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Divergent effects of TGF- β inhibition in bone metastases in breast and lung cancer

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Date of receipt: 27/02/2013

Date of acceptance: 29/04/2013

Work scholarship from the SEIOMM to attend the 33 Congress of the ASBMR (Toronto, Canada. 2010).

Summary

Background: The objective of this study lies in the determination of the validity of transforming growth factor β (TGF- β) as a therapeutic target in models of metastasis deriving from different histological types of lung cancer.

Material and methods: 4-week-old immunodeficient mice inoculated with lung and breast cancer lines were treated with cytokine inhibitor peptide, control peptide or placebo. Weekly bioluminescence and microradiographic measurements were taken to determine the effects of the treatment on tumor burden and metastatic lesions in the long bones.

Results: Treatment with the specific peptide against TGF- β has a protector effect in the bone of animals inoculated with the breast cancer lines, unlike what happens in the control peptide and placebo groups. However, the anti-TGF- β treatment lacks the significant therapeutic effects on the bone metastases which develop in lung cancer bearing animals.

Conclusions: The role of TGF- β as a potential therapeutic target in bone metastasis is highly dependent on the histopathological type and subtype of tumor.

Key words: TGF- β , bone, metastasis, animal models.

Introduction

The skeleton is one of the preferred targets for tumour cells. Neoplasia of the breast, prostate, lung and myeloma very frequently generate bone metastases¹. The prognosis for survival from a diagnosis of bone metastasis varies depending on the type of tumour. In patients with lung cancer with this type of metastasis, the average survival is measured frequently in months, being the lowest of all types of tumour with bone tropism². Furthermore, on many occasions this phenomenon is detected at the time of diagnosis of the disease, a fact which contributes to this neoplasia being the primary cause of death from cancer³.

The high frequency of bone metastasis may be explained by the endothelial fenestrations of the bone marrow, which could facilitate the establishment of the metastatic cells⁴. The bone is also a propitious environment for the development of these cells, since it is a medium rich in growth factors such as transforming growth factor β (TGF- β)⁵. The intracellular signalling route for this cytokine involves the phosphorylation of the Smad proteins, which ultimately allows the expression of the target genes in the nucleus. TGF- β has opposing functions in carcinogenesis. On the one hand, signalling through its receptor triggers an anti-proliferative response in conditions of oncogenic stress. This takes place through the induction of the expression of tumour suppressor genes such as the cyclin-dependent kinase inhibitors (CDKI) or the repression of oncogenes such as c-Myc and members of the ID family⁶. On the other, there are neoplasms which keep this pathway intact, evading this cytostatic response and simultaneously favouring tumour progression. Notable among other mechanisms is its contribution to the evasion of immunity mediated by the T CD8+lymphocytes⁷ and angiogenesis by the induction of vascular endothelial growth factor (VEGF) and MMPs (matrix metalloproteinases) in the tumour and the endothelium⁸. The tumour cell may also use TGF- β signalling for the progression of the metastasis in the bone microenvironment, given that this cytokine favours the expression of the protein related to the parathyroid hormone (PTHrP) or IL1^{9,10}. These factors induce the expression of RANKL (ligand for receptor activator for nuclear factor κ B) in the membrane of the osteoblasts. The recognition of the ligand for the corresponding receptor in the mononuclear precursors entails their activation and the formation of the mature osteoclasts. The action of these cells promotes the release from the bone matrix of cytokines which prime the metastatic colonisation. Thus a positive feedback process or "vicious circle" is generated which magnifies the osteolytic effects.

The genetic or pharmacological inhibition of the signalling pathway of TGF- β has been shown to have therapeutic benefits in preclinical models. The expression of the dominant negative form of the TGFBR2 receptor in breast tumour cells or the use of the TGFRI SD-208 inhibitor in animals inoculated with melanoma cells with bone tropism has shown a reduction in metastatic lesions in the

bone which these cell lines occasion^{11,12}. Similarly, our group demonstrated this effect in a model of bone metastasis derived from a large cell pulmonary carcinoma line¹³.

The main objective of this study is to assess the contribution of TGF- β to bone metastasis in models derived from other histopathological models of lung cancer such as adenocarcinoma or carcinoid. A model of bone metastasis derived from breast cancer was used as a control.

Material and methods

Cell culture

Tumour cell lines A549, H727 and MDA-MB-231 from lung adenocarcinoma, lung carcinoid and breast cancer, respectively, were used. The lines A549 and H727 were transfected with the retroviral vector SFG-NES-TGL (kindly donated by Dr Ponomarev), which contains the luciferase reporter gene. The cells were cultivated at 37°C and 5% CO₂ in sterile conditions in RPMI (A549 and H727) or DMEM (MDA-MB-231) medium, supplemented with 10% FBS, 100 units/ml of penicillin and 100 μ g/ml of streptomycin (Invitrogen®).

Animals and intracardiac inoculation (I.C)

An injection of tumour cells was made into the left ventricle of 24 immunosuppressed mice, 4 weeks of age (Harlan Laboratories) in accordance with previous descriptions^{14,15}. The cells were resuspended in PBS at a concentration of 2x10⁶ cells/ml. The animals were anaesthetised intraperitoneally prior to the inoculation with ketamine (65 mg/kg) and xylazine (2.5 mg/kg). 2x10⁵ cells (100 μ l) were injected using a 29G calibrated needle.

All the protocols for working with laboratory animals were approved by the Ethics Committee for Animal Experimentation of the University of Navarra (CEEAA).

Therapeutic regimen

Control peptides p41, or anti-TGF- β p17 or p144 (kindly donated by Digna Biotech as available) were used to assess this cytokine as a therapeutic target. Both had similar activity *in vivo*^{13,16}. Five days after the inoculation of the tumour cell lines the animals were divided in 3 groups of 8 mice, each of which was to receive a daily dose, intraperitoneally, of 3.75 mg/kg of p144, control peptide (p41) or vehicle (physiological serum). The mice inoculated with the MDA-MB-231 line were divided equally, and from day 7 treated on alternate days with 2.5 mg/kg of p17 (similar activity to p144), control peptide p41 or vehicle.

The dose of p17 was chosen on the basis of its activity in previous studies^{13,17}, while that of p144 was higher to ensure that the results obtained *in vivo* with the lung cancer cell lines were not attributable to a low concentration of the peptide. The time period between the intracardiac inoculation and the start of the treatment had been established in earlier experiments, in which the time necessary for the cells to be detected in bone was determined, through bioluminescence or isolation of the metastatic cells.

The effect on the tumour was determined on a weekly basis through bioluminescence and/or osteolysis through X-ray analysis, looking for possible differences.

The duration of each experiment depended on the development of metastasis for each cell line.

Bioluminescence

D-luciferin (PROMEGA) was administered intraperitoneally at a concentration of 150 mg/kg to the previously-anaesthetised animals. After 5 minutes different images of bioluminescence were taken in a CCD chamber using the Living Image (IVIS® system, Xenogen) programme. The same programme was used to quantify the bioluminescent signal defining the lower extremities. From the data obtained was subtracted the luminometric value of a mouse not injected with cells (control). The values obtained were divided by the luminometric value of each extremity obtained before the start of treatment. The luminometric signal obtained appears superimposed on the rodent.

The cells transfected with SFG-NES-TGL, A549 y H727 vectors had a detectable signal. No tumour-related luminometric data from the animals inoculated with the MDA-MB-231 line was obtained, given that the vector with the luciferase gene with which it was transfected did not generate a signal, probably due to the methylation of the promoter¹³.

Radiographic analysis

The X-rays were carried out under anaesthetic using a Faxitron® (MX-20) X-ray model. Film sensitive to this radiation (MIN-R, Kodak®) was used at 20kV for 20 seconds at 2x magnification. The radiographs were digitized at a resolution of 1200 ppi (Epson® Expression 1680 Pro). The area of osteolytic lesions was analysed with the computerised image analysis programme AnalySIS® (GmbH). The relative quantification of the metastatic areas was expressed as the percentage of the sum of the areas of lesion in the femur and tibia with respect to the total surface area of these long bones on the films.

Computerised microtomography (µCT)

Femoral and tibial joints representative of each experimental group were analysed in a microCT device (micro CAT II, Siemens® Preclinical Solutions) at 75.0 kVp and 250 uA. Each scan was carried out at a resolution of 10 µm. The two-dimensional images were reconstructed using a standard deconvolution procedure with a Shepp-Logan filter. For the reconstruction of the images the COBRA®_Exxim programme was used. The images were stored in three-dimensional frames with a voxel size of 19*19*23 µm.

Statistical analysis

The SPSS 15.0 software programme was used to determine the statistical value of the results. A significance level of $\alpha=0.05$ was used. Values of p lower than this limit were considered as significant

(*). A single factor ANOVA analysis was carried out to study the metastatic area of the radiographs with multiple Tukey comparisons and the Kruskal-Wallis test followed by multiple comparisons with Mann-Whitney for bioluminescence. The p values obtained were adjusted with the Bonferroni method.

Results

The radiological analysis of the femurs and tibias of the animals inoculated with the MDA-MB-231 line showed a drastic reduction in the metastases of the animals treated with p17 in comparison with those treated with control peptide or placebo three weeks from the initiation of the experiment (Figure 1). In the animals inoculated with the H727 line, the bioluminescence images at day 22 showed a slight effect of p144 on the tumour load in the lower extremities of the animals treated with this peptide (*p< 0.05, Figure 2A). This effect lost its significance in the days following the experiment (Figure 2B). Similarly, the radiographical analysis of the long bones revealed a slight reduction in metastatic lesions in these mice which was not significant (Figures 3A and 3B). On the other hand, in the animals inoculated with the A549 line the p144 did not demonstrate any protective effect on the tumour load determined by bioluminescence (Figures 4A and 4B) or on the development of osteolytic lesions (Figures 5A and 5B). These results indicate that the prometastatic activity of TGF- β is highly dependent on each cell line.

Discussion

Metastasis involves the acquisition of new functions on the part of the cell in the primary tumour. These include motility and invasion, intravasation, survival in circulation, adhesion to the endothelium, extravasation and growth or colonisation of the target organs¹⁸. This requires a genetic and/or epigenetic programme –little understood at present– influenced by the selective pressure established in the tumour itself and its microenvironment. Therefore, the identification of the targets involved in metastasis is critical. Knowledge of the factors effecting this process could allow the development of new antimetastatic therapies which would have an impact on the quality of life of cancer patients. This fact justifies the development of models of metastasis which reproduce clinical reality. The model based on intracardiac inoculation recapitulates the final stages of metastasis: extravasation, homing, and colonisation of target organs. With this technique a high and rapid incidence of metastasis with reproducible results is achieved. This approximation has been used before with tumour lines of melanoma, prostate and breast^{10,19}. The main limitation is the exclusion of the initial events of the metastatic cascade such as invasion, motility and intravasation towards the pulmonary parenchyma. The models which recapitulate these initial stages of metastasis such as orthopaedic injection have been successful for breast cancer. Here, an extirpation of the primary tumour after its

Figure 1. (Left) Radiographic analysis at day 21 of the metastatic area in the long bones of animals inoculated with breast cancer line MDA-MB-231. The osteolytic lesions (arrows) were evaluated after treatment with placebo (PBS), p17 (anti-TGF- β peptide) or p41 (control peptide). * $p < 0.05$. (Right) representative X-rays of long bones of animals from each group

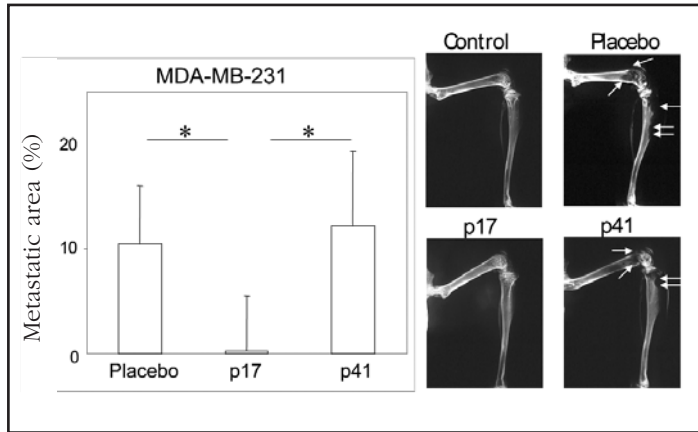
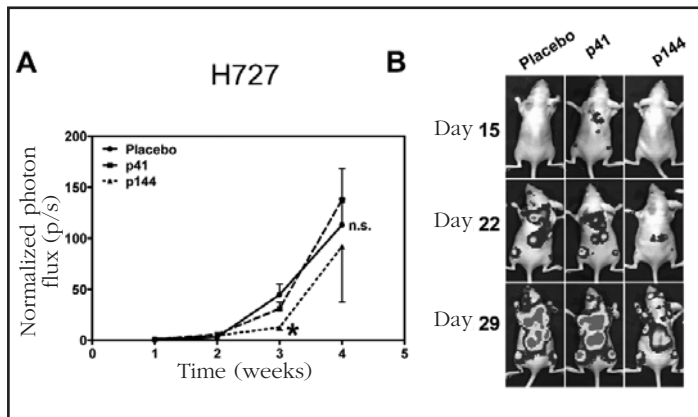


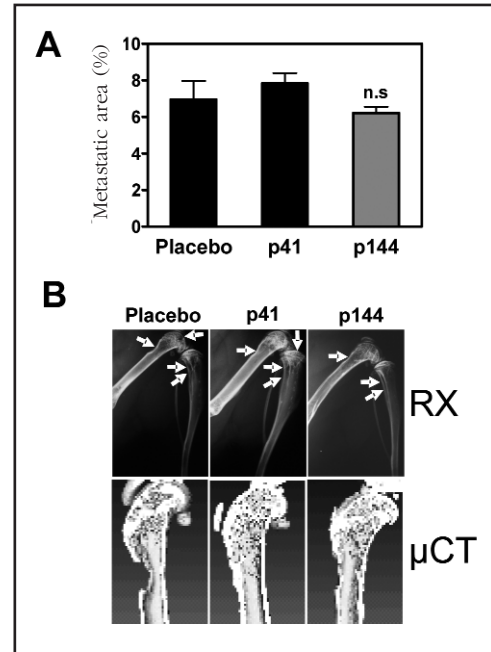
Figure 2 A. Analysis of bioluminescence in mice inoculated with the H727 line and treated with placebo, p144 (anti-TGF- β peptide) or p41. A moderate reduction in the tumour load 22 days after inoculation is observed in animals treated with p144 (* $p < 0.05$; n.s., not significant in the later week). B. Images of bioluminescence representative of each group during the course of the experiment



growth is carried out to facilitate the later appearance of metastases at a distance²⁰. However, this approximation is not viable in the case of lung cancer, since the death of the animals frequently ensues before the development of the metastasis, due to the rapid growth of the tumour. Furthermore, the frequency of bone metastasis in orthotopic models is low.

In spite of the importance of TGF- β in the bone microenvironment and in metastasis, its therapeutic potential should be treated with caution. Firstly, the systemic inhibition of this pathway may generate collateral affects, since signalling through this pathway is of vital importance in the homeostatic process of tissues. Secondly, the results obtained show that the effects of this pathway depend on the cellular context. Breast tumour cells evade

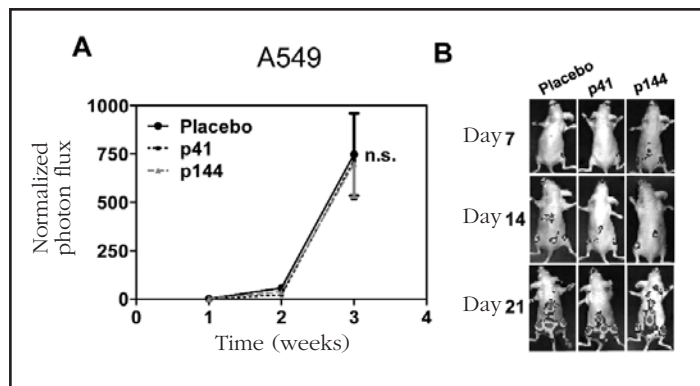
Figure 3. A. Radiographic analysis of the metastatic area in the long bones of animals inoculated with the pulmonary carcinoid line H727. B. Histomorphometric analysis: X rays (XR) and computerised microtomography (μ CT)



TGF- β 's inhibitory signals, keeping intact the route to the limbs which favours the prometastatic function of the cytokines. The great abundance of transcriptional inhibitors of the genes involved in the cytostatic response in these cells may account for this phenomenon²¹. This may explain the therapeutic improvement obtained in the model of bone metastasis of the MDA-MB-213 cell line. As occurs with breast cancer, pulmonary oncogenesis may involve the loss of the tumour-suppressant effects TGF- β ²². Therefore, this

cytokine may be of therapeutic interest for the treatment of bone metastasis of lung cancer. In agreement with these results, the use of a peptide inhibitor of TGF- β in a model of large cell carcinoma showed a reduction in bone metastasis¹³. However, the results shown in this work demonstrate that this effect on adenocarcinoma and carcinoid of the lung are of little or no significance. These experiments substantiate therefore the cell-dependent context of the TGF- β pathway. Furthermore, it is possible that other cytokines present in abundance in the bone, such as IGF-1²³, may constitute key elements for the progression of the vicious circle and consequent metastatic colonisation of bone. Future studies may determine other ideal targets for the development of antimetastatic therapies.

Figure 4. A. Analysis of bioluminescence in those mice inoculated with the A549 line treated with placebo, p144 or p41. (n.s., not significant). B. Images of bioluminescence representative of each group during the course of the experiment

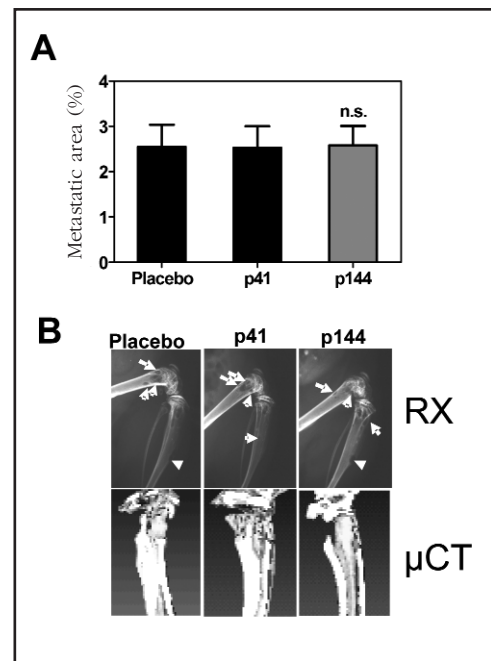


This work was financed by the "FIMA UTE project" agreement, RTICC RD06/0020/0066, PI042282, FIT-090100-2005-46, SAF-2009-11280 and the "Ortiz de Landáruzi" Award (67/2055, Government of Navarra) and "Fundación La Caixa" obtained by F.L.D. L-R by FIMA and the FPU National Programme and I.A by FIMA and the Government of País Vasco. F.L is a researcher in the Programme 13.

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Figure 5. A. Radiographic analysis of the metastatic area in the long bones of animals inoculated with the pulmonary adenocarcinoma line A549. B. Histomorphometric analysis: XR and μ CT



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