



## CLART® CMA

CLART® CMA is an *in vitro* diagnostics test line of products for the detection of mutations in genes associated with response to therapy in cancer patients.

### FEATURES OF CMA DIAGNOSTIC KITS :

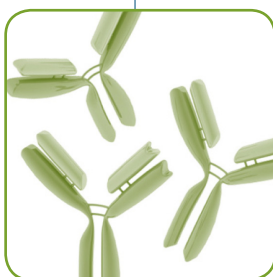
- All kits have been validated for automatic and manual DNA extraction from FFPE samples and cell lines.
- Mutational status can be detected for single or multiple genes just combining references.
- High sensitivity and specificity.
- Avoids unnecessary toxicity caused by improper selected antitumor therapy, as well as its associated costs.
- Each mutation is detected in triplicate avoiding unspecific bindings.
- Three internal quality controls included per sample:
  - **Genomic DNA control:** validates the extraction performance.
  - **Amplification control:** avoids false negative results.
  - **Biotin markers:** check the proper performance of the visualization reagents provided with the kit.

- Short turnaround time (5 hours).
- Reduces the amount of sample required. All mutations from any kit can be detected in a single array.
- Compatible with any GENOMICA automation system.

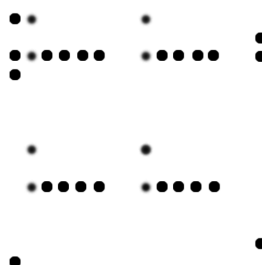
### DATA MANAGEMENT :

- Automatic reading and interpretation of results (CAR®).
- User-friendly report format (html, bmp).
- Printable, exportable and storable files.

### REPORTING RESULTS :



► Report and image obtained by CAR® reader.



Genomica S&T CAR, CAP System		Working list Run ID Run ID: 2013_07_18_11_40_57_153 Date and time: Thu Jul 18 11:41:02 2013		GENOMICA	
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Pin	Array ID	Array	Array ID	Wavelength	Sample reference	Result
A1	10913	CLART CMA	00000000010913	1	1	Out alignment failed. The image could not be analyzed. The result is not valid.
B1	10913	CLART CMA	00000000010913	2	2	Out alignment failed. The image could not be analyzed. The result is not valid.
C1	10913	CLART CMA	00000000010913	3	3	Out alignment failed. The image could not be analyzed. The result is not valid.
D1	10913	CLART CMA	00000000010913	4	4	Out alignment failed. The image could not be analyzed. The result is not valid.
E1	10913	CLART CMA	00000000010913	5	5	Out alignment failed. The image could not be analyzed. The result is not valid.
F1	10913	CLART CMA	00000000010913	6	6	Out alignment failed. The image could not be analyzed. The result is not valid.
G1	10913	CLART CMA	00000000010913	7	7	Out alignment failed. The image could not be analyzed. The result is not valid.
...	...	...	...	...	...	Out alignment failed. The image could not be analyzed. The result is not valid.



# CLART<sup>®</sup> CMA KRAS · BRAF · PI3K & NRAS · iKRAS

Specific detection of somatic mutations in oncogenes determining response to therapy in colorectal cancer patients.

## KRAS

- G12A
- G12D
- G12R
- G12C
- G12S
- G12V
- G13D
- Q61H (A>T)
- Q61L

## BRAF

- V600E
- V600K

## PI3K

- E542K
- E545D
- E545K
- H1047R

## NRAS

- G12D
- G16R
- G61K
- G61L
- Q61H (A>T)
- K117N (G>C)
- A146T (G>A)

## iKRAS

- Q61H (A>C)
- A146V
- A146T
- K117N (A>C)

## ● MAIN FEATURES :

### KRAS · BRAF · PI3K :

- Diagnostic specificity close to 100% in all point mutations.
- Diagnostic sensitivity from 87% to 100% in BRAF and PI3K

### NRAS · iKRAS :

- Detects the presence of the most prevalent mutations of NRAS and infrequent KRAS with a diagnostic sensitivity and specificity ≥98% .

## ● ORDERING REFERENCES :

### KRAS · BRAF · PI3K :

#### CLART<sup>®</sup> CMA KRAS

Amplification 24 tests: CS-0412-24

#### CLART<sup>®</sup> CMA BRAF

Amplification 24 tests: CS-0512-24

#### CLART<sup>®</sup> CMA PI3K

Amplification 24 tests: CS-0612-24

#### CLART<sup>®</sup> CMA KBP Array

Genotyping 24 test: CS 0712-24

### NRAS · iKRAS :

#### CLART<sup>®</sup> CMA NRAS · iKRAS

Amplification 24 tests: CS-0114-24

#### CLART<sup>®</sup> CMA NiK Array

Genotyping 24 tests: CS-0214-24

\* Panels can be run and purchased separately.

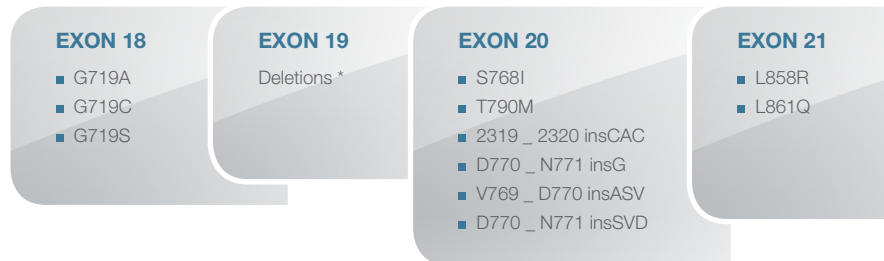
## ● BIBLIOGRAPHY :

1. "Biomarkers Predicting Clinical Outcome of Epidermal Growth Factor Receptor-Targeted Therapy in Metastatic Colorectal Cancer". *Review J Natl Cancer Inst* 2009;101:1308-1324.
2. KRAS, BRAF, PIK3CA, and PTEN mutations: implications for targeted therapies in metastatic colorectal cancer". *Lancet Oncol* 2011; 12: 594-603
3. "NRAS and KRAS testing by a new diagnostic method to detect point mutations in colorectal cancer specimens: CLART NRAS-iKRAS. 2014 ASCO Annual Meeting Proceedings



# CLART® CMA EGFR

Specific detection of somatic mutations, deletions and insertions in EGFR determining response to therapy in non-small-cell lung cancer patients.



\*6223, 12370, 12370, 6255, 12384, 12382, 6225, 12678, 6218, 12728, 6220, 12419, 6210, 13556, 12386, 12385, 18427, 12403, 12383, 6254, 13551, 12367, 12422, 12387, 26038, 13552, 12416, 23571. (According to COSMIC ID nomenclature).

## ● MAIN FEATURES :

- An average diagnostic sensitivity  $\geq 92\%$  in the most of the mutations.
- Diagnostic specificity  $\geq 99\%$  .

## ● ORDERING REFERENCES :

### CLART® CMA EGFR Amplification

24 tests: CS-1014 –24

### CLART® CMA EGFR Genotyping

24 tests: CS-1114-24

## ● BIBLIOGRAPHY :

1. ZHANG, Z., STIEGLER, A. L., BOGGON, T. J., KOBAYASHI, S. & HALMOS, B. (2010) EGFRmu-tated lung cancer: a paradigm of molecular oncology. *Oncotarget*, 1, 497-514. PAO, W., MILLER, V.A., POLITI, K.A., RIELY, G.J., SOMWAR, R.,
2. ZAKOWSKI, M.F., KRIS, M.G., VARMUS, H. (2005) Acquired resistance of lung adenocarci-nomas to gefitinib or erlotinib is associated with a second mutation in the EGFR kinase domain. *PLoS Medicine*, 2, 3, e73.
3. YASUDA, H., KOBAYASHI, S., COSTA, D.B. (2012) EGFR exon 20 insertion mutations in non-smallcell lung cancer: preclinical data and clinical implications. *Lancet Oncol*, 13: e23–31.
4. PEREZ-MORENO, P., BAMBRILLA E., THOMAS, R., SORIA, J.C. (2012) Squamous Cell Carci-noma of the Lung: Molecular Subtypes and Therapeutic Opportunities. *Clin Cancer Res*. 2012 May 1;18(9):2443-2451. ROSENZWEIG, S.A., ATREYA, H.S. (2010) Defining the pathway to insulin-like growth factor system targeting in cancer. *Biochem Pharmacol*, 15;80(8):1115-24.



# CLART<sup>®</sup> CMA BRAF · MEK1 · AKT1

GENOMICA.COM



Detection of specific somatic mutations in oncogenes  
determining response to therapy in melanoma patients.

## BRAF

- V600E
- V600K

## MEK1

- I111S
- P124S
- E203K

## AKT1

- Q79K

## ● MAIN FEATURES :

- Diagnostic sensitivity in BRAF > 93% .
- The obtained diagnostic specificity for all the mutations in the MEK1 and AKT1 genes is 100% .
- Diagnostic specificity ≥99% .

## ● ORDERING REFERENCES :

### BRAF · MEK1 · AKT1 :

#### CLART<sup>®</sup> CMA BRAF

Amplification 24 tests: CS-0216-24

Genotyping 24 tests: CS-0716-24

#### CLART<sup>®</sup> CMA BRAF · MEK1 · AKT1

Amplification 24 tests: CS-0316-24

Genotyping 24 tests: CS-0716-24

## ● BIBLIOGRAPHY :

1. "Advances in personalized targeted treatment of metastatic melanoma and non-invasive tumor monitoring." Klinac D, Gray ES, Millward M, Ziman M. *Front Oncol.* 2013 Mar 19;3:54. doi: 10.3389/fonc.2013.00054. eCollection 2013.
2. "Effects of AKT inhibitor therapy in response and resistance to BRAF inhibition in melanoma". Lassen A, Atefi M, Robert L, Wong DJ, Cerniglia M, Comin-Anduix B, Ribas A. *Mol Cancer.* 2014 Apr 16;13:83. doi: 10.1186/1476-4598-13-83.
3. Detection of BRAF V600 mutations in melanoma: evaluation of concordance between the Cobas<sup>®</sup> 4800 BRAF V600 mutation test and the methods used in French National Cancer Institute (INCa) platforms in a real-life setting.

