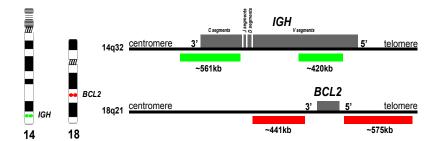


IGH/ BCL2

Intended Use

The *IGH/BCL2* DNA-FISH Probe is designed to detect the translocation between the *IGH* gene located on 14q32 and the *BCL2* gene located on 18q21, using fluorescence *in situ* hybridization (FISH). The translocation between the *IGH* and *BCL2* gene is designated as t(14;18)(q32;q21) and is the hallmark of follicular lymphoma (FL). The t(14;18) translocation is more frequent in lower FL grades, such as FL grades 1 and 2 (88%), than in higher grades, such as FL 3b (4-13%).^[1] The translocation is also detected in 30% of diffuse large B-cell lymphoma (DLBCL) cases, but is less frequently detected in other non-Hodgkin lymphomas.^[1,2] By conventional cytogenetics, the t(14;18)(q32;q21) translocation that involves the *IGH* and *BCL2* genes is indistinguishable from the t(14;18)(q32;q21) translocation that involves the *IGH* and *MALT1* genes. Such translocations can be distinguished by using the respective DNA-FISH probes.^[1]



Schematic of the IGH/BCL2 DNA-FISH Probe:

Horizontal red and green bars indicate the regions covered by the probes (approximate to scale, GRCh37/Hg19/2009). The directly labeled *IGH* (green) and *BCL2* (red) probes flank the breakpoints in the *IGH* and *BCL2* genes, respectively.

Signal Interpretation

In normal diploid metaphase and interphase nucleus, two green and two red signals would be observed corresponding to the two normal homologous chromosomes 14 and 18, respectively (Figures 1 and 2). Upon translocation, the most commonly observed pattern is a single green and red signal, representing the normal chromosomes 14 and 18, and two fusion signals (red/green or yellow) representing the two translocated chromosomes. It is recommended to confirm variant pattern or atypical signal patterns by metaphase analysis whenever possible.

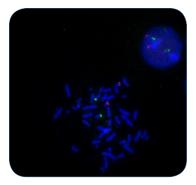


Figure 1: Normal diploid metaphase and interphase nucleus (from normal peripheral blood specimen) with 2 red (*BCL2*) and 2 green (*IGH*) signals.

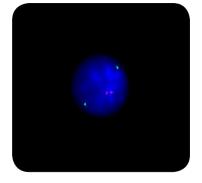


Figure 2: Normal diploid interphase nucleus (from bone marrow specimen) with 2 red (*BCL2*) and 2 green (*IGH*) signals.

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Fluorophore	Excitation _{max}	Emission max
Green	496 nm	520 nm
Red	580 nm	603 nm
DAPI	360 nm	460 nm

Fluorescence Microscopy Filter Requirements

References

1. Heim, S., Mitelman, F. (Ed). Cancer Cytogenetics, 2009 (3rd Edition). Wiley-Blackwell, New Jersey. P. 297-358.

2. Fan, Y.S., Rizkalla, K. Cancer Genet Cytogenet, 2003. 143(1): p. 73-9.

Instructions for use are available at www.cancergeneticsitalia.com

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