Cat Nr/REF: KBI-10730

Critical region 1 (red):

Interpretation:

For professional use only **English**

Poseidon™ Repeat Free™ BCL6 (3g27) Break probe - Optimized for Tissue Hybridization -

Introduction: Chromosomal translocation at band 3g27 affecting the BCL6 locus are among the most frequent

changes in B-NHL. A FISH strategy using two differently labeled flanking BCL6 probes provides a robust, sensitive, and reproducible method for the detection of common and uncommon

abnormalities of BCL6 gene in interphase nuclei.

Intended use: The BCL6 (3q27) Break Probe is optimized to detect translocations involving the BCL6 gene

region at 3g27 in a dual-color, split assay on metaphase/interphase spreads, blood smears and

bone marrow cells

The probe is especially developed for use on paraffin sections and recommended to be used in combination with a Poseidon FISH Kit providing necessary reagents to perform FISH (KBI-60002. KBI-60003 or KBI-60001) for optimal results. For applications on metaphase/interphase

spreads, blood smears and bone marrow cells it is advised to use KBI-10607. The distal BCL6 gene region probe is direct-labeled with PlatinumBright550.

The **proximal BCL6** gene region probe is direct-labeled with Platinum *Bright* 495. Critical region 2 (green):

Reagent Poseidon probes are direct-labeled DNA probes provided in a ready-to-use format. Apply 10 µl of

probe to a sample area of approximately 22 x 22 mm.

Please refer to the Instructions for Use for the entire Poseidon FISH protocol.

Poseidon Repeat Free probes do not contain Cot-1 DNA. Hybridization efficiency is

therefore increased and background, due to unspecific binding, is highly reduced.

The BCL6 (3g27) Break probe is designed as a dual-color split probe to detect translocations at 3g27. A break is defined when a red/green or yellow fusion signals (F) splits into separate red and green signals. Only red and green signals which are more than one signal diameter apart from each other are counted as a break. Co-localized red/green or yellow signals identify the normal chromosome(s) 3.

Signal patterns other than those described above may indicate variant translocations or other complex rearrangements. Investigators are advised to analyze metaphase cells for the

interpretation of atypical signal patterns.

	Normal Signal Pattern	3q27 Split
Expected Signals	2F	1F1R1G

References: Butler et al, 2002, Cancer Res, 62; 4089-4094.

Sanchez-Izquierdo, 2001, Leukemia, 15; 1475-1484.

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Application Manual

KBI-10730 ON BCL-6 (3g27), Break (tissue)











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