DRAFT: PathVysion™ HER-2 DNA Probe Kit
(LSI® HER-2/*neu* SpectrumOrange™ / CEP® 17 SpectrumGreen™)
Order Number 30-161060

Proprietary Name: PathVysion HER-2 DNA Probe Kit

Common or Usual Name: Fluorescence *in situ* hybridization (FISH) reagents

Intended Use

The PathVysion HER-2 DNA Probe Kit (PathVysion Kit) is designed to detect amplification of the HER-2/*neu* gene via fluorescence *in situ* hybridization (FISH) in formalin-fixed, paraffin-embedded human breast cancer tissue specimens. Results from the PathVysion Kit are intended for use as an adjunct to existing clinical and pathologic information currently used as prognostic factors in stage II, node-positive breast cancer patients. The PathVysion Kit is further indicated as an aid to predict disease-free and overall survival in patients with stage II, node positive breast cancer treated with adjuvant cyclophosphamide, doxorubicin, and 5-fluorouracil (CAF) chemotherapy.

The PathVysion Kit is indicated as an aid in the assessment of patients for whom HERCEPTIN® (Trastuzumab) treatment is being considered (see HERCEPTIN package insert).

Warning

HERCEPTIN therapy selection
NOTE: All of the patients in the HERCEPTIN clinical trials were selected using an investigational immunohistochemical assay (CTA). None of the patients in those trials were selected using the PathVysion assay. The PathVysion assay was compared to the CTA on a subset of clinical trial samples and found to provide acceptably concordant results. The actual correlation of the PathVysion assay to HERCEPTIN clinical outcome in prospective clinical trials has not been established.

Adjuvant therapy selection
The Vysis PathVysion kit is not intended for use to screen for or diagnose breast cancer. It is intended to be used as an adjunct to other prognostic factors currently used to predict disease-free and overall survival in stage II, node-positive breast cancer patients and no treatment decision for stage II, node-positive breast cancer patients should be basedon HER-2/*neu* gene amplification status alone. Selected patients with breast cancers shown to lack amplification of HER-2/*neu* may still benefit from CAF (cyclophosphamide, doxorubicin, 5-fluorouracil) adjuvant therapy on the basis of other prognostic factors which predict poor outcome (e.g. tumor size, number of involved lymph nodes, and hormone receptor status). Conversely, selected patients with breast cancers shown to contain gene amplification may not be candidates for CAF therapy due to pre-existing or intercurrent medical illnesses.

Required Training
Vysis will provide training in specimen preparation, assay procedure, and interpretation of FISH testing of the HER-2/*neu* gene for inexperienced users. It is also recommended that a laboratory which has previously received training but now has new personnel performing the assay, request training for the new users.

Summary and Explanation

Among all cancers in the U.S., breast cancer is expected to be the most common cancer (32% / 182,000) in women and to be the second most common cause of death from cancer (18% / 46,000) in 1995 [1]. After surgery, breast cancers with positive axillary nodes, which account for 30% of all breast cancers [2], are associated with a shorter disease-free survival [3,4] and a shorter overall survival [5] than node negative breast cancers. It has been generally accepted that patients with breast cancer and positive axillary nodes at diagnosis, should be offered adjuvant systemic treatment.

Amplification or overexpression of the HER-2/*neu* gene has been shown to be an indicator of poor prognosis in node-positive breast cancer [6-10]. In one study, the prognostic value of HER-2/*neu* appears to be stronger among patients treated with chemotherapy [7]. However, in predicting disease-free and overall survival in individual patients, other established prognostic factors such as tumor size, number of positive lymph nodes, and steroid receptor status must also be taken into consideration.

The fluorescence *in situ* hybridization (FISH) technique has been used to detect HER-2/*neu* gene amplification in human breast carcinoma cell lines in both interphase and metaphase cells [11-14]. FISH appears to be an alternative technique capable of overcoming many of the inherent technical and interpretative limitations of other techniques, such as immunohistochemistry [15]. For quantification of HER-2/*neu* gene amplification, FISH assesses not only the level of HER-2/*neu* gene amplification directly in the tumor cells while retaining the characteristic morphology of the tissue studied, but also the spatial distribution of oncogene copies in individual uncultured primary breast carcinomas.

Principles of the Procedure

In situ hybridization is a technique that allows the visualization of specific nucleic acid sequences within a cellular preparation. Specifically, DNA fluorescence *in situ* hybridization (FISH) involves the precise annealing of a single stranded fluorescently labeled DNA probe to complementary target sequences. The hybridization of the probe with the cellular DNA site is visible by direct detection using fluorescence microscopy.

The Locus Specific Identifier (LSI®) HER-2/*neu* DNA probe is a 190 Kb SpectrumOrange™ directly labeled fluorescent DNA probe specific for the HER-2/*neu* gene locus (17q11.2-q12). The Chromosome Enumeration Probe (CEP®) 17 DNA probe is a 5.4 Kb SpectrumGreen directly labeled fluorescent DNA probe specific for the alpha satellite DNA sequence at the centromeric region of chromosome 17 (17p11.1-q11.1). The probes are pre-mixed and pre-denatured in hybridization buffer for ease of use. Unlabeled blocking DNA is also included with the probes to suppress sequences contained within the target loci that are common to other chromosomes. This PathVysion Kit is designed for the detection of HER-2/*neu* gene amplification in formalin-fixed, paraffin-embedded human breast tissue specimens by FISH. The assay is rapid, non-radioactive, requires little tumor material, and is capable of detecting as few as 2 to 8 copies of the oncogene.

Formalin-fixed paraffin embedded tissue specimens are placed on slides. The DNA is denatured to its single-stranded form and subsequently allowed to hybridize with the

PathVysion probes. Following hybridization, the unbound probe is removed by a series of washes, and the nuclei are counterstained with DAPI (4.6 diamidino-2-phenylindole), a DNA-specific stain that fluoresces blue. Hybridization of the PathVysion probes is viewed using a fluorescence microscope equipped with appropriate excitation and emission filters allowing visualization of the intense orange and green fluorescent signals. Enumeration of the LSI HER-2/*neu* and CEP 17 signals is conducted by microscopic examination of the nucleus, which yields a ratio of HER-2/*neu* gene to chromosome 17 copy number.

Reagents and Instruments

Materials Provided

This kit contains sufficient reagents to process approximately 20 assays. An assay is defined as one 22 mm x 22 mm target area.

1) LSI HER-2/*neu* SpectrumOrange (low copy number *E. coli* vector) / CEP 17 SpectrumGreen DNA Probe (*E. coli* plasmid)
   Vysis P.N.: 30-171060
   Quantity: 200 µL
   Storage: -20°C in the dark
   Composition: SpectrumGreen fluorophore-labeled alpha satellite DNA probe for chromosome 17, SpectrumOrange fluorophore-labeled DNA probe for the HER-2/*neu* gene locus, and blocking DNA, pre-denatured in hybridization buffer.

2) DAPI Counterstain
   Vysis P.N.: 30-804860
   Quantity: 300 µL
   Storage: -20°C in the dark
   Composition: 1000 ng/mL DAPI (4,6-diamidino-2-phenylindole) in phenylenediamine dihydrochloride, glycerol, and buffer

3) NP-40
   Vysis P.N.: 30-804820
   Quantity: 4 mL (2 vials)
   Storage: -20 to 25°C
   Composition: NP-40

4) 20X SSC salts
   Vysis P.N.: 30-805850
   Quantity: 66 g for up to 250 mL of 20X SSC solution
   Storage: -20 to 25°C
   Composition: sodium chloride and sodium citrate

Note: Material Safety Data Sheets (MSDS) for all reagents provided in the kits are available upon request from the Vysis Technical Service Department.

Storage and Handling

Store the unopened PathVysion Kit as a unit at -20°C protected from light and humidity. The 20X SSC salts and NP-40 may be stored separately at room temperature. Expiration dates for each of the components are indicated on the individual component labels. These storage conditions apply to both opened and unopened components.

Exposure to light, heat or humidity may affect the shelf life of some of the kit components and should be avoided. Components stored under conditions other than those stated on the labels may not perform properly and may adversely affect the assay results.

Materials Required but Not Provided

Laboratory Reagents

- ProbeChek HER-2/*neu* Normal Control Slides (Normal Signal Ratio) Order No. 30-805093; Formalin-fixed, paraffin-embedded, cultured human breast cancer cell line (MDA-MB-231; normal LSI HER-2/*neu*:CEP 17 ratio) applied to glass microscope slides. Quantity: 5 slides Store the control slides at 15-30°C in a sealed container with desiccant to protect them from humidity.
- ProbeChek HER-2/*neu* Cutoff Control Slides (Weakly Amplified Signal Ratio) Order No. 30-805042; Formalin-fixed, paraffin-embedded, cultured human breast cancer cell line (Hs 578T; low level HER-2/*neu* amplification) applied to glass microscope slides Quantity: 5 slides Store the control slides at 15-30°C in a sealed container with desiccant to protect them from humidity.
- Paraffin Pretreatment Reagent Kit (Vysis Cat. # 32-801200), which includes:
  - Pretreatment Solution (NaSCN) Quantity: 5 x 50 mL
  - Protease (Pepsin (2500-3000 units/mg)) Quantity: 5 x 25 mg
  - Protease Buffer (NaCl solution, pH 2) Quantity: 5 x 50 mL
  - Wash Buffer (2X SSC, pH 7) Quantity: 2 x 250 mL
- Neutral buffered formalin solution (4% formaldehyde in PBS)
- Hemo-De clearing agent (Scientific Safety Solvents #HD-150)
- Hematoxalin and eosin (H & E)
- Immersion oil appropriate for fluorescence microscopy. Store at room temperature (15-30°C).
- Ultra-pure, formamide. Store at 4°C for up to one month from delivery (See manufacturer's recommendations for detailed information).
- Ethanol (100%). Store at room temperature.
- Concentrated (12N) HCl
- 1N NaOH
- Purified water (distilled or deionized or Milli-Q). Store at room temperature.
- Rubber cement
- Drierite

Laboratory Equipment

- Precleaned silanized or positively charged glass microscope slides
- Slide warmer (45 - 50°C)
- 22 mm x 22 mm glass coverslips
- Microliter pipettor (1-10 µL) and sterile tips
- Polypropylene microcentrifuge tubes (0.5 or 1.5 mL)
- Timer
- Microtome
- Magnetic stirrer
- Vortex mixer
- Microcentrifuge
- Graduated cylinder
- Water baths (37±1°C, 72±1°C, and 80±1°C)

- Protein-free water bath (40°C)
- Air incubators (37°C and 56°C)
- Diamond-tipped scribe
- Humidified hybridization chamber
- Forceps
- Disposable syringe (5 mL)
- Coplin jars (6) Suggested type: Wheaton Product No. 900620 vertical staining jar
- Fluorescent microscope equipped with recommended filters (see next section)
- pH meter and pH paper
- Calibrated thermometer
- Microscope slide box with lid
- 0.45 µm pore filtration unit

Microscope Equipment and Accessories

Microscope: An epi-illumination fluorescence microscope is required for viewing the hybridization results. *If an existing fluorescence microscope is available, it should be checked to be sure it is operating properly to ensure optimum viewing of fluorescence in situ hybridization assay specimens.* A microscope used with general DNA stains such as DAPI, Propidium Iodide, and quinacrine may not function adequately for FISH assays. Routine microscope cleaning and periodic "tune-ups" by the manufacturer's technical representative are advisable.

Note: *Often, a presumed failure of reagents in an in situ assay may actually indicate that a malfunctioning or sub-optimal fluorescence microscope or incorrect filter set is being used to view a successful hybridization assay.*

Excitation Light Source: A 100 watt mercury lamp with life maximum of about 200 hours is the recommended excitation source. Record the number of hours that the bulb has been used and replace the bulb before it exceeds the rated time. Ensure that the lamp is properly aligned.

Objectives: Use oil immersion fluorescence objectives with numeric apertures ≥ 0.75 when using a microscope with a 100-watt mercury lamp. A 40X objective, in conjunction with 10X eyepieces, is suitable for scanning. For FISH analysis, satisfactory results can be obtained with a 63X or 100X oil immersion achromat type objective.

Immersion Oil: The immersion oil used with oil immersion objectives should be one formulated for low auto fluorescence and specifically for use in fluorescence microscopy.

Filters: Multi-bandpass fluorescence microscope filter sets optimized for use with the CEP and LSI DNA probe kits are available from Vysis for most microscope models. The recommended filter sets for the PathVysion Kit are the DAPI/9-Orange dual bandpass, DAPI/Green dual bandpass and DAPI/Green/Orange (V.2) triple bandpass. Hybridization of the LSI HER-2/*neu* and CEP 17 probes to their target regions is marked by orange and green fluorescence, respectively. All of the other DNA will fluoresce blue with the DAPI stain.

Preparation of Working Reagents

20X SSC (3M sodium chloride, 0.3M sodium citrate, pH 5.3)

To prepare 20X SSC pH 5.3, add together:

66 g	20X SSC
200 mL	purified water
250 mL	final volume

Mix thoroughly. Measure pH at room temperature with a pH meter. Adjust pH to 5.3 with concentrated HCl. Bring the total volume to 250 mL with purified water. Filter through a 0.45 µm pore filtration unit. Store at room temperature for up to 6 months.

Denaturing Solution (70% formamide / 2X SSC, pH 7.0-8.0)

To prepare denaturing solution, add together:

49 mL	formamide
7 mL	20X SSC, pH 5.3
14 mL	purified water
70 mL	final volume

Mix thoroughly. Measure pH at room temperature using a pH meter with glass pH electrode to verify that the pH is between 7.0 - 8.0. This solution can be used for up to one week. Check pH prior to each use. Store at 2 - 8°C in a tightly capped container when not in use.

Ethanol Solutions

Prepare v/v dilutions of 70%, 85%, and 100% using 100% ethanol and purified water. Dilutions may be used for one week unless evaporation occurs or the solution becomes diluted due to excessive use. Store at room temperature in tightly capped containers when not in use.

Post-Hybridization Wash Buffer (2X SSC/0.3% NP-40)

To prepare, add together:

100 mL	20X SSC, pH 5.3
847 mL	Purified water
3 mL	NP-40
1000 mL	Final Volume

Mix thoroughly. Measure pH at room temperature using a pH meter. Adjust pH to 7.0 - 7.5 with 1N NaOH. Adjust volume to 1 liter with purified water. Filter through 0.45 µm pore filtration unit. Discard used solution at the end of each day. Store unused solution at room temperature for up to 6 months.

Warnings and Precautions

1. For *In Vitro* Diagnostic Use.
2. The PathVysion Kit is intended for use only on formalin-fixed, paraffin-embedded breast cancer tissue; it is not intended for use on fresh or non-breast cancer tissue.
3. All biological specimens should be treated as if capable of transmitting infectious agents. The control slides provided with this kit are manufactured from human cell lines that have been fixed in 10% formalin. *Because it is often impossible to know which might be infectious, all human specimens and control slides should be treated with universal precautions. Guidelines for specimen handling are available from the U.S. Centers for Disease Control and Prevention* [16].
4. Exposure of the specimens to acids, strong bases, or extreme heat, should be avoided. Such conditions are known to damage DNA and may result in FISH assay failure.
5. Failure to follow all procedures for slide denaturation, hybridization, and detection may cause unacceptable or erroneous results.
6. To identify target areas, H & E staining should be conducted on every 10th slide of the same tissue block.
7. Hybridization conditions may be adversely affected by the use of reagents other than those provided by Vysis, Inc.
8. Proper storage of kit components is essential to ensure the labeled shelf life. Assay results may be adversely effected by kit components stored under other conditions.
9. If stored at low temperatures, 20X SSC may crystallize. If the crystals cannot be redissolved at room temperature, the solution should be discarded.
10. If any other working reagents precipitate or become cloudy, they should be discarded and fresh solutions prepared.
11. The DAPI Counterstain contains DAPI (4,6-diamidino-2-phenylindole) and 1,4-phenylenediamine.
    - DAPI is a possible mutagen based on positive genotoxic effects. Avoid inhalation, ingestion, or contact with skin.
    - 1,4 phenylenediamine is a known dermal sensitizer and a possible respiratory sensitizer. Avoid inhalation, ingestion, or contact with skin. Refer to MSDS for specific warnings.
12. Fluorophores are readily photobleached by exposure to light. To limit this degradation, handle all solutions containing fluorophores in reduced light. This includes all steps involved in handling the hybridized slide. Carry out all steps which do not require light for manipulation (incubation periods, washes, etc.) in the dark.
13. LSI HER-2/*neu* & CEP 17 DNA probe mixture contains formamide, a teratogen. Avoid contact with skin and mucous membranes.
14. Calibrated thermometers are required for measuring temperatures of solutions, water baths, and incubators.
15. Always verify the temperature of the pretreatment solution, denaturation solution and wash buffers prior to each use by measuring the temperature of the solution in the Coplin jar with a calibrated thermometer.
16. All hazardous materials should be disposed of according to your institution's guidelines for hazardous disposal.

Specimen Processing and Slide Preparation

Specimen Collection and Processing

The PathVysion Kit is designed for use on formalin-fixed, paraffin-embedded tissue specimens. Tissue collections should be performed according to the laboratory's standard procedures. **Selection of tissue for PathVysion assay should be performed by the pathologist**. Exposure of the specimens to acids, strong bases, or extreme heat, should be avoided. Such conditions are known to damage DNA and may result in FISH assay failure.

Breast tissue should be prepared in sections between 4 and 6 microns thick. Formalin-fixed, paraffin-embedded tissue may be handled and stored according the laboratory's routine procedures. To ensure optimum results from the PathVysion Kit, these methods should be consistent for all specimens analyzed. To identify target areas, H & E staining should be conducted on every 10th slide of the same tissue block.

Tissue sections should be mounted on the positive side of an organosilane-coated slide in order to minimize detachment of the tissue from the slide during FISH assay. The PathVysion Kit contains reagents sufficient for approximately 20 assays; one assay for the PathVysion Kit is defined as a 22 mm x 22 mm area. Larger specimen sections will require more than 10 μL of probe per assay.

Slide Preparation from Formalin-Fixed, Paraffin-Embedded Tissue

The following method may be used for preparing slides from formalin-fixed, paraffin-embedded tissue specimens:

1. Cut 4-6 μm thick paraffin sections using a microtome.
2. Float the sections in a protein-free water bath at 40°C.
3. Mount the section on the positive side of an organosilane-coated slide.
4. Allow slides to air dry.

(Start processing ProbeChek control slides here)

5. Bake slides overnight at 56°C.

Slide Pretreatment

Slides must be deparaffinized and the specimens fixed prior to assay with the PathVysion Kit. The package insert for the Vysis Paraffin Pretreatment Reagent Kit (Product No. 32-801200) contains detailed instructions. The following is a brief description of the procedure.

Deparaffinizing Slides

- Immerse slides in Hemo-De for 10 minutes at room temperature.
- Repeat twice using new Hemo-De each time.
- Dehydrate slides in 100% EtOH for 5 minutes at room temperature. Repeat.
- Air dry slides or place slides on a 45-50°C slide warmer.

Pretreating Slides

- Immerse slides in 0.2N HCl for 20 minutes.
- Immerse slides in purified water for 3 minutes.
- Immerse slides in Wash Buffer for 3 minutes.
- Immerse slides in Pretreatment Solution at 80°C for 30 minutes.
- Immerse slides in purified water for 1 minute.
- Immerse slides in Wash Buffer for 5 minutes. Repeat.

Protease Treatment

- Remove excess buffer by blotting edges of the slides on a paper towel.
- Immerse slides in Protease Solution at 37°C for 10 minutes.
- Immerse slides in Wash Buffer for 5 minutes. Repeat.
- Dry slides on a 45-50°C slide warmer for 2-5 minutes.

Fixing the Specimen

- Immerse slides in 10% buffered formalin at room temperature for 10 minutes.
- Immerse slides in wash buffer for 5 minutes. Repeat.
- Dry slides on a 45-50°C slide warmer for 2-5 minutes.
- Proceed with the PathVysion HER-2 assay protocol.

Assay Procedure

Fluorescence *In Situ* Hybridization Procedure Summary

Probe Preparation

1. Allow the probe to warm to room temperature so that the viscosity decreases sufficiently to allow accurate pipetting.
2. Vortex to mix. Centrifuge each tube for 2-3 seconds in a bench-top microcentrifuge to bring the contents to the bottom of the tube. Gently vortex again to mix.

Denaturation of Specimen DNA

The timing for preparing the probe solutions should be carefully coordinated with denaturing the specimen DNA so that both will be ready for the hybridization step at the same time.

1. Prewarm the humidified hybridization chamber (an airtight container with a piece of damp blotting paper or paper towel approximately 1 in. x 3 in. taped to the side of the container) to 37°C by placing it in the 37°C incubator prior to slide preparation. Moisten the blotting paper or paper towel with water before each use of the hybridization chamber.
2. Verify that the pH of the denaturing solution is 7.0 - 8.0 at room temperature before use. Add denaturing solution to Coplin jar and place in a 72±1°C water bath for at least 30 minutes, or until the solution temperature reaches 72±1°C. Verify the solution temperature before use.
3. Mark the areas to be hybridized with a diamond-tipped scribe.
4. Denature the specimen DNA by immersing the prepared slides in the denaturing solution at 72±1°C (≤6 slides per jar) for 5 minutes. Do not denature more than 6 slides at one time per Coplin jar. ***Note: Verify the solution temperature before each use.***
5. Using forceps, remove the slide(s) from the denaturing solution and immediately place into a 70% ethanol wash solution at room temperature. Agitate the slide to remove the formamide. Allow the slide(s) to stand in the ethanol wash for 1 minute.
6. Remove the slide(s) from 70% ethanol. Repeat step 5 with 85% ethanol, followed by 100% ethanol.

7. Drain the excess ethanol from the slide by touching the bottom edge of the slide to a blotter, and wipe the underside of the slide dry with a laboratory wipe.
8. Dry the slide(s) on a 45-50°C slide warmer for 2-5 minutes.

Hybridization

1. Apply 10 µL of probe mixture to target area of slide. Immediately place a 22 mm x 22 mm glass coverslip over the probe and allow it to spread evenly under the coverslip. Air bubbles will interfere with hybridization and should be avoided. The remaining probe solution should be refrozen immediately after use.
2. Seal coverslip with rubber cement as follows: Draw the rubber cement into a 5 mL syringe. Eject a small amount of rubber cement around the periphery of the coverslip overlapping the coverslip and the slide, forming a seal around the coverslip.
3. Place slides in the pre-warmed humidified hybridization chamber. Cover the chamber with a tight lid and incubate at 37°C overnight (14-18 hours).
4. Skip to Post-Hybridization Washes.

Post-Hybridization Washes

1. Add post-hybridization wash buffer (2X SSC/0.3% NP-40) to a Coplin jar. Prewarm the post-hybridization wash buffer by placing the Coplin jar in the 72±1°C water bath for at least 30 minutes or until solution temperature has reached 72±1°C. ***Note: The temperature of the wash solution must return to 72±1°C before washing each batch.***
2. Add post-hybridization wash buffer to a second Coplin jar and place at room temperature. Discard both wash solutions after 1 day of use.
3. Remove the rubber cement seal from the first slide by gently pulling up on the sealant with forceps.
4. Immerse slide(s) in post-hybridization wash buffer at room temperature and float off coverslip.
5. After coverslip has been carefully removed, remove excess liquid by wicking off the edge of the slide and immerse slide in post-hybridization wash buffer at 72±1°C for 2 minutes (≤6 slides/jar).
6. Remove each slide from the wash bath and air dry in the dark in an upright position. (A closed drawer or a shelf inside a closed cabinet is sufficient.)
7. Apply 10 µL of DAPI counterstain to the target area of the slide and apply a glass coverslip. Store the slide(s) in the dark prior to signal enumeration.

Slide Storage

Store hybridized slides (with coverslips) at -20°C in the dark. After removing from -20°C storage, allow slide(s) to reach room temperature prior to viewing using fluorescence microscopy.

Signal Enumeration

Assessing Slide Adequacy

Evaluate slide adequacy using the following criteria:

- Probe Signal Intensity: The signal should be bright, distinct, and easily evaluable. Signals should be in either bright, compact, oval shapes or stringy, diffuse, oval shapes.
- Background: The background should appear dark or black and relatively free of fluorescence particles or haziness.

If any of the above features are unsatisfactory, consult the troubleshooting guide (Table 2) and process a fresh slide.

Recognition of Target Signals

Use the prescribed filter. Adjust the depth of the focus, and become familiar with the size and shape of the target signals and noise (debris). Enumerate hybridization signals only among tumor cells. Tumor cells in general are larger than normal cells, lymphocytes, and epithelial cells. Identify target areas by H & E stain on every 10th slide of the same tissue block. Identify these areas on the coverslip after the FISH assay is performed.

Selection of Optimum Viewing Area and Evaluable Nuclei

Use a 25X objective to view the hybridized area and locate the target of interest (tumor cells as identified by H & E stain). Avoid areas of necrosis and where the nuclear borders are ambiguous. Skip those nuclei with signals that require subjective judgment. Skip signals with weak intensity and non-specificity, or with noisy background. Skip nuclei with insufficient counterstain to determine the nuclear border. Enumerate only those nuclei with discrete signals.

Signal Enumeration Guidelines

Using a 40X objective, scan several areas of tumor cells to account for possible heterogeneity. Select an area of good nuclei distribution; avoid areas of the target where hybridization signals are weak. Using a 63X or 100X objective, begin analysis in the upper left quadrant of the selected area and, scanning from left to right, count the number of signals within the nuclear boundary of each evaluable interphase cell according to the guidelines provided below and in Figure 1.

- Focus up and down to find all of the signals present in the nucleus.
- Count two signals that are the same size and separated by a distance equal or less than the diameter of the signal as one signal.
- Do not score nuclei with no signals or with signals of only one color. Score only those nuclei with one or more FISH signals of each color.
- For each nucleus, count the number of LSI HER-2/*neu* signals and the number of CEP 17 signals. (Note: It may be necessary to alternate between the DAPI/9-orange, DAPI/green and DAPI/Green/Orange (V.2) filter sets to view both color signals). Record counts (see Recommended Method for HER-2 to CEP 17 Ratio Determination, below)..

Figure 1
Dual Color Signal Counting Guide

Key: ○ = green probe, CEP 17

● = orange probe, LSI HER-2/neu

1		Nuclei are overlapping and all areas of both of the nuclei are not visible but signals are not in overlapping area. Count as two orange and two green in each nucleus.
2		Count as two orange signals and two green signals. One orange signal is diffuse.
3		Don't count. Nuclei are overlapping, all areas of nuclei are not visible and some signals are in overlapping area.
4		Count as two orange signals and two green signals. One orange signal is split.
5		Count as one orange signal and two green signals. One green signal is split and the orange signal is split.
6		Count as two orange signals and one green signal.
7		Count as three orange signals and one green signal.
8		Count as four orange signals and two green signals.

Recommended Method for LSI HER-2 to CEP 17Ratio Determination

The recommended method for LSI HER-2/*neu* to CEP 17 ratio determination is by dividing the total number of LSI HER-2/neu signals by the total number of CEP 17 signals in counting the same 20 nuclei.

1. Determine and record the number of LSI HER-2/*neu* and CEP 17 counts in 20 nuclei (see sample data sheet, below).
2. Add all LSI HER-2/*neu* signals. This sum represents the Total LSI HER-2/*neu* signals, e.g., 252.
3. Add all CEP 17signals. This sum represents the Total CEP 17 signals, e.g., 44.
4. To calculate the final result, use the following ratio.

   Total LSI HER-2/*neu* signals (step 2)/Total CEP 17 signals (step 3), e.g., 252/44 equals a ratio of 5.73, which is positive for HER-2/*neu* amplification.

   If the LSI HER-2/*neu* to CEP 17 ratio is borderline (1.8-2.2), count an additional 20 nuclei and recalculate the ratio.

5. Report results as follows:
   If the ratio is <2, HER-2/*neu* gene amplification was not observed;
   if the ratio is ≥2, HER-2/*neu* gene amplification was observed.
   ***Note*: A ratio at or near the cutoff (1.8 - 2.2) should be interpreted with caution, as described above.**

	HER-2/*neu*	CEP 17
1	5	1
2	6	1
3	10	1
4	11	1
5	14	1
6	17	1
7	6	2
8	11	2
9	15	2
10	14	2
11	16	2
12	20	2
13	8	3
14	10	3
15	12	3
16	14	3
17	18	3
18	19	3
19	11	4
20	15	4
Total	**252**	**44**

Quality Control

Use of Control Slides

Control slides must be run concurrently with patient slides to monitor assay performance and to assess the accuracy of signal enumeration. Control slides should be used beginning with the Slide Preparation procedure (see Specimen Processing and Slide Preparation section above). Controls should be run on each day of FISH testing and with each new kit lot. Vysis ProbeChek control slides are required with each run of patient slides processed. In addition, individual users may choose to use their own control material, providing it is characterized and validated in accordance with CLIA high-complexity requirements.

Assess control slide adequacy and perform signal enumeration according to the instructions in the Signal Enumeration section above. The criteria for slide adequacy must be satisfied and the LSI HER-2/*neu*:CEP 17 ratio results should be within the established ranges for acceptable test performance. See Table 1 for acceptable ProbeChek slide results.

In no case should FISH test results be reported if assay controls fail. If control slides fail to meet the slide acceptance criteria, the assay may not have been performed properly or the PathVysion Kit component(s) may have performed inadequately. A repeat analysis with fresh control slides and patient specimen slide(s) will be necessary. If control slides meet the acceptance criteria but the results are outside the specified range, the enumeration may not have been performed correctly and an independent, repeat analysis of the same slide may be appropriate. In the event of hybridization failure, with either the specimen or control slide(s), consult the troubleshooting guide in Table 2.

For clinical specimens, when interpretation of the hybridization signal is difficult and there is insufficient specimen sample for re-assay, the test is uninformative. If there are insufficient cells for analysis, the test is uninformative.

Patient specimens should be controlled according to standard laboratory procedure requirements. Hybridization quality and enumeration results should be documented on an appropriate form. Hybridization quality and efficiency should be considered when evaluating results.

Table 1
Acceptable HER-2/*neu* : CEP 17 Ratio Ranges for ProbeChek HER-2/neu Control Slide

Control	HER-2/*neu*:CEP 17 Acceptable Range
Normal Ratio Control Slide (Cat. No 30-805093)	0.75-1.25
Cutoff (Weakly Amplified) Ratio Control Slide (Cat. No. 30-805042)	1.60-2.00

Table 2
Troubleshooting Guide

Problem	Probable Cause	Solution
• No signal or weak signals	• Inappropriate filter set used to view slides	• Use recommended filters
	• Microscope not functioning properly	• Call microscope manufacturer's technical representative
	• Improper lamps (i.e. Xenon or Tungsten)	• Use a mercury lamp (100 watt recommended)
	• Mercury lamp too old	• Replace with a new lamp
	• Mercury lamp misaligned	• Realign lamp
	• Dirty and/or cracked collector lenses	• Clean and replace lens
	• Dirty or broken mirror in lamp house	• Clean or replace mirror
	• Hybridization conditions inappropriate	• Check temp of 37±1°C incubator • Increase hybridization time to at least 14 hours
	• Inappropriate post-hybridization wash temperature	• Check temp. of 72±1°C water bath
	• Air bubbles trapped under coverslip and prevented probe access	• Apply coverslip by first touching the surface of the hybridization mixture
	• Insufficient amount of hybridization solution for section	• Increase amount of hybridization solution to 20 µL per section
	• Inadequate protease digestion	• Check temp. of 37±1°C bath • Check that pH of buffer is 2.0±0.2 • Increase digestion time, up to 60 min.
	• Section overfixed	• Eliminate fixation in pretreatment • Optimal fixation time in preparation of paraffin embedded slides is 24-48 hrs. Longer fixation times will lead to progressive degradation of morphology.
	• DNA loss (poor DAPI staining)	• Check fixation conditions
• Variation of signal intensity across tissue section	• Inherent in many tissue sections	• Check DAPI staining. If DAPI staining in poor area is good, then score slide. If DAPI staining is poor in poor area, increase fixation time
	• Probe unevenly distributed on slide due to air bubbles under coverslip	• Repeat hybridization on next adjacent section or same slide and make sure no air bubbles are trapped under coverslip
	• Oversized section	• Increase volume of hybridization solution to 20 µL on large tissue sections
• Noisy background	• Inadequate wash stringency	• Check pH of 7.2-7.5 wash buffer • Check temperature of 72±1°C bath • Provide gentle agitation during wash • Increase wash time to 5 minutes
• Tissue loss or tissue morphology degraded	• Tissue section underfixed (poor DAPI staining)	• Check fixation time/conditions
	• Inappropriate slides used	• Use positively charged slides
	• Improper slide baking	• Check temp. of 56°C oven
	• Over-pretreatment	• Check temp. of 80±1°C pretreatment • Decrease pretreatment time • Decrease protease digestion time
	• Over-denaturation	• Check temp. of 72±1°C denaturation bath • Decrease denaturation time
	• Tissue section overfixed (cell boundaries will be distinct)	• Eliminate fixation step • Optimal fixation time in preparation of paraffin embedded slides is 24-48 hrs. Longer fixation times will lead to progressive degradation of morphology
	• Tissue section was torn removing coverslip after hybridization	• Allow coverslip to soak off in wash buffer

Contact the Vysis Technical Services Department at 800-553-7042 for further assistance

Interpretation of Results

The number of LSI HER-2/*neu* and CEP 17 signals per nucleus are recorded in columns. Results on enumeration of 20 interphase nuclei from tumor cells per target are reported as the ratio the total HER-2/*neu* signals to those of CEP 17. Our clinical study found that specimens with amplification showed a LSI HER-2/*neu*:CEP 17 signal ratio of ≥2.0; normal specimens showed a ratio of <2.0.

Results at or near the cutoff point (1.8 - 2.2) should be interpreted with caution. In the event of a borderline result (1.8 – 2.2), particularly if there also appears to be variability of the counts from nucleus to nucleus, 20 additional nuclei should be enumerated. The specimen slide should be re-enumerated by another technician to verify the results. If still in doubt, the assay should be repeated with a fresh specimen slide. If the test results are not consistent with the clinical findings, a consultation between the pathologist and the treating physician is warranted.

<u>Reasons to Repeat the Assay</u>

The following are situations requiring repeat assays with fresh specimen slides and the appropriate control slides. Consult the troubleshooting guide (Table 2) for probable causes and the actions needed to correct specific problems.

1. If one or both of the control slides fail to meet the slide acceptance criteria, the specimen slide results are not reliable, and the assay must be repeated.

2. If there are fewer than 20 evaluable nuclei, the test is uninformative and the assay should be repeated.

3. If, upon assessing the slide quality as described in the Signal Enumeration section, any of the aspects (signal intensity, background, or cross-hybridization) are unsatisfactory, the assay must be repeated.

Limitations

1. The PathVysion Kit has been optimized only for identifying and quantifying chromosome 17 and the HER-2/*neu* gene in interphase nuclei from formalin-fixed, paraffin-embedded human breast tissue specimens. Other types of specimens or fixatives should not be used.
2. The performance of the PathVysion Kit was validated using the procedures provided in this package insert only. Modifications to these procedures may alter the performance of the assay.
3. Performance characteristics of the PathVysion Kit have been established only for node positive patients receiving the designated regimens of CAF and for metastatic breast cancer patients being considered for HERCEPTIN therapy. Performance with other treatment regimens has not been established.
4. The clinical interpretation of any test results should be evaluated within the context of the patient's medical history and other diagnostic laboratory test results.
5. FISH assay results may not be informative if the specimen quality and/or specimen slide preparation is inadequate.
6. Technologists performing the FISH signal enumeration must be capable of visually distinguishing between the orange and green signals.

Expected Values

FISH interphase analysis was performed on human breast tissue specimens from 524 breast cancer patients to verify the cutoff point and to assess the expected ratio of LSI HER-2/*neu* to CEP 17.

Sixty nuclei were enumerated per specimen. Based on a cutoff point of 2.0 for assessing HER-2/*neu* gene amplification in breast tissue specimens (see "Establishment of Cut-off Point", below, for details on determining cutoff point), 433 of the specimens were negative and 91 positive for HER-2/*neu* gene amplification. The distribution of ratios of HER-2/*neu* to CEP 17 signals for the 433 non-amplified specimens is summarized in Table 3.

Table 3
Distribution of Ratio of HER-2/*neu* to CEP 17 Signals in Non-Amplified Breast Tissue Specimens

Statistics	Range		
	0.1-1.0	1.1-1.5	1.6-1.99
mean	0.86	1.15	1.72
S.D.	0.14	0.13	0.11
n	185	226	22

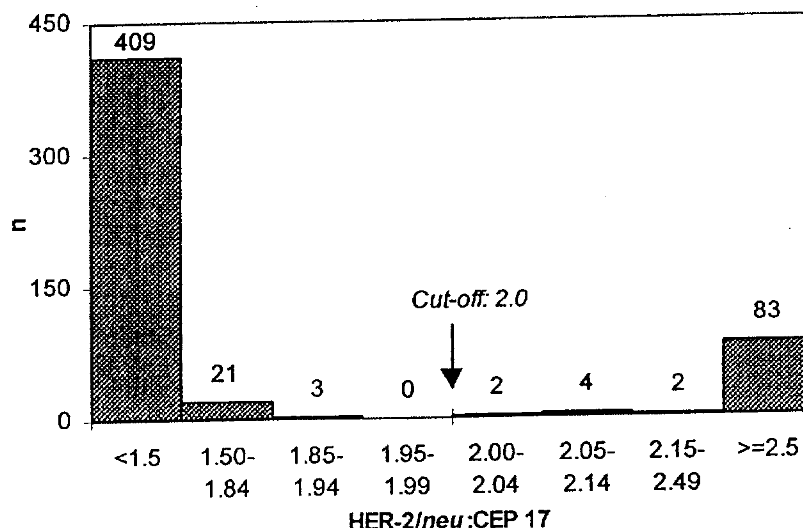
The distribution of ratios of HER-2/*neu* to CEP 17 signals for the 91 HER-2/*neu* amplified specimens is summarized in Table 4.

Table 4
Distribution of Ratio of HER-2/*neu* to CEP 17 Signals in Amplified Breast Tissue Specimens

Statistics	Range		
	2.0-5.0	5.1-10.0	>10.0
mean	3.35	7.39	12.77
S.D.	0.95	1.41	1.80
n	33	42	16

Establishment of Cut-off Point
In the pivotal CALGB 8869 study, the cutoff point for determining HER-2/*neu* gene amplification was determined to be 2.0, based on best-fit analysis of clinical outcome of CAF treatment. Among the 433 non-amplified specimens, the largest ratio of LSI HER-2/*neu* to CEP 17 signals was 1.95, and among the 91 amplified samples, the smallest ratio of LSI HER-2/*neu* to CEP 17 signals was 2.0. This gap between the largest value among normal specimens and the smallest value among amplified specimens reduces the chance of misclassification, with 2.0 as the cutoff point. The distribution of the ratio of LSI HER-2/*neu*:CEP 17 in the 524 specimens from the study described above are shown in Figure 2.

Figure 2
Signal Distribution



Performance Characteristics

Analytical Sensitivity and Specificity

Hybridization Efficiency
On the ProbeChek™ quality control slides, the average percentage of cells with no hybridization signal was 0.0 to 2.0%. These slides are prepared from formalin-fixed, paraffin-embedded breast cancer cell lines, and represent the best case scenario for hybridization efficiency. Thus, under these conditions, the hybridization efficiency is expected to be 98%, with <2% cells having no signal for either probe.

Analytical Sensitivity
The analytical sensitivity of the PathVysion Kit probes was tested in the reproducibility study described below. In that study, the 1.0 - 1.2 HER-2/*neu*:CEP 17 ratio specimen was estimated with a mean of 1.05 (±0.03), and the 1.6 - 2.0 HER-2/*neu*:CEP 17 ratio specimen was estimated with a mean of 1.81 (±0.08). The upper 95% CI was 1.11 for the 1.0 - 1.2 ratio specimen and the lower 95% CI for the 1.6 - 2.0 specimen was 1.65. Thus, the limit of detection for the PathVysion Kit in interphase cells is estimated to be a ratio of 1.5.

Analytical Specificity

Locus specificity studies were performed with metaphase spreads from normal lymphocytes according to standard Vysis QC protocols. A total of 254 metaphase spreads were examined sequentially by G-banding to identify chromosome 17, and the HER-2/*neu* gene locus, followed by FISH. No cross-hybridization to other chromosome loci was observed in any of the 254 cells examined; hybridization was limited to the intended target regions of the two probes.

Stringency studies were also performed, according to standard Vysis protocols, on formalin-fixed, paraffin-embedded tissue specimens to determine the optimum denaturation time and temperature; hybridization time and temperature; post-hybridization wash time and temperature; and post-hybridization wash buffer composition. For the denaturation step, three temperatures (65°C, 73°C, and 80°C) were tested for 2 minutes, 5 minutes and 8 minutes each. The results showed no statistical difference in the overall rating among all denaturation temperatures and durations; all combinations passed the quality evaluation. Stringency of the hybridization step was tested in two parts; first, hybridizations were conducted at 5 different temperatures (27°C, 32°C, 37°C, 42°C, and 47°C) for 18 hours, then for 5 different durations (10 hr, 14 hr, 18 hr, 22 hr, and 26 hr) at the recommended temperature (37°C). Hybridization quality was significantly affected by both hybridization temperature and time. ,Hybridizations at 37°C for 18 hours showed the highest overall quality ratings, although there were no statistically significant differences in hybridization quality between 14 and 18 hours. Thus, an incubation time of 14-18 hours is recommended.

The post-hybridization wash step was tested in a similar manner; first assays were conducted at 5 different temperatures (69°C, 71°C, 73°C, 76°C, and 80°C), then for different durations, ranging from 2 to 8 minutes at 73°C. Wash temperature was a significant factor, with 73°C resulting in the highest ratings. Wash times between 2 and 5 minutes all produced acceptable results, but increasing the wash time to 8 minutes significantly lowered the overall quality ratings in some samples. Therefore the recommended post-hybridization wash conditions are 72±1°C for 2 minutes.

The wash buffer composition was also analyzed to determine the effect on signal intensity and probe specificity. Increasing the salt concentration from 0.4X SSC to 2X SSC increased the signal intensity, but did not appear to compromise the probe specificity. Thus, a wash buffer composition of 2X SSC / 0.3% NP-40 is recommended.

Reproducibility and Repeatability Pre-clinical Studies

The repeatability of the FISH assay for HER-2/*neu* was determined on consecutive sections of normal and amplified breast tissue, as well as on different thicknesses of the same tissue. On 10 consecutive tissue sections from one normal breast tissue, the average ratio of HER-2/*neu* to CEP 17 copy number was 1.19 (S.D. = 0.05); the results are shown in Table 5.

Table 5
Average Number of Signals per Cell and Ratio of HER-2/*neu*:CEP 17 Copy Number in Consecutive Sections (with normal HER-2/*neu*)

	Section Number									
	1	2	3	4	5	6	7	8	9	10
HER-2	3.8	3.2	3.6	3.5	3.6	3.5	3.4	3.5	3.3	3.1
CEP 17	3.1	3.1	3.0	2.7	3.0	3.0	2.8	2.1	3.0	2.7
Ratio	1.2	1.2	1.2	1.3	1.2	1.2	1.2	1.6	1.1	1.1

On 10 consecutive tissue sections from one specimen with amplified HER-2/*neu*, the average ratio of HER-2/*neu* to CEP 17 copy number was 3.61 (S.D. = 0.50); the results are shown in Table 6.

Table 6
Average Number of Signals per Cell and Ratio of HER-2/*neu*:CEP 17 Copy Number in Consecutive Sections (with amplified HER-2/*neu*)

	Section Number									
	1	2	3	4	5	6	7	8	9	10
HER-2	4.7	4.9	5.9	4.5	3.6	4.6	4.6	4.8	4.5	4.2
CEP 17	1.2	1.3	1.3	1.3	1.3	1.3	1.4	1.3	1.4	1.3
Ratio	3.9	3.7	4.7	3.6	2.8	3.7	3.3	3.8	3.3	3.3

Similarly, on 8 consecutive normal tissue sections of different thickness (2-8 microns), the average ratio of HER-2/*neu* to CEP 17 copy number was 1.15 (S.D. = 0.16); the results are shown in Table 7. These results demonstrated an acceptable degree of reproducibility of the HER-2/*neu* FISH assay in tissue sections with thicknesses between 4 and 8 microns.

Table 7
Average Number of Signals per Cell and Ratio of HER-2/*neu* :CEP 17 Copy Number in Consecutive Sections of Different Thickness

	Thickness of Section (microns)							
	2	2	4	4	6	6	8	8
HER-2	2.3	2.4	2.4	2.7	2.7	2.8	2.6	3.3
CEP 17	1.7	1.8	2.3	2.5	2.7	2.7	2.5	3.2
Ratio	1.4	1.4	1.1	1.1	1.0	1.1	1.1	1.0

Reproducibility Using Control Slides

To assess the reproducibility of the HER-2/*neu* and CEP 17 assay, analyses for the ratio of HER-2/*neu* to CEP 17 were assessed for inter-site, inter-lot, inter-day, and inter-observer reproducibility on **control slides** with differing levels of HER-2/*neu* gene amplification. Four specimens consisting of formalin-fixed, paraffin-embedded tissue from human breast tumor cell lines with normal (1.0-1.2) and amplified (1.6-2.0, 3-5, 7-11) ratios of HER-2/*neu* to CEP 17 were evaluated for HER-2/*neu* and CEP 17 according to the instructions for signal enumeration in the package insert. The overall hybridization success rate was 98.3% (118/120) on the first try. Hybridization of the two replacement slides was successful.

Using ANOVA, statistically significant variations were observed between observers, which reflects the subjectivity in signal interpretation and enumeration. No statistically significant variations were observed in any of the other study parameters. The mean, standard deviation, and percent CV of the observed ratios of HER-2/*neu* to CEP 17 are shown in Tables 8-11.

Table 8
Site-to-Site Reproducibility

Ratio of HER-2/*neu* to CEP 17	Statistics	Site #1	Site #2	Site #3
1.0-1.2	Mean	1.08	1.01	1.07
	S.D	0.03	0.04	0.07
	C.V.(%)	2.66	3.58	6.77
	n	8	8	8
1.6-2.0	Mean	1.81	1.71	1.78
	S.D.	0.05	0.05	0.19
	C.V.(%)	2.88	2.78	10.50
	n	8	8	8
3.0-5.0	Mean	4.39	3.65	4.49
	S.D.	0.22	0.18	0.79
	C.V.(%)	4.99	4.93	17.64
	n	8	8	8
7.0-11	Mean	7.21	8.26	8.23
	S.D.	0.15	0.83	0.87
	C.V.(%)	2.07	10.10	10.55
	n	8	8	8

S.D. (Standard Deviation), C.V.(%) (Coefficient of Variation).

Table 9
Lot-to-Lot Reproducibility

Ratio of HER-2/*neu* to CEP 17	Statistics	Lot #1	Lot #2	Lot #3	Lot #4
1.0-1.2	Mean	1.05	1.07	1.02	1.04
	S.D	0.07	0.06	0.03	0.05
	C.V.(%)	6.48	6.06	3.21	4.87
	n	6	6	6	6
1.6-2.0	Mean	1.78	1.77	1.77	1.75
	S.D.	0.10	0.13	0.15	0.09
	C.V.(%)	5.65	7.49	8.54	5.07
	n	6	6	6	6
3.0-5.0	Mean	4.08	3.92	4.57	4.14
	S.D.	0.44	0.34	0.96	0.40
	C.V.(%)	10.78	8.74	20.92	9.56
	n	6	6	6	6
7.0-11	Mean	7.67	7.72	7.89	8.33
	S.D.	0.69	0.72	0.88	1.06
	C.V.(%)	8.97	9.36	11.16	12.68
	n	6	6	6	6

S.D. (Standard Deviation), C.V.(%) (Coefficient of Variation).

Table 10
Day-to-Day Reproducibility

Ratio of HER-2/*neu* to CEP 17	Statistics	Assay Day #1	Assay Day #2	Assay Day #3	Assay Day #4
1.0-1.2	Mean	1.06	1.07	1.02	1.04
	S.D	0.06	0.07	0.05	0.04
	C.V.(%)	5.65	6.61	4.58	4.03
	n	6	6	6	6
1.6-2.0	Mean	1.76	1.77	1.77	1.77
	S.D.	0.17	0.14	0.08	0.10
	C.V.(%)	9.62	7.99	4.31	5.65
	n	6	6	6	6
3.0-5.0	Mean	4.24	4.48	4.10	3.89
	S.D.	0.48	0.97	0.36	0.38
	C.V.(%)	11.25	21.56	8.89	9.71
	n	6	6	6	6
7.0-11	Mean	7.91	8.01	7.72	7.97
	S.D.	1.11	0.90	0.57	0.89
	C.V.(%)	13.99	11.22	7.39	11.20
	n	6	6	6	6

S.D. (Standard Deviation), C.V.(%) (Coefficient of Variation).

Table 11
Observer-to-Observer Reproducibility

Ratio of HER-2/*neu* to CEP 17	Statistics	Observer #1	Observer #2
1.0-1.2	Mean	1.06	1.04
	S.D	0.07	0.03
	C.V.(%)	7.00	2.85
	n	12	12
1.6-2.0	Mean	1.71	1.82
	S.D.	0.10	0.11
	C.V.(%)	6.01	6.20
	n	12	12
3.0-5.0	Mean	4.05	4.31
	S.D.	0.44	0.73
	C.V.(%)	10.80	16.84
	n	12	12
7.0-11	Mean	7.52	8.28
	S.D.	0.49	0.95
	C.V.(%)	6.55	11.44
	n	12	12

S.D. (Standard Deviation), C.V.(%) (Coefficient of Variation).

Assay Portability

A five-center, blinded, randomized, comparative study using formalin-fixed, paraffin-embedded **human breast cancer specimens** was conducted to assess assay portability [17]. Study specimens consisted of formalin-fixed, paraffin-embedded human breast cancer tissue sections with varying levels of HER-2/*neu* gene amplification. The specimens included one normal (no amplification), two with low level, and one with moderate level HER-2/*neu* gene amplification, as determined by FISH.

Day-to-Day Reproducibility

Table 12 shows the mean observed ratios of LSI HER-2/*neu* to CEP 17 for the three assay days. As shown by the p-values, there were no statistically significant variations in ratio values across the three study days (all p-values >0.05). The results of this study demonstrated that the PathVysion assay is reproducible from day to day.

Table 12
Summary Statistics of LSI HER-2/*neu* to CEP 17 by Assay Day

Expected Ratio	Statistics	Assay Day #1	Assay Day #2	Assay Day #3	P-value
1.0-1.2	**Mean**	**1.01**	**1.06**	**1.05**	0.6826
	S.D.	0.10	0.12	0.08	
	C.V.(%)	9.90	11.32	7.62	
	n	15	15	15	
2.1-2.8	**Mean**	**2.54**	**2.43**	**2.32**	0.5535
	S.D.	0.19	0.32	0.22	
	C.V.(%)	7.48	13.17	9.52	
	n	15	15	15	
2.5-3.5	**Mean**	**3.18**	**2.98**	**3.03**	0.2083
	S.D.	0.30	0.31	0.32	
	C.V.(%)	9.43	10.40	10.56	
	n	15	15	15	
5.0-7.0	**Mean**	**5.69**	**5.63**	**5.69**	0.9620
	S.D.	0.53	0.49	0.86	
	C.V.(%)	9.31	8.70	15.11	
	n	15	15	15	

Site-to-Site Reproducibility

Table 13 shows the mean observed ratios of LSI HER-2/*neu* to CEP 17 for the five study sites.. There was some statistically significant variation in the ratio value across the five study sites (p=<0.05 for the normal and 2.5-3.5 specimen). However, these differences were not clinically relevant; 99% of the specimens were correctly classified as positive or negative for HER-2/*neu* gene amplification.

Table 13
Site-to-Site Reproducibility

Expected Ratio	Statistics	Site #1	Site #2	Site #3	Site #4	Site #5	P-value
1.0-1.2	**Mean**	**1.00**	**1.16**	**1.01**	**1.04**	**0.97**	0.0001
	S.D.	0.09	0.09	0.07	0.09	0.04	
	C.V.(%)	9.00	7.76	6.93	8.65	4.12	
	n	9	9	9	9	9	
2.1-2.8	**Mean**	**2.40**	**2.46**	**2.57**	**2.26**	**2.65**	0.0965
	S.D.	0.19	0.26	0.52	0.22	0.24	
	C.V.(%)	7.92	10.60	20.20	9.73	9.06	
	n	9	9	9	9	9	
2.5-3.5	**Mean**	**3.01**	**3.09**	**3.41**	**2.74**	**3.08**	<0.0001
	S.D.	0.21	0.35	0.20	0.23	0.16	
	C.V.(%)	6.98	11.3	5.87	8.39	5.19	
	n	9	9	9	9	9	
5.0-7.0	**Mean**	**5.48**	**5.22**	**5.94**	**5.82**	**5.91**	0.0568
	S.D.	0.66	0.43	0.56	0.89	0.18	
	C.V.(%)	12.0	8.24	9.43	15.30	3.05	
	n	9	9	9	9	9	

The summary of assay variations for all five sites are presented in Table 14. The standard deviation (S.D.) and the coefficient of variation (C.V.) were relatively small and stable across all ratios of LSI HER-2/*neu* to CEP 17.

Table 14
Summary of Site-to-Site Reproducibility

Ratio of HER-2/*neu* to CEP 17	Mean	Standard Deviation	C.V. (%)	N
1.0-1.2	1.04	0.10	9.60	45
2.1-2.8	2.47	0.32	12.96	45
2.5-3.5	3.07	0.31	10.10	45
5.0-7.0	5.67	0.63	11.11	45

This study had a 100% hybridization success rate, validating the ease of use of the PathVysion HER-2 Kit.

Clinical Studies

Dose of Cyclophosphamide, Adriamycin and 5-Fluorouracil (CAF) (CALGB 8869 Study)

The interaction between HER-2/*neu* gene amplification and dose of CAF was evaluated in a retrospective analysis of a single randomized clinical trial, CALGB 8869. This was a large, prospective, randomized trial in stage II, node-positive breast cancer patients that evaluated three different doses of adjuvant CAF chemotherapy: a high dose (cyclophosphamide at 600 mg/m^2, doxorubicin at 60 mg/m^2, and 5-fluorouracil at 600 mg/m^2 for four cycles), a moderate dose (cyclophosphamide at 400 mg/m^2, doxorubicin at 40 mg/m^2, and 5-fluorouracil at 400 mg/m^2 for six cycles), or a low dose (cyclophosphamide at 300 mg/m^2, doxorubicin at 30 mg/m^2, and 5-fluorouracil at 300 mg/m^2 for four cycles). Doses were administered on a 28-day cycle.

Archived tissue specimens from 572 patients, randomly selected from the original study population, were included for analysis by FISH assay with DNA probe. The objectives of this study were to determine whether amplification of the HER-2/*neu* gene provides statistically significant and independent prognostic information pertaining to disease-free survival and overall survival in stage II, node positive breast cancer patients receiving adjuvant CAF therapy; and to explore the relationship between HER-2/*neu* gene amplification and clinical data, including such factors as tumor grade and steroid receptor status.

Among these 572 tumor specimens, 45 were excluded due to FISH assay failures, and 3 were duplicate assays. This left 524 cases for analysis. Using the Vysis PathVysion DNA Probe Kit, HER-2/*neu* gene amplification was defined as ≥2 (i.e., the ratio of average HER-2/*neu* to average CEP 17 signals with 60 nuclei counted). A total of 433 patient samples were found to be HER-2/*neu*-negative and 91 HER-2/*neu*-positive.

The table below lists the several baseline characteristics of the 524 patients whose archived tumor specimens were selected for evaluation by this assay, as well as, details of the adjuvant treatments received on the original CALGB 8869 study.

Table 15
Comparison of Patient Characteristics at Baseline and Details of Adjuvant Treatment

	HER-2/*neu* amplification* n=91	No HER-2/*neu* amplification* n=433
Age		
<40	17.6	14.5
40-50	39.6	40.0
>50	42.9	48.5
Premenopausal	46.2	39.5
Peri/Postmenopausal	53.8	60.5
Tumor size		
≤2 cm	31.9	37.2
>2 - ≤5	57.1	58.4
>5	9.9	3.9
unknown	1	0.5
Positive nodes		
≤3	59.3	55.9
4-9	27.5	34.9
≥10	13.2	9.2
ER (+)	49.5	71.4
PR(+)	35.2	61.7
ER (+) or PR (+)	60.4	77.8
CAF dose regimen received		
High	33.0	34.4
Moderate	34.1	31.4
Low	33.0	34.2

*percent of patients

The results of analysis with Cox proportional hazard model for disease-free survival using FISH measurement of HER-2/*neu* gene amplification showed a statistically significant interaction between HER-2/*neu* gene amplification and the CAF dose regimen received (p=0.033, likelihood test, see Table 16). Similarly, the results of Cox proportional hazard model for overall survival also showed a statistically significant interaction between HER-2/*neu* gene amplification and the CAF dose regimen received (p=0.028, likelihood test, see Table 16)

Table 16
Cox Proportional Hazard Model Showing Likelihood-Ratio Tests for Disease-free and Overall Survival

	Disease-Free Survival			Overall Survival		
Source	DF	ChiSq	P value	DF	ChiSq	P value
CAF	2	5.56	0.06	2	4.57	0.10
Square root: # positive nodes	1	72.87	0.0000	1	56.32	0.0000
Tumor > 2 cm	1	13.77	0.0002	1	12.93	0.0003
Premenopausal	1	1.96	0.16	1	0.10	0.76
HER-2 ratio	1	10.05	0.0015	1	10.52	0.0012
HER-2 ratio interaction of CAF dose	2	6.84	0.033	2	7.15	0.028

Disease-free survival probabilities (Table 17, Figure 3a) are comparable among the three dose groups of patients with HER-2/*neu* -negative tumors. For example, at 7 years post-randomization the estimated disease-free survival probabilities are 55%, 63%, and 61% for low (L), moderate (M), and high (H) CAF dose groups, respectively. The dose effect is greater for patients with HER-2/*neu*- positive tumors (Table 17, Figure 3b), with disease-free survival at 7 years of 36%, 44%, and 66% for L, M, and H CAF dose groups, respectively. The corresponding figures for overall survival at 7 years (Table 18, Figure 3c) have a similar relationship: 64%, 75%, and 70% for patients with HER-2/*neu* -negative tumors, and 48%, 50%, and 76% for patients with HER-2/*neu*- positive tumors, again for L, M, and H CAF dose groups, respectively (Table 18, Figure 3d).

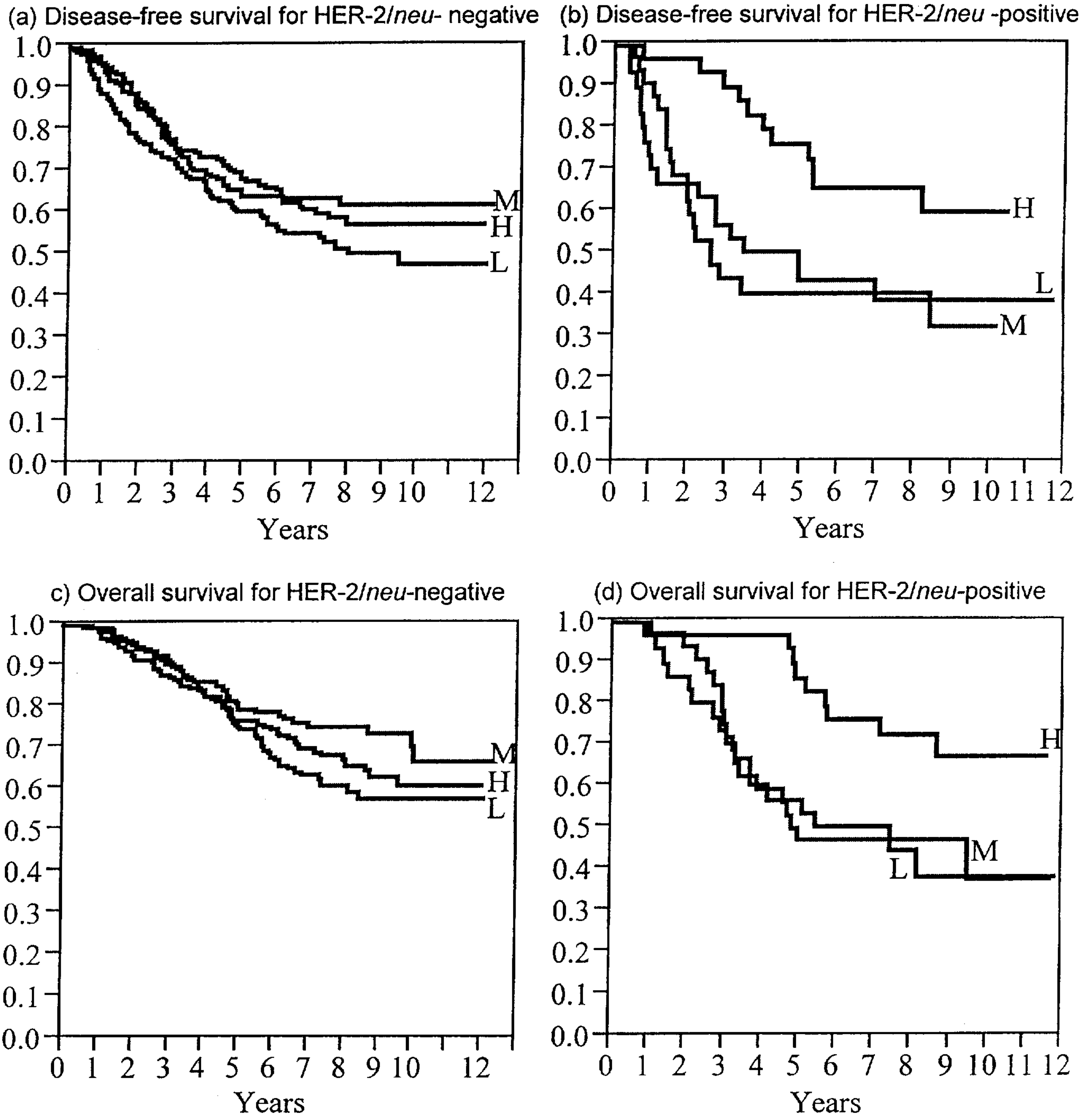
Table 17
Disease-free Survival Probabilities

CAF Dose	HER-2/*neu* negative	HER-2/*neu* positive
Low	55%	36%
Moderate	63%	44%
High	61%	66%

Table 18
Overall Survival Probabilities

CAF Dose	HER-2/*neu* negative	HER-2/*neu* positive
Low	64%	48%
Moderate	75%	50%
High	70%	76%

Figure 3
Disease-free (a, b) and overall (c, d) survival for patients with HER-2/*neu*-negative (a, c) and positive (b, d) tumors for the three CAF dose groups, H, M, and L*



*HER-2/*neu* positivity means HER/CEP ≥ 2. Sample sizes in (a, c) are 149, 136, and 148 (for H, M, and L) and in (b, d) are 30, 31, and 30. The significance levels for the HER-2/*neu* by CAF interaction from the proportional hazards models (Table 16) are 0.033 for disease-free survival—(a) vs. (b)—and 0.028 for overall survival—(c) vs. (d).

FISH analysis of the study specimens showed that there was a significant dose-response effect of adjuvant chemotherapy with CAF in patients with HER-2/*neu* gene amplification, but not in patients with no or minimal HER-2/*neu* amplification. This association was found in both disease-free and overall survival. In addition, this study found no correlation between HER-2/*neu* copy number, as assessed by FISH, and patient age, menopausal status, tumor size or the number of positive nodes. A statistically significant negative correlation was observed between HER-2/*neu* copy number and both estrogen (ER) and progesterone (PR) receptor status.

Concordance with the Clinical Trial Assay (CTA)

The primary mechanism of HER-2 protein overexpression in human breast cancer appears to be via gene amplification [18,19]. Fluorescence *in situ* hybridization (FISH) detection of HER-2/*neu* gene amplification provides an additional diagnostic method to define HER-2 overexpression.

The PathVysion Kit was compared to the Clinical Trial Assay (CTA), which was used to enroll patients into the Genentech-sponsored pivotal HERCEPTIN trials (H0648g, H0649g, H0650g) [19]. To establish concordance between FISH and the CTA, a subset of 623 specimens (317 positive and 306 negative, as determined by the CTA), were randomly selected in an intended 1:1 ratio from the specimens screened for enrollment in the HERCEPTIN trials. FISH assays were performed on all specimens, with informative results achieved on 529 specimens. The results from the analysis of the 529 informative cases are presented in Table 19.

Table 19
CTA versus FISH

FISH	CTA Score				Total
	0	1+	2+	3+	
Negative	207	28	67	21	323
Positive	7 (3.2%)	2 (6.7%)	21 (23.9%)	176 (89.3%)	206
Total	214	30	88	197	529

The results showed a 2x2 concordance of 82% (95% CI 79% - 85%), where concordance was defined as the proportion of samples rated 0 or 1+ by CTA and not amplified by FISH plus the proportion of samples rated 2+ or 3+ by CTA and amplified by FISH. These data are consistent with a high concordance between protein overexpression [as determined by immunohistochemistry (CTA)] and gene amplification [as determined by FISH (Vysis PathVysion)].

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