Cat Nr/REF: KBI-10003

English For professional use only

Poseidon™ Repeat Free™ FIP1L1 – CHIC2 – PDGFRA (4q12) Deletion, Break probe

Introduction: The deletion of the CHIC2 locus generates a fusion FIP1L1-PDGFRA gene giving

raise to a novel tyrosine kinase. This deletion has been observed in patients with idiophatic hypereosinophilic syndrome (HES), chronic eosinophilic leukemia (CEL), systemic mast cell disease, and chronic myeloproiferative disorders (CMPD).

Intended use: The FIP1L1-Chic2-PDGFRA probe is optimized to detect the Chic2 deletion at 4q12

associated with the FIP1L1/PDGFRA fusion in a dual-color, dual-fusion assay on metaphase/interphase spreads, blood smears and bone marrow cells, It also

detects translocation involving the FIP1L1 and PDGFRA region.

The probe is 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.

Critical region 1 (red): The CHIC2 (4q12) gene region is direct-labeled with Platinum Bright 550.

Critical region 2 (green): The FIP1L1 and PDGFRA (4q12) gene regions are direct-labeled with

PlatinumBright495.

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.

Interpretation: The FIP1L1-Chic2-PDGFRA probe is designed as deletion probe, where loss of

CHIC2 region is observed as loss of a red signal leaving a green signal at 4q12. Split of the probe in case of a translocation at 4q12 results in a break of one fusion signal, observed as a 2F1G signal pattern. Single color fusion (F) signals will identify

the normal chromosomes 4.

Signal patterns other than those described above may indicate variant translocations or other complex rearrangements, such as hyperdipoloidy for chromosome 4. Investigators are advised to analyze metaphase cells or use additional probes (e.g. SE Chromosome 4) for the interpretation of atypical signal

patterns.

	Normal Signal Pattern	Del(4q12)	Translocation at 4q12	Translocation + Deletion at 4q12*
Expected Signals	2F	1F1G	2F1G	1F2G

^{*} hyperdiploidy for Chromosome 4 has to be verified for this signal constellation

References: Cools et al, N Engl J Med, 2003, 348, 1201-1214. Godlib et al. Blood. 2004. 103. 2879-2891.

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REPEAT-FREET FISH PROBES

Application Manual

KBI-10003 ON FIP1L1-CHIC2-PDGFRA (4q12) Del, Break











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