Commentary

Recent translational research: circulating tumor cells in breast cancer patients

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Abstract

In breast cancer patients, hematogenous tumor cell dissemination can be detected, even at the single cell level, by applying immunocytochemical and molecular assays. Various methods for the detection of circulating tumor cells in the peripheral blood have been described. Results from recently reported studies suggest that circulating tumor cell levels may serve as a prognostic marker and for the early assessment of therapeutic response in patients with metastatic breast cancer. However, in early-stage breast cancer, the impact of circulating tumor cells is less well established than the presence of disseminated tumor cells in bone marrow; several clinical studies have demonstrated that cells of the latter type are an independent prognostic factor at primary diagnosis. In this article we briefly summarize recent studies examining the presence of circulating tumor cells in the blood and discuss further clinical applications.

Introduction

Adjuvant systemic therapy has contributed substantially to the decline in mortality from breast cancer in the Western world [1]. However, adjuvant systemic therapy is associated with substantial cost and toxicity. With the use of traditional prognostic factors, it is still not possible to reliably identify those breast cancer patients who will relapse with metastatic disease [2]. Research efforts have therefore been undertaken to identify additional parameters enabling individual risk assessment. In addition to their role as prognostic factors, these parameters may also serve as targets for new therapeutic approaches (for example HER-2/neu trastuzumab) [3].

Classically, prognostic factor research has been focused on molecular characteristics of the primary tumor. However, it seems more logical to examine micrometastases in distant sites as an indication of future cancer behavior. In that regard, immunocytochemical and molecular assays enable the specific detection of systemically detected metastatic tumor cells, even at the single-cell stage, in the blood and bone marrow (BM). These technical advances provide an opportunity to study one of the first crucial steps in the metastatic cascade.

Because BM collection is invasive and inconvenient, many investigators have studied techniques to detect and characterize tumor cells in blood. Recently reported studies indicate that, with modern detection techniques, circulating tumor cells (CTCs) could provide a reliable and convenient source for clinically relevant analysis. Pretreatment CTC levels seem to be prognostic in women with metastatic disease starting a new therapy. Moreover, CTC levels may also be important in determining therapeutic efficacy. Fewer data are available for CTCs in patients with early stage disease.

In this review we consider the current aspects in the field of CTC detection in breast cancer patients and discuss perspectives for clinical and research applications.

Techniques for detecting CTCs: new developments for tumor cell enrichment and cell detection

Enumeration of CTCs in human blood is fraught with many challenges. First, one must separate these rare cancer cells from billions of erythrocytes and millions of leucocytes present in 10 ml of blood, and then one must confirm that the identified 'event' is, indeed, a CTC. Many strategies have been used to separate CTCs from normal blood cells (e.g. centrifugal migration through Ficoll gradient or filtration), and

BM, bone marrow; CTCs, circulating tumor cells; DTCs, disseminated tumor cells; FISH, fluorescence in situ hybridization; RT-PCR, reverse transcriptase polymerase chain reaction.

to confirm the malignant origin of the cells on the basis of the identification of epithelial or breast-specific markers.

For rigorous multicenter studies and/or routine clinical use, it is of importance to demonstrate reproducibility of evaluations and stability of blood samples, to allow shipment of samples. With modern assay systems, such requirements can be fulfilled [4]. In this regard, recent developments of new enrichment systems for CTCs seem to provide an accurate and reliable method to identify and characterize these rare events [5,6]. For example, the nearly entirely automated system, designated CellSearch™, has been cleared by the US Federal Drug Administration. This system combines immunomagnetic enrichment and fluorescent immunocytochemical characterization of these rare events. In a prospective, multi-institutional clinical trial, CTC detection with this system provided significant prognostic information both before and soon (4 weeks) after initiation of chemotherapy in patients with measurable metastatic breast cancer [6,7]. A prospective randomized clinical trial is planned by the US Breast Cancer Intergroup, led by the Southwest Oncology Group, to evaluate whether patients with metastatic breast cancer who have persistently elevated CTC levels after one cycle of first-line chemotherapy benefit from an early change of therapy.

PCR-based assays for CTC detection

Expression of epithelial and/or breast cancer-specific genes can be exploited to provide evidence of disseminated tumor cells (DTCs). In addition to immunologic techniques, PCR techniques have been employed to detect these rare events. The major limiting factor for the detection of single tumor cells by RT-PCR is illegitimate low-level transcription of tumor-associated or epithelium-specific genes in normal cells [8]. Quantitative RT-PCR may increase discrimination between mRNA expression in normal cells and tumor cells, thereby increasing the specificity of the RT-PCR approach. In one study, Ring and colleagues [9] compared immunomagnetic separation or cell filtration combined with laser scanning cytometry with a multimarker real-time RT-PCR assay for the detection of breast cancer cells. The multimarker real-time RT-PCR assay was found to be the most sensitive technique for the detection of CTCs [9]. However, it is not clear which of the methods provides clinical relevant information.

Two separate studies have suggested that RT-PCR detection of CTCs is associated with poor prognosis. One of these employed cytokeratin 19 RT-PCR in the blood of non-metastatic breast cancer patients before adjuvant therapy [10], whereas in the other a multimarker PCR assay was used to detect CTCs in metastatic patients [11]. CTC detection by PCR for cytokeratin 19 before neoadjuvant chemotherapy has also been reported to be associated with an increased risk for recurrence [12]. These intriguing but preliminary results require validation in prospective multicenter studies,

because assay conditions vary between studies and there is no consensus regarding these conditions.

Can blood examination replace BM aspiration?

Results from individual studies, and a recently published pooled analysis of several studies, have consistently demonstrated that the presence of DTCs in BM has a strong prognostic impact on patient survival [13]. Because BM aspiration and biopsy are invasive and inconvenient, several investigations have addressed whether one might garner the same information from CTCs. Preliminary studies suggest that CellSearch™ can be used to detect DTCs in BM and CTCs in blood of patients with primary breast cancer [14,15]. Optimal cutoff values for different patient groups are still under investigation. At the San Antonio Breast Cancer Symposium in 2005, it was reported that prognostic information from CTC determination can also be obtained in patients treated with endocrine therapy but that the optimal time point for the evaluation of CTCs in these patients might be later during the course of treatment than for patients treated with chemotherapy [16]. In addition, for certain patient groups (for example, patients with estrogen receptornegative tumors or patients treated with chemotherapy) the clinically relevant number of CTCs might be lower than the five cells previously described [16].

Recent studies suggest that the frequency of CTCs in blood seems to be lower than that of DTCs in BM. Conventional Ficoll gradient centrifugation might therefore not provide sufficient enrichment of malignant cells. This could explain the lack of clinical relevance of CTC detection described in earlier studies.

Other methods of detection of CTCs and DTCs have also been compared [5,17,18]. Using immunocytochemistry, Piegra and colleagues [18] showed a correlation between the detection of CTCs in the blood and DTCs in BM of patients with primary breast cancer at the time of surgery. However, although they did detect an association between the two, only the presence of DTCs in BM provided significant prognostic information. In a study by Wiedswang and colleagues [17], CTCs and DTCs from BM were examined by immunocytochemistry from breast cancer patients with a median time of 40 months after operation. Both the presence of CTCs and that of DTCs in BM was significantly associated with disease-free survival and breast cancer-specific survival, and the presence of both CTCs and DTCs resulted in an especially poor prognosis. However, this study did not exclude patients who had relapsed before the sampling of blood or BM. Excluding the patients with breast cancer-related events before the sample collections, CTC detection was not significantly associated with disease-free survival/distant-disease-free survival, in contrast to DTC detection [17]. Kataoka and colleagues [19] examined blood and BM at the time of surgery for the presence of tumor cells with a quantitative RT-PCR assay for cytokeratin 7 mRNA.

The presence of CTCs in blood, but not DTCs in BM, was correlated with an shorter distant metastasis-free survival. In multivariate analysis, the presence of CTCs was an independent prognostic factor in lymph-node-negative patients [19]. This is in contrast to a study of Benoy and colleagues [20], who found a prognostic impact only for BM but not for blood with the use of RT-PCT for cytokeratin 19 and mammaglobin. This study also included patients with metastatic breast cancer.

Currently, these findings do not support a change from DTC detection from BM to CTC analysis. However, they do suggest that the presence of CTCs could provide information additional to that gained from DTC analysis.

Molecular characterization of CTCs

Recent technical developments permit an examination of the genome of even single DTCs. Using immunocytochemistry, immunofluorescence, and/or fluorescence *in situ* hybridization (FISH), several groups have reported tumor-antigen expression and/or chromosomal aberrations of DTCs in BM and CTCs, indicating the malignant origin of these disseminated cells and indicating genetic heterogeneity [2,21].

The biologic properties of CTCs are not yet well understood. It seems that these cells have little proliferative potential [5]. With current techniques, it is also possible to perform gene expression profiling from CTCs [22,23]. This enables studies that should allow deeper insights into the mechanisms allowing CTCs to form manifest metastasis. For example, it seems that, for breast cancer, only a small proportion of tumor cells, so-called 'cancer stem cells', have the capacity for self-renewal and unlimited growth [24]. In view of the resistance of DTCs and CTCs to chemotherapy it is possible that some of these cells have stem-cell properties. Identifying these 'metastatic stem cells' will be the one of the challenges for future research.

One aspect of clinical importance is the identification of therapeutic targets such as HER-2/neu on CTCs. HER-2 can be detected on CTCs with cytogenetic [21], immunocytochemical [25,26] and PCR methods [27]. There are several possible implications of these findings. For example, it has been reported that nearly one-third of patients whose primary tumors were reportedly negative for HER-2 had CTCs with amplified HER-2 (as determined by FISH). Given the extraordinary impact of trastuzumab (Herceptin), a humanized monoclonal antibody directed against HER-2, in both the metastatic and adjuvant settings, monitoring CTCs for HER-2 could have an enormous influence on the application of this therapy. In addition, the presence of HER-2-positive CTCs seems to be associated with an impaired prognosis [26,27].

Conclusions and perspectives

Besides the prognostic information gained from CTCs, they may also provide a tool for predicting or monitoring the

efficacy of systemic therapy. The study with the highest level of evidence performed so far has been in patients with metastatic disease. However, CTC measurements might also have particular clinical relevance in the context of adjuvant systemic therapy. Indeed, adjuvant therapeutic efficacy can currently be assessed only in large-scale clinical trials after observation periods of several years. An assay that permits the assessment of treatment effects in this context is of high clinical importance. It was previously shown that CTCs can be detected during and after adjuvant or neoadjuvant chemotherapy [5,28], but the clinical impact of this finding is still unclear.

Prospective clinical studies are now ongoing to evaluate whether the eradication of CTCs in the blood (and also DTCs in BM) is correlated with a longer disease-free period and overall survival in the adjuvant setting and whether therapy guided by CTC testing is able to improve the outcome for metastatic patients. An additional important goal is the possibility of identifying metastatic-tumor-specific targets so as to improve therapy regimens.

Competing interests

VM declares no competing interests. DH received research funding from Immunicon and occasional honoraria for speaking engagements from Veridex. KP is co-founder of Micromet AG, Munich, Germany.

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