

# Zyto Light SPEC ERBB2/CEN 17 Dual Color Probe Kit

**REF** Z-2020-5

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**REF** Z-2020-20

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For the qualitative detection of human ERBB2 gene amplifications and alpha satellites of chromosome 17 by fluorescence *in situ* hybridization (FISH)



In vitro diagnostic medical device according to EU directive 98/79/EC

# 1. Intended use

The <u>ZytoLight SPEC ERBB2/CEN 17 Dual Color Probe Kit</u> is intended to be used for the qualitative detection of human ERBB2 gene amplifications as well as the detection of chromosome 17 alpha satellites in formalin-fixed, paraffin-embedded specimens such as human breast cancer or gastric cancer tissues by fluorescence *in situ* hybridization (FISH).

Interpretation of the results must be made within the context of the patient's clinical history with respect to further clinical and pathologic data of the patient by a qualified pathologist.

#### 2. Clinical relevance

The ERBB2 gene (a.k.a. HER2 and NEU) is located in the chromosomal region 17q12 and encodes a 185-190 kDa transmembrane glycoprotein, p185, acting as a cellular growth factor receptor. The p185 protein belongs to the EGFR (epidermal growth factor receptor) subgroup of the RTK (receptor tyrosine kinase) superfamily also including EGFR (ERBB1), ERBB3 (HER3), and ERBB4 (HER4). Amplification of the proto-oncogene ERBB2, observed in approximately 20% of all breast cancer samples, has been correlated with a poor prognosis of the disease. Similar results have been obtained for a variety of other malignant neoplasms, e.g., ovarian cancer, stomach cancer, and carcinomas of the salivary gland.

# 3. Test principle

The fluorescence *in situ* hybridization (FISH) technique allows for the detection and visualization of specific nucleic acid sequences in cell preparations. Fluorescently-labeled DNA fragments, so called FISH probes, and their complementary target DNA strands in the preparations are codenatured and subsequently allowed to anneal during hybridization. Afterwards, unspecific and unbound probe fragments are removed by stringency washing steps. After counterstaining the DNA with DAPI, hybridized probe fragments are visualized using a fluorescence microscope equipped with excitation and emission filters specific for the fluorochromes with which the FISH probe fragments have been directly labeled.

# 4. Reagents provided

The <u>ZytoLight SPEC ERBB2/CEN 17 Dual Color Probe Kit</u> is available in two sizes and is composed of:

C	Component	Quantity		
Code		5 \\	20	Container
PT1	Heat Pretreatment Solution Citric	150 ml	500 ml	Screw-cap bottle (large)
ES1	Pepsin Solution	1 ml	4 ml	Dropper bottle, white cap
WB1	Wash Buffer SSC	210 ml	560 ml	Screw-cap bottle (large)
PL8	ZytoLight SPEC ERBB2/CEN 17 Dual Color Probe	0.05 ml	0.2 ml	Reaction vessel, red lid
WB2	25x Wash Buffer A	50 ml	2x50 ml	Screw-cap bottle (medium)
MT7	DAPI/DuraTect- Solution	0.2 ml	0.8 ml	Reaction vessel, blue lid
	Instructions for use	1	1	

<u>Z-2020-5 (5 tests)</u>: Components **ES1**, **PL8**, and **MT7** are sufficient for 5 reactions. Component **WB2** is sufficient for 5x 3 staining jars of 70 ml each. Component **PT1** is sufficient for 2 staining jars of 70 ml each. Component **WB1** is sufficient for 3 staining jars of 70 ml each.

<u>Z-2020-20 (20 tests)</u>: Components **ES1**, **PL8**, and **MT7** are sufficient for 20 reactions. Component **WB2** is sufficient for 11x 3 staining jars of 70 ml each. Component **PT1** is sufficient for 7 staining jars of 70 ml each. Component **WB1** is sufficient for 8 staining jars of 70 ml each.

The ZytoLight SPEC ERBB2/CEN 17 Dual Color Probe (PL8) is composed of:

- ZyGreen (excitation 503 nm/emission 528 nm) labeled polynucleotides ( $\sim$ 10.0 ng/ $\mu$ l), which target sequences mapping in 17q12-q21.1\* (chr17:37,572,531-38,181,308) harboring the ERBB2 gene region (see Fig. 1).
- ZyOrange (excitation 547 nm/emission 572 nm) labeled polynucleotides (~1.5 ng/μl), which target sequences mapping in 17p11.1-q11.1 specific for the alpha satellite centromeric region D17Z1 of chromosome 17.
- Formamide based hybridization buffer

<sup>\*</sup>according to Human Genome Assembly GRCh37/hg19

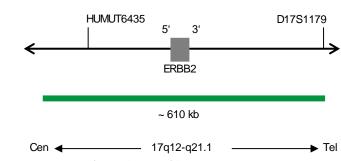


Fig. 1: SPEC ERBB2 Probe map (not to scale)

# 5. Materials required but not provided

- Positive and negative control specimens
- Microscope slides, positively charged
- Water bath (37°C, 98°C)
- Hybridizer or hot plate
- Hybridizer or humidity chamber in hybridization oven
- Adjustable pipettes (10 μl, 25 μl)
- Staining jars or baths
- Timer
- Calibrated thermometer
- Ethanol or reagent alcohol
- Xvlene
- Deionized or distilled water
- Coverslips (22 mm x 22 mm, 24 mm x 60 mm)
- Rubber cement, e.g., <u>Fixogum Rubber Cement</u> (Prod. No. E-4005-50/-125) or similar
- Adequately maintained fluorescence microscope (400-1000x)

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- Immersion oil approved for fluorescence microscopy
- Appropriate filter sets

## 6. Storage and handling

The components of the kit must be stored at 2-8°C. Additionally, the <u>DAPI/DuraTect-Solution</u> (MT7) and the probe solution (PL8) must be stored protected from light. Return components to storage conditions immediately after use. If these storage conditions are followed, the kit will function, without loss of performance, at least until the expiry date printed on the label. Do not use reagents beyond expiry date indicated on the label.

# 7. Warnings and precautions

- Read the instruction for use prior to use!
- Do not use the reagents after the expiry date has been reached!
- This product contains substances (in low concentrations and volumes) that are harmful to health and potentially infectious. Avoid any direct contact with the reagents. Take appropriate protective measures (use disposable gloves, protective glasses, and lab garments)!
- If reagents come into contact with skin, rinse skin immediately with copious quantities of water!
- A material safety data sheet is available on request for the professional user.
- Do not reuse reagents.
- Avoid cross-contamination of samples as this may lead to erroneous results.
- The specimens must not be allowed to dry during the hybridization and washing steps!
- The probe (PL8) and the <u>DAPI/DuraTect-Solution</u> (MT7) should not be exposed to light, especially strong light, for a longer period of time, i.e., all steps should be accomplished, where possible, in the dark and/or using lightproof containers!

# Hazard and precautionary statements for PL8:

The hazard determining component is Formamide.



## Danger

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H351	Suspected of causing cancer.
H360FD	May damage fertility. May damage the unborn child.
H373	May cause damage to organs through prolonged or repeated exposure.
P201	Obtain special instructions before use.
P202	Do not handle until all safety precautions have been read and understood.
P260	Do not breathe dust/fume/gas/mist/vapours/spray.
P280	Wear protective gloves/protective clothing/eye protection/face protection.
P308+P313	IF exposed or concerned: Get medical advice/attention.
P405	Store locked up.

## Hazard and precautionary statements for PT1, WB1, and WB2:

The hazard determining component is a mixture of: 5-chloro-2-methyl-4-isothiazolin-3-one [EC no. 247-500-7] and 2-methyl-2H-isothiazol-3-one [EC no. 220-239-6] (3:1).

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#### Warning

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H317	May cause an allergic skin reaction.
P261	Avoid breathing dust/fume/gas/mist/vapours/spray
P272	Contaminated work clothing should not be allowed out of the workplace.
P280	Wear protective gloves/protective clothing/eye protection/face protection.
P302+P352	IF ON SKIN: Wash with plenty of water.
P333+P313	IF skin irritation or rash occurs: Get medical advice/attention.
P362+P364	Take off contaminated clothing and wash it before reuse.

#### 8. Limitations

- For in vitro diagnostic use.
- For professional use only.
- The clinical interpretation of any positive staining, or its absence, must be done within the context of clinical history, morphology, other histopathological criteria as well as other diagnostic tests. It is the responsibility of a qualified pathologist to be familiar with the FISH probes, reagents, diagnostic panels, and methods used to produce the stained preparation. Staining must be performed in a certified, licensed laboratory under the supervision of a pathologist who is responsible for reviewing the stained slides and assuring the adequacy of positive and negative controls.
- Specimen staining, especially signal intensity and background staining, is dependent on the handling and processing of the specimen prior to staining. Improper fixation, freezing, thawing, washing, drying, heating, sectioning, or contamination with other specimens or fluids may produce artefacts or false results. Inconsistent results may result from variations in fixation and embedding methods, as well as from inherent irregularities within the specimen.
- $\bullet$   $\,$  The probe should be used only for detecting loci described in 4. "Reagents provided".
- The performance was validated using the procedures described in this instruction for use. Modifications to these procedures might alter the performance and have to be validated by the user.

# 9. Interfering substances

Red blood cells present in the specimen might exhibit autofluorescence which hinders signal recognition.

The following fixatives are incompatible with FISH:

- Bouin's fixative
- B5 fixative
- Acidic fixatives (e.g., picric acid)
- Zenker's fixative
- Alcohols (when used alone)
- Mercuric chloride
- Formaldehyde/zinc fixative
- Hollande's fixative
- Non-buffered formalin

# 10. Preparation of specimens

#### Recommendations:

- Fixation in 10% neutrally buffered formalin for 24 h at room temperature (18-25°C).
- Sample size  $\leq 0.5$  cm<sup>3</sup>.
- Use premium quality paraffin.
- Embedding should be carried out at temperatures lower than 65°C.
- Prepare 2-4 μm microtome sections.
- Use positively charged microscope slides.
- Fix for 2-16 h at 50-60°C.

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# 11. Preparatory treatment of the device

25x Wash Buffer (WB2) is to be pretreated according to the instructions in 12. "Assay procedure". All other kit reagents are ready-to-use. No reconstitution, mixing, or dilution is required. Bring probe to room temperature (18-25°C) before use, protect from light. Prior to opening the vial, mix by vortexing and spin down briefly.

# 12. Assay procedure

# 12.1 Day 1

#### Preparatory steps

- (1) Prepare two ethanol series (70%, 90%, and 100% ethanol solutions):
  Dilute 100% ethanol with deionized or distilled water. These solutions can be stored in suitable containers and can be re-used.
- (2) Heat Pretreatment Solution Citric (PT1): Warm to 98°C.
- (3) Wash Buffer SSC (WB1): Bring to room temperature (RT). WB1 may form precipitates at 2-8°C, which do not affect the quality and should dissolve when heated.
- (4) ZytoLight FISH Probe: Bring to RT before use, protect from light.

# Optional, when performing post-fixation step:

(strongly recommended if tissue fixation is not optimal)

Prepare a 1% Formaldehyde solution using the Formaldehyde Dilution Buffer Set (PT-0006-100)

# Pretreatment (dewax/proteolysis)

- (1) Incubate slides for 10 min at 70°C (e.g., on a hot plate).
- (2) Incubate slides for 2x 10 min in xylene.
- (3) Incubate in 100%, 100%, 90%, and 70% ethanol, each for 5 min.
- (4) Wash 2x 2 min in deionized or distilled water.
- (5) Incubate for 15 min in pre-warmed <u>Heat Pretreatment Solution Citric</u> (PT1) at 98°C.

We recommend not to use more than eight slides per staining jar.

- (6) Transfer slides immediately to deionized or distilled water, wash for 2x 2 min and drain off or blot off the water.
- (7) Apply (dropwise) Pepsin Solution (ES1) to the specimens and incubate for 15 min at 37°C in a humidity chamber.

ES1 may form precipitates, which do not affect the quality.

Depending on multiple factors, e.g. nature and duration of fixing, thickness of sections, and nature of tissue/cells, different incubation times may be required. As an incubation guideline, we recommend an incubation time of 2-30 min for tissue samples and 2-15 min for cell samples. As a general rule, we recommend to ascertain the optimum time for proteolysis in pretests.

(8) Wash for 5 min in Wash Buffer SSC (WB1).

# Optional, when performing post-fixation step:

Incubate slides for 15 min in 1% Formaldehyde solution and wash subsequently for 5 min in Wash Buffer SSC (WB1)

- (9) Wash for 1 min in deionized or distilled water
- (10) Dehydration: in 70%, 90%, and 100% ethanol, each for 1 min
- (11) Air dry sections.

Note: Make sure to completely dry sections prior to probe application since residual moisture may reduce signal intensity and/or affect tissue morphology.

#### Denaturation and hybridization

Pipette 10 µl of the <u>Zyto Light SPEC ERBB2/CEN 17 Dual Color Probe</u>
 (PL8) onto each pretreated specimen.

Avoid long exposure of the probe to light.

(2) Cover specimens with a 22 mm x 22 mm coverslip (avoid trapped bubbles) and seal the coverslip.

We recommend using rubber cement (e.g., Fixogum Rubber Cement) for sealing.

- (3) Place slides on a hot plate or hybridizer and denature specimens for 10 min at 75°C.
- (4) Transfer the slides to a humidity chamber and hybridize overnight at 37°C (e.g., in a hybridization oven).

It is essential that the tissue/cell samples do not dry out during the hybridization step.

#### 12.2 Day 2

#### Preparatory steps

(1) Preparation of 1x Wash Buffer A: Dilute 1 part 25x Wash Buffer A (WB2) with 24 parts deionized or distilled water. Fill three staining jars with the 1x Wash Buffer A and pre-warm it to 37°C.

Diluted 1x Wash Buffer A is stable for one week when stored at 2-8°C.

(2) <u>DAPI/DuraTect-Solution</u> (MT7): Bring to room temperature before use, protect from light.

#### Post-hybridization and detection

- (1) Carefully remove the rubber cement or glue.
- (2) Remove the coverslip by submerging in 1x Wash Buffer A at 37°C for
- (3) Wash using 1x Wash Buffer A for 2x 5 min at 37°C.

The 1x Wash Buffer A should be pre-warmed. Check with a thermometer if necessary.

- (4) Incubate the slides in 70%, 90%, and 100% ethanol, each for 1 min.
- (5) Air dry the samples protected from light.
- (6) Pipette 25 µl <u>DAPI/DuraTect-Solution</u> (MT7) onto the slides. Avoiding trapped bubbles, cover the samples with a coverslip (24 mm x 60 mm). Incubate in the dark for 15 min.

Using a pipette tip which has been cut off to increase the size of the opening, can make the pipetting process easier. Avoid long exposure to light.

- (7) Store the slide in the dark. For longer storage periods, this should take place at 2-8°C.
- (8) Evaluation of the sample material is carried out by fluorescence microscopy. Filter sets for the following wavelength ranges are required:

Fluorescent dye	Excitation	Emission
ZyGreen	503 nm	528 nm
ZyOrange	547 nm	572 nm

#### 13. Interpretation of results

With the use of appropriate filter sets, the hybridization signals of the probe appear green (ERBB2 gene region) and orange (CEN 17).

**Normal situation**: In interphases of normal cells or cells without an amplification involving the ERBB2 gene region, two green signals and two orange signals appear (see Fig. 2).

**Aberrant situation:** In cells with an amplification of the ERBB2 gene region, an increased number of green signals or green signal clusters will be observed (see Fig. 2).

Overlapping signals may appear as yellow signals.

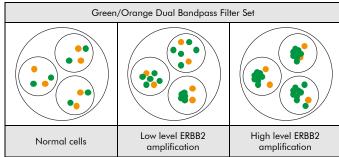


Fig. 2: Expected results in normal and aberrant interphase nuclei

Other signal distribution may be observed in some abnormal samples which might result in a different signal pattern than described above, indicating variant rearrangements. Unexpected signal patterns should be further investigated.

## Please note:

- Due to decondensed chromatin, single FISH signals can appear as small signal clusters. Thus, two or three signals of the same size, separated by a distance ≤ 1 signal diameter, should be counted as one signal.
- Do not evaluate overlapping nuclei.
- Do not count over-digested nuclei (recognized by dark areas visible inside of the nuclei).

- Do not count nuclei with strong auto-fluorescence, which hinders signal recognition.
- A negative or unspecific result can be caused by multiple factors (see chapter 17).
- In order to correctly interpret the results, the user must validate this
  product prior to use in diagnostic procedures according to national
  and/or international guidelines.

#### 14. Recommended quality control procedures

In order to monitor correct performance of processed specimens and test reagents, each assay should be accompanied by internal and external controls. If internal and/or external controls fail to demonstrate appropriate staining, results with patient specimens must be considered invalid.

**Internal control:** Non-neoplastic cells within the specimen that exhibit normal signal pattern, e.g., fibroblasts.

External control: Validated positive and negative control specimens.

#### 15. Performance characteristics

**Accuracy:** The location of hybridization of the probe was evaluated on metaphase spreads of a karyotypically normal male. In all tested specimens the probe hybridized solely to the expected loci. No additional signals or cross-hybridizations were observed. Therefore, the accuracy was calculated to be 100%.

**Analytical sensitivity:** For the analytical sensitivity assessment, the probe was evaluated on metaphase spreads of karyotypically normal males. All nuclei showed the expected normal signal pattern in all tested specimens. Therefore, the analytical sensitivity was calculated to be 100%.

**Analytical specificity:** For the analytical specificity assessment, the probe was evaluated on metaphase spreads of karyotypically normal males. In all tested specimens, all signals hybridized solely to the expected target loci and no other loci. Therefore, the analytical specificity was calculated to be 100%.

#### 16. Disposal

The disposal of reagents must be carried out in accordance with local regulations.

#### 17. Troubleshooting

Any deviation from the operating instructions can lead to inferior staining results or to no staining at all.

Weak signals or no signals at all

Possible cause	Action
No target sequences available	Use appropriate controls
Cell or tissue sample not fixed properly	Optimize fixing time and fixative or apply a post-fixation step as described in "12. Assay procedure".
Heat pretreatment, proteolysis, denaturation, hybridization, or stringency wash temperature incorrect	Check temperature of all technical devices used, using a calibrated thermometer
Proteolytic pretreatment not carried out properly	Optimize pepsin incubation time, increase or decrease if necessary
Probe evaporation	When using a hybridizer, the use of the wet stripes/water filled tanks is mandatory. When using a hybridization oven, the use of a humidity chamber is required. In addition, the coverslip should be sealed completely, e.g., with Fixogum, to prevent drying-out of the sample during hybridization
Too low concentrated stringency wash buffer	Check concentration of Wash Buffer A
Old dehydration solutions	Prepare fresh dehydration solutions

Fluorescence microscope adjusted wrongly	Adjust correctly
Inappropriate filter sets used	Use filter sets appropriate for the fluochromes of the probe.  Triple-bandpass filter sets provide less light compared to single or dualbandpass filter sets. Consequently, the signals may appear fainter using these triple-bandpass filter sets
Photo-damage of the probes/fluorophores	Accomplish hybridization and washing steps in the dark

Cross hybridization signals; noisy background

Possible cause	Action
Incomplete dewaxing	Use fresh solutions; check duration of dewaxing
Proteolytic pretreatment too strong	Reduce pepsin incubation time
Probe volume per area too high	Reduce probe volume per section/area, distribute probe dropwise to avoid local concentration
Slides cooled to room temperature before hybridization	Transfer the slides quickly to 37°C
Too high concentrated stringency wash buffer	Check concentration of Wash Buffer A
Washing temperature following hybridization too low	Check temperature; increase if necessary
Dehydration of specimens between the individual incubation steps	Prevent dehydration by sealing the slides and performing incubation in a humid environment

Tissue morphology degraded

Possible cause	Action
Cell or tissue sample has not been fixed properly	Optimize fixing time and fixative or apply a post-fixation step as described in "12. Assay procedure".
Proteolytic pretreatment not carried out properly	Optimize pepsin incubation time, increase or decrease if necessary
Insufficient drying before probe application	Extend air-drying

Overlapping nuclei

Possible cause	Action
Inappropriate thickness of tissue sections	Prepare 2-4 $\mu$ m microtome sections

Specimen floats off the slide

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Possible cause	Action	
Unsuitable slide coating	Use appropriate slides	
Proteolytic pretreatment too strong	Reduce pepsin incubation time	

#### Weak counterstain

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Possible cause	Action	
Low concentrated DAPI solution	Use <u>DAPI/DuraTect-Solution (ultra)</u> (Prod. No. MT-0008-0.8) instead	
DAPI incubation time too short	Adjust DAPI incubation time	

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#### 18. Literature

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Our experts are available to answer your questions. Please contact helptech@zytovision.com



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