

An overview of the MLPA workflow

Α Day 1 Sample DNA denaturation **Probe Hybridization** Overnight incubation Day 2 Neighboring probes ligation PCR amplification Capillary electrophoresis After Day 2 QC and Data analysis by Coffalyser.Net software

B BRCA1 and BRCA2 relevant probe sets

Discovery sets:
P002-D1-0918 BRCA1-v01
P090-C1-0319 BRCA2-v01
P239-C1-0319 BRCA1 Region-v01 (Extended to BRCA1 up- and downstream genes)

C Control samples

Control samples	Purposes
3 normal DNA samples with intact BRCA1 and BRCA2	To evaluate the overall assay performance and as normalization control for calculating the final ratio (FR)
SALSA Artificial duplication DNA: SD024-S01	As positive duplication control for P002:BRCA1 and P090:BRCA2 MLPA assays
BRCA009 Blood DNA (BRCA009-B)	As positive duplication and deletion control for P239:BRCA1 up- and downstream MLPA assay

BRCA010

MLPA assay QC checklist

QC1.Electrophoresis data

Strength of capillary electrophoresis (CE) signal:

PASSED

Size standard and CE quality: **PASSED**

Regularity, distribution and shape of peaks:

PASSED

MLPA peaks fall into the optimal range of our CE platform: **PASSED**

QC2.QC Fragments check

Benchmark fragment at 92nt: DETECTED

Four DNA quantity fragments:

DETECTED and

ABOVE THRESHOLD

Two DNA denaturation fragments:

DETECTED and

ABOVE THRESHOLD

Chromosome X-specific fragment (For sex checking):

MATCHED

Chromosome Y-specific fragment (For sex checking):

MATCHED

P002:BRCA1 (QC PASSED) P090:BRCA2 (QC PASSED)

Derived from the MLPA checklists generated by Daisy

QC3.Control samples

A. Positive control

(SD024) Artificial duplication in P002:BRCA1 and BRCA2:P090:

DETECTED

(BRCA009-B) In-house positive control for P239: DETECTED

Two chromosome X-specific

fragment: DETECTED

One chromosome Y-specific

fragment: DETECTED

B. No DNA control

Only five major peaks <121nt:

DETECTED

C. Reference samples

Three normal references:

INCLUDED

Variation of individual probe among three references (SD<=0.10):

PASSED

Normal BRCA1/2 copy number:

DETECTED

D. Non-BRCA specific reference probes

Variation of reference probes

(SD<=0.10): PASSED

Final ratio of each reference probe fall within normal copy number

range: PASSED

QC4.Data quality check in Coffalyzer.Net

Maximum FRSS, FMRS and CAS quality scores:

PASSED

No missing probe:

PASSED

Bin size correct and no peak overlapping:

PASSED

Lot number of the probes was accurately shown:

PASSED

P239:BRCA1 up- and downstreams (QC PASSED)