

## EDITORIAL

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# Pancreatic Neoplasm in 2012: An Update. Tissue Is an Issue!

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Approximately 42,470 new cases of pancreatic cancer are diagnosed per year in USA, which represents approximately 3% of all newly diagnosed cancers [1]. Pancreatic cancer remains the fourth most common cause of cancer-related mortality in the United States. This close incidence to mortality ratio depicts the considerable diagnostic and therapeutic challenges faced by the patients as well as the care takers treating these patients. Sadly, more than 90% of patients diagnosed with pancreatic cancer succumb to their disease within 5 years; 75% within one year [2]. The major explanation for this poor prognosis is the lack of a therapeutic time window [3]. Pre- and early cancerous lesions are beyond our threshold of detection. Pancreatic cancer is diagnosed at an advanced disease in majority of the cases, and is further characterized with a high rate of local and distant recurrence following surgical resection, and relative chemo-resistance. Even an early stage tumor at the time of initial diagnosis can be metastatic and resistant to conventional therapies.

In the era of personalized medicine, understanding about premalignant lesions, knowledge of genetic abnormalities and development of targeted therapies we are often limited by the lack of tissue [4]. The tissue is extremely important to unveil the nature of the disease as well as to identify new molecular markers to predict outcome and response to treatment, and to aid in the detection of premalignant pancreatic lesions. This is of utmost importance as we have witnessed the integration of molecular markers and active therapies in other solid tumors, such as lung cancer. It is the prime time to do so for pancreatic adenocarcinoma.

The best chance of sampling tissue specimen is at the time of surgical resection. Unfortunately, only 10-20% patients are surgical candidates while the majority of patients with advanced disease are often diagnosed by

fine-needle aspiration either via an EUS- or CT-guided biopsy, hence provide small sample. Due to this limitation only few cells are available which handicap any further investigation on the collected tissue.

Several cytotoxic and biological agents targeting epithelial tumor cells show promising results in pre-clinical and preliminary human studies but have failed to show relevant effects in larger randomized clinical studies. This discrepancy between experimental results and clinical results seem to be at least partly a result of the tumor microenvironment. Pancreatic ductal adenocarcinoma is characterized by remarkable desmoplasia, forming more than 80% of the tumor mass [5]. The desmoplasia is composed of extracellular matrix proteins, myofibroblastic pancreatic stellate cells, and immune cells associated with a multitude of cytokines, growth factors, and extracellular matrix metabolizing enzymes. We are just learning to appreciate the role of this complex process in carcinogenesis and resistance to chemotherapy. Investigators have shown that stellate cells produce extracellular matrix proteins, cytokines and growth factors that promote the growth of the cancer cells. Recent studies also suggest that interactions between extracellular matrix proteins and desmoplastic secreted growth factors with the cancer cells of pancreatic ductal adenocarcinoma activate intracellular signals including reactive oxygen species that act to make the cancer cells resistant to dying [6]. These findings suggest that the desmoplasia of pancreatic ductal adenocarcinoma is a key factor in regulating carcinogenesis of pancreatic ductal adenocarcinoma as well as responses to therapies.

It is clear that we are entering a new era of cancer therapy, in which molecular profiling of tumor specimens is likely to become routinely performed. This is made easier as the technology is more readily available. The incorporation of well-designed correlative studies into the design of therapeutic trials in pancreatic cancer therefore remains crucial to the advancement of this field. However, we have shown our failure to adequately collect tissue in major randomized phase III studies as well as cooperative group trials. Two such examples are:

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**Table 1.** Definitions of various types of biomarkers.

<b>Prognostic biomarkers</b>	Biomarkers that provide information about the patients overall cancer outcome, regardless of therapy [9]
<b>Predictive biomarkers</b>	Biomarkers that can be used in advance of therapy to estimate response or survival of a specific patient on a specific treatment compared with another treatment [10]

- Radiation Therapy Oncology Group (RTOG) 9704. Phase III randomized study of adjuvant fluorouracil-based chemoradiotherapy preceded and followed by fluorouracil *versus* gemcitabine in patients with resected adenocarcinoma of the pancreas: only 225 samples were found adequate from 538 patients after pancreatic resection [7].

- National Cancer Institute of Canada (NCIC) Clinical Trials Group (CTG) PA.3. Erlotinib plus gemcitabine compared with gemcitabine alone in patients with advanced pancreatic cancer: only 26% patients out of 569 had adequate tissue sampled [8].

Biomarkers are either prognostic or predictive (Table 1). Prognostic biomarkers are intrinsic indicators for tumor's aggressiveness and patients' final clinical outcome, regardless of the therapy received. Their clinical relevance is significant as they allow for better risk stratifications as well as rapid assessment of likelihood of disease progression or recurrence.

On the other hand, predictive markers are parameters used to predict treatment responses. Customized chemotherapies based on certain biomarkers have been shown to have better efficacy and result in improved outcome in cancer patients.

Several potential predictive biomarkers for cytotoxic therapy in pancreatic cancer have been identified, such as secreted protein rich in cysteine (SPARC) expression, *KRAS*, human equilibrative nucleoside transporter 1 (hENT-1), cytidine deaminase CDA, and cyclin-dependent kinase activity [11, 12, 13, 14, 15, 16, 17, 18] (Table 2). MicroRNA and cancer stem cells have also been identified as predictive biomarkers as well as potential therapeutic targets in this setting [19].

Predictive and prognostic markers identified in pancreatic cancer patients are summarized in Table 3.

Prospective collection of tissues mandated by the protocol in patients with pancreatic cancer is cumbersome and complicated for many reasons: fine needle aspiration offers a safe diagnostic test and provided ample tissue for confirmation of the

**Table 2.** Potential predictive markers of efficacy and/or resistance to therapy in pancreatic adenocarcinoma.

Therapy	Marker
DNA damaging chemotherapy	Breast cancer gene (BRCA) 1/2 mutation
Radiotherapy	
Poly ADP ribose polymerase (PARP) inhibitor	
Nab-paclitaxel	Secreted protein rich in cysteine (SPARC) expression
Gemcitabine	Human equilibrative nucleoside transporter (hENT)
5-fluorouracil (5-FU)	Cyclin dependent kinase (CDK)
	Cytidine deaminase (CDA) expression
5-fluorouracil (5-FU)	Thymidine phosphorylase (TP)
Capecitabine	Thymidylate synthase (TS)
	Dihydropyrimidine dehydrogenase (DPD)
Erlotinib	<i>KRAS</i> wild-type status
C-met inhibition	C-met expression
Chemoradiation	<i>SMAD4 (DPC4)</i> retention

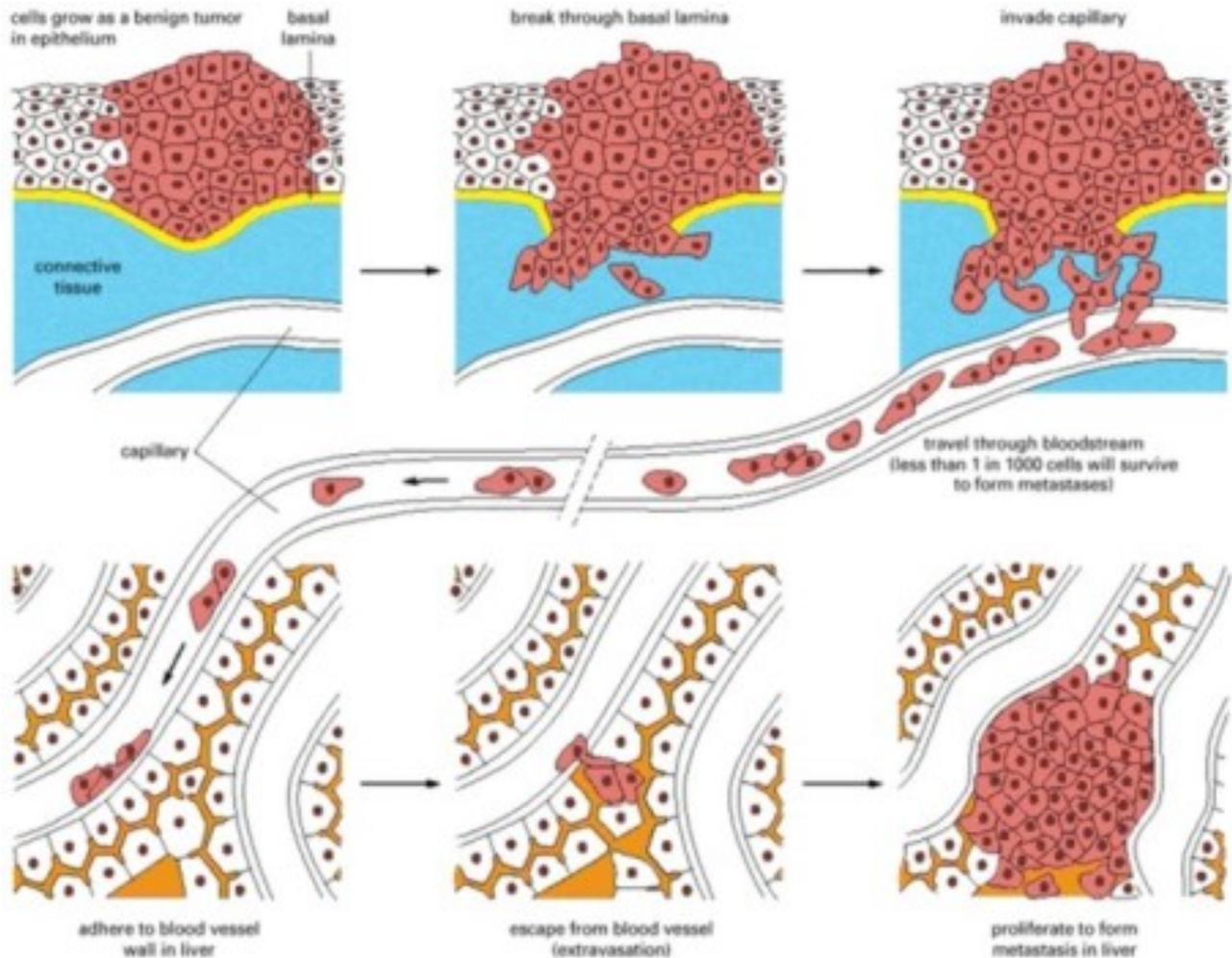
diagnosis, but not enough to be further utilized for research purposes; core biopsies may add more risk to the patients; these procedures can add extra cost and if a repeat biopsy is needed then the cost can be doubled. Though these issues seem as a hindrance, we must overcome these barriers to move forward in the field of pancreatic cancer.

The ability to perform circulating tumor cells collection and proteomic biomarker profiling from serum samples may help us overcoming the traditional barriers to performing correlative studies. Circulating tumor cells are cancer cells that are detached from primary tumor sites and travel in the peripheral blood circulation system, leading to distant metastasis [20] (Figure 1). CTCs are typically enriched and detected via immunomagnetic separation system [21] or via microfluidic circulating tumor cell-chip system [22, 23].

De Albuquerque *et al.* [24] reported the prognostic values of CTCs detection in pancreatic adenocarcinoma. By using the high affinity antibodies BM7 (MUC 1) in addition to conventional VU1D9 (EpCAM), circulating tumor cell detection was reported in 49.3% of 144 peripheral blood samples from 39 patients with advanced pancreatic adenocarcinoma. The detection of such circulating tumor cells portended poor prognosis (median progression free survival: 60.7 days *vs.* 163.6 days in patients with positive and negative circulating tumor

**Table 3.** Summary of predictive and prognostic markers identified in pancreatic cancer patients.

Higher levels of CA 19-9 in a resected pancreatic cancer patient suggests micrometastases
CA 19-9 is a highly significant predictor of overall survival in patients with resected pancreatic cancer
<i>SMAD4 (DPC4)</i> is a predictive biomarker in patients with localized pancreatic cancer
VEGF is a prognostic marker in resected pancreatic cancer
miR-10b is a predictive marker of response to neo-adjuvant therapy in pancreatic cancer
REG4 protein overexpression is an unfavorable response to preoperative chemoradiotherapy in patients with pancreatic cancer
Alpha 1-antichymotrypsin (AACT) may be a useful prognostic marker in patients with advanced stage pancreatic cancer
Human equilibrative nucleoside transporter 1 (hENT1) and deoxycytidine kinase (dCK) are useful predictive markers to response to gemcitabine in patients with pancreatic cancer
<i>EGFR</i> and <i>KRAS</i> mutation status were not identified as markers predictive of a survival benefit from the combination of erlotinib with gemcitabine for the first-line treatment of advanced pancreatic cancer



**Figure 1.** Circulating tumor cells and process of metastasis (copyright ©2002 from Molecular Biology of the Cell by Alberts *et al.* [21]. Reproduced by permission of Garland Science/Taylor & Francis LLC).

cell detections, respectively;  $P < 0.0001$ ). As such, authors concluded that circulating tumor cells can act as an independent prognostic biomarker.

The prognostic and predictive values of circulating tumor cells have been well established in breast and prostate cancer, though their utility in pancreatic cancer is very limited. As the technologies further advance, it is possible that circulating tumor cells may emerge as a critical prognostic as well as predictive biomarkers in pancreatic cancer [25].

Genome-wide analysis using high-throughput DNA method for potential molecular biomarker identifications and analysis is an attractive strategy in pharmacogenetics. Investigators have shown a promising set of single nucleotide polymorphisms, such as PYCARD and MACRE2, which appears to have strong positive correlation with efficacy from gemcitabine-based chemotherapy in pancreatic cancer [26].

Therefore, in view of data available, our clinical practice remains unchanged, though some of aforementioned biomarkers appear to have a potential prognostic and predictive role and have to be explored further. Given these promising preliminary data, future clinical trials using hybrid chemotherapy design [27],

tailored towards standardized biomarker assay, may bring forward more insight and confirmatory data for this interesting concept.

Consensus report of the National Cancer Institute (NCI) Clinical Trials Planning Meeting on pancreas cancer treatment [28] pays emphasis on the enhancement of research to identify and validate the relevant targets and molecular pathways in pancreatic cancer, cancer stem cells, and the microenvironment. In addition, emphasis was also placed on developing rational combinations of targeted agents and the development of predictive biomarkers to assist selection of patient subsets. The report also recommends that phase III clinical trials should be implemented only if there is a meaningful clinical signal of efficacy and safety in the phase II setting. Therefore, the emphasis must be on performing well-designed phase II studies with uniform sets of basic entry and evaluation criteria with survival as a primary endpoint. Patients with either metastatic or locally advanced pancreatic ductal adenocarcinoma must be studied separately.

A better understanding of the biology of desmoplasia in the mechanism of pancreatic ductal adenocarcinoma will likely provide significant opportunities for better

treatments for this devastating cancer. Development of biorepositories in the conduct of randomized phase III trials in this disease is mandatory. This cannot be achieved without more funding and change in our approach and attitude towards research practice in the field of pancreatic cancer. Furthermore, improved awareness and understanding of hereditary genetic abnormalities predisposing to pancreatic adenocarcinoma present us the potential for both screening of at-risk individuals and development of molecularly-targeted treatment modalities. The pharmacogenomic studies have identified biomarkers of efficacy to established chemotherapy but prospective validation of these predictive and prognostic biomarkers need to be achieved immediately followed by their incorporation into clinical decision making.

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