

SALSA® digitalMLPA™ Probemix D007 Acute Lymphoblastic Leukemia



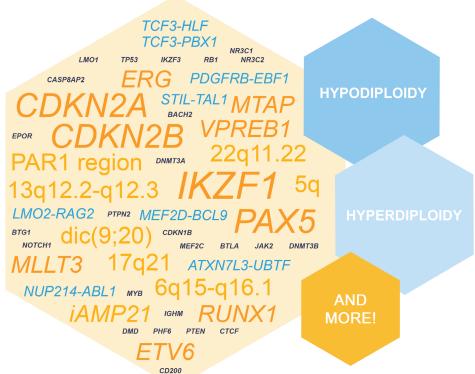
- ✓ High resolution coverage of 73 ALL-related genes and 8 regions
- ✓ Gross copy number detection with 250 probes across the genome
- ✓ High dynamic range for CNA detection
- ✓ Quick turnaround time of 48-72 hours

D007 Acute Lymphoblastic Leukemia is a new generation panel detecting an extensive number of ALL-associated genes and regions, as well as gross copy number alterations (CNAs). Bring down lab-handling time and optimise your resources with its targeted approach.

Recurrent and clonal genetic alterations in different subtypes of Acute Lymphoblastic Leukemia (ALL) are well-characterised, and associated with differences in disease outcome. MRC Holland's SALSA® MLPA® technology has become a prime method in studying CNAs in ALL, especially *IKZF1*^{del}. D007 Acute Lymphoblastic Leukemia is based on the groundbreaking SALSA® digitalMLPA™ technology, which combines the broad scale of next generation sequencing with the unparalleled sensitivity in CNA detection of MLPA. In a single reaction, D007 Acute Lymphoblastic Leukemia can detect:

- Partial chromosome gains, losses and high-level amplifications
- Hyperdiploidy and hypodiploidy
- Intrachromosomal gene fusions
- Intragenic CNAs

digitalMLPA can be run on any Illumina platform, can be combined with NGS libraries on the same flow cell, and is analysed using our free Coffalyser digitalMLPATM software, meaning no bioinformatic skills are required.



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- ➤ Combines genes and regions included in the well-established MLPA probemixes P335 ALL-IKZF1, P202 IKZF1-ERG, P327 iAMP21-ERG, P329 CRLF2-CSF2RA-IL3RA, P383 T-ALL, P056 TP53.
- ➤ Allows for the examination of B-cell differentiation and cell cycle control genes, T-ALL-associated alterations, iAMP21 and CNAs of the PAR1 region, and much more.
- ➤ Examines subtelomeric, pericentromeric and middle regions of the chromosomal arms to detect gross CNAs and hyper-/hypodiploidy by using 250 karyotyping probes.

D007 Acute Lymphoblastic Leukemia: target genes & regions Karyotyping probe CNA probe ■ Chr. region (# of probes) CNA & fusion probe Probe targeting region border BCL9 × q16.1 **5q** (26) PBX1* PHA1 ETV6 13q12.2 CDKN1B * -q12.3 * (32) ■ LMO2 FLT3 * BAG2 CTCF ABL1 NUP214 NOTCH1 IL3RA P2RY8 GYG2 BCL2L1 * IKZF3 ATXN7L3 * VPREB1 IGLV7-43 * UBTF * HNRNPUL1*

Required materials

- 20 ng of sample DNA, peripheral blood or bone marrow-derived
- SALSA® digitalMLPA™ probemix, reagents and barcode plates
- · Thermocycler with heated lid
- Illumina sequencing platform (all types), flow cell and reagents

References

Benard-Slagter A et al. (2017). J Mol Diagn. 19:659-72.

PHF6

For a full list of references, see assay page on mrcholland.com.

