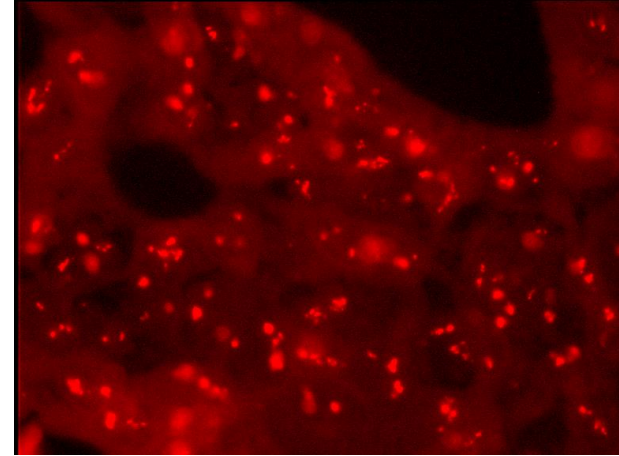
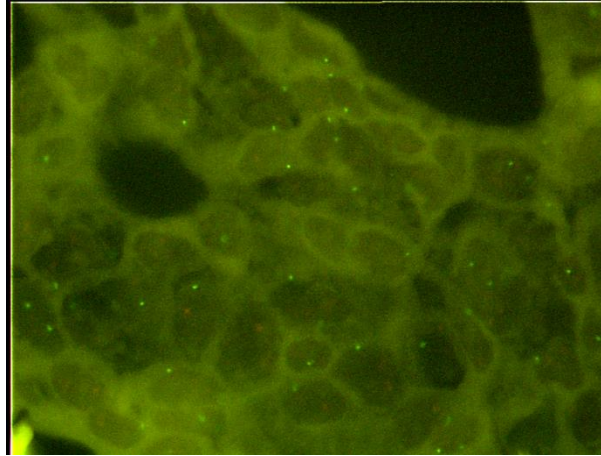
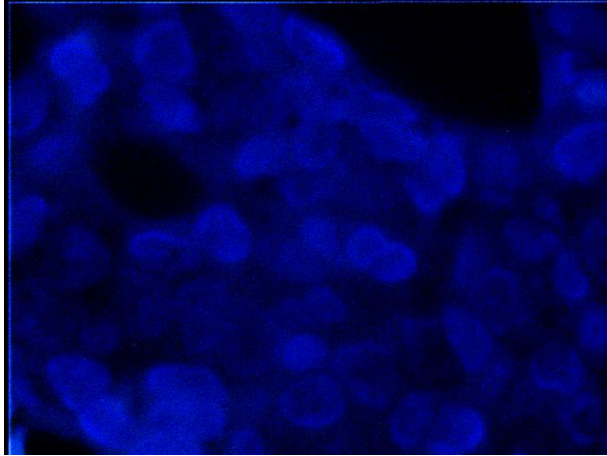


FITC
TR

**Excitation
wave length**
495nm
596nm

**Emission
wave length**
520nm
615nm

FISH (HER2)



DAPI

DAPI

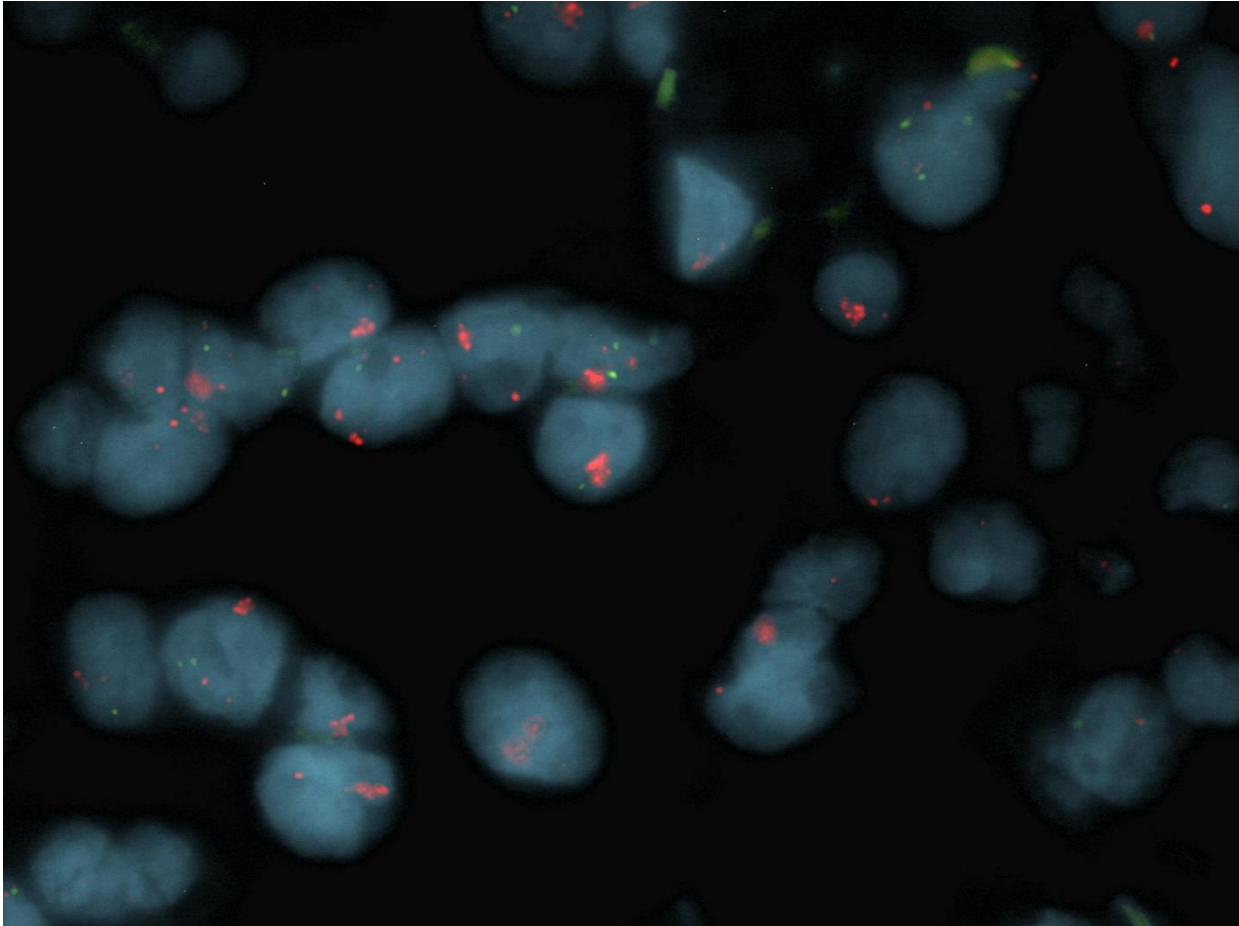
FITC

CEP17

TR

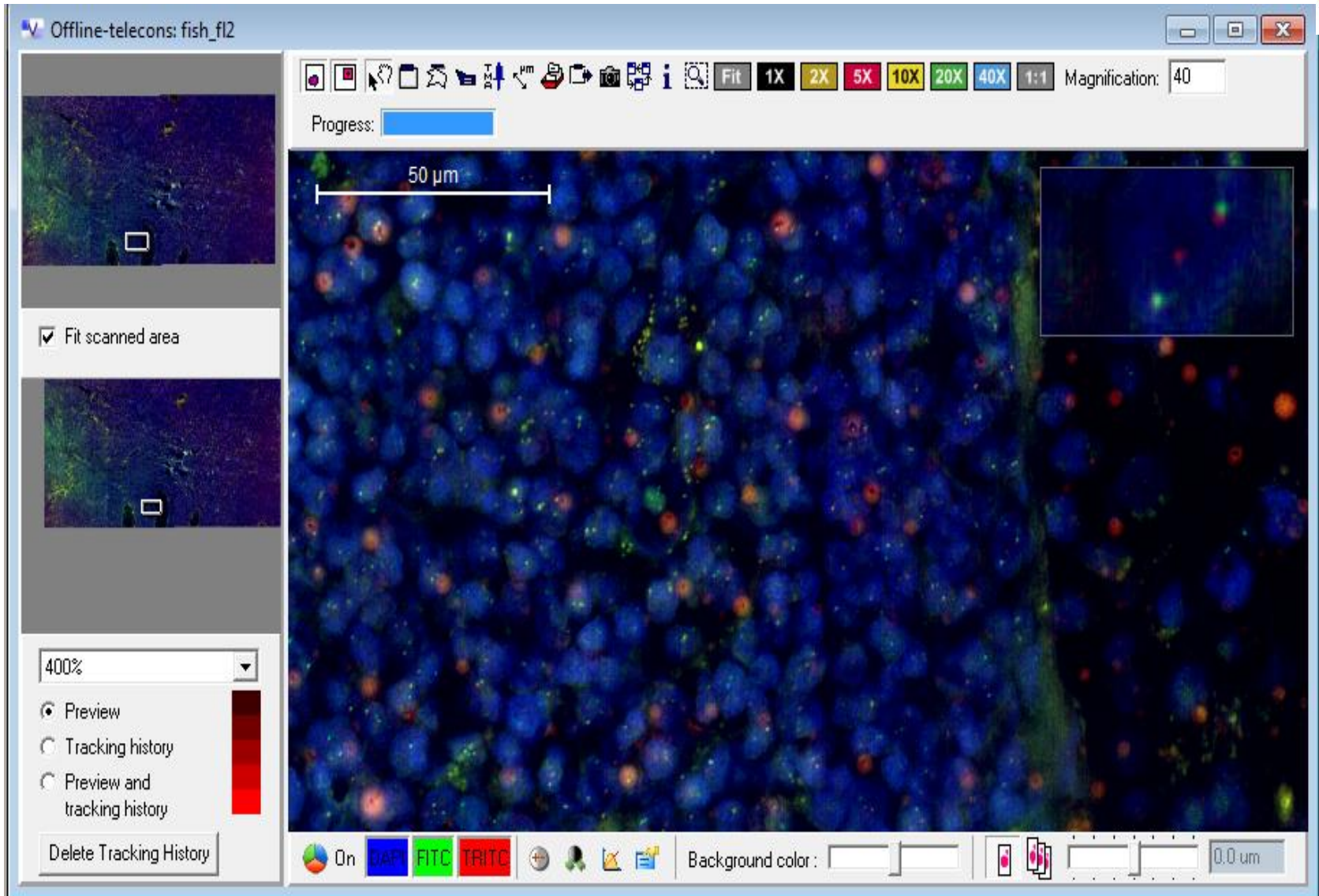
HER2

FISH (HER2)



DAPI / FITC/TR Triple filter image

EXAMINATION



EXAMINATION



Offline-telecons: fish_f12

Progress:

50 µm

Magnification: 40

Fit 1X 2X 5X 10X 20X 40X 1:1

400%

Preview
 Tracking history
 Preview and tracking history

Delete Tracking History

On DAPI FITC TRITC Background color: 0.0 um

**BEFORE EXAMINING FISH SLIDES
YOU SHOULD REMEMBER THE MORPHOLOGY
KEEP H&E SLIDES WITH YOU**

INTERPHASE FISH ON PARAFFIN TISSUE SECTIONS



Normal tissue

Tumour tissue

Intact
nuclei



Journal of Pathology 2002,198 (2):163-170

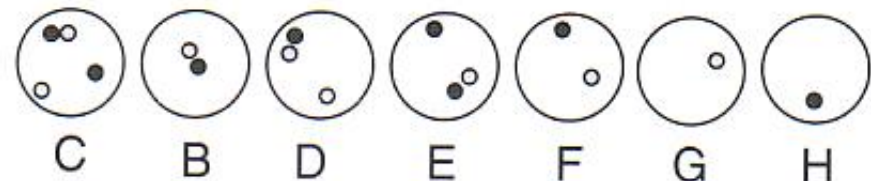
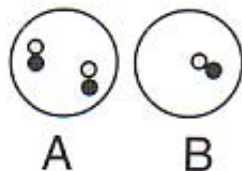
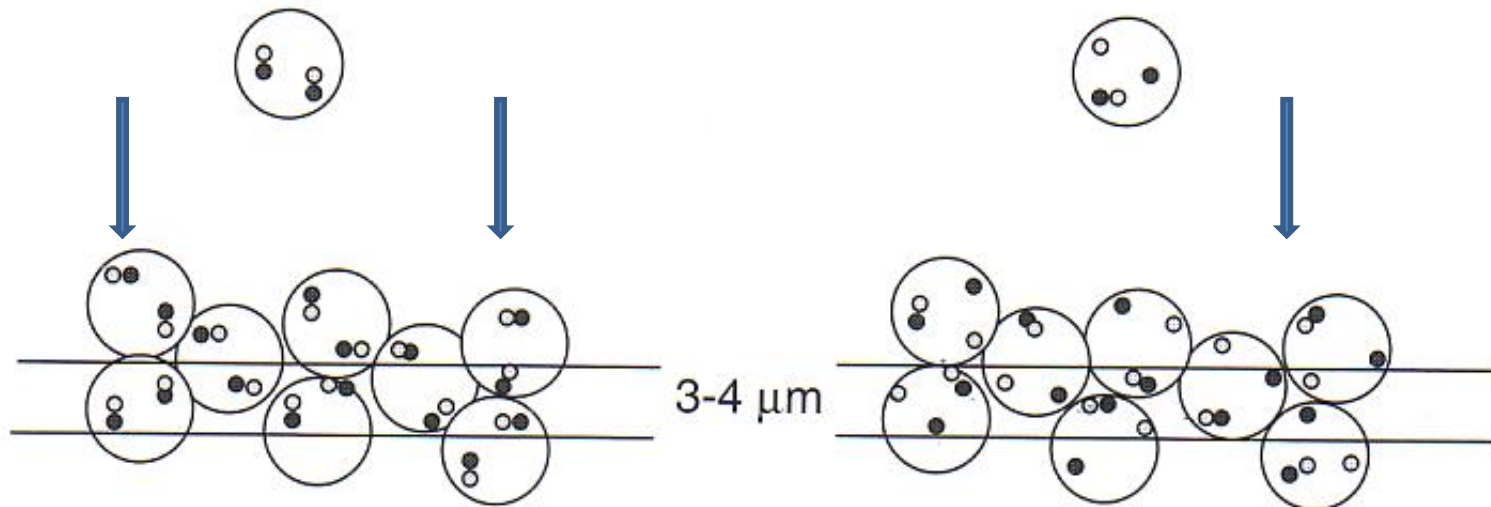
INTERPHASE FISH ON PARAFFIN TISSUE SECTIONS

Normal tissue

Tumour tissue

Intact nuclei

Tissue section



WHAT COULD BE SEEN BY USING ISH?



GENE COPY NUMBER INCREASE (AMPLIFICATIONS)

GENE OR CHROMOSOME LOSS (DELETIONS)

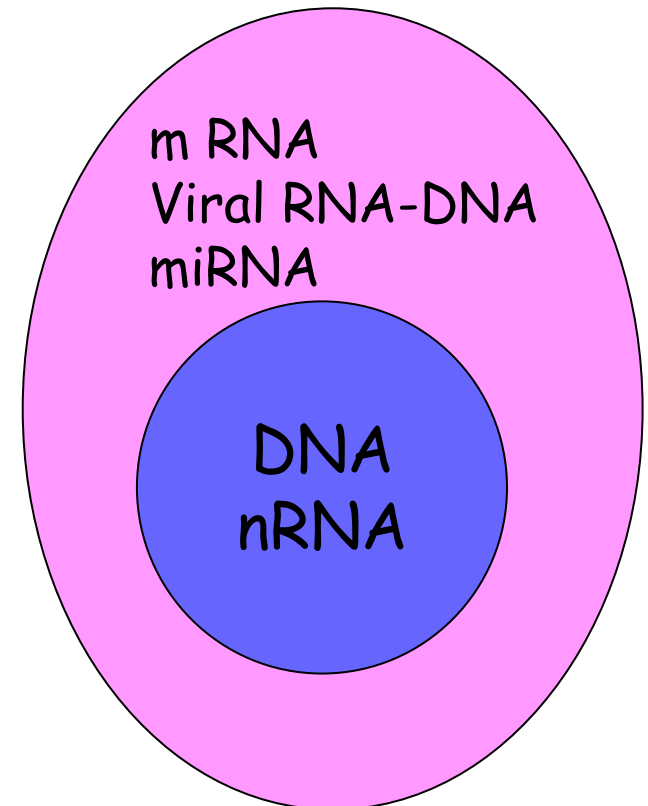
GENE REARRANGEMENTS (TRANSLOCATIONS)

INFECTIOUS AGENTS

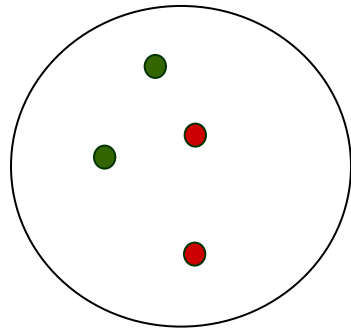
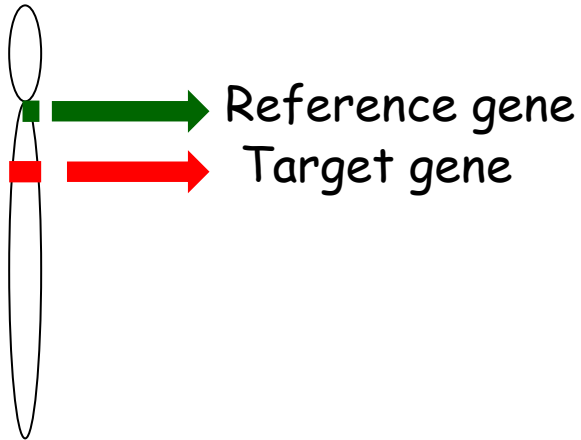
PROTEIN EXPRESSION (mRNA)

CHIMERISM (X, Y)

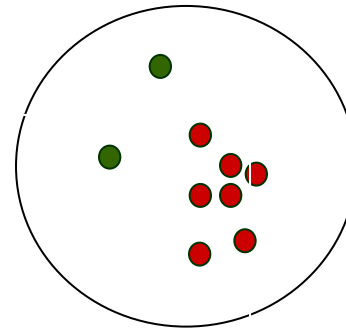
miRNA's



AMPLIFICATION SIGNALS

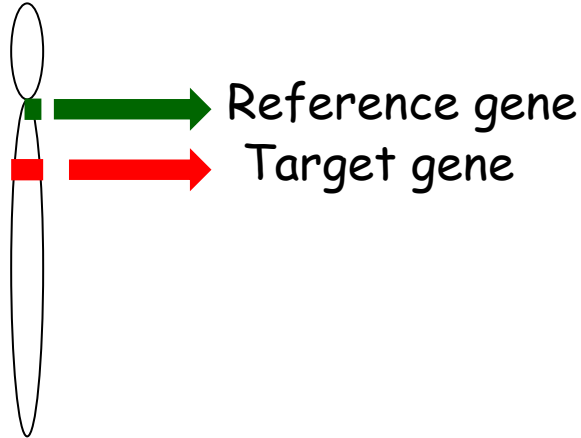


NORMAL

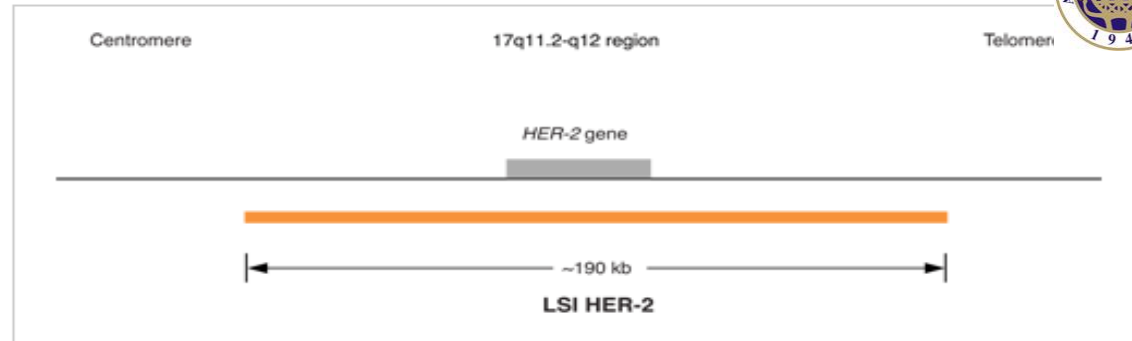


AMPLIFICATION

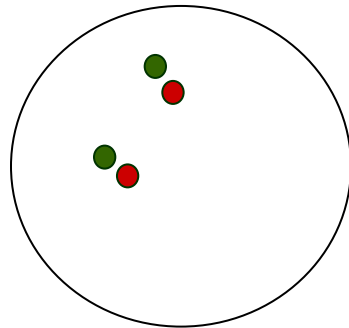
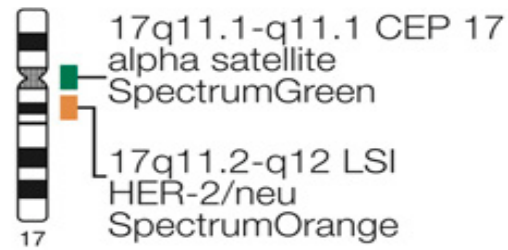
AMPLIFI



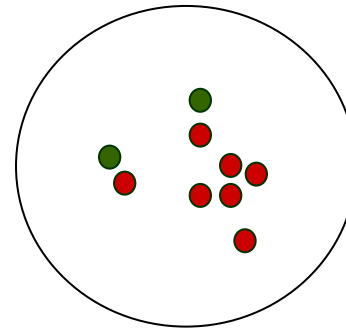
Probe Map



Ideogram

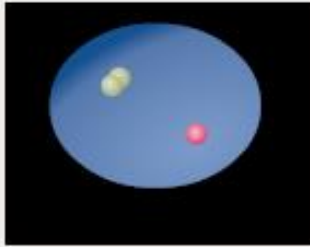
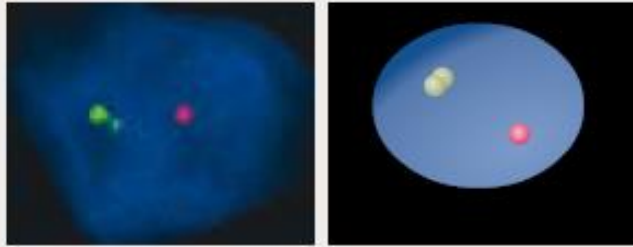


NORMAL



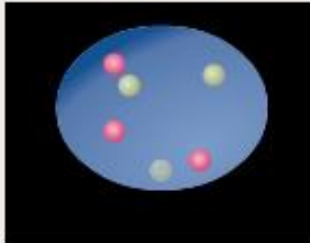
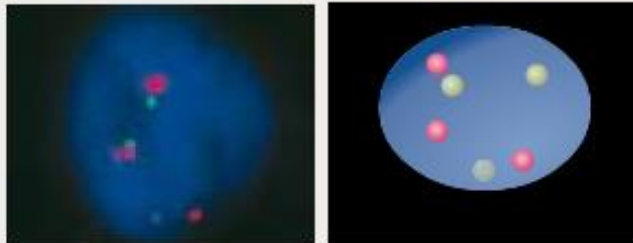
AMPLIFICATION

Counting Guide



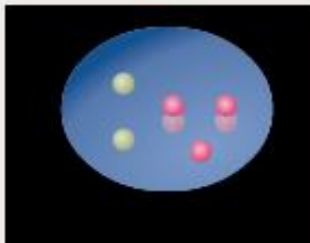
One green signal (split) indicates the presence of one copy of chromosome 17*. One red signal indicates the presence of one copy of the *HER2* gene.

The ratio of *HER2* to CEN-17 is $1/1 = 1$; non-amplified.



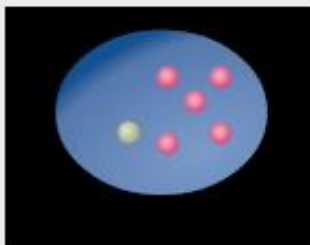
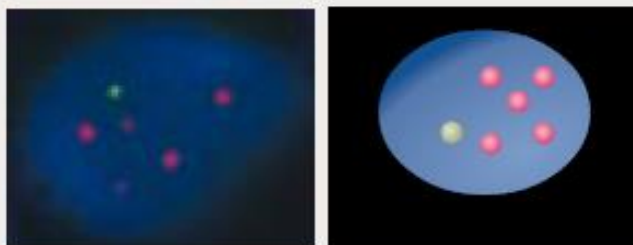
Three green signals (one out of focus) indicate the presence of three copies of chromosome 17. Three red signals indicate the presence of three copies of the *HER2* gene.

The ratio of *HER2* to CEN-17 is $3/3 = 1$; non-amplified.



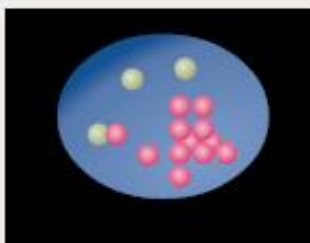
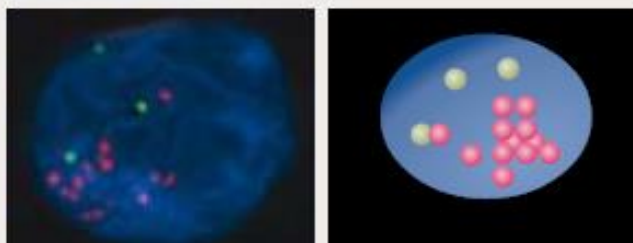
Two green signals indicate the presence of two copies of chromosome 17. Three red signals (two split signals) indicate the presence of three copies of the *HER2* gene*.

The ratio of *HER2* to CEN-17 is $3/2 = 1.5$; non-amplified.



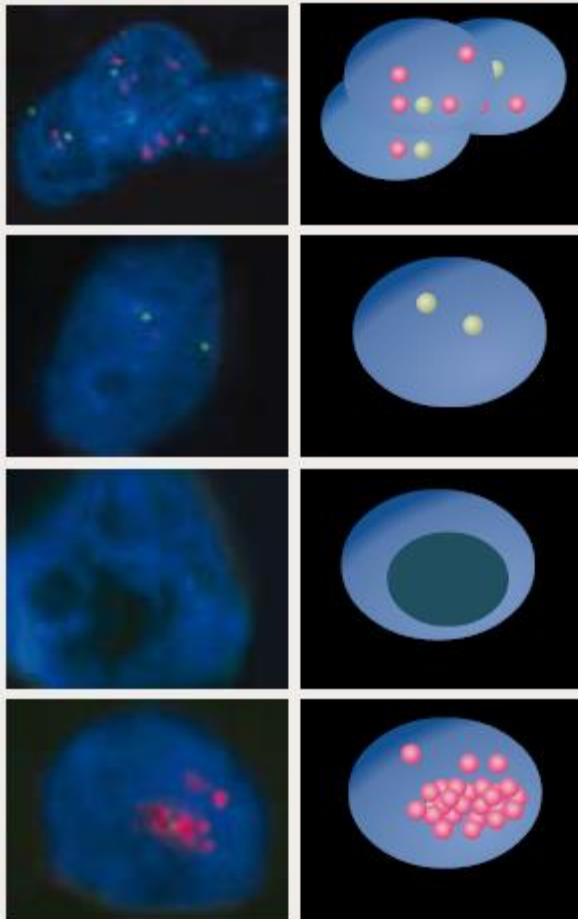
One green signal indicates the presence of one copy of chromosome 17. Five red signals indicate the presence of five copies of the *HER2* gene.

The ratio of *HER2* to CEN-17 is $5/1 = 5$; amplified.



Three green signals indicate the presence of three copies of chromosome 17. Approximately 12 red signals indicate the presence of 12 copies of the *HER2* gene (cluster estimation).

The ratio of *HER2* to CEN-17 is $12/3 = 4$; amplified.



Do not score (nuclei are overlapping, not all areas of nuclei are visible).

Do not score nuclei with signals of only one color (two green signals).

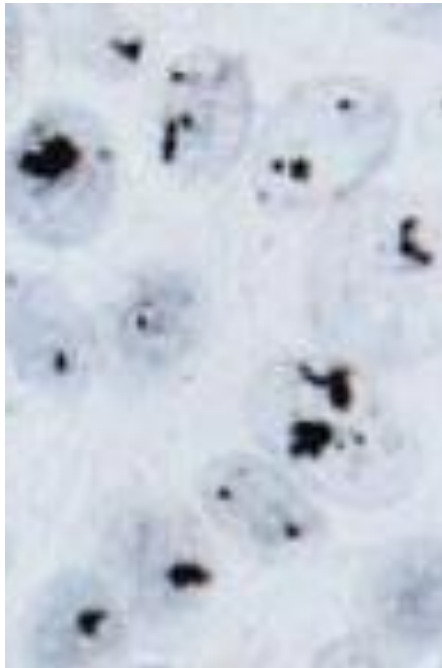
Do not score (overdigested nuclei).

Cluster of red signals hiding green signals. Check the green signals with a specific FITC filter, or do not score.

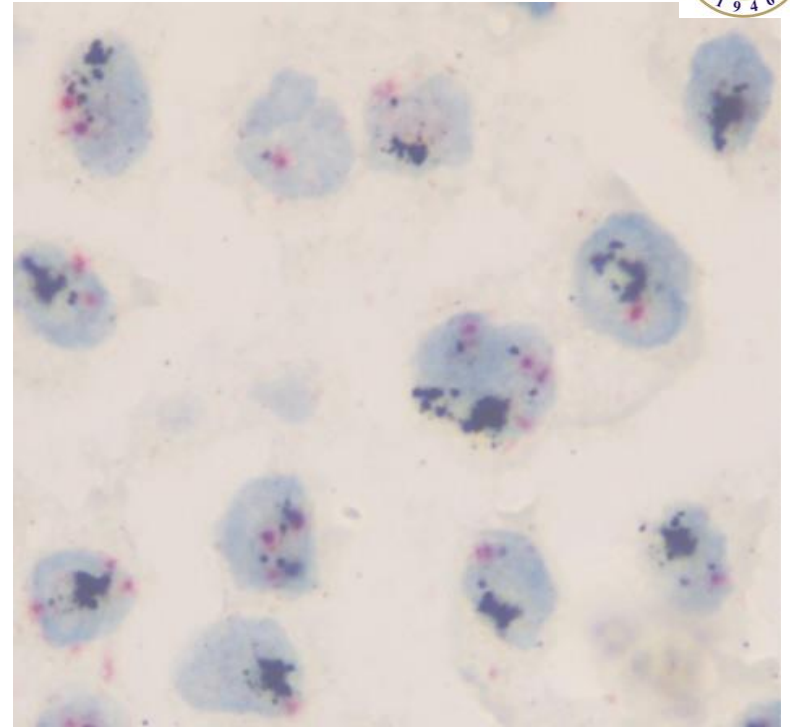
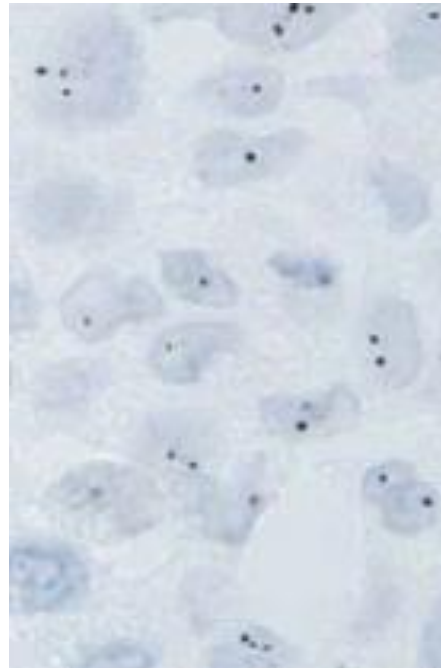
**Two signals of the same size, separated by a distance equal to or less than the diameter of one signal, are counted as one signal.*

SISH

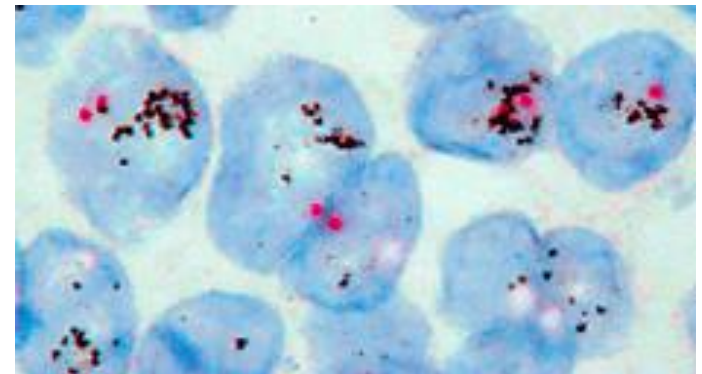
Her 2 neu



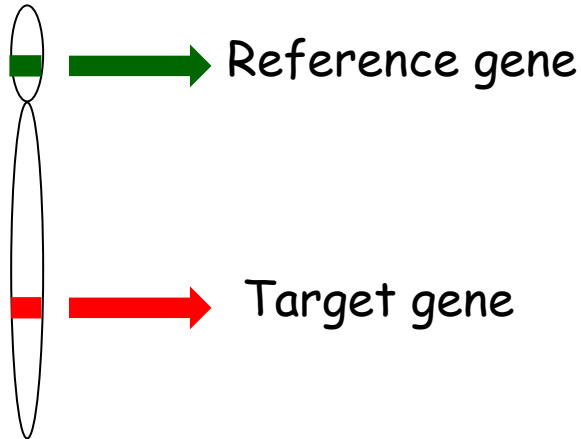
Chr 17



Her 2 neu / Chr 17



DELETION SIGNALS



Ideograms

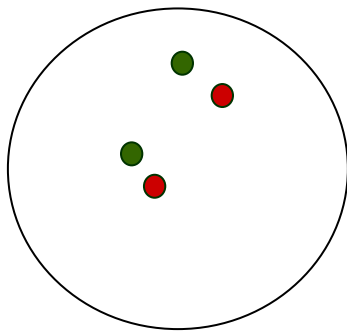
LSI 1p36
SpectrumOrange



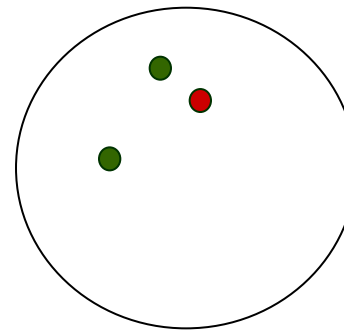
LSI 1q25
SpectrumGreen

LSI 19p13
SpectrumGreen

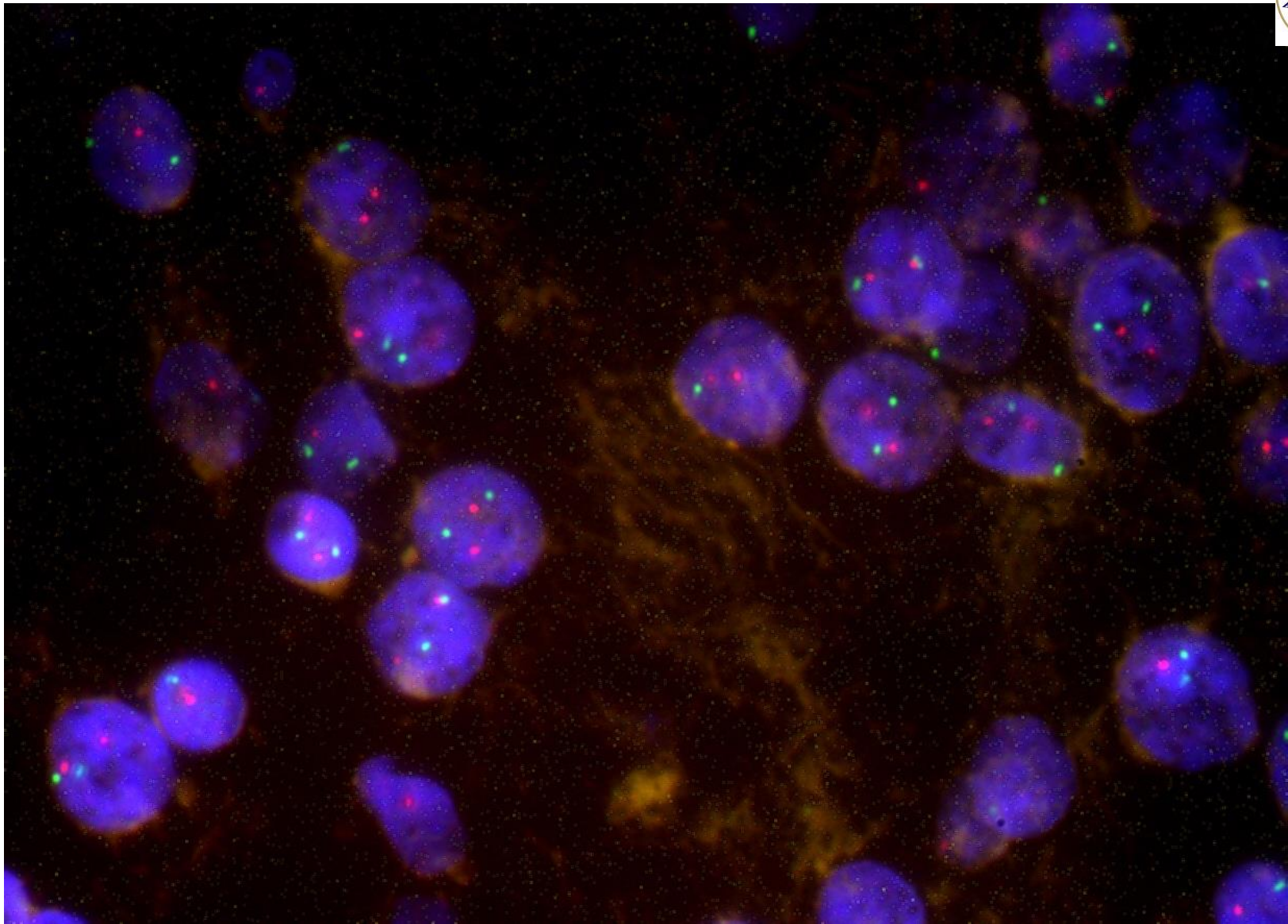
LSI 19q13
SpectrumOrange



NORMAL



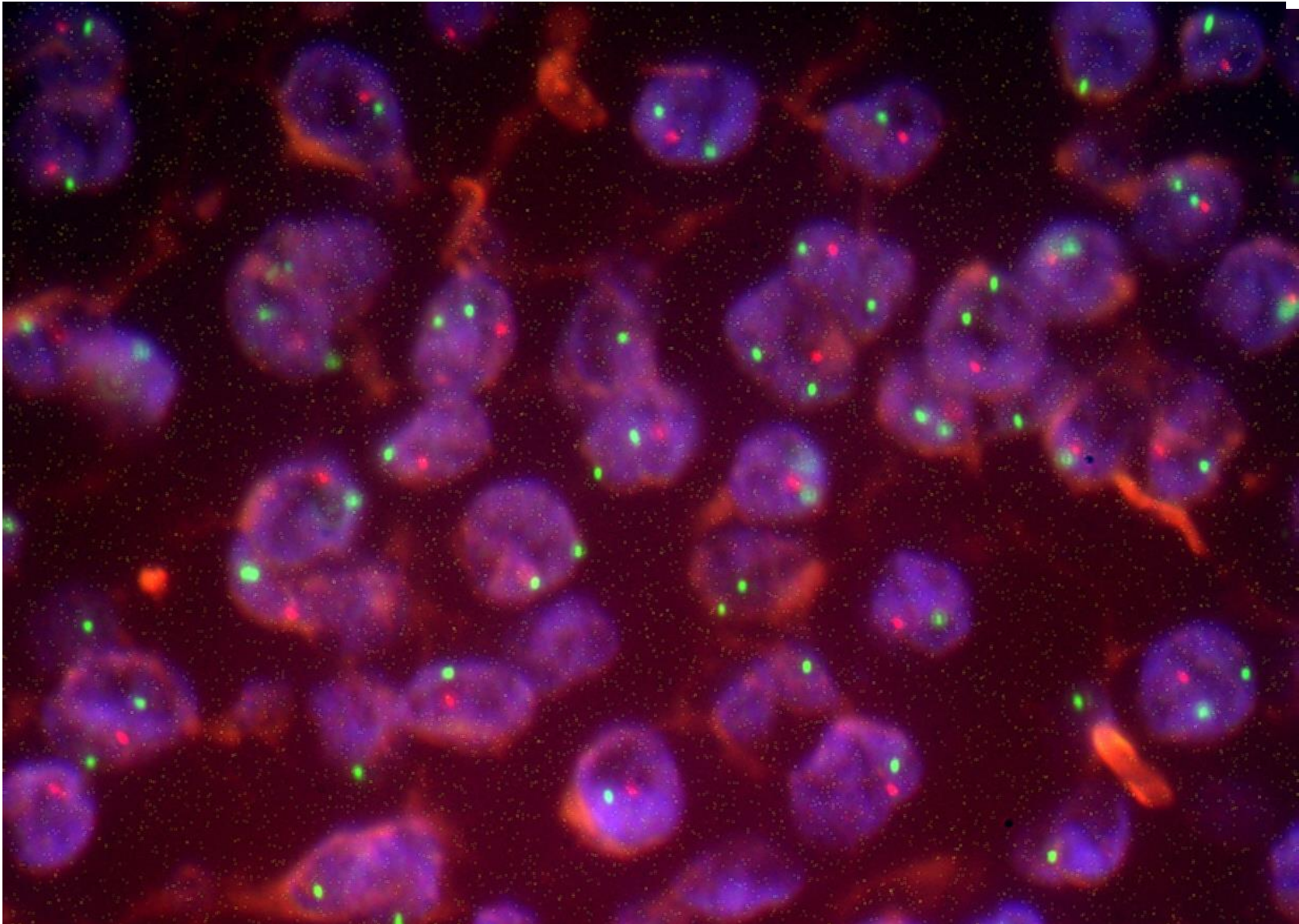
DELETED



Chr 1P

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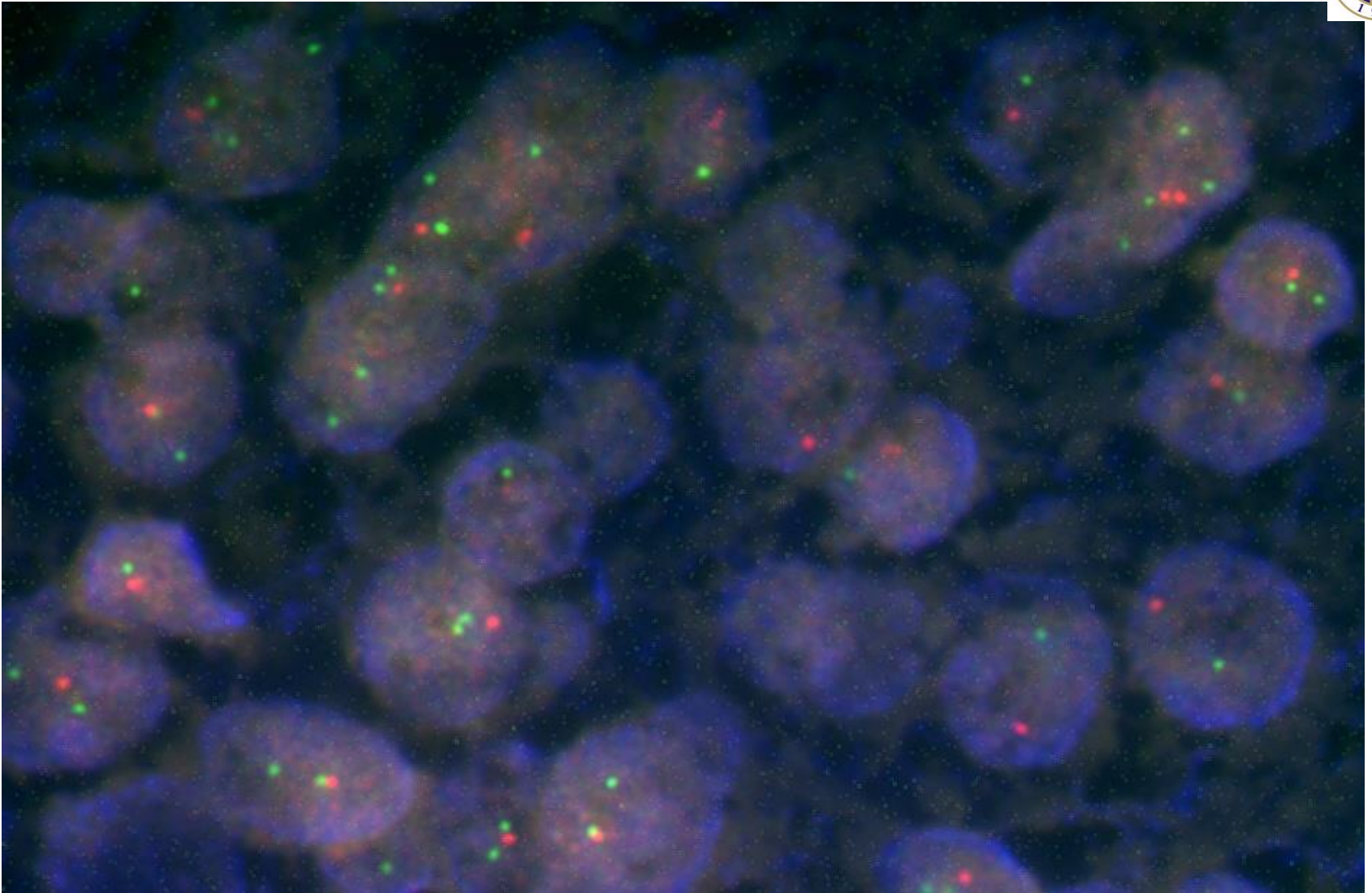
Normal signals



Chr 1P

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Deletion Signals



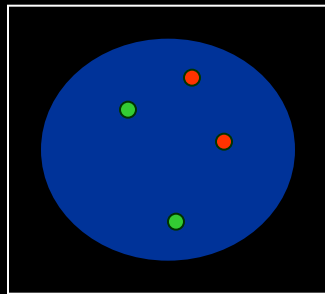
Chr 19q

Deletion signals



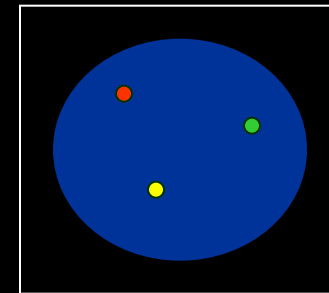
TRANSLOCATION (REARRANGEMENT) SIGNALS

FUSION PROBE



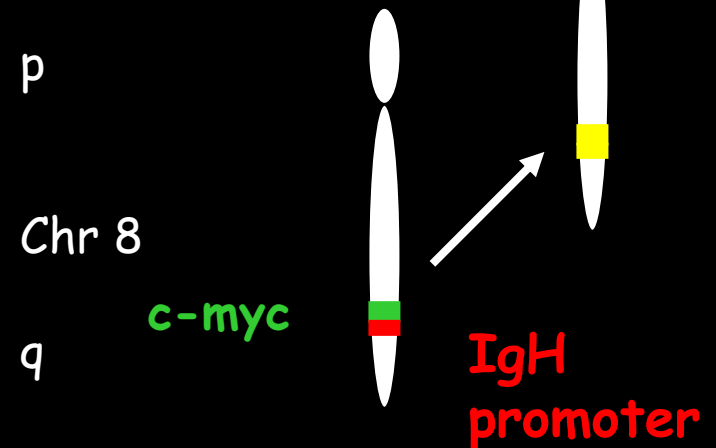
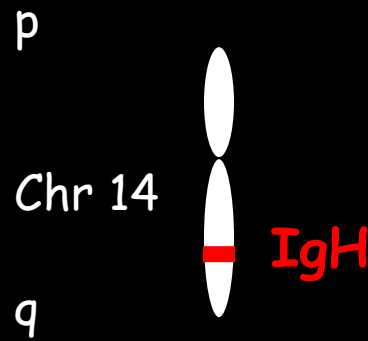
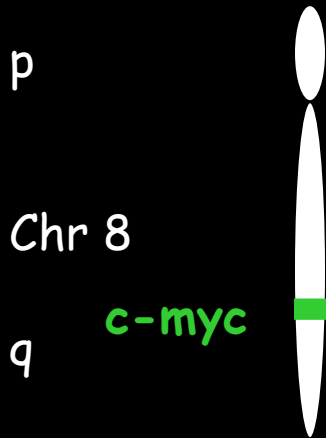
c-myc; IgH

t(8;14)(q24;q32)



Normal

Abnormal



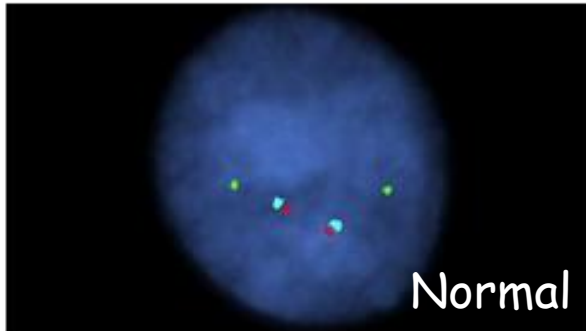
Pitfalls:

Does not recognize translocations between alternative partner genes
 %5-10 false positive fusion signals due to the cellular superposition on cell smears
 False positive signals are increase on tissue sections.

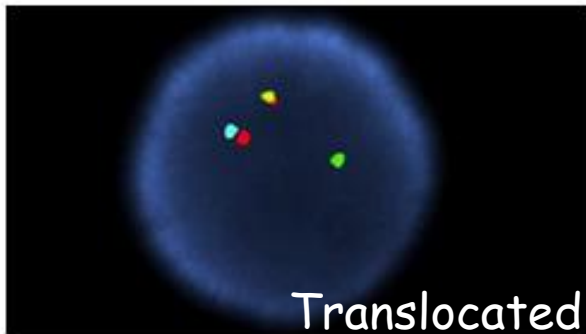
Triple signal fusion probes



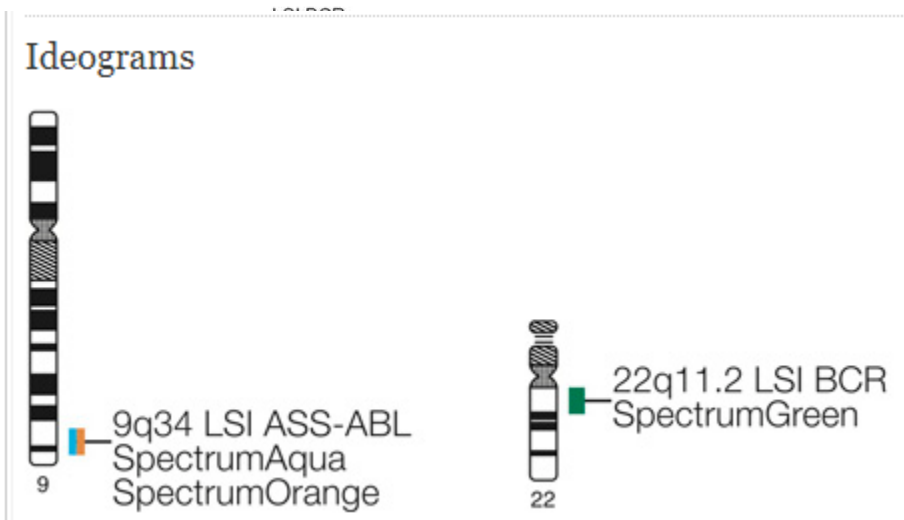
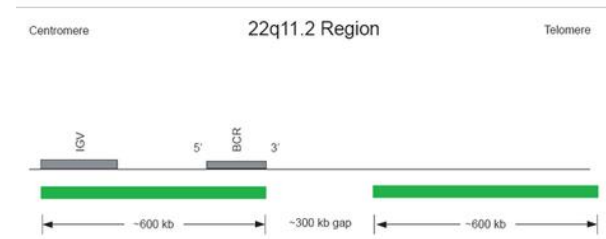
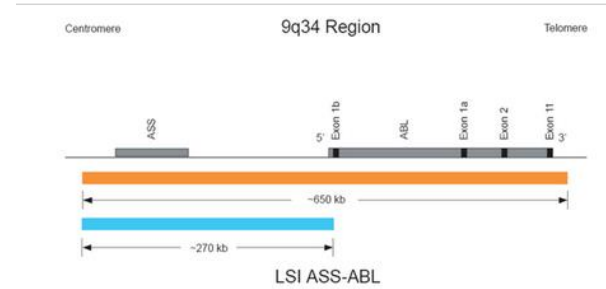
Bcr-Abl



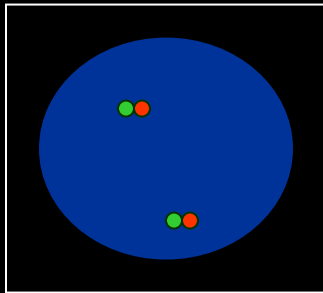
Nucleus showing the two aqua/orange and two green signal pattern.



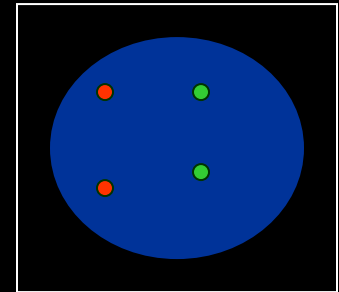
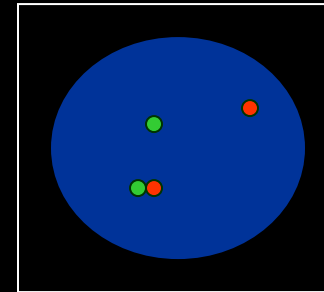
Nucleus showing the one aqua/orange, one green, and one orange/green fusion (yellow) signal pattern.



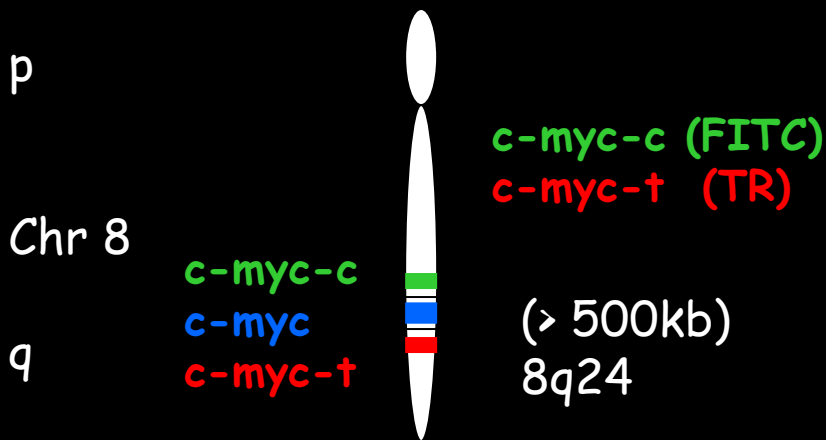
SPLIT PROBE (SEGREGATION ASSAY)



Normal



Abnormal



Probe sensitivity spectrum

c-myc; IgH	t(8;14)(q24;q32)
c-myc; Igk	t(2;8)(p12;q24)
c-myc; Igl	t(8;22)(q24;q11)
c-myc; TCR-a	t(8;14)(q24;q11)

Advantages:

All the alternative translocations independent from the partner genes can be detected
Decreases the false positive signals due to the superposition

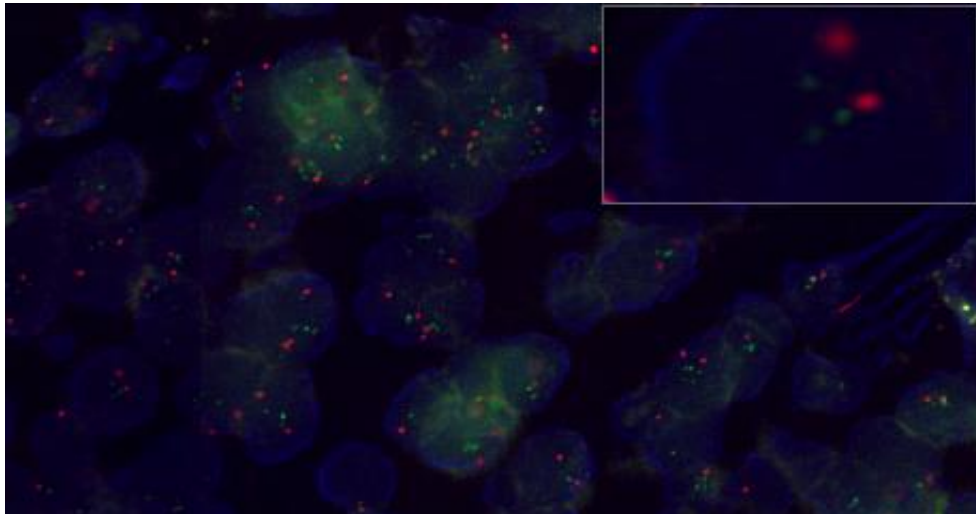
Split signal probes



2p23 LSI ALK
SpectrumOrange
SpectrumGreen



LSI ALK Dual Color, Break Apart
Rearrangement Probe



EML4-ALK testing in non-small cell carcinomas of the lung: a review with recommendations

Erik Thunnissen · Lukas Bubendorf · Manfred Dietel ·
Göran Elmberger · Keith Kerr · Fernando Lopez-Rios ·
Holger Moch · Włodzimierz Olszewski ·
Patrick Pauwels · Frédérique Penault-Llorca ·
Giulio Rossi

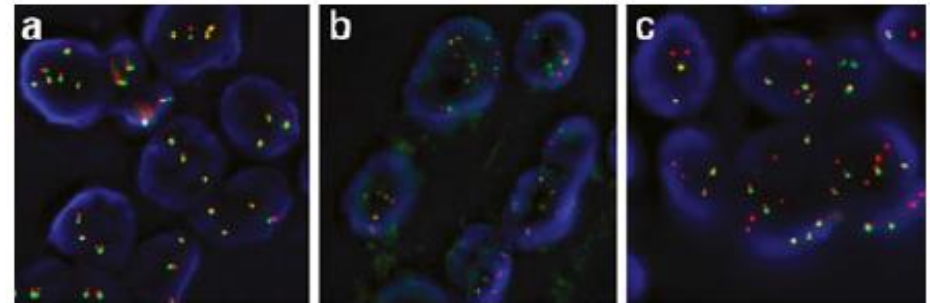
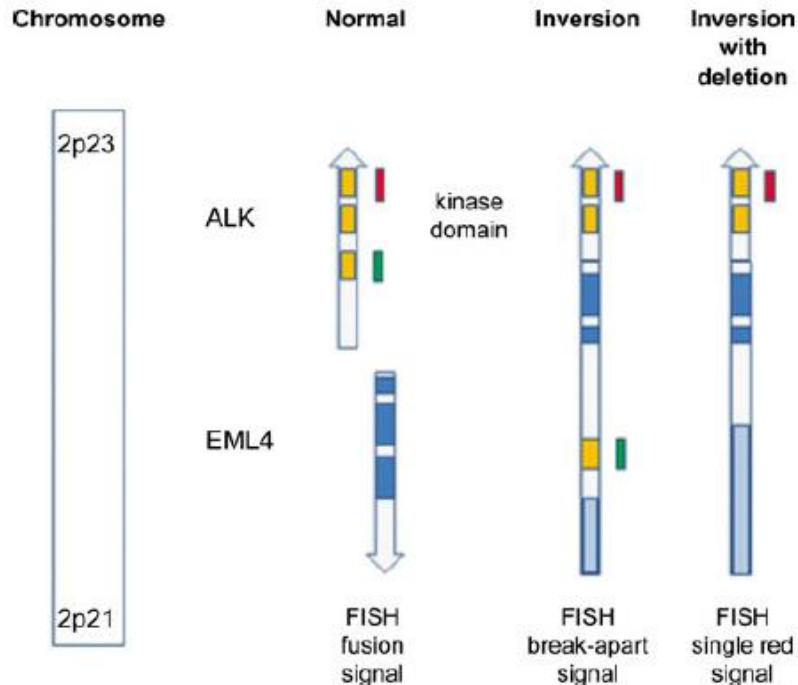
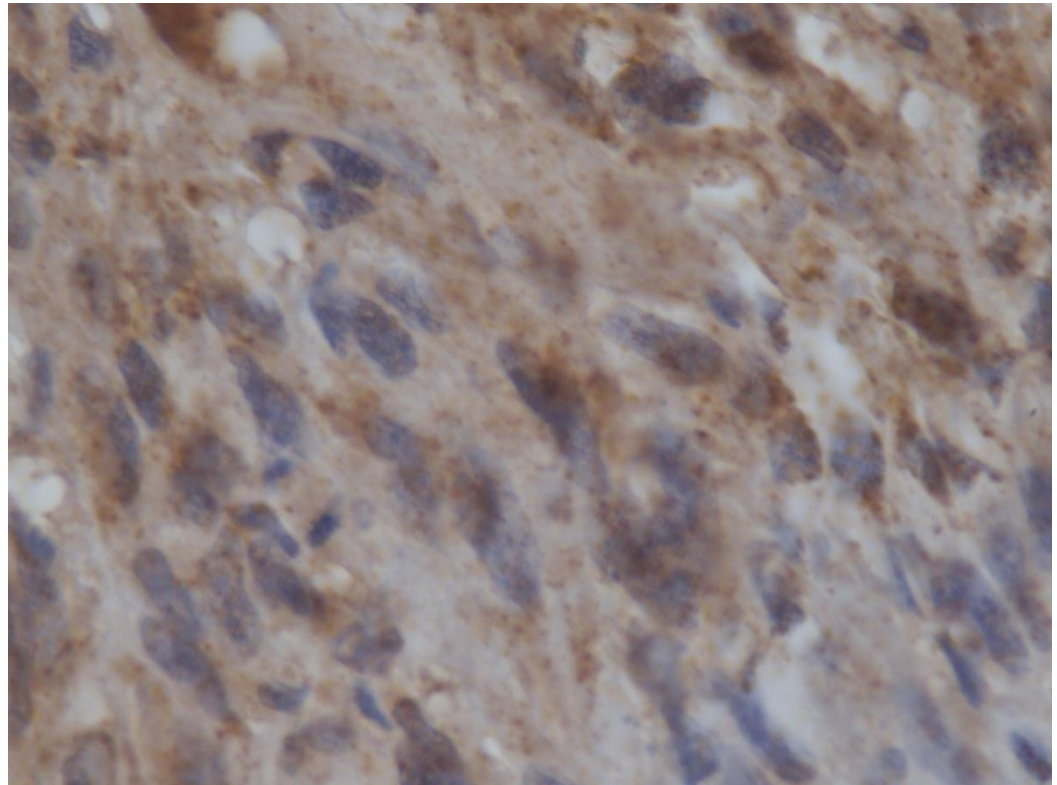
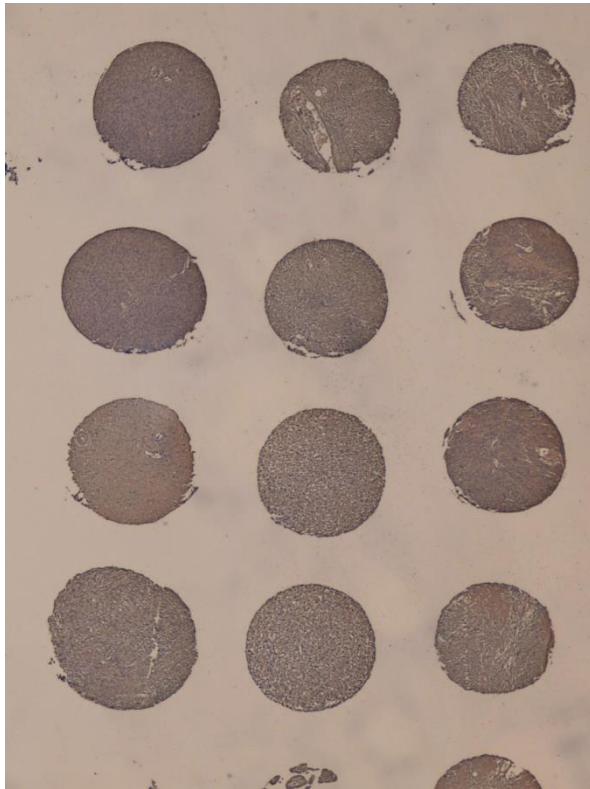


Fig. 2 Representative examples of ALK FISH findings in three pulmonary adenocarcinomas (Vysis ALK Break Apart FISH probe). All three carcinomas show increased ALK copy number. **a** Normal signals, no rearrangement. Note that some of the signals are fused and produce a yellow signal, while others have *green* and *red* signals in close proximity. **b** One or two break apart signals per nucleus, indicative of inversion. **c** Single red signals, indicative of inversion and deletion. Note that the cancer cells in **b** and **c** contain both rearranged and normal ALK signals. Cells are considered ALK FISH positive when there is: (1) ≥ 1 set of red and green signals that are ≥ 2 signal diameters apart, or (2) a single red signal without a corresponding green signal in addition to fused (normal) signals. A sample is considered negative if < 5 cells ($< 10\%$) are positive and positive if > 25 cells ($> 50\%$) are positive. A sample is considered equivocal if 5–25 cells (10–50%) are positive

RNA ISH for PDGFRA expression



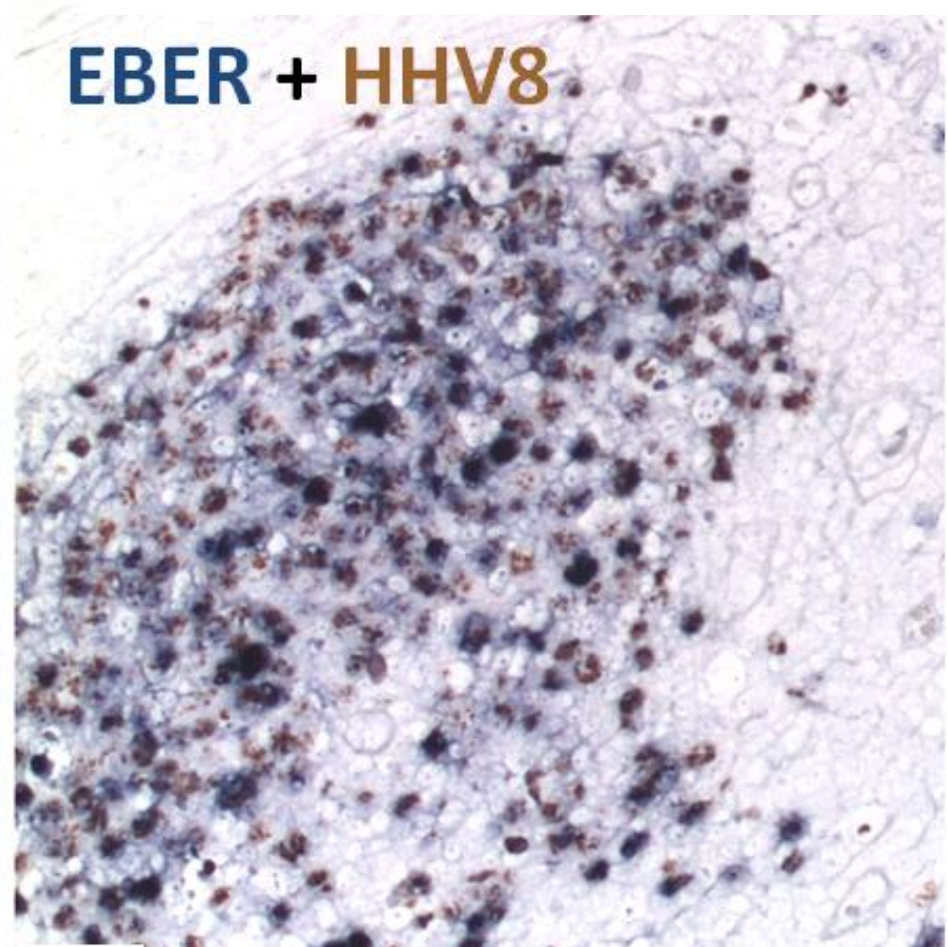
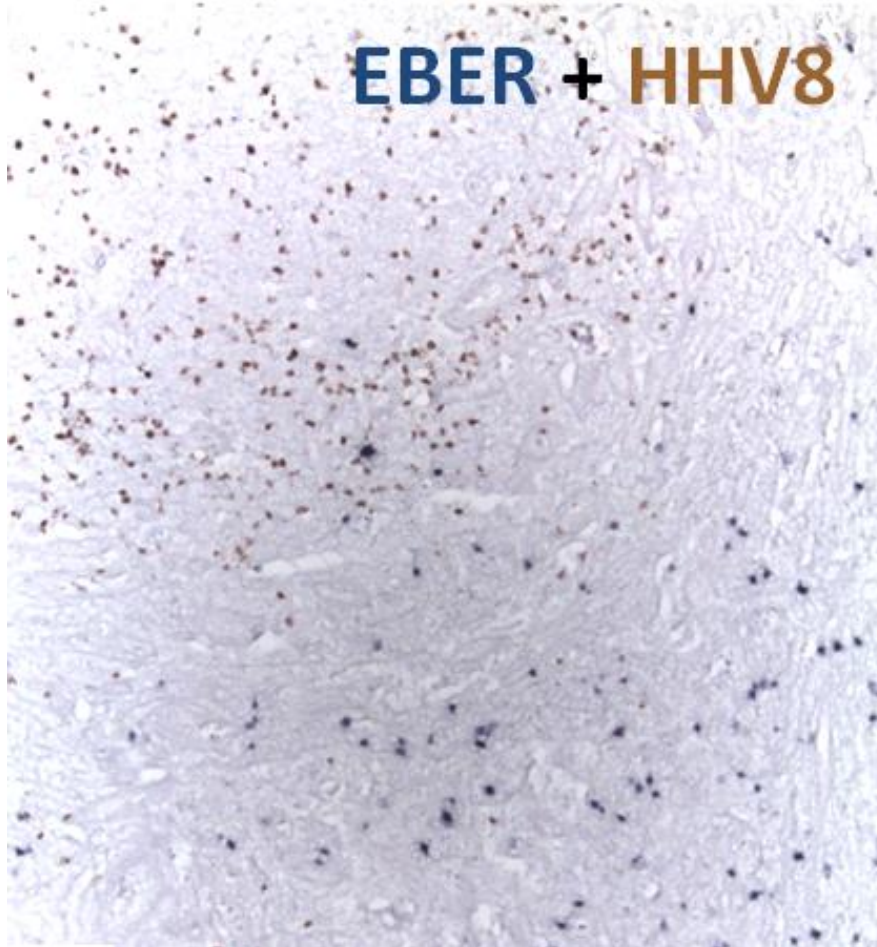


Early lesions in lymphoid neoplasia

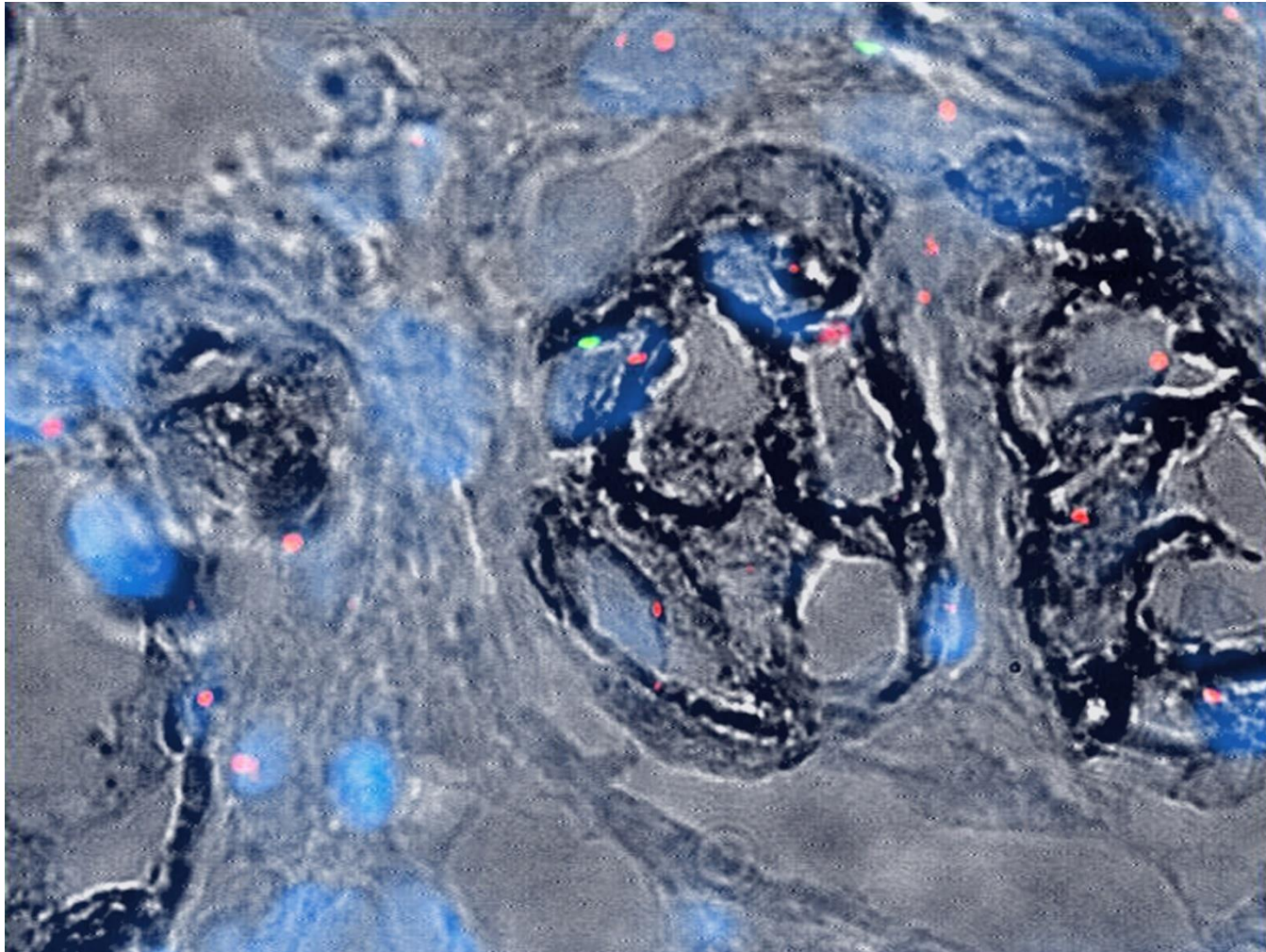
Conclusions based on the Workshop of the XV. Meeting of the European Association of Hematopathology and the Society of Hematopathology in Uppsala, Sweden

Falko Fend · José Cabecadas · Philippe Gaulard ·
Elaine S. Jaffe · Philip Kluin · Isinsu Kuzu ·
LoAnn Peterson · Andrew Wotherspoon ·
Christer Sundström

DOUBLE LABELLING IMUNOHISTOCHEMISTRY + ISH



DOUBLE LABELLING IMUNOHISTOCHEMISTRY + FISH





17-22 October 2014

Istanbul

Hilton Convention Center

17th Meeting of the European Association for Haematopathology



www.eahp2014.org

TOPIC

"Redefining the spectrum of small B-cell lymphomas in light of current technology".

GOAL

To integrate modern biotechnological developments and morphological pathology to better understand small B cell lymphomas.

EUROPEAN ASSOCIATION
FOR HAEMATOPATHOLOGY



Society for Hematopathology



FEDERATION OF
PATHOLOGY
SOCIETIES OF
TURKEY



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HEMATOLOGY



THANK YOU FOR YOUR ATTENTION

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