

Instructions for Use

CE IVD

IdentiClone® *BCL2/J_H* Translocation Assay

For identification of follicular cell lymphoma and other lymphomas and leukemias.

IVD For *In Vitro* Diagnostic Use



 Storage Conditions: **-85°C to -65°C**
(DNA controls may be separated from assay kits and stored at 2°C to 8°C)

Catalog#	Products	Quantity
REF 93090020	IdentiClone <i>BCL2/J_H</i> Translocation Assay – Gel Detection	33 Reactions
REF 93090040	IdentiClone <i>BCL2/J_H</i> Translocation Assay MegaKit – Gel Detection	330 Reactions

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1. Intended Use

The IdentiClone *BCL2/J_H* Translocation Assay is an *in vitro* diagnostic product intended for PCR-based detection of *BCL2/J_H* t(14;18) gene translocations in patients with suspect lymphoproliferations and can be used to:

- Distinguish lymphoma from benign lymphoid hyperplasia
- Distinguish follicular lymphoma from other B-cell lymphomas that may have a similar appearance
- Monitor and evaluate disease recurrence

2. Summary and Explanation of the Test

2.1. Background

The *BCL2* t(14;18) (q32;q21) translocation is found in 80-90% of follicular lymphomas and 30% of diffuse large cell lymphomas; however, this translocation is rarely present in other lymphoproliferative diseases. The t(14;18) brings about a juxtaposition of the *BCL2* gene with the immunoglobulin heavy chain (*IGH*) gene joining segment. This leads to a marked increase in expression of bcl-2 driven by the immunoglobulin heavy chain gene enhancer. The bcl-2 protein inhibits programmed cell death (apoptosis) leading to cell accumulation.

The majority of breakpoints on 18q21-22 occur within the major breakpoint region (Mbr) in the 3' untranslated region of exon 3 (60-70% of the cases), and the minor cluster region (mcr) located 3' to *BCL2* exon 3 (20-25% of the cases). Some breakpoints occur at distant loci and will not be identified by this particular test. Therefore a negative result does not completely exclude the presence of a *BCL2/IGH* gene rearrangement in the sample.¹

Invivoscribe's IdentiClone assays represent a new approach to PCR-based clonality testing. These standardized assays were carefully optimized testing positive and negative control samples using multiplex master mixes. Assay development was followed by extensive validation including the testing of more than 400 clinical samples using Revised European/American Lymphoma (REAL) Classification. Testing was performed at more than thirty prominent independent testing centers throughout Europe in a collaborative study known as the BIOMED-2 Concerted Action.¹

The gel detection based assays cannot reliably detect clonal populations comprising less than 1% of the total lymphocyte cell population. **Always interpret the results of this test in the context of morphologic and other relevant data and do not use alone for a diagnosis of malignancy.**

2.2. Summary

This test kit includes four (4) master mixes. The *BCL2/J_H* translocation master mixes (*BCL2/J_H* Tube A, B, and C) target the joining (J) region of the immunoglobulin heavy (*IGH*) chain gene and distinct regions of the *BCL2* gene. These master mixes are used to detect major breakpoint region (Mbr) and minor cluster region (mcr) of the *BCL2* t(14;18) translocations. The fourth master mix, the Specimen Control Size Ladder, targets multiple genes and generates a series of amplicons of approximately 100, 200, 300, 400, and 600 base pairs to ensure that the quality and quantity of input DNA is adequate to yield a valid result. A single thermal cycler program and similar detection methodologies are used with many of our assays which improves consistency and facilitates cross training on a broad range of different assays.

This assay is based on the EuroClonality/BIOMED-2 Concerted Action BMH4-CT98-3936.



3. Principles of the Procedure

3.1. Polymerase Chain Reaction (PCR)

PCR assays are routinely used for the identification of chromosome translocations. This test targets the Mbr and mcr regions of the *BCL2*/*J_H* translocations and amplifies genomic DNA between primers that target the *BCL2* gene and the conserved joining (*J*) regions of the *IGH* gene (*BCL2*/*J_H* Tubes A, B, and C). Breakpoints that occur outside the Mbr and mcr regions will not be identified by this particular test. Therefore, a negative result does not completely exclude the presence of a *BCL2*/*J_H* gene rearrangement in the sample.¹ DNA from a normal lymphocyte population will also produce a negative result.



- t(14;18) Tube A: two (2) *BCL2* MBR primers + one (1) *J_H* primer
 t(14;18) Tube B: four (4) *BCL2* 3'MBR primers + one (1) *J_H* primer
 t(14;18) Tube C: three (3) *BCL2* mcr primers + one (1) *J_H* primer

Figure 1. Depicted is a schematic diagram of the *BCL2*/*J_H* t(14;18) translocation showing the *BCL2* gene on the left and the Ig heavy chain (*IGH*) gene on the right. Shown are the relative positions and orientations for the major breakpoint region (Mbr) primers, the minor cluster region (mcr) primers, and the *J_H* primer, which are included in the 3 *BCL2*/*J_H* master mix tubes.

3.2. Gel Detection

Gel electrophoresis, such as agarose gel electrophoresis or non-denaturing polyacrylamide gel electrophoresis (PAGE), is commonly used to resolve amplicon products based on their size, charge, and conformation. Since DNA is negatively charged, when an electrical potential (voltage) is applied across the gel containing PCR products, the electrical field causes the amplicons to migrate through the gel. Smaller DNA fragments are able to easily migrate through the gel matrix, whereas larger DNA fragments migrate more slowly. This causes a separation of the amplicon products based on size. Ethidium bromide or other DNA intercalating dyes can then be used to stain and detect these products in the gel.


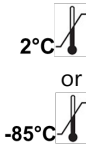
4. Reagents

4.1. Reagent Components

Table 1: Available Assays

Catalog #	Description	Quantity
REF 93090020	IdentiClone <i>BCL2</i> / <i>J_H</i> Translocation Assay – Gel Detection	33 Reactions
REF 93090040	IdentiClone <i>BCL2</i> / <i>J_H</i> Translocation Assay MegaKit – Gel Detection	330 Reactions

Table 2: Kit Components

Reagent	Catalog # (REF)	Reagent Components (active ingredients)	Unit Quantity	93090020 # of Units	93090040 # of Units	Storage Temp.
Master Mixes	23090050CE	<i>BCL2</i>/<i>J_H</i> Tube A – Unlabeled Multiple oligonucleotides targeting the major breakpoint region (Mbr) of the <i>BCL2</i> gene and the J region of the <i>IGH</i> gene in a buffered salt solution.	1500 µL	1	10	 -65°C
	23090060CE	<i>BCL2</i>/<i>J_H</i> Tube B – Unlabeled Multiple oligonucleotides targeting the 3' major breakpoint region (3' Mbr) of the <i>BCL2</i> gene and the J region of the <i>IGH</i> gene in a buffered salt solution.	1500 µL	1	10	
	23090070CE	<i>BCL2</i>/<i>J_H</i> Tube C – Unlabeled Multiple oligonucleotides targeting the minor cluster region (mcr) of the <i>BCL2</i> gene and the J region of the <i>IGH</i> gene in a buffered salt solution.	1500 µL	1	10	
Template Amplification Control Master Mix	20960020	Specimen Control Size Ladder – Unlabeled Multiple oligonucleotides targeting housekeeping genes.	1500 µL	1	10	
Positive Control DNAs	40881750	IVS-0030 Clonal Control DNA 200 µg/mL of DNA in 1/10 th TE solution	100 µL	1	5	 2°C or -65°C
	40900070	IVS-P002 Clonal Control DNA 1600 pg/mL of plasmid DNA diluted in IVS-0000 Polyclonal Control DNA in 1/10 th TE solution	100 µL	1	5	
	40881810	IVS-0031 Clonal Control DNA 200 µg/mL of DNA in 1/10 th TE solution	100 µL	1	5	
Negative (Normal) Control DNA	40920010	IVS-0000 Polyclonal Control DNA 200 µg/mL of DNA in 1/10 th TE solution	100 µL	1	5	

Note: There are no preservatives used in the manufacture of this kit.

4.2. Warnings and Precautions

- **IVD** This product is for *in vitro* diagnostic use.
- Use this assay kit as a system. Do not substitute other manufacturer's reagents. Dilution, reducing amplification reaction volumes, or other deviation in this protocol may affect the performance of this test and/or nullify any limited sublicense that comes with the purchase of this testing kit.
- Materials are stable until the labeled expiration date when stored and handled as directed. Do not use kits beyond their expiration date.
- Close adherence to the protocol will assure optimal performance and reproducibility. Use care to ensure use of correct thermal cycler program, as other programs may provide inaccurate/faulty data, such as false positive and false negative results.
- Do not mix or combine reagents from kits with different lot numbers.
- Wear appropriate personal protective equipment and follow good laboratory practices and universal precautions when working with specimens. Handle specimens in approved biological safety containment facilities and open only in certified biological safety cabinets. Use molecular biology grade water for the preparation of specimen DNA.
- Due to the analytical sensitivity of this test, use extreme care to avoid the contamination of reagents or amplification mixtures with samples, controls or amplified materials. Closely monitor all reagents for signs of contamination (*e.g.*, negative controls giving positive signals). Discard reagents suspected of contamination.
- To minimize contamination, wear clean gloves when handling samples and reagents and routinely clean work areas and pipettes prior to doing PCR.
- Autoclaving does not eliminate DNA contamination. Follow uni-directional workflow in the PCR laboratory; begin with master mix preparation, move to specimen preparation, then to amplification, and finally to detection. Do not bring amplified DNA into the areas designated for master mix or specimen preparation.
- Dedicate all pipettes, pipette tips, and any equipment used in a particular area to that area of the laboratory.
- Use sterile, disposable plastic ware whenever possible to avoid RNase, DNase, or cross-contamination.

4.3. Storage and Handling

- For any duration other than immediate use, **store assay kits at -85°C to -65°C.**
- The optimum storage temperature for DNA controls is 2°C to 8°C, but DNA controls can be stored long term at -85°C to -65°C.
- All reagents and controls must be thawed and vortexed or mixed thoroughly prior to use to ensure that they are resuspended completely. Excessive vortexing may shear DNA and cause labeled primers to lose their fluorophores.
- Materials are stable until the labeled expiration date when stored and handled as directed. Do not use kits beyond the expiration date.
- Due to high salt concentrations, PCR master mixes are sensitive to freeze/thaw cycles. Aliquot master mixes into sterile o-ring screw-cap tubes if necessary.

5. Instruments

5.1. Thermal cycler

- Use or function: Amplification of DNA samples
- Performance characteristics and specification:
 - Minimum Thermal Range: 15°C to 96°C
 - Minimum Ramping Speed: 0.8°C/sec
- Follow manufacturer's installation, operation, calibration, and maintenance procedures.
- See section 7.4: *Amplification* for thermal cycler program.

5.2. Electrophoresis Unit

- Use or function: DNA fragment separation
- Performance characteristics and specification:
 - Capable of running at 35V to 135V for extended times
- Follow manufacturer's installation, operation, calibration, and maintenance procedures.

5.3. UV Illumination Unit

- Use or function: DNA detection
- Performance characteristics and specification:
 - Capable of emitting light at a wavelength of ~302 nm
- Follow manufacturer's installation, operation, calibration, and maintenance procedures.

6. Specimen Collection and Preparation

6.1. Precautions

Biological specimens from humans may contain potentially infectious materials. Handle all specimens in accordance with the OSHA Standard on Bloodborne Pathogens or Biosafety Level 2.

6.2. Interfering Substances

The following substances are known to interfere with PCR:

- Divalent cation chelators
- Low retention pipette tips
- EDTA (not significant at low concentrations)
- Heparin

6.3. Specimen Requirements and Handling

This assay tests **genomic DNA** (gDNA) from the following sources:

- 5 cc of peripheral blood, bone marrow biopsy, or bone marrow aspirate anti-coagulated with heparin or EDTA (stored at 2°C to 8°C and shipped at ambient temperature)
- Minimum 5 mm cube of tissue (stored and shipped frozen; or stored and shipped in RPMI 1640 at ambient temperature or on ice)
- 2 µg of gDNA (stored at 2°C to 8°C and shipped at ambient temperature)
- Formalin-fixed paraffin embedded tissue or slides (stored and shipped at ambient temperature)

6.4. Sample Preparation

Extract the gDNA from patient specimens as soon as possible. Resuspend DNA to a final concentration of 100 µg to 400 µg per mL in 1/10th TE (1 mM Tris-HCl, pH 8.0; 0.1 mM EDTA) or in molecular biology grade or USP water. This is a robust assay system. A wide range of DNA concentrations will generate a valid result. Therefore, quantifying and adjusting DNA concentrations is generally not necessary. Testing sample DNA with the Specimen Control Size Ladder master mix will ensure that DNA of sufficient quality and quantity was present to yield a valid result.









6.5. Sample Storage

Store gDNA at 2°C to 8°C or at -85°C to -65°C for long term storage.

7. Assay Procedure

7.1. Materials Provided

Table 3: Kit components

Catalog #	Description
 23090050CE	<i>BCL2</i> / <i>J_H</i> Tube A – Unlabeled
 23090060CE	<i>BCL2</i> / <i>J_H</i> Tube B – Unlabeled
 23090070CE	<i>BCL2</i> / <i>J_H</i> Tube C – Unlabeled
 20960020	Specimen Control Size Ladder – Unlabeled
 40881750	IVS-0030 Clonal Control DNA
 40900070	IVS-P002 Clonal Control DNA
 40881810	IVS-0031 Clonal Control DNA
 40920010	IVS-0000 Polyclonal Control DNA

7.2. Materials Required (not provided)

Table 4: Materials Required (not provided)

Reagent/Material	Recommended Reagents/Materials and Suppliers	Catalog # (REF)	Notes
DNA Polymerase	Roche: <ul style="list-style-type: none"> EagleTaq DNA Polymerase Invivoscribe, Inc.: <ul style="list-style-type: none"> EagleTaq DNA Polymerase¹ or equivalent 	05206944190 60970100	N/A
Glass Distilled De-ionized Molecular Biology Grade or USP Water	N/A	N/A	Sterile and free of DNase and RNase.
Calibrated Pipettes	Rainin: <ul style="list-style-type: none"> P-2, P-20, P-200, and P-1000 pipettes Or SL-2, SL-20, SL-200, and SL-1000 pipettes 	N/A	Must be able to accurately measure volumes between 1 μ L and 1000 μ L.
Thermal cycler	Bio-Rad: <ul style="list-style-type: none"> MJ Research PTC-100 or PTC-200, PTC-220, PTC-240 Perkin-Elmer <ul style="list-style-type: none"> PE 9600 or PE 9700 	N/A	N/A
Vortex Mixer	N/A	N/A	N/A
PCR plates or tubes	N/A	N/A	Sterile
Filter barrier pipette tips	N/A	N/A	Sterile, RNase/DNase/Pyrogen-free
Microcentrifuge tubes	N/A	N/A	Sterile
Gel Electrophoresis Unit	N/A	N/A	For polyacrylamide gels
Ethidium Bromide	Thermo Fisher Scientific: <ul style="list-style-type: none"> UltraPure™ 10 mg/mL Ethidium Bromide 	15585-011	N/A
6% Polyacrylamide Gels	Thermo Fisher Scientific: <ul style="list-style-type: none"> Novex® TBE Gels (6%, 12 well) 	EC62652Box	N/A
TBE Running Buffer	Thermo Fisher Scientific: <ul style="list-style-type: none"> Novex TBE Running Buffer (5X) 	LC6675	Dilute 1:5 prior to use.
Gel Loading Buffer	Thermo Fisher Scientific: <ul style="list-style-type: none"> 10X BlueJuice™ Gel Loading Buffer Novex Hi-Density TBE Sample Buffer (5X) 	10816-015 LC6678	N/A
100 bp DNA Ladder	Thermo Fisher Scientific: <ul style="list-style-type: none"> TrackIt™ 100 bp DNA Ladder 	10488-058	N/A

¹**Note:** This product is for sale and use in the European Economic Area only. It is not to be resold or transferred to another party. See also Legal Notice in section 15.

7.3. Reagent Preparation

- Test all unknown samples with the Specimen Control Size Ladder master mix to ensure that no inhibitors of amplification are present and there is DNA of sufficient quality and quantity to generate a valid result.
- Singlicate test results are valid; however, **duplicate** testing is recommended when possible. If duplicate testing provides inconsistent results, re-testing or re-evaluation of the sample is necessary.
- Test **positive**, **negative** and **no template** controls for each master mix.
- It is recommended to batch multiple samples in a run to avoid running out of the negative control (IVS-0000 Polyclonal Control DNA). If batching samples is not practical, IVS-0000 Polyclonal Control DNA is also available for purchase separately.

7.3.1. Using gloved hands, remove the master mixes from the freezer. Allow the tubes to thaw completely; then gently vortex to mix.

7.3.2. In containment hood or dead air box, remove an appropriate aliquot from each master mix to individual clean, sterile microcentrifuge tubes.

- Aliquot volumes are 45 μL for each reaction.
- Include an additional reaction volume for every 15 reactions to correct for pipetting errors.
- Thus, for each master mix (except for the Specimen Control Size Ladder), the number of reactions (**n**) is:

n = 2 × # of samples	(run each sample in duplicate)
+ 1	positive control DNA (See section 7.7 <i>Recommended Positive Controls</i>)
+ 1	negative control DNA (IVS-0000 Polyclonal Control DNA)
+ 1	no template control (water)
+ 1	to correct for pipetting errors
<hr/>	
n = 2 × # of samples + 4	Total

- Therefore the total aliquot volume for each master mix is **n × 45 μL** .
- For the Specimen Control Size Ladder master mix, the number of reactions (**m**) is:

m = # of samples	(run each sample in duplicate)
+ 1	positive control DNA (IVS-0000 Polyclonal Control DNA)
+ 1	no template control (water)
+ 1	to correct for pipetting errors
<hr/>	
m = # of samples + 3	Total

- Therefore the total aliquot volume for the Specimen Control Size Ladder master mix is **m × 45 μL** .

7.3.3. Add 1.25 units (or 0.25 μL at 5 U/ μL) of Taq DNA polymerase per reaction to each master mix.

- The total Taq DNA polymerase added to each master mix is **n × 0.25 μL** , and **m × 0.25 μL** for the Specimen Control Size Ladder master mix.
- Gently vortex to mix.

7.3.4. For each reaction, aliquot 45 μL of the appropriate master mix + DNA polymerase solution into individual wells in a PCR plate or tube.

7.3.5. Add 5 μL of appropriate template (sample DNA, positive control DNA, negative control DNA, or water) to the individual wells containing the respective master mix solutions.

- Pipette up and down several times to mix.

7.3.6. Cap or cover the PCR plate.

- Samples are now ready to be amplified on a thermal cycler.
- If amplification cannot be performed immediately following reagent preparation, store PCR plate or tubes at 2°C to 8°C for up to 24 hours.

Quick Guide:

For each master mix and n reactions, mix:

n × 45 μL Master Mix

n × 0.25 μL Taq DNA polymerase

Vortex gently to mix.

Aliquot **45 μL** of master mix + DNA polymerase solution into each reaction well.

Add **5 μL** of appropriate Template to each well

Total reaction volume = **50 μL**

7.4. Amplification

7.4.1. Amplify the samples using the following PCR program:

- Use the **calculated** option for temperature measurement with the BioRad MJ Research PTC thermal cyclers.

Table 5: Thermal cycling conditions

Step	Temperature	Duration	Cycles
1	95°C	7 minutes	1
2	95°C	45 seconds	35
3	60°C	45 seconds	
4	72°C	90 seconds	
5	72°C	10 minutes	1
6	15°C	∞	1

7.4.2. Remove the amplification plate or tubes from the thermal cycler.

- Although amplified DNA is stable at room temperature for extended periods of time, store PCR products at 2°C to 8°C until detection.
- Detection must be within 30 days of amplification.

7.5. Gel Detection – Polyacrylamide TBE Gels

7.5.1. Prepare a 2% MetaPhor or NuSieve 3:1 agarose/1X TBE gel.

7.5.2. Place gel in electrophoresis unit and cover with 1X TBE buffer.

7.5.3. Mix 20 µL of each PCR product with 4 µL of 6X gel loading buffer.

7.5.4. Load 20 µL of this mixture into separate wells of the gel, then load 4 µL of the 100 bp DNA Ladder flanking the samples.

7.5.5. Run at 100V for 90 minutes.

- Voltage and electrophoresis time depend on the PCR amplicon size, gel length, and % of agarose in the gel.
- Voltage and run time can be adapted accordingly.

7.5.6. Stain gel with ethidium bromide or other equivalent dye.

7.5.7. Place gel over UV illuminator to visualize bands.

7.5.8. Photograph and interpret resulting data. (see sections 8: *Interpretation of Results* and 10: *Expected Values*)

7.6. Quality Control

Positive and negative (or normal) controls are furnished with the kit and can be run in singlicate each time the assay is performed to ensure proper performance of the assay. In addition, include a no template control (*e.g.*, water) to test for contamination of the master mix or cross-contamination of reactions. A buffer control may also be added to ensure that no contamination of the buffer used to resuspend the samples has occurred. The values for the positive controls are provided under section 10.1: *Expected Size of Amplified Products*. Additional controls and sensitivity controls (dilutions of positive controls into negative control) are available from Invivoscribe.

7.7. Recommended Positive Controls

- The amplicon sizes listed were determined using an ABI platform or by gel electrophoresis.

Table 6: Recommended Positive Controls

Master Mix	Target	Control DNA	Catalog #	Product Size in base pairs (bp)
<i>BCL2/J_H</i> Tube A	Mbr of <i>BCL2/J_H</i>	Valid Size Range IVS-0030 Clonal Control DNA	---	100 - 2500 254, ~750 ^a
<i>BCL2/J_H</i> Tube B	3' Mbr of <i>BCL2/J_H</i>	Valid Size Range IVS-P002 Clonal Control DNA	---	100 - 2500 ~140
<i>BCL2/J_H</i> Tube C	mcr of <i>BCL2/J_H</i>	Valid Size Range IVS-0031 Clonal Control DNA	---	100 - 2500 382, ~800 ^a
Specimen Control Size Ladder	Multiple Genes	Valid Size Range IVS-0000 Polyclonal Control DNA	---	100, 200, 300, 400, 600^b 100, 200, 300, 400, 600 ^b

^aNote: The ~750 bp band (*BCL2/J_H* Tube A) and the ~800 bp band (*BCL2/J_H* Tube C) are generally weak bands.

^bNote: Because smaller PCR fragments are preferentially amplified, it is not unusual for the 600 bp fragment to have a diminished signal or to be missing entirely.

8. Interpretation of Results

Although positive results are highly suggestive of malignancy, interpret both positive and negative results in the context of all clinical information and laboratory test results. The size range for each master mix has been determined by testing positive and negative control samples. For accurate and meaningful interpretation it is important to ignore peaks that occur outside of the valid size range for each master mix.

8.1. Analysis

- 8.1.1. Report samples that fail to amplify following repeat testing as “**A result cannot be reported on this specimen because there was DNA of insufficient quantity or quality for analysis**”.
- 8.1.2. Repeat testing of samples that test negative if the positive control reaction failed.
- 8.1.3. If samples run in duplicate yield differing results, re-test or re-evaluate samples for sample switching.
- 8.1.4. All assay controls must be examined prior to interpretation of sample results. If the controls do not yield the correct results, the assay is not valid and the samples cannot be interpreted.

Table 7. The following describes the analysis of each of the controls, and the decisions necessary based upon the results.

Type of Control	Expected Result	Aberrant Result
No Template Control	No amplification present, continue with analysis	Amplification present, repeat the assay.
Polyclonal Control	Product size is consistent with expected size listed in section 10.1: <i>Expected Size of Amplified Products</i> . No clonal rearrangements are present. Continue with analysis.	Clonal rearrangements are present. Repeat the assay.
Positive Control (This can also be an extraction control if positive control material is taken through extraction processes)	Product size is consistent with expected size listed in section 10.1: <i>Expected Size of Amplified Products</i> . Continue with analysis.	Repeat the assay.
Specimen Control Size Ladder (This amplification control is <u>essential</u> for samples of unknown quantity and quality.)	If all of the 100, 200, 300, 400, and 600 bp peaks are seen, continue with analysis. Because smaller PCR fragments are preferentially amplified, it is not unusual for the 600 bp fragment to have a diminished signal or to be missing entirely. Continue with analysis.	If no bands are seen, repeat the assay <u>unless specimen tests positive</u> . If only 1, 2, or 3 bands are seen, re-evaluate sample for DNA degradation <u>unless specimen tests positive</u> .

8.2. Sample Interpretation

Given that the controls produce expected results, interpret the clinical samples as follows:

- 8.2.1. One or more prominent positive bands^a within the valid size range are reported as:
 - “Positive for the detection of a *BCL2/J_H* t(14;18) translocation consistent with the presence of a clonal cell population. In the context of overall diagnostic criteria, clonal cell populations can indicate the presence of a hematologic malignancy.”
- 8.2.2. An absence of positive bands^a within the valid size range is reported as:
 - “Negative for the detection of a *BCL2/J_H* t(14;18) translocation.”

^aNote: Criteria for defining a positive band are as follows:

- Products generated from samples that fall within the valid size range and produce a distinct and discrete band(s) are consistent with a positive band.

9. Limitations of Procedure

- This assay does not identify 100% of clonal cell populations.
- This assay cannot reliably detect less than one (1) positive cell per 100 normal cells.
- Always interpret the results of molecular clonality tests in the context of clinical, histological and immunophenotypic data.
- PCR-based assays are subject to interference by degradation of DNA or to inhibition of PCR due to EDTA, heparin, and other agents.

10. Expected Values

10.1. Expected Size of Amplified Products

- The amplicon sizes listed were determined using an ABI platform or by gel electrophoresis.

Table 8: Expected Size of amplified Products

Master Mix	Target	Control DNA	Catalog #	Product Size in base pairs (bp)
<i>BCL2/J_H</i> Tube A	Mbr of <i>BCL2/J_H</i>	Valid Size Range	---	100 - 2500
		IVS-0000 Polyclonal Control DNA	40920010	---
		IVS-0030 Clonal Control DNA	40881750	254, ~750 ^a
		IVS-0031 Clonal Control DNA	40881810	---
<i>BCL2/J_H</i> Tube B	3' Mbr of <i>BCL2/J_H</i>	Valid Size Range	---	100 - 2500
		IVS-0000 Polyclonal Control DNA	40920010	---
		IVS-P002 Clonal Control DNA	40900070	~140
		IVS-0030 Clonal Control DNA	40881750	---
<i>BCL2/J_H</i> Tube C	mcr of <i>BCL2/J_H</i>	Valid Size Range	---	100 - 2500
		IVS-0000 Polyclonal Control DNA	40920010	---
		IVS-0030 Clonal Control DNA	40881750	---
		IVS-0031 Clonal Control DNA	40881810	382, ~800 ^a
Specimen Control Size Ladder	Multiple Genes	Valid Size Range	---	100, 200, 300, 400, 600^b
		IVS-0000 Polyclonal Control DNA	40920010	100, 200, 300, 400, 600 ^b

^aNote: The ~750 bp band (*BCL2/J_H* Tube A) and the ~800 bp band (*BCL2/J_H* Tube C) are generally weak bands.

^bNote: Because smaller PCR fragments are preferentially amplified, it is not unusual for the 600 bp fragment to have a diminished signal or to be missing entirely.

10.2. Sample Data

The data shown below were generated using the master mix indicated. Amplified products were run on a 2% agarose gel.

BCL2/JH Tube A

Lane 1 = 100% IVS-0007
Lane 2 = 100% IVS-0030
Lane 3 = 1% IVS-0030
Lane 4 = 100% IVS-0000

Valid Size Range =
100-2500 bp

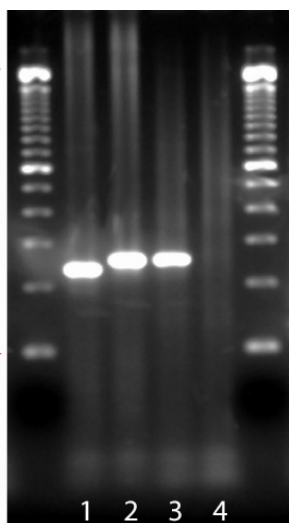


Figure 2. *BCL2/J_H* Tube A master mix.

BCL2/JH Tube B

Lane 1 = 100% IVS-0007
Lane 2 = 100% IVS-P002
Lane 3 = 1% IVS-P002
Lane 4 = 100% IVS-0000

Valid Size Range =
100-2500 bp

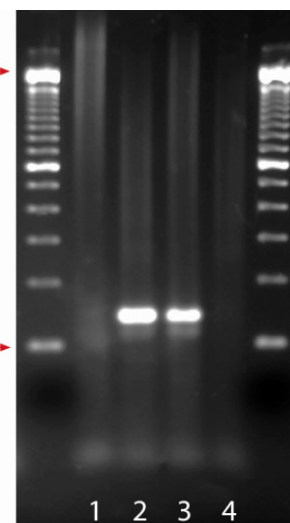


Figure 3. *BCL2/J_H* Tube B master mix.

BCL2/JH Tube C

Lane 1 = 100% IVS-0001
Lane 2 = 100% IVS-0031
Lane 3 = 1% IVS-0031
Lane 4 = 100% IVS-0000

Valid Size Range =
100-2500 bp

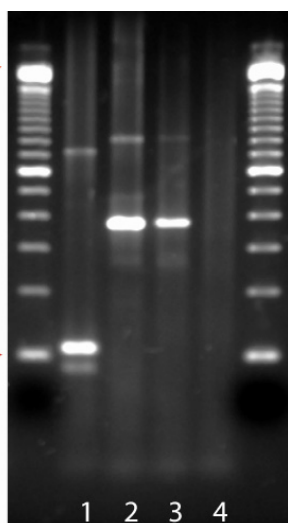


Figure 4. *BCL2/J_H* Tube C master mix.

Specimen Control Size Ladder - Unlabeled

Lane 1 = 100 bp ladder
Lane 2 = 50 bp ladder
Lane 3 = Genomic DNA
Lane 4 = Genomic DNA

~600 bp →
~400 bp →
~300 bp →
~200 bp →
Amplicon Sizes: ~100 bp →

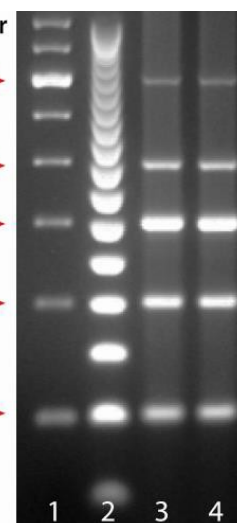


Figure 5. Specimen Control Size Ladder master mix.

11. Performance Characteristics

The initial evaluation of this assay was performed in three laboratories on DNA derived from 124 cases of follicular cell lymphoma (FCL) known to carry the t(14;18) translocation. 109 cases were identified with the *BCL2/J_H* fusion gene (88%) using this PCR assay. The final testing and evaluation was performed on samples in 11 independent laboratories. False positive results (0.4%) were observed in only 12 of 3036 analyses.

This IdentiClone *BCL2/J_H* Translocation Assay was found to be more sensitive than Southern blot analysis. Sensitivity differed slightly between the master mixes. However, overall sensitivity for the assay was determined to be between one (1) positive cell in 10^2 normal cells (1.0%) and one (1) positive cell in 10^3 normal cells (0.1%).

In conclusion, the performance characteristics of this robust three tube multiplex PCR assay were evaluated in order to maximize the detection of the t(14;18) breakpoint. This strategy is capable of amplifying across the breakpoint region in the majority of cases of follicular lymphoma with a cytogenetically defined translocations.

12. References

1. Miller, JE, Wilson, SS, Jaye, DJ, Kronenberg, M. (1999). An automated semiquantitative B and T cell clonality assay. *Molecular Diagnostics*. 4:101-117.

13. Technical and Customer Service

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	For <i>In Vitro</i> Diagnostic Use		Expiration Date
	Catalog Number		Authorized Representative in the European Community
	Reagent Volume		Manufacturer
	Lot Number		Consult Instructions for Use
	Storage Conditions		

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