Diagnostic Utility of Liquid Biopsy to Analyze Circulating Tumor DNA Via Next-Generation Sequencing

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Recent oncology advancements in next-generation sequencing (NGS) technologies for analyzing circulating tumor DNA (ctDNA) from liquid biopsy (LB) samples have garnered significant interest due to their potential applications across various stages of the disease: cancer screening, diagnosis at early stage, profiling tumor mutations, monitoring treatment response, detecting minimal residual disease, and selecting the most appropriate treatments based on genomic alterations.

Unlike tissue biopsy, the current standard method for cancer diagnosis, LB is a minimally invasive. It allows for real-time evaluation of the tumor's genomic profile without invasive and costly procedures. The most commonly studied sample type for LB is plasma from blood.





This report evaluates the potential of LB for diagnostic purpose by analyzing ctDNA via NGS in breast, lung, and colorectal cancer, aiming to assess its inclusion in the Catalan Health System (CHS) benefit package. Specific objectives are: **1**. To describe the diagnostic utility of ctDNA from LB by NGS in breast, lung and colorectal cancer, **2**. To describe the diagnostic utility of commercial ctDNA LB-based NGS panels, **3**. To identify the clinical scenarios in which LB to analyse ctDNA could be performed as the first choice in breast, lung and colorectal cancer, and **4**. To plan a budgetary impact analysis (BIA) of the inclusion of ctDNA LB-based NGS panels in breast, lung and colorectal cancer.

1 Umbrella review of the diagnostic utility of ctDNA from LB by NGS in breast, lung and colorectal cancer



3 Narrative review of targeted Scientific Societies (SS) recommendations and expert opinions



RESULTS



Diagnostic utility of ctDNA analysis from LB by NGS vs. tissue biopsy in breast, lung and colorectal cancer

| Breast cancer | | Lung cancer (NSCLC) | | Colorectal cancer | |
|--|---|---|---|---|--|
| SR with MA included | Galvano et al. 2022 | SR with MA/HTA report included | Esagian 2020, Sebastiao et al. 2020, Ontario Health 2020 | SR with MA included | Galvano et al. 2019, Ye et al. 2021 |
| Use of the ctDNA LB-based NGS test | Tumor profiling in advanced cancer stage | Use of the ctDNA LB- based NGS test | Tumor profiling in advanced cancer stage | Use of the ctDNA LB- based NGS test | Tumor profiling in advanced cancer stage |
| Quality assessment | Very low | Quality assessment | Low to Very low | Quality assessment | Very low |
| Diagnostic utility data vs. tissue biopsy | <i>PIK3CA</i> Sensitivity: 0.83, Specificity: 0.98 | Diagnostic utility of ctDNA LB-based NGS | <i>EGFR, ALK, ROS-1, BRAF, RET, MET</i> exon 14 | Diagnostic utility of ctDNA LB-based NGS | RAS Sensitivity: 0.75, Specificity: 0.91, AUC of |

AUC of SROC: 0.98

2 Commercial NGS panels

ctDNA LB-based NGS panels reviewed: FoundationOne Liquid CDx, Oncomine[™] Lung cfDNA Assay Oncomine[™] Colon cfDNA Assay, Oncomine[™] Lung Cell-Free Total Nucleic Acid Assay, Oncomine[™] Pan-Cancer Cell-Free Assay, Oncomine[™] Precision Assay, Oncomine[™] Precision Assay GX (Ion Torrent, Guardant 360, AVENIO ctDNA Panel Targeted Kit, AVENIO ctDNA Expanded Kit, AVENIO ctDNA Surveillance Kit, TruSight Oncology 500 ctDNA

None of the reviewed panels have an intended use for cancer screening or early stage diagnosis

| NGS test with CE man | FoundationOne Liquid CDx (Roche), Guardant 360 CDx k (Guardant Health). Intended use: Tumor mutation profiling for treatment selection |
|----------------------------|--|
| Mutations detection | Oncomine[™] Lung cfDNA Assay and Oncomine[™] Colon cfDNA Assay: only single nucleotide variants (SNV) and indels. Other panels: also copy number variations (CNV) and gene fusions. |
| Diagnostic utility data | Heterogeneity between tests in diagnostic utility data. Sensitivity (7 panels): 89.2-99.9%; Specificity (4 panels): 99.6-99.9%; Positive Predictive Value (PPV) (7 panels): 99-100%; Negative PV (NPV) (2 panels): 94-100%; PPA (2 panels): 98-100%; Negative percent agreement (NPA) (2 panels): 93.9-100%; Overall Predictive value (1 panel): 93.4% |

data vs. tissue biopsySensitivity: 0.76, Specificity: 0.99, AUC ofSROC: 0.99

EGFR T790M Sensitivity: 0.78, Specificity: 0.85

ALK, BRAF, ERBB2, EGFR, KRAS, MET, RET *i ROS 1*Positive Percent Agreement, PPA: 53-68%
depending on the analysed gene

3 SS recommendations and expert opinions about LB in breast, lung and colorectal cancer

SS reviewed: SEOM (Sociedad Española de Oncología Médica), ESMO (European Society of Medical Oncology), ASCO (American Society of Clinical Oncology)

| Cancer creening or early liagnosis | Not recommended |
|---|--|
| umor profiling | Recommended Analysis of ctDNA from LB in advanced disease for cancer genotyping and guide treatment decisions. The use of BL as the first choice is only recommended in specific cases, such as not having tissue samples or when urgent |

cases, such as not having tissue samples or when urgent therapeutic decisions need to be made.

In the specific case of breast cancer, it is recommended to analyze ESR1 in an advanced state to the progression of hormonal therapy,

data vs. tissue biopsy SROC: 0.81

KRAS Sensitivity: 0.65, Specificity: 0.88, AUC of SROC: 0.86

4 Budgetary impact analysis protocol

| BIA features | BIA features | | | | |
|---------------------------------------|--|--|--|--|--|
| Data obtention | Narrative review | | | | |
| Objective | To estimate the monetary units that should be disbursed by the CHS | | | | |
| Perspective | Healthcare payer | | | | |
| Population | Patients likely to be diagnosed or already diagnosed with lung, breast or colorectal cancer of any age and any stage of the disease. | | | | |
| Temporal horizon | 1 year | | | | |
| Comparison scenarios | Different scenarios based on different % expansion of LB use vs. tissue biopsy. A complementary or alternative use of the LB use will also be considered. Results will be differentiated according to the use of LB in diagnostic/screening or tumor profiling. | | | | |

preferably through the determination of ctDNA.

E / CONCLUSIONS

- Existing evidence of LB to analyse ctDNA via NGS from SR with MA is limited to the molecular profile application and for certain target genes.
- Most NGS panels allow the detection of SNV, indels, CNV, and fusions.
- SS recommendations suggest that LB may be complementary to tissue biopsy.
- The BIA will help decision-makers understand the potential costs and economic benefits in different scenarios, facilitating better resource management.



Declaration of funding: This HTA report was performed with funding from the Catalan Ministry of Health

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