# Liquid gold: blood-based biopsies make headway

Jane de Lartigue, PhD

Pathologic and, increasingly, molecular analysis of tumor tissue biopsies is the gold standard in initial diagnosis of cancer, but liquid biopsies, which analyze tumor-derived material circulating in the bloodstream are gaining traction. Here, we discuss the current state of development of this complementary and potentially alternative approach to tumor analysis.

#### Liquid biopsy gaining traction

Biopsies enable oncologists to gather information about a potential or established tumor, including confirmation of the presence of cancerous tissue and determination of its histological characteristics, such as tumor grade and stage, as well as its molecular features, such as the presence of certain gene mutations. Ultimately, this information can be put to use in determining the most appropriate course of treatment.

The current gold standard is a tissue biopsy that typically involves an invasive procedure to permit the collection of a piece of tumor tissue. Yet, tissue biopsies are not always feasible because of the location of the tumor or the poor performance status of many patients with advanced disease. They also provide only a snapshot of the disease at the time at which they were taken and don't necessarily reflect the genetic heterogeneity or evolution of a tumor over time.

The detection of components that are derived from the tumor circulating in the blood of cancer patients had fueled the idea of blood-based diagnostics in oncology – so-called liquid biopsies. These have rapidly gained traction in the past several decades as a less expensive (the cost of performing genomic analyses on blood samples is at least an order of magnitude less than on tissue samples), less invasive (requiring only a simple blood draw) alternative source of information about tumors.<sup>1</sup>

As researchers have refined the ability to exploit liquid biopsies, commercial interest has been piqued. More than 35 companies within the United States alone are developing liquid biopsies, and it's easy to see why with a market projected to be in the many billions of dollars.<sup>2</sup>

#### Seeking out tumor clues in the blood

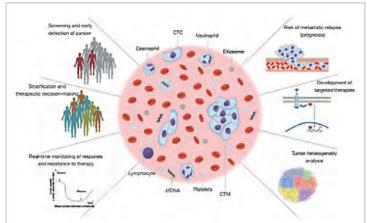
Liquid biopsies consist of a 10-15 mL blood sample drawn into a tube that contains an anticoagulant and it can contain several different types of tumorassociated material. Thus far, two components – circulating tumor cells (CTCs) and circulating tumor DNA (ctDNA) – have formed the cornerstone of liquid biopsies. At present, it is not clear whether these components are released randomly, as a byproduct of tumor cell death or if they are released as part of a specific biologic process, such as for the colonization of metastatic sites. It reality, it may be a little of both, and active dissemination may be particularly relevant for CTCs, among which are postulated to be a population of cancer stem cells that can initiate distant metastases.<sup>3,4</sup>

The discovery of CTCs dates back to the 1860s, when cells that were morphologically identical to the tumor were identified in the blood of a patient with metastatic cancer. Their potential significance was not fully realized until a few decades ago, when they were found to exist from early on in the course of disease.<sup>3,4</sup>

CTCs, which can be either single cells or clusters of cells known as microemboli, have a short half-life in the bloodstream – less than 2 ½ hours – and are also extremely rare (1 mL of blood contains 1-10 CTCs) against a background of many millions of normal cells. Thus the detection and isolation of CTCs presents a significant challenge. More than 40 different platforms are being developed for the isolation and enrichment of CTCs. For the most part, these use a method called positive selection to pick out CTCs.<sup>1,3,4</sup>

Positive selection exploits the biological or physical properties that are specific to CTCs and absent in normal cells, for example, the presence of a specific tumor-associated antigen on their surface or differences in size, density or electric charge. The limitations of this method are that, not only do you

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**FIGURE 1** Potential clinical utility of CTC and ctDNA analyses in cancer care. Reproduced with permission: Calabuig-Farinãs S, Jantus-Lewintre E, Herreros-Pomares A, Camps C. Circulating tumor cells versus circulating tumor DNA in lung cancer – which one will win? Reproduced with permission. Calabuig-Fariñas et al. Transl Lung Cancer Res. 2016;5(5):466-482.

CTC, circulating tumor cell; cfDNA, circulating free DNA; CTM, circulating tumor microemboli

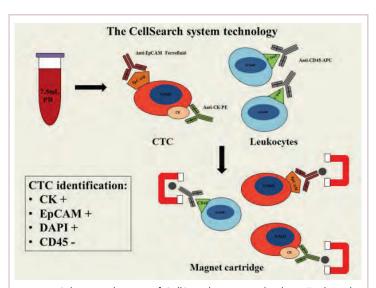


FIGURE 2 Schematic diagram of CellSearch system technology. To date, the CellSearch (Janssen Diagnostics/Veridex LLC) system is the only CTC assay approved by the US Food and Drug Administration. It is a semi-automated system able to detect and enumerate CTCs. Analysis is carried out in 7.5 mL of whole blood using three different antibodies against the epithelial cell adhesion molecule (EpCAM), cytokeratin (CK) and CD45. Peripheral blood is mixed with magnetic iron nanoparticles coated with anti-EpCAM antibody to confer magnetic properties to the epithelial cells. To detect these cells and exclude the presence of leukocytes, anti-CK and anti-CD45 fluorescent antibodies are used. In addition, cell nuclei are labelled with the DAPI nuclear dye to allow microscopic identification of the relevant cell fraction. A strong magnetic field is then applied to separate the cell population that can be detected and counted by digital fluorescent microscopy. Reproduced with permission. Truini et al. Clinical applications of circulating tumor cells in lung cancer patients by CellSearch system. Front Oncol. 2014;4(242):1-5.

PB, peripheral blood; CTC, circulating tumor cell; CK-PE, cytokeratins-phytoerythrin; CD45-APC, cluster of differentiation 45-allophycocyanin; N-DAPI, nucleus stained with DAPI. need to know something about CTCs to begin to understand what makes them truly unique and ensure only isolation of CTCs, but their phenotype is also thought to be continually changing.<sup>1,3,4</sup>

In recent years, the focus has shifted toward technologies that use negative depletion, meaning that they target the other types of cells in the blood sample and filter those away until only the CTCs are left behind. The most advanced are devices that use microfluidic technology to sort the cells, such as the CTC-iChip system being developed by researchers at Massachusetts General Hospital in Boston.<sup>5</sup>

ctDNA consists of small fragments of nucleic acids that are not contained within a cell or associated with cell fragments and is thought to be present in 50%-90% of patients, depending on the type of cancer they have. ctDNA has a similarly short half-life in the circulation to CTCs and, like CTCs, ctDNA is present at very low levels in the bloodstream. Although levels of ctDNA have been shown to increase with increasing tumor burden, it is still often obscured by the presence of other cell-free DNA derived from non-tumor cells.

ctDNA can be distinguished from other cell-free DNA by the presence of somatic mutations and a number of highly sensitive methods have been developed to detect them, including the amplification-refractory mutation system (ARMS); digital polymerase chain reaction; and the beads, emulsification, amplification, and magnetics (BEAMing) system. Next-generation sequencing technologies, including tagged-amplicon deep sequencing (TAm-Seq), the Safe-Sequencing System (Safe-SeqS), and cancer personalized profiling by deep sequencing (CAPP-seq), can also be used and the race for ever more sensitive analytical tools is ongoing.<sup>1,3,4,6</sup>

## Applying liquid biopsies now and in the future

There are a plethora of potential applications for liquid biopsies<sup>3,7</sup> (Figure 1), and probably the most exciting among them is the potential for screening for and early detection of cancer. The fact that ctDNA and CTCs have both been shown to be present from the earliest stages of disease has sparked interest in the possibility of developing simple blood tests to identify tumors before they become detectable by other methods and at a point at which they may be curable.

Given that both are present at such low levels within the circulation and are particularly sparse at earlier stages of disease, current technologies may lack the specificity and sensitivity for this application at present. However, numerous clinical trials are ongoing.

For CTCs, simple enumeration has been the most extensively investigated application to date. Numerous studies have shown that the number of CTCs in the bloodstream

iquid biopsy type	Tumor type	Key finding	Data
CTC (CellSearch)	Metastatic breast cancer <sup>8</sup>	Elevated and increasing levels of CTCs at baseline and after therapy were independent prognostic indi- cators of PFS and OS	Pretreatment Median PFS 6.6 mo (≥5CTCs/7.5mL blood) vs 11.4 mo (<5CTCs/7.5 mL blood; P < .0001) Median OS 15.5 mo vs 37.1 mo (P < .0001) Postreatment Median PFS 4.8-6.7 mo vs 9.6-10.7 mo (P < .0001) Median OS 13.1-22.4 mo vs 27-41.5 mo (P < .0001)
CTC (CellSearch)	Metastatic castration- resistant prostate cancer <sup>9</sup>	CTC level was the most accurate and independent predictor of OS	Pretreatment Median OS 4.5 mo (≥5CTCs/7.5 mL blood) vs 21.7 mo (<5CTCs/7.5 mL blood; P < .0001) Post-reatment Median OS 6.7-9.5 mo vs 19.6-20.7 mo (P < .0001)
CTC (CellSearch)	Metastatic colorectal cancer <sup>10</sup>	CTC count before and during treat- ment is an independent predictor of PFS and OS	Pretreatment Median PFS 4.5 mo (≥3CTCs/7.5 mL blood) vs 7.9 mo (<3CTCs/7.5 mL blood; P = .0002) Median OS 9.4 mo vs 18.5 mo (P < .0001) Posttreatment Median PFS 1.2-3.8 mo vs 6.3-7.3 mo (P < .0001) Median OS 3.3-6.1 mo vs 14.6-15.7 mo (P < .0001)
ctDNA (Cobas EGFR muta- tion test v2)	Non–small-cell lung cancer	Strong concordance between EGFR mutation rates identified by plasma ctDNA and tumor biopsy	ENSURE study evaluating treatment with erlotinib vs gem- citabine/cisplatin as first-line treatment in patients with EGFR mutation-positive NSCLC. 98.6% enrolled patients had available plasma ctDNA samples. Concordance, 76.7% (EGFR mutation positive cases) and 98.2% (EGFR mutation negative cases). EGFR positivity according to plasma ctDNA correlated with improved PFS following treatment with erlotinib

CTC, circulating tumor cell; ctDNA, circulating tumor DNA; PFS, progression-free survival; OS, overall survival; EGFR, epidermal growth factor receptor

has prognostic significance in various different tumor types. Three such studies led to the first regulatory approval for a CTC detection system (Table 1).<sup>8-10</sup>

TABLE 1 Kov clinical trials of liquid biopsion

CellSearch (Janssen Diagnostics/Veridex LLC) is a semi-automated system that captures CTCs on the basis of their expression of an epithelial antigen, epithelial cell adhesion molecule (EpCAM). To do this, it uses magnetic particles coated with EpCAM antibodies that should positively select CTCs. The cells are then stained with a variety of fluorescent antibodies that help to further distinguish them as CTCs (Figure 2).<sup>4,11</sup>

This assay is approved by the FDA for monitoring patients with metastatic breast, colorectal (CRC), or prostate cancers and, in conjunction with information garnered from other diagnostic tests, allows assessment of patient prognosis. The presence of CTCs above a certain threshold ( $\geq 5$  CTCs/7.5 mL blood for prostate cancer and breast cancer, and  $\geq 3$  CTCs/7.5 mL blood for CRC) were independent and accurate predictors of poorer survival.<sup>8-10,12</sup>

One area in which liquid biopsies could really come into their own is in providing more real-time analysis of tumors. This is something that has proven particularly challenging with tissue biopsies because repeating these invasive procedures is problematic. But the ease of repeat blood draws means that serial liquid biopsies could be performed and might offer the possibility of monitoring disease progression and evolution over the course of disease and particularly in response to treatment.

Indeed, studies have shown that in addition to baseline CTC counts, changes in CTC number during treatment are also prognostic. There was improved survival among patients whose CTC counts decreased below a threshold value during treatment and vice versa. This is also an approved use for CellSearch though at present it is not widely clinically implemented.<sup>12</sup>

#### **Clinical utility remains elusive**

The ultimate goal would be for liquid biopsies to have an impact on treatment decisions, allowing oncologists to change management strategy based on predicted sensitivity or resistance to therapy, so-called clinical utility. Thus far, clinical utility has proved elusive, though liquid biopsies using ctDNA to evaluate tumor genotype have come closest.

The Cobas EGFR Mutation Test v2 recently became the first ctDNA-based liquid biopsy to receive regulatory approval. It was approved as a companion diagnostic to identify patients with advanced non–small-cell lung can
 TABLE 2 Select ongoing studies of CTCs

Clinicaltrials.gov identifier/trial name	Description	
Predictive, prognostic and diagnostic value		
NCT01710605/STIC-CTC	Investigating the prognostic value of baseline CTC count in hormone receptor-positive breast cancer	
NCT02119559/CIRCUTEC	Examining the potential of CTCs as an early predictive marker of response to first-line treatment with cetuximab in patients with inoperable, recurrent and/or metastatic HNSCC	
NCT02554448	Detection of CTCs in stage III renal cancer patients undergoing neoadjuvant therapy	
NCT02610764/ESO-CTC	Evaluating the influence of multimodal therapy on CTC count in resectable esophageal adenocarcinoma	
NCT02887612	Utility of ctDNA for prediction of relapse in gastric cancer	
NCT01848015	Assessing the predictive value of CTCs for recurrence of advanced gastric cancer after resection	
NCT02072616/CTC-Pancreas	Role of CTCs in the diagnosis and therapeutic management of pancreatic adenocarcinoma	
NCT02155426	Prognostic role of changes in CTC counts in patients with advanced NSCLC treated with first- or second-line chemotherapy and targeted therapy	
NCT01961713	CTC analysis in patients with localized prostate cancer undergoing prostatectomy	
Guiding therapeutic decision making		
NCT01548677/TREAT-CTC	Role of trastuzumab in HER2-negative early breast cancer with HER2-positive CTCs	
NCT01975142/CirCe	Examining the use of HER2-targeting therapy in breast cancer patients with HER2-amplified CTCs	
NCT01349842/CirCe01	Evaluation of CTCs for guiding third-line chemotherapy for metastatic breast cancer patients	
NCT01619111/DETECT III	Comparing lapatinib in combination with standard therapy vs standard therapy alone in patients with HER2-negative, hormone receptor-positive metastatic breast cancer with HER2-positive CTCs	
NCT02035813/DETECT IV	Evaluating treatment with standard endocrine therapy and mTOR inhibitor everolimus in patients with HER2-negative, hormone receptor-positive metastatic breast cancer with persistent HER2-negative CTCs	

CTC, circulating tumor cell; ctDNA, circulating tumor DNA; HER2, human epidermal growth factor receptor 2; HNSCC, head and neck squamous cell carcinoma; mTOR, mammalian target of rapamycin

cer (NSCLC) who have specific mutations in the epidermal growth factor receptor (*EGFR*) gene and are therefore eligible for treatment with the EGFR inhibitor erlotinib.<sup>13</sup>

Approval was based on comparison of *EGFR* mutation identification rates using plasma ctDNA samples and tumor tissue samples from patients enrolled in the phase 3 ENSURE trial, which compared the efficacy of erlotinib with chemotherapy as first-line therapy in patients with advanced NSCLC. Of the 217 patients enrolled in the trial, 98.6% of patients had both tumor biopsy and plasma ctDNA samples available for testing. Concordance between the two types of biopsy in identifying patients with *EGFR* mutations was high and patients with *EGFR* positivity according to liquid biopsy results demonstrated improved progression-free survival when treated with erlotinib.<sup>14</sup>

The results of a large-scale genomic analysis of various different types of tumors using ctDNA were also recently presented at the 2016 American Society of Clinical Oncology meeting. Blood samples from more than 15,000 patients with 50 different tumor types, including advanced lung cancer (37%), breast cancer (14%), and CRC (10%), were collected and compared with either available tumor biopsy samples from the same cases (n = 398) or, in the majority of cases, with The Cancer Genome Atlas database, which uses tumor biopsies to perform genome-wide sequencing studies. Both types of biopsy revealed very similar mutation patterns when the Guardant360 next-generation sequencing test, which targets 70 genes, was applied. In particular, when *EGFR*, *BRAF*, *KRAS*, *ALK*, *RET*, and *ROS1* mutations were identified by tumor tissue biopsy, the same mutations were reported in 94%-100% of plasma samples.<sup>15</sup>

Studies of the clinical utility of ctDNA and CTCs are among ongoing clinical trials of liquid biopsies (Tables 2 and 3). The potential for using CTCs to guide treatment decisions has become particularly relevant in breast cancer in light of results showing that patients with primary tumors that are negative for human epidermal growth factor receptor 2 (HER2) amplification, an important biomarker in breast cancer, may have CTCs that are HER2positive, in up to 30% of cases. These patients may therefore TABLE 3 Select ongoing studies of ctDNA **Clinicaltrials.gov identifier** Description Early-stage detection of cancer NCT02612350 Utility of plasma ctDNA in asymptomatic subjects at high risk of cancer for the detection of neoplastic disease NCT02715102 Utility of plasma ctDNA for detection of lung cancer in patients undergoing diagnostic or therapeutic bronchoscopy or thoracotomy NCT02778412 Utility of plasma ctDNA for detection of thyroid cancer in patients undergoing diagnostic fine needle aspirations of the thyroid Monitoring relapse/recurrence NCT02842203 Evaluating ctDNA as a prognostic marker and monitor of disease recurrence in stage III colorectal cancer NCT02887612 Utility of ctDNA for prediction of relapse in gastric cancer Biomarker/guiding therapy NCT02623257 EGFR mutations in ctDNA in patients with advanced NSCLC Utility of ctDNA as biomarkers in treatment of biliary tract cancer NCT02809898 ctDNA, circulating tumor DNA; EGFR, epidermal growth factor receptor; NSCLC, non-small-cell lung cancer

still benefit from HER2-targeted therapy.<sup>16</sup>

The DETECT studies are the first phase 3 trials in which treatment decisions are being based on the phenotypic characteristics of CTCs. DETECT III (NCT01619111) is comparing lapatinib in combination with standard therapy with standard therapy alone in patients with HER2-negative metastatic breast cancer who have HER2-positive CTCs, whereas DETECT IV (NCT02035813) is enrolling patients with HER2-negative, hormone receptor-positive metastatic breast cancer and persistent HER2-negative CTCs to receive standard endocrine therapy and the mammalian target of rapamycin inhibitor everolimus.

#### Other targets and sources for liquid biopsy

Another approach to liquid biopsies that is also beginning to take off is to collect tumor-derived exosomes from the bloodstream. Exosomes are tiny, fluid-filled, membranebound sacks that bud off from the surface of a cell to expel waste or to transport cargo from one cell to another. DNA, RNA, and protein can be extracted from tumor-derived exosomes and could also serve as molecular biomarkers relating to the cancer cells from which they came.<sup>6,7</sup>

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Exosome Diagnostics is bringing the first exosomebased diagnostic tests to the market and recently teamed up with Amgen for the development of these liquid biopsies.<sup>17</sup> In January 2016, they launched ExoDx Lung (ALK), for detection of *EML4-ALK* gene fusions in patients with NSCLC, using a proprietary platform for the isolation of RNA from exosomes. Data that was presented at several different conferences in 2015 demonstrated a sensitivity of 88% and specificity of 100% for this diagnostic when compared with tissue *ALK* status in NSCLC patients receiving a second-generation ALK inhibitor following progression on prior ALK inhibitor therapy.<sup>18</sup>

In September, they subsequently announced the launch of a test that analyses genetic information from exosomes collected from a urine sample taken from prostate cancer patients. Using a 3-gene signature, in combination with a proprietary algorithm, this diagnostic generates a score assessing a prostate cancer patient's risk for higher grade, more aggressive disease. It is designed to complement the prostate-specific antigen score and has demonstrated accuracy in ruling out the presence of high-grade cancer before an initial biopsy in more than 1500 patients.<sup>19</sup>

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