

Novel tumor markers in the serum of testicular germ cell cancer patients: a review

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Abstract: Serum tumor markers have an important role in the management of patients with testicular cancer. They are useful for diagnosis, staging and risk assessment, follow-up, evaluation of response, and early detection of relapse. Alpha-fetoprotein, human chorionic gonadotropin, and lactate dehydrogenase are established serum markers in testicular cancer, but they have a limited sensitivity. Ongoing research may lead to the identification of novel biomarkers. Therefore, we review the experimental analyses for nucleic acids, circulating tumor cells, and proteins as potential biomarkers in the serum of testicular germ cell cancer patients.

Keywords: biomarker, serum, testicular germ cell cancer

Introduction

Testicular germ cell cancer (TGCT) is the most common malignant tumor in men aged 15–44 years, with its incidence rising in many countries; however, mortality rates remain low and many men are cured.¹ The main factors contributing to the excellent cure rates are careful staging at the time of diagnosis, adequate early treatment based on polychemotherapy, radiotherapy, and surgery, as well as strict follow-up and salvage therapies. The introduction of the highly specific serum tumor markers alpha-fetoprotein (AFP) and human chorionic gonadotropin (HCG) especially improved the clinical management of patients with TGCT, and these markers nowadays play important roles in diagnosis, risk stratification, treatment monitoring, and surveillance.² However, about 50% of TGCT patients have normally ranged tumor markers;³ thus, additional markers could be helpful for the clinician. Several researchers are working on improved noninvasive biomarkers. We will discuss recent advances in the field of circulating tumor cells (CTCs), cell-free nucleic acids, and proteins.

Nucleic acids microRNA

microRNAs (miRNAs) are small RNA molecules involved in several essential biological processes⁴ that cover embryogenic development, cell differentiation, apoptosis, and tumorigenesis.⁵ miRNAs have the potential to qualify as biomarkers in various malignancies because they mostly reveal a high stability in body fluids.^{5,6} Palmer et al⁷ recently demonstrated that miR-371-373 and miR-302 clusters are highly overexpressed in all malignant germ cell tumors (compared to normal control tissues and benign tumors), irrespective of patient age, histologic subtype, or anatomic site of the tumor.⁴ High expression of the miR-371-3 and miR-302 clusters

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was demonstrated in the serum of a 4-year-old boy with a yolk sac tumor and miRNA levels declined after chemotherapy.⁴ This finding stimulated research enormously, and Belge et al confirmed high levels of serum miR-371-3 with a standardized quantitative real-time polymerase chain reaction technique in a small cohort (n=11) of TGCT patients;⁶ unfortunately, HCG/AFP levels were not indicated in the report. Serum levels of each of the three miRNAs (miR-371, miR-372, and miR-373) decreased significantly after treatment.⁶ Dieckmann et al confirmed these results in a subsequent study and, most importantly, showed that the diagnostic information of the novel markers was superior to the classical tumor markers AFP and HCG (increased in four and two out of 24 TGCT patients, respectively).⁵ Patients in clinical stage I also showed a decline of circulating miRNAs following orchidectomy. Further research in a series with 80 TGCT patients confirmed the excellent sensitivity of serum miRNAs:⁸ an increase of miR-371, miR-372, miR-373, or miR-367 was observed in 98% of the examined TGCT patients; the miRNAs thus outperformed AFP (36%) and HCG (57%). Our own data also confirm the improved sensitivity (85%) and specificity (99%) of miR-371 when compared to AFP (14%) and HCG (37%).⁹ In summary, several independent studies indicate an increase of miR-371–373 in TGCT patients irrespective of their histological subtypes and, therefore, result in a better diagnostic performance than AFP/HCG.

DNA

DNA fragments also circulate in a patient's blood, and its quantification seems to be a universal cancer biomarker.¹⁰ Earlier, we demonstrated an increase of short, presumably apoptotic DNA fragments in a cohort of 74 patients with TGCT (compared to 35 control subjects).¹¹ The increase of genomic DNA allowed us to identify TGCT patients with a sensitivity of 84% and a specificity of 97%.¹¹ Whereas mitochondrial DNA is also released into circulation, the underlying mechanism seems to be different from genomic DNA: the quantification of mitochondrial DNA was less sensitive (60%) at a similar specificity (94%) in the same study cohort.¹² Notably, the diagnostic performance of genomic and mitochondrial DNA was superior to that of AFP and HCG (24% and 43% sensitivity, respectively), and cell-free serum DNA levels were similar in seminoma and non-seminoma TGCT.

The analysis of circulating DNA allows for the detection of tumor-specific alterations. DNA hypermethylation is frequent during carcinogenesis, and also occurs in TGCT. Serum DNA

hypermethylation in the above mentioned cohorts was detected more frequently in patients with TGCT than in healthy individuals, including *APC* = adenomatosis polyposis coli 57% and 6%, *p16(INK)* 53% and 17%, *p14(ARF)* = alternate reading frame) 53% and 0%, *RASSF1A* = ras association domain family member 1A 47% and 0%, *PTGS2* = prostaglandin-endoperoxide synthase 2 45% and 0%, and *GSTP1* = glutathione-S-transferase π 1 25% and 0%, respectively. The amount of diagnostic information increased when multiple gene sites were analyzed in combination (67% sensitivity and 97% specificity), and was therefore superior to AFP and HCG (24% and 43% sensitivity, respectively).¹³ The X-inactive-specific transcript, which is located on the X-chromosome, is methylated in male somatic cells. The detection of unmethylated X-inactive-specific transcript DNA in plasma enabled Kawakami et al to specifically identify TGCT patients with 64% sensitivity in a cohort of 25 TCGT patients (the diagnostic accuracy of AFP/HCG was not mentioned).¹⁴ DNA methylation patterns were similar in seminoma and non-seminoma patients.^{13,14} So far, studies on serum/plasma DNA still require independent validation; the possible combination of DNA quantification and the assessment of qualitative alterations render cell-free DNA a study object of particular interest.

CTCs

The detection of CTCs is feasible in many malignancies, especially in patients with an advanced disease.¹⁵ Using the CellSearch assay (Veridex, Raritan, NJ, USA), CTCs were detected in the peripheral blood of 17.5% of the tested TGCT patients. Patients with advanced metastasized (clinical stage III) TGCT and non-seminoma TGCT especially showed an increase of CTCs. CTCs further correlated with AFP, HCG, and lactate dehydrogenase (LDH) levels.¹⁶ An earlier study aimed to identify CTCs via the detection of AFP and HCG messenger RNA (mRNA) using real-time polymerase chain reaction in peripheral blood cells: overall, 27% of patients had an increase of these mRNA, and there was also a correlation of AFP/HCG mRNA positivity with clinical stage IIC/III.¹⁷ Given the low frequency of CTCs in patients with minimal tumor burden, there seems to be only a limited potential for CTCs in the clinical management of patients with TGCT. Furthermore, serum AFP (25%) and HCG (49%) provided a similar or even better sensitivity.

Proteins

TRA-I-60

A report in 1991 suggested that terato-related antigen monoclonal antibody (TRA-I-60) could serve as a tumor marker for

TGCT patients with an embryonal carcinoma component.¹⁸ The analysis of TRA-1-60 in stage I non-seminoma TGCT indicated similar diagnostic performance as the conventional markers. However, during follow-up, TRA-1-60 levels were increased 1 month before radiological signs of recurrence and decreased during treatment.¹⁹ A subsequent study by Lajer et al confirmed an increase of TRA-1-60 in TGCT patients, but the clinical utility was limited by a high rate of false-positive elevations.²⁰

Neuron-specific enolase (NSE)

NSE, an isoenzyme of the glycolytic enzyme 2-phospho-D-glycerate-hydrolase, is a marker for tumors of neuroendocrine origin. It is recognized as a useful tumor marker in small-cell lung cancer and melanoma.²¹ Early studies indicated an increase of NSE in seminoma TGCT, and the diagnostic sensitivity and specificity were in the same order as those of HCG.^{22,23} However, large-scale studies have shown frequent false-positive or false-negative results of NSE during follow-up examinations, and thus do not support NSE to be a valuable tumor marker in germ-cell tumors.^{24,25} Furthermore, hemolysis leads to an elevation of serum NSE.²⁵ Accordingly, Sturgeon et al point out that in spite of promising results (30%–50% positivity of patients with seminomas and less often in non-seminomatous germ cell tumor patients), the use of NSE is limited and, so far, only in an experimental stage of development.²⁶

Lectin (LCA)-reactive AFP

It is sometimes difficult to decide on a treatment plan for seminoma patients with slightly increased AFP levels as these could also be due to liver disease. LCA-reactive AFP, determined by LCA-affinity electrophoresis, might be a specific tool to identify AFP derived from undetected embryonal cell carcinoma or yolk sac tumor elements.²⁷ Furthermore, Kawai et al showed that LCA-reactive AFP was detected after the normalization of total AFP levels in stage I TGCT patients (n=3) who suffered from tumor recurrence.²⁷

Fas and Fas ligand

Fas/Apo-1, a member of the tumor necrosis factor receptor superfamily, can transduce a signal that leads to the rapid induction of apoptosis by binding the Fas ligand (FasL). Serum levels of soluble FasL are increased in patients with various malignancies.²⁸ In TGCT patients with a seminoma component, Fas and FasL are expressed in 73% of tumor tissues, whereas its expression is only 56% for Fas and 11% for FasL in non-seminoma tissue.²⁹ An elevation of soluble

FasL was noticed in 38% of patients with seminomatous TGCT elements (n=24).²⁹ Unfortunately, the sensitivity of HCG was not provided in the study cohort.

CD30

CD30 (TNFRSF8) is a protein of the tumor necrosis factor receptor family. It can be detected in extracellular fluids (eg, anaplastic large-cell lymphoma)³⁰ and it correlates with the tumor burden. CD30 is expressed exclusively in embryonal carcinoma components of TGCT.³¹ Latza et al reported soluble CD30 in the serum of all patients (n=8) with embryonal cell carcinoma, but only in 20% of patients with other TGCT components.³⁰ Moreover, serum levels of soluble CD30 antigen seem to be a promising parameter for monitoring patients with embryonal cell carcinoma.³⁰ Persistent CD30 expression in embryonal carcinoma cells after cisplatin-based chemotherapy is indicative of recurrent and progressive TGCT,³² and monitoring this protein in the serum may be helpful for clinical decision making.

C-reactive protein (CRP)

CRP is an acute-phase reactant and a useful marker of systemic inflammation.³³ Systemic inflammation has been linked to cancer progression, and, in patients with urological cancers, the presence of a systemic inflammatory response is thought to be indicative of a poor prognosis. However, CRP has not been studied as a prognostic biomarker in TGCT. Furthermore, CRP levels have been proven to increase after radiotherapy in contrast to chemotherapy or surgery alone, and it is speculated that a CRP increase could be predictive of late complications like cardiovascular disease³⁴ and chronic cancer-related fatigue.³³

Free β -subunit of HCG (HCG β)

In the serum of patients with seminoma, HCG is increased in only about 15%–20% of the cases. However, 30%–40% of seminoma patients have been found to have increased serum concentrations of HCG β .³⁵ The determination of HCG β increases the frequency of marker-positive seminoma from 17% to 57% and of marker-positive relapses from 32% to 59%. But still, about 40% of marker-positive seminoma and relapses would have been missed with an assay measuring HCG and HCG β together. In patients with seminomatous testicular cancer, HCG β is superior to HCG, and in some non-seminomatous germ cell tumor patients it provides additional information.³⁵ Notably, an increase of both HCG β and HCG was in most cases limited to patients with a choriocarcinomatous component, whereas

the majority of seminoma patients had only an increase of HCG β but not HCG.³⁵ The lack of HCG β may therefore be indicative of choriocarcinoma.

Hyperglycosylated HCG (HCG-h)

HCG-h contains larger and more complex carbohydrate chains than regular HCG.³⁶ Lempiäinen et al showed that a major proportion of HCG was hyperglycosylated preoperatively, at relapse, and shortly after treatment.³⁶ The serum concentrations of HCG-h and HCG correlated strongly with each other and had similar diagnostic value. HCG-h does not appear to provide clinical information additional to that obtained by conventional HCG assays.³⁶

Angiogenic growth factors (pleiotrophin [PTN] and fibroblast growth factor-2 [FGF-2])

Increased serum concentrations of various angiogenic growth factors have been described as potential markers for tumor progression and metastasis in other tumor entities.³⁷ Various angiogenic growth factors (ie, FGF-2, vascular endothelial growth factor, FGF-4, transforming growth factor-beta, epidermal growth factor, and PTN) have been analyzed by Aigner et al in TGCT.³⁷ Significantly elevated levels of FGF-2 and PTN were observed in the serum of testicular cancer patients, and FGF-2 serum levels especially allowed TGCT patients (n=22) and normal individuals (n=21) to be distinguished very accurately (sensitivity/specificity was not reported). FGF-2 serum levels were also elevated in early tumor stages. In comparison, AFP was elevated in 66% of non-seminoma patients, and HCG was increased in 23% of seminoma patients and 56% of non-seminoma patients. The data indicate that serum levels of FGF-2 and PTN might be useful novel tumor markers but confirmative studies are required.

Circulating full-length and caspase-cleaved cytokeratin 18 (CK18)

Circulating full-length and caspase-cleaved CK18 are considered biomarkers of chemotherapy-induced cell death measured using a combination of the M30 and M65 enzyme-linked immunosorbent assays. M30 measures caspase-cleaved CK18 produced during apoptosis and M65 measures the levels of both caspase-cleaved and intact CK18, the latter of which is released from cells undergoing necrosis.³⁸ During necrotic and apoptotic cell death, CK18 and other cytokeratins are released into the blood in either their intact or their caspase-cleaved forms. There they remain relatively stable in the circulation of patients with cancer. Previous studies have highlighted their

potential as prognostic, predictive, and pharmacological tools in the treatment of cancer. M65 and M30 levels appear to reflect chemotherapy-induced changes that correlate with changes in markers routinely used in the clinic for managing patients with non-seminomatous TGCT.³⁸ The overall decreases in M65 and M30 are indicative of treatment response, as they appear to reflect a decrease in tumor load due to chemotherapy-induced tumor cell death. The correlation between M65/M30 levels and International Germ Cell Consensus Classification (IGCCC) prognosis groups, as well as their overall agreement with LDH, AFP, and HCG levels, suggest that M65/M30 may also have prognostic value in patients with non-seminomatous TGCT.³⁸ Although some patients were recruited prospectively in this study, additional studies are needed to underline the benefit of M65/M30 measurements during chemotherapeutic treatment.

Conclusion

The established serum tumor markers AFP, HCG, and LDH play an important role in the clinical management of TGCT patients; but because of their limited sensitivity, additional markers are needed. Very promising novel markers such as circulating microRNAs or FGF-2, which are detectable at similar frequencies in seminoma/non-seminoma TGCT patients and are absent in healthy subjects, are of particular interest as diagnostic tools. The analysis of M65/M30 may provide additional information as a treatment monitoring marker. Nevertheless, validating studies in large-scale, prospective, and clinically relevant patient cohorts are urgently needed. The detection technique of serum miR-371 is being optimized by a startup at the University of Bremen (miRdetect), and prospective studies to prove the clinical use of this marker are to be expected within the next few years.

Disclosure

The authors report no conflicts of interest in this work.

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