

Interactions between radiotherapy and endocrine therapy in breast cancer

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Abstract

Whenever radiation therapy is given with curative intent there is the risk of serious damage to normal tissue. This risk increases with the dose of radiation, as does the probability of local tumour control. In the attempt to cure, the doses reach a level that inevitably causes some undesirable adverse effects, ranging from undetectable, or minimal, to unacceptably severe. Over the last few years, a number of reports have suggested that the prediction of normal tissue response after radiotherapy may be achieved by assays on samples withdrawn from the patients prior to treatment, although recent reports have described mixed results. The ability to predict tumour response to anti-hormones in patients with breast cancer has important implications with regard to treatment. Recent discoveries promise to provide individualized treatment options. However, there are no data to support that, used jointly, the combination of radiotherapy and hormone therapy may achieve an enhancement of breast cancer tumour response. Nowadays, development in cancer therapy is increasingly arising out of studies in basic science; its implementation in the hands of clinicians is improving the management of patients with cancer. In addition, as the biological aspects of irradiation and hormonal therapy offer an explanation, at least in part, for the outcome observed in patients with breast cancer after therapy, we have focused this review on trying to analyse the most relevant experimental research about the relative roles of radiotherapy and hormonal therapy, the corresponding side-effects and, taking into account recent advances, future areas of research that we consider of major importance in the field.

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Introduction

Breast cancer is a very heterogeneous disease in which oestrogen acts as an endocrine growth factor in at least one-third of breast cancer cases and the effects are mediated via the oestrogen-receptor (ER) pathway. The huge differences in the biological characteristics found in the study of patients with breast cancer, including data from clinical, pathological, cellular and molecular studies, are sufficient to explain the dramatic changes, and the diversity, in the treatments recommended over the past two decades. It is well known that prognosis factors influence therapeutic conduct and, based on them, clinical trials have been performed in order to decide treatment, dose and the optimum sequence of different therapeutic agents to be used. The problems of defining the best therapy are immense and clearly this is an important area for research in future.

Tumour size, lymph node status, histological type and

grade, oestrogen and progesterone receptors and patient age are accepted as established prognostic and predictive factors valid for the selection of treatment of patients with breast cancer (Albain *et al.* 1994, Zavagno *et al.* 2000). The role of other promising new factors, such as p53, bcl-2, bax, the truncated form of epidermal growth factor receptor gene (ERB-2), alone and/or in association with other factors to define the patho-biological profile of the tumours, could represent an initial framework for a biologically tailored therapy (Ross & Fletcher 1999, Daidone *et al.* 2000, Ziyaie *et al.* 2000, Bièche *et al.* 2001), the importance of which will be determined in ongoing prospective studies.

The natural history of breast cancer is characterized by an initial stage of sensitivity to medical treatments whereas, at later stages, sometimes even without the previous use of anti-oestrogen or chemotherapeutic agents, the tumour becomes resistant to these growth-inhibitory agents.

Although mastectomy remains the most frequently used treatment in the early stages, the demand for conservative surgical measures, with or without adjuvant therapy, is growing among women with breast cancer, and many studies indicate that such an approach is adequate for a large subgroup of these patients (Jha *et al.* 2001). Since there are no clear differences in survival, especially in non-invasive breast cancer in which 95% or more of the patients are cured no matter how they are treated, we need to consider the fact that individual patients differ in the importance they place on the risk and benefits of adjuvant treatments. In addition, given that the gradually increasing success of cancer treatment has led to longer patient survival which unfortunately implies a greater opportunity for late effects to appear, increasing in severity and impact on quality of life, this parameter might be evaluated after treatment (Amichetti *et al.* 1999) to provide better ways to support shared decision making between patients and their physicians (Eifel *et al.* 2001).

Women with primary invasive breast cancer receive both local and systemic treatment. Surgery and radiation therapy are local treatments to reduce the risk of recurrent cancer in the breast, chest wall and regional lymph nodes. Cytotoxic chemotherapy and hormonal therapy are systemic treatments given after local treatment to reduce systemic recurrences and overall mortality from breast cancer. Nonetheless, the management of this class of tumour is a continuing challenge because of the high local recurrence rates and poor survival due to metastatic disease (Tan *et al.* 2001).

A number of patient and tumour characteristics, which are present at the time of primary cancer diagnosis, affect the subsequent risk of local and distal recurrences. However, the predictive use of one individual characteristic is not straightforward because of interaction with other patients features, tumour and treatment factors but it is clear that locoregionally recurrent breast cancer is an independent predictor of subsequent distant metastasis (Clemons *et al.* 2001). To identify properly, by means of multivariate analysis, the major factors determining the risk of tumour progression, and to include them in a prognosis index for survival would indeed be useful.

Nowadays, development in cancer therapy is increasingly arising out of studies in basic science; its implementation in the hands of clinicians is improving the management of patients with cancer. In addition, as the biological aspects of irradiation and hormonal therapy offer an explanation, at least in part, for the outcome observed in patients with breast cancer after therapy, we have focused this review on trying to analyse the most relevant experimental research on the relative roles of radiotherapy and hormonal therapy, the corresponding side-effects and, taking into account recent advances, future areas of research that we consider of major importance in the field.

Radiation damage to DNA

DNA represents the most important target for radiation action on a cell, and subsequently on the tumour and its surrounding normal tissues. A therapeutic dose of low-linear energy transfer (LET) radiation causes a large number of ionizations in every cell. Some of these directly damage DNA; others generate free radicals that react with DNA. Free radicals are extensively scavenged by compounds containing (–SH) sulphhydryl groups. Lesions in DNA are very effectively repaired by enzymatic reactions. Some of them fail to repair and become fixed, and any differences in induction and/or repair among double-strand breaks will lead to differences in cell lethality (Steel *et al.* 1989, Ruiz de Almodóvar *et al.* 1994, Núñez *et al.* 1995), but there are many sources of evidence in the sense that DNA damage is the critical event in radiation-induced cell death (Yasui *et al.* 2001).

The cellular microenvironment of tumours influences their responsiveness to therapeutic agents. None of them is simpler than oxygen, although a host of chemical and pharmacological agents that modify the biological effects of ionizing radiation have been discovered. Considerable controversy surrounds the interpretation of radiosensitivity data from steroid hormones and anti-hormones on oestrogen-sensitive breast cancer (Sarkaria *et al.* 1994, Villalobos *et al.* 1996, Amorino *et al.* 2000). Currently, our understanding about the mechanisms of radiation-induced cell death in hormone-dependent tumour cells is limited (Siles *et al.* 1998).

Apoptosis is obviously an important process in biology but, in tumour cells, which have lost the normal inter-relationship of cell proliferation and cell death, an in-depth analysis of these two processes would facilitate the design of more effective treatment strategies using radiation therapy alone or combined with other therapeutic modalities. Some authors have pointed out that hormonal treatment might alter the radiation sensitivity in breast cancer regardless of the receptor status, suggesting that hormonal agents may act both via receptor- and non-receptor-binding mechanisms (Wazer *et al.* 1991, Paulsen *et al.* 1996).

The importance of cell cycle control in the cell response to DNA-damaging agents is widely accepted and many genes, or the pharmacological modulation of their functionality, might be a critical determinant of cellular radiosensitivity, with a possible differential effect in tumours with non-functional p53 and normal tissues (Valenzuela *et al.* 2000).

The tumour suppressor protein p53 plays a key role in the cell's decision to arrest the cell cycle or undergo apoptosis following a genotoxic insult (Siles *et al.* 1996). The presence of damaged DNA in the cell activates the repair mechanisms as well as the signal transduction pathways, leading to cell cycle arrest and repair or apoptosis. Several of these responses are mediated by the stabilization and activation of p53. It is there-

fore crucial to understand the events from the initial radiation-induced DNA damage to p53 activation.

bcl-2 protein is a potent anti-apoptotic protein that inhibits a mitochondria-operated pathway of apoptosis in many cells. DNA damage agents and death receptor ligands can activate this mitochondrial mechanism and it has been suggested that the enhanced expression levels of bcl-2 present in tumour cells can render these cells less sensitive, not only to chemotherapeutic drugs but also to tumour necrosis factor-related apoptosis-inducing ligand (Ruiz de Almodóvar *et al.* 2001). Poly (ADP-ribosylation) of different nuclear acceptors by poly (ADP-ribose) polymerase-1 (PARP-1) is an early event in that cascade; although the nature and the consequence of this interaction are still very controversial, it seems that PARP-1 partially contributes to radiation-induced cell death (Valenzuela *et al.* 2002).

Tumour and normal tissue response to radiation treatment

Genetic, epigenetic and nutritional variables have been invoked to explain variations in the intrinsic radiosensitivity of both human tumour and normal tissue cells. Nowadays, it is recognized that these differences exist and that they may be related to the clinical ability to cure and tolerance to treatment. The success of radiotherapy depends on the total radiation dose. In an attempt to cure, the doses reach a level that inevitably cause some collateral side-effects. With each new patient, the radiation oncologist pays attention to those aspects that can be controlled as plans for the treatment of the malignant disease are made; however, there are no ways of accounting for tumour and normal tissue sensitivities which might be the major determinant of successful treatment.

The dilemma for the physician is how to protract the treatment in a way that maximizes control of the tumour and minimizes normal tissue injury. Thus, the tolerance of the normal tissues encompassed within the treatment volume is the factor that limits the dose (Withers 1992). Radiation therapy is associated with a broad spectrum of normal tissue acute and late reactions, ranging from undetectable or minimal to unacceptably severe (Tucker *et al.* 1992, Burnet *et al.* 1998). Although a significant proportion of this variation can be attributed to treatment-related factors, there is an increasing evidence showing that the major determining differences are related to intrinsic biological factors (Safwat *et al.* 2002).

Over the last few years, a number of important publications have suggested that the prediction of normal tissue response after radiotherapy may be achieved by assays performed on cells or tissue samples withdrawn from the patients prior to radiation treatment, although recent reports have described mixed results (Ozsahin *et al.* 1997, Barber *et al.* 2000, Peacock *et al.* 2000, West *et al.* 2001).

The ability to predict these differences in radiation sensitivity would have important implications with regard to cancer treatment. For example, a strategy based on testing human normal tissue radiosensitivity might permit the individualization of treatment by dose escalation in resistant patients without increasing normal tissue complications (Bentzen 1997). But advocating dose escalation is advocating research that 'is state-of-the-art, not standard of practice' (Glatstein 2001). However, even if an approach to predict *in vivo* response prior to radiotherapy treatment was validated, clinical implementation would be difficult.

We have previously reported the potential of assays of DNA damage in lymphocytes for this purpose (Núñez *et al.* 1998), and the existence of an inter-individual variation in the radiosensitivity of normal tissue, measured on lymphocytes as the initial number of DNA double-strand breaks induced by Gy per DNA unit (dsb/Gy per DNA unit). Our results have shown that the radiosensitivity parameter in normal cells can differ by a factor of 10, ranging from 0.5 to 4.7 dsb/Gy per DNA unit (Ruiz de Almodóvar *et al.* 2002). Among the patients studied we detected 15 women who developed severe skin reactions. Whether DNA initial radiation-induced damage is related or not to the severity of normal tissue effects after radiation treatment might be better clarified with further research that is currently underway in several centres including our own.

As cancer treatment becomes more effective, more patients survive longer and the importance of short- and long-term morbidity is increased. The categorization of possible side-effects according to a scoring system already shows an improvement in the quality assurance of care and thus in the patients' quality of life. Different scoring protocols are found in the literature. Among them, the European Organization for Research and Treatment of Cancer/Radiation Therapy Oncology group (RTOG) classification (Cox *et al.* 1995) is a very comprehensive system that deals with both acute and late reactions. Our previous results (López *et al.* 2002) point out the advantage of the RTOG system over others, at least for evaluating the acute effects produced by radiation treatment on the skin of women with breast cancer.

Clinical oncologists are aware of the fact that late damage is indeed the most important component of morbidity. However, so far, no consistent relationship has been proved between acute and late reactions. Late injury, in particular, represents a threat to normal tissues, either by direct cell death or delayed apoptosis. In both events, the ultimate expression is cell loss and atrophy. Moreover, if enough cells are depleted and not repopulated, there will be a functional organ compromise. Recognition and clearance of apoptotic cells after exposure to radiation produces both a persistent macrophage activation and an inflammatory type response, effects that persist long after the initial radiation insult (Lorimore *et al.* 2001).

In breast cancer, the time between the first introduction of new advanced medical tools and the full realization of the potential benefits in the clinic is often quite long, especially considering the long follow-up period required to assess the outcome adequately. The maximum severity of acute effects can be judged during the course of radiotherapy. Late effects would be better judged after 5 years or more. This allows sufficient time for complications to develop fully in some normal tissues, though not in all. The log-normal variation in cellular radiosensitivity among individuals (Ruiz de Almodóvar *et al.* 2002) could probably be due to polymorphisms of normal genes responsible for processing normal and damaged DNA and/or for chromatin conformation status. In these senses, a single mutated copy of the ataxia telangiectasia gene (ATM) gene occurs in approximately 1% of the general population; these individuals may be at an increased risk of carcinogenesis or radiosensitivity. It has been described very recently that possession of an ATM mutation may be predictive of an increased rate of subcutaneous late tissue effects after radiotherapy for breast cancer (Iannuzzi *et al.* 2002).

It seems clear that in order to quantify the relative merits of new treatment modalities or treatment techniques accurately, clinically relevant radiobiological models to predict response are needed. Once accurate genetic- and/or cell survival-based predictive assays become available, radiation therapy will become an exact science allowing truly individual optimization and allow consideration of the various side-effects that the patient is willing to accept (Brahme 2001).

Mechanism and tolerability of hormone therapies in breast cancer

The normal breast epithelium undergoes cyclical waves of proliferation, differentiation and apoptosis during the menstrual cycles, and after lactation arrest it undergoes remodelling by apoptosis. Factors including oestradiol, progesterone, prolactin, glucocorticoids, insulin, aldosterone, growth factors, as well as contact with the extra-cellular matrix and with the immune system's cells, are all coordinate events that lead finally to mammary epithelium renewal. In the last two decades, studies on breast cancer biology, and relevant information about the biological characteristics of the tumours, have made us aware that in pathological situations many proteins are differently expressed and that they, via cell-type-specific signalling pathways, may contribute to cancer proliferation and tumour progression through the inhibition of apoptosis.

One of the aims of anti-tumour therapy is to obtain tumour regression by inhibiting proliferation and stimulating apoptosis in cancer cells as specifically as possible. One possible way to induce apoptosis in cancer cells is by blocking signal transduction of growth factors and hormone receptors. Apoptosis is a biological event which has been described in

experimental systems to be under the control of two main categories of proteins: (a) the bcl-2 family members and (b) the family of cysteinyl aspartate-specific proteases, the caspases. Up to now, at least 13 caspases have been identified in mammalian cells. In experimental models, it has been shown that anti-oestrogen-induced apoptosis is mediated by both caspase-3 and c-jun-NH₂-terminal kinase, this mechanism being effective against certain ER-negative tumours (Mandlekar *et al.* 2000).

Tamoxifen (TAM), a non-steroidal anti-oestrogen, has been used as a treatment for breast cancer for a quarter of a century. A general issue regarding the use of TAM as endocrine therapy is the fact that during long-term treatment drug resistance may occur and that the compound bears the risk of producing second malignancies (Assikis *et al.* 1996). It is believed that one of the main reasons for these disadvantages is based on the fact that this compound has, besides its antagonist properties, also agonist properties.

In breast cancer, TAM has cytostatic effects mediated through the ER, although functional heterogeneity within populations of cancer cells contributes to seemingly paradoxical effects of anti-oestrogens and the development of anti-oestrogen resistance. In addition, TAM has also been shown to be cytotoxic to both ER-positive and ER-negative cells and such an effect is believed to be mediated by the induction of apoptosis. The cell survival threshold is determined by the balance between cell death-suppressor and cell death-promoter signals provided by external factors or stimuli as well as by intracellular molecules. Proteins of the caspase family are typically present in inactive forms in the cytosol, and are activated by binding to apoptosis-related death receptors or by sequential cleavages. On the other hand, the bcl-2 family of genes has a central role in the control of the response acting either as an apoptosis inhibitor or promoting the cell death programme (Daidone *et al.* 1999, Lilling *et al.* 2000, Ruiz de Almodóvar *et al.* 2001). Despite knowledge about the ER status, it is not always possible to predict which breast cancer will respond to TAM. However, it seems obvious that a relationship between effective TAM therapy and its known anti-proliferative (Danova *et al.* 1993) and pro-apoptotic actions (Chen *et al.* 1996) are representative of different mechanisms of tumour response (Cameron *et al.* 2000). The induction of apoptosis by the compound ICI 128780 (ZM) is associated with a dramatic decrease in bcl-2 expression (Diel *et al.* 1999). In cells lacking ER, it has been demonstrated that TAM is able to induced oxidative stress, thereby causing thiol depletion and activating the transcriptional factor nuclear factor (NF)κB through the modulation of redox metabolism. Importantly, the induction of apoptosis by TAM is not linked to down-regulation or functional inactivation by phosphorylation of the anti-apoptotic bcl-2 protein (Ferlini *et al.* 1999). An early event in the induction of apoptosis by TAM involves mitochondrial depolarization and caspase activation that ultimately lead to cellu-

lar disassembly, and that mechanism occurs independently of p53 (Diëtze *et al.* 2001).

In tumour biopsies, increased bcl-2 expression correlates with the expression of ER, whereas a low expression of bcl-2 characterizes most of Brca-1-associated breast cancer and its down-regulation may account for the increased apoptosis and the high proliferative rate observed in Brca-1-associated carcinomas (Freneaux *et al.* 2000).

In summary, effective anti-oestrogen therapy appears to cause tumour regression by two, possibly separate, mechanisms: (a) reducing bcl-2 expression that is associated with increased apoptosis and (b) reducing proliferation with no significant change in Bcl-2 expression. In addition, TAM has recently been suggested to exert an anti-angiogenic effect that contributes to reducing tumour bulk when this drug is used in postmenopausal women with large, operable or locally advanced ER-rich primary breast cancers (Marson *et al.* 2001).

There have been several major developments concerning the use of hormonal agents for the treatment of breast cancer over the past few years which impact on current endocrine therapy. Not only do we have a larger number of agents in the clinic, but there has also been a trend to reduced drug toxicity. New treatments are being assessed after surgery, but are also being tested pre-operatively and in combination with radiotherapy and chemotherapy. Among them, endocrine treatments are effective in the following situations. (a) As adjuvant systemic therapies in premenopausal women: these include reducing circulating oestrogen by ablation or by inhibiting ovarian oestrogen production. (b) As adjuvant systemic therapy in postmenopausal women: the mainstay of therapy is the prevention of oestrogen binding to its receptor using an anti-oestrogen or lowering oestrogen levels with aromatase inhibitors. (c) As neoadjuvant systemic therapy in locally advanced breast cancer. (d) As primary treatment for elderly breast cancer patients with locoregional disease and as palliative treatment. (e) For prevention in women with a high risk of developing breast cancer.

Most of our endocrine treatments deprive the tumour cells of oestradiol. New data and the newest treatments available indicate the need to re-think our general approach to endocrine therapy and endocrine prevention (Howell *et al.* 2001). This is of particular interest today because the results of new combined therapies, including chemotherapy, endocrine therapy and radiotherapy modalities, have also been reported (Crump *et al.* 1997, Brain *et al.* 1999, Zambetti *et al.* 1999, Tan *et al.* 2001, Vicini *et al.* 2001).

TAM was introduced more than 20 years ago for the palliative treatment of advanced breast cancer. Response rate to treatment ranges from 20% to 60% depending on patient and tumour characteristic and on evaluation stringency. Although this therapy is generally seen as less toxic than many other cancer treatments, it is true that quantifiable data on the side-effects of hormonal therapies such as TAM or

goserelin are scarce. A critical review of the published reports describes morbidity in women treated in such a way as minimal or well tolerated, even when accompanied by data suggesting that the majority of patients do experience some undesirable symptoms. Furthermore, TAM is associated with an increased risk of uterine cancer, thromboembolism, hot flushes, sexual dysfunction and cataracts (Fabian & Kimler 2001) and we need to point out that without good comprehensive morbidity profiles for hormone therapies, prospective patients cannot make informed judgements on proposed treatments (Fellowes *et al.* 2001).

The objectives of combined therapy

The biological problems presented by combined radiotherapy and hormone therapy are considerable. Radiotherapy by itself induces complex changes both in tumours and in the adjacent normal tissues. The response to hormone therapy and chemotherapy are similarly complex: in addition to some of the factors that determine the response to radiation treatment we also have problems of drug delivery, drug resistance and metabolism. The combination of radiotherapy and endocrine therapy involves the combined complexity of both modalities, plus the interaction between them. A combination of drugs and radiation can result as non-interactive if each modality appears to exert its own individual effect; interactive refers to a situation where there is evidence that one modality modifies the effect of the other at comparable levels of overall toxicity.

A combination of radiotherapy and hormonal therapy administered sequentially is indeed useful in the management of breast cancer patients and it is supported because (a) radiation is used to treat the primary tumour and endocrine therapy is added as an adjuvant to deal with systemic spread of disease; there might be spatial co-operation between them; (b) although cellular apoptosis may be a common outcome, each one of the two therapeutic modalities might reach this effect through independent mechanisms; (c) both therapies produce changes in the cell cycle distribution that seems to occur by different mechanisms; and (d) collateral side-effects from one or other modality appear different and the possibilities of their being cumulative are poor. A number of clinical trials and a great deal of empirical experience have shown, without any doubt, that both treatments play major roles in breast cancer treatment.

For instance, local control is generally obtained by modified radical mastectomy and radiotherapy on the chest wall, but in a selected subgroup of women, i.e. those with a tumour of limited size without oedema or erythema, conservative surgery may be offered. Apart from the details of dose fractionation, many other factors can influence the incidence and severity of normal tissue reactions. These include concomitant treatment with chemotherapy (Smith & Lipton 2001). However, other data have pointed out a significant reduction

in the surgical requirements for mastectomy after treatment with neoadjuvant chemoendocrine therapy, without deterioration in local or distal relapse (Makris *et al.* 1998), although this treatment has not provided obvious survival benefit to women with breast cancer (Gazet *et al.* 2001). The need for mastectomy is reduced but not abolished; in some studies this effect is associated with a small increase in the risk of local recurrence (Smith & Lipton 2001). Post-operative radiotherapy forms an intrinsic part of breast conservation therapy, substantially reducing the risk of breast relapse.

Early complications of radiotherapy include tiredness, skin erythema and moist desquamation but the radiation-induced toxicity on the patients' skin is found with approximately the same frequency regardless of the breast cancer surgical treatment applied. In addition, in our experience, the administration of chemotherapy and concomitant irradiation does not increase the intensity, or its frequency, of the acute adverse effects observed on the skin included in the treatment field (López *et al.* 2002). However, it is generally accepted that some chemotherapy drugs, including taxanes, may enhance the effectiveness of radiation therapy although they may increase the incidence of radiation pneumonitis (Taghian *et al.* 2001). Because 5 weeks of whole-breast radiotherapy results in one of the most significant impediments to the widespread use of breast-conserving techniques, acceleration of treatment by confining radiotherapy to the tumour bed has recently been explored. The result of the accelerated treatments using an interstitial implant to deliver radiation to the tumour bed alone over 4–5 days seemed to produce 5-year results equivalent to those achieved with conventional treatment (Vicini *et al.* 2001). In Vicini *et al.* (2001), selection criteria were highly restrictive and follow-up was relatively short. Extended follow-up will therefore be required to determine the long-term efficacy of this treatment approach.

Most of the acute side-effects of treatment are reversible, and there is little or no detectable increase in long-term risk of cardiac or lung toxicity or second cancers with the use of current regimens (Lind *et al.* 2001). An increased incidence of radiation pneumonitis among patients treated with local regional radiotherapy compared with those receiving local radiotherapy only has been reported recently (Lind *et al.* 2002). Moreover, women treated with breast conservation and post-operative radiotherapy showed a good quality of life with a preserved favourable body image and a lack of negative impact on sexuality. Radiation therapy does not lead to any significant additional problems capable of affecting the quality of life (Amichetti *et al.* 1999). However, unconventional and more aggressive irradiation protocols are usually associated with the aggravation of acute reactions and it has been noted that a non-healing acute response can directly progress into a late effect; that consequential effects might be minimized by means of the amelioration of the acute response to irradiation (Dörr & Hendry 2001) and an

extended follow-up will be required to determine the long-term efficacy of each new treatment.

Breast cancer research has developed at a rapid pace over the last decades and new options have been evaluated (Tubiana-Mathieu *et al.* 2001). Recent discoveries promise to provide individualized treatment options. However, there are no data to support that using different treatments jointly might obtain an enhancement of tumour response quantifiable in terms of increase of overall survival with no impairment in the quality of life of the patients. We believe that this, conceptually and experimentally, is a very difficult problem that must first be conducted in experimental models.

Perspectives

A possible way to progress in basic research with a clear projection to clinical practice is the study of chromatin's conformation and its pharmacological modulation. In the mammalian genome, methylation takes place only at cytosine bases that are located 5' to a guanine in a CpG dinucleotide (Bird 2002). These epigenetic changes, in particular aberrant promoter hypermethylation that is associated with inappropriate gene silencing, affect virtually every step in tumour progression (Jones & Baylin 2002). DNA methylation is a biochemical modification of chromatin status that can influence gene expression. Our ideas of how DNA methylation is established in the mammalian genome are changing as rapidly as is our understanding of how chromatin organization modulates gene expression.

Very strong evidence accumulated in the past years suggests that cancer cells usurp and use this mechanism to their benefit by inactivating tumour suppressor genes and causing loss of proteins involved in cell cycle regulation, DNA transcription, DNA repair and apoptosis (Santini *et al.* 2001). This has led to the pursuit of DNA methyl transferase inhibition as a drug target for therapeutic intervention (Bender *et al.* 1998). Not surprisingly, targeting these pathways often also results in radiosensitization (Pervan *et al.* 2001), perhaps through the activation of p53 DNA damage response pathway (Karpf *et al.* 2001) and increases in the amount of DNA radiation-induced damage and impairing cellular repair and recovery.

Loss of expression for both the ER α and E-cadherin genes has been linked to disease progression in human ductal breast carcinomas and has been associated with aberrant 5' CpG island methylation (Nass *et al.* 2000). Restoration of ER expression could be a strategy to improve the response of the breast negative receptor tumours to treatment with anti-oestrogen. One DNA methyltransferase inhibitor undergoing clinical evaluation is 5-aza-2'-deoxycytidine (clinical name, decitabine). Taking into account that aberrant methylation occurs in tumours but not in the normal surrounding tissue, this characteristic might be used to improve the clinical outcome of women with breast cancer without increasing

the associated side-effects. Another drug that inhibits methylation is procainamide, which is used for the treatment of cardiac arrhythmias. This drug is a non-competitive inhibitor of the methyltransferase enzymes and, given the therapeutic promise of reversing methylation changes in cancer, several companies are pursuing this important goal to identify better inhibitors of DNA methylation to be used in clinical studies.

Growing evidence suggests a relationship between alterations in chromatin structure by histone deacetylation and DNA methylation, the development of cancer and higher resistance to treatment. Hopefully, the restoration of chromatin status could provide a modulation in the radiation and hormonal response of tumours (Allan *et al.* 2000). This approach could be important because combined demethylating agents and histone deacetylase inhibitors induce a 300- to 400-fold increase in the transcription of ER, suggesting that the activities of both DNA methyl transferase and histone deacetylase are key regulators of methylation-mediated ER silencing (Yang *et al.* 2001). Because ER is a critical growth-regulatory gene in breast cancer, and also a target for anti-oestrogen therapy, is it important to understand its transcriptional regulation better. Whether the pre-treatment of breast cancer cells with 5-aza-2'-deoxycytidine sensitizes cells against anti-oestrogen and radiotherapy, enhancing antiproliferative, necrotic and apoptotic cell death, needs to be further investigated.

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