Product Description
SALSA® Binning DNA SD096-S01

Version S01

Catalogue number
- **SD096:** SALSA Binning DNA, 6 reactions

Precautions and warnings
For professional use only. Always consult the most recent product description AND the corresponding probemix product description AND the MLPA General Protocol before use: www.mrcholland.com. Binning DNA is not known to contain any harmful agents.

Safety data sheet
Based on the concentrations present, none of the ingredients are hazardous as defined by the Hazard Communication Standard. **A Safety Data Sheet (SDS) is not required for these products**: none of the preparations contain dangerous substances (as per Regulation (EC) No 1272/2008 [EU-GHS/CLP] and amendments) at concentrations requiring distribution of an SDS (as per Regulation (EC) No 1272/2008 [EU-GHS/CLP] and 1907/2006 [REACH] and amendments). If spills occur, clean with water and follow appropriate site procedures.

General information
The SALSA Binning DNA SD096 is a research use only (RUO) reagent to be used in combination with SALSA MLPA probemix P496-A1 KMT2A, a SALSA MLPA Reagent Kit and Coffalyser.Net™ analysis software for the processes of linking all probe signals to their identity by use of the probe lengths. SD096 contains the targets of all probes included in the above-listed probemix, including the mutation-specific probe target *ASXL1* c.1934dupG.

Binning DNA should never be used as a reference sample in the MLPA data analysis. Neither should it be used in quantification of mutation signals.

Experimental set up
MLPA reactions for binning purposes should be performed with 5 µl of Binning DNA. Inclusion of one reaction with SALSA Binning DNA SD096 in the initial MLPA experiment is essential as it can aid in data binning of the peak pattern when using Coffalyser.Net software. Furthermore, Binning DNA should be included in the experiment whenever changes have been applied to the set-up of the capillary electrophoresis device (e.g. when a different polymer type is used).

Data analysis
Coffalyser.Net software should be used for analysis of MLPA experiments. When performing the fragment analysis step in Coffalyser.Net, select SD096 in the *bin smpl* –column. By selecting the SD096 sample as your binning sample, probes will be correctly identified in the peak pattern across all samples. Coffalyser.Net software is freely downloadable at www.mrcholland.com.

Binning DNA content
SD096 consists of a mixture of female genomic DNA from healthy individuals and a titrated amount of plasmid DNA that contains a partial sequence of the *ASXL1* gene. This partial sequence includes one mutation that will be detected by the mutation-specific probe present in the above-listed probemix. See Table 1 and the corresponding probemix product description for more details on the mutation-specific probe target present. The indicated mutation-specific probe will generate a signal on SD096. The indicated mutation-specific probe will generate a signal on SD096.

Please note that the plasmid DNA also contains the target sequence of the 105 nt chromosome Y specific control fragment. As a result, the 100 and 105 nt control fragments indicate the presence of two copies chromosome X and one copy chromosome Y.

Table 1. Mutation-specific probe target in Binning DNA SD096-S01

Probemix	Gene/Exon	Probe length (nt)	Probe ID	Probemix version	Details
P496	*ASXL1* exon 13 ‡	160	18261-SP0848-L26260	A1	c.1934dupG

Note: Please consult the corresponding probemix product description for more information about exon numbering, mutation nomenclature and gene transcripts used.

‡ An unspecific peak might be detected at one nucleotide shorter length from the expected length of this *ASXL1* c.1934dupG mutation-specific probe at 160 nt. Please analyse the peak pattern carefully when making calls for this mutation-specific probe.

More information: www.mrcholland.com; www.mrcholland.eu	
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Implemented changes in the product description
Version S01-01 - 10 November 2023 (03) - Not applicable, new document.