

# Bone marrow angiogenesis and mast cell density increase simultaneously with progression of human multiple myeloma

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**Summary** Immunohistochemical, cytochemical and ultrastructural data showing vivid angiogenesis and numerous mast cells (MCs) in the bone marrow of 24 patients with active multiple myeloma (MM) compared with 34 patients with non-active MM and 22 patients with monoclonal gammopathy of undetermined significance (MGUS) led us to hypothesize that angiogenesis parallels progression of MM and that MCs participate in its induction via angiogenic factors in their secretory granules.

**Keywords** angiogenesis; mast cell; multiple myeloma; tumour progression

Angiogenesis is obligatory in the enhancement of progression (growth, invasion and metastasis) of solid tumours (Folkman, 1995). New vessels promote growth by conveying oxygen and nutrients and removing catabolites, whereas endothelial cells secrete growth factors for tumour cells (Hamada et al, 1992; Folkman, 1995). Endothelial cells also secrete a variety of matrix-degrading proteinases which facilitate invasion (Mignatti and Rifkin, 1993). Lastly, an expanding endothelial surface increases opportunities for tumour cells to enter the circulation and metastasize (Aznavorian et al, 1993).

Tumour cells may not be the only source of angiogenic factors within a tumour. Host inflammatory cells, including fibroblasts, macrophages and mast cells (MCs), which are recruited and activated by tumour cells via paracrine mechanisms act synergistically with these cells by secreting the same or other factors (Polverini, 1996). MCs play a decisive role in the synergism (Norrby and Whooley, 1993). Also, experimentally induced tumours display MC accumulation close to the tumour cells before the onset of angiogenesis (Kessler et al, 1976), and those induced in MC-deficient mice display both reduced angiogenesis and ability to metastasize (Starkey et al, 1988; Dethlefsen et al, 1994).

Knowledge on these relations in haematological tumours is circumstantial. Angiogenesis is correlated with tumour growth (S-phase fraction) in monoclonal gammopathies (Vacca et al, 1994), and with progression stages in B-cell non-Hodgkin's lymphomas (Ribatti et al, 1998) and in mycosis fungoides (Vacca et al, 1997).

This paper presents the results of an investigation on angiogenesis and MC counts in the bone marrow of patients with monoclonal gammopathy of undetermined significance (MGUS) and multiple myeloma (MM) grouped according to a pathway of progression.

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## MATERIALS AND METHODS

### Patients

A total of 80 Caucasian patients who fulfilled the South West Oncology Group diagnostic criteria for MM and MGUS (Durie, 1991) were studied (Table 1). Myeloma patients were defined as active or non-active, according to clinical performance and

**Table 1** Patient clinical and immunological data

Total no.	80
Multiple myeloma	58
Active	24
Average age (median, range)	64 years (66.5, 42–87)
Men/women	15/9
M-component IgG/IgA/IgD $\kappa$ or $\lambda$	16/6/1/1
Diagnosis	10
Stage I/II/III; A/B <sup>a</sup>	1/2/7; 6/4
Relapse <sup>b</sup>	8
Progression	6
Non-active	34
Average age (median, range)	66 years (68, 45–80)
Men/women	20/14
M-component IgG/IgA $\kappa$ or $\lambda$	22/8/4
Response	20
Plateau <sup>c</sup>	14
Monoclonal gammopathy of undetermined significance	22
Average age (median, range)	62 years (63.8, 45–86)
Men/women	12/10
M-component IgG/IgA/IgM	18/2/2

<sup>a</sup>According to Durie and Salmon. <sup>b</sup>Relapse defined as M-component increase >50% from the lowest value, or clinical and bone marrow relapse when the M-component did not reflect tumour load and disease activity. <sup>c</sup>Plateau phase defined as post-treatment M-component decrease >50%, and lasting for at least 6 months without treatment.

**Table 2** Microvessel area and mast cell counts in the bone marrow of patients

	MGUS (n = 22)	Non-active MM (n = 34)	Active MM (n = 24)
Microvessel area ( $\mu\text{m}^2$ )	1.1 $\pm$ 0.5 (0.9; 0.2–2.5)	1.2 $\pm$ 0.6 (1.3; 0.2–2.2)	5.7 $\pm$ 3* (5.2; 1.2–12.8)
Number of mast cells	1.3 $\pm$ 1* (1; 0–3)	1.6 $\pm$ 1.2 (1.5; 0–4)	4.8 $\pm$ 2* (5; 1–8)

Results are expressed as means  $\pm$  1 standard deviation (median; range) in 250 $\times$  microscopic fields (125  $\mu\text{m}^2$ ). The cellular area in MGUS, non-active and active MM was 42.1  $\pm$  8.8  $\mu\text{m}^2$ , 46.6  $\pm$  11.2  $\mu\text{m}^2$  and 52.4  $\pm$  9.6  $\mu\text{m}^2$ . \* $P < 0.01$  compared with non-active MM and MGUS

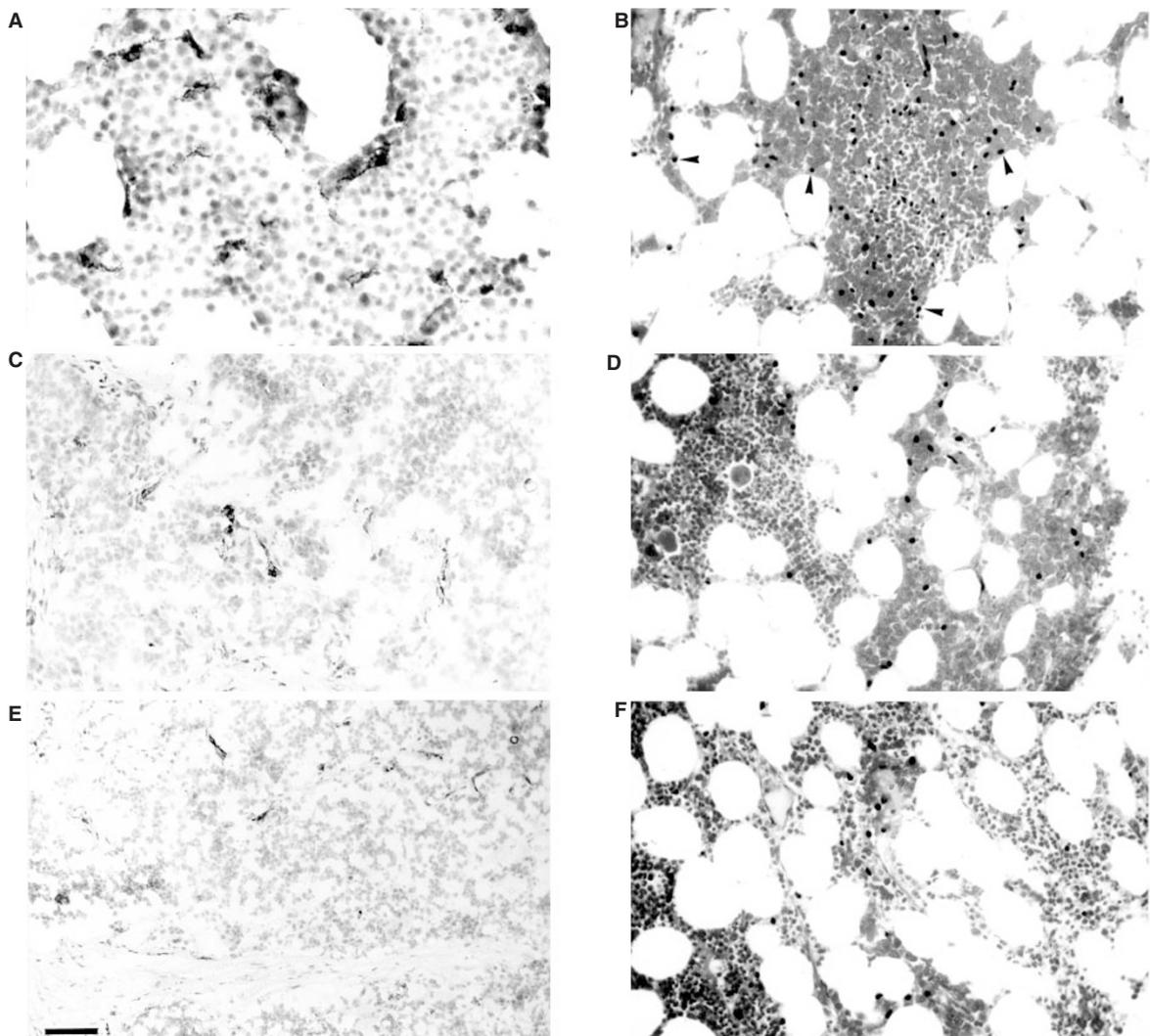
M-component level (Durie, 1991). Active patients were those: (a) at diagnosis, with symptomatic disease and an increase in the M-component level in the 3 months before analysis; (b) at relapse; (c) with unresponsive and rapidly progressive disease (leukaemic

progression), characterized by severe bone pain, hypercalcaemia and pancytopenia. Non-active patients were those in: (a) post-treatment complete/objective response; (b) the off-treatment plateau phase. MGUS, non-active-MM and active MM constitute a progression pathway because: (i) the clinical evolution from one step to the next is typical; (ii) the plasma cell S-phase fraction and tumour mass rise significantly in the transition from one step to the next (Durie, 1991).

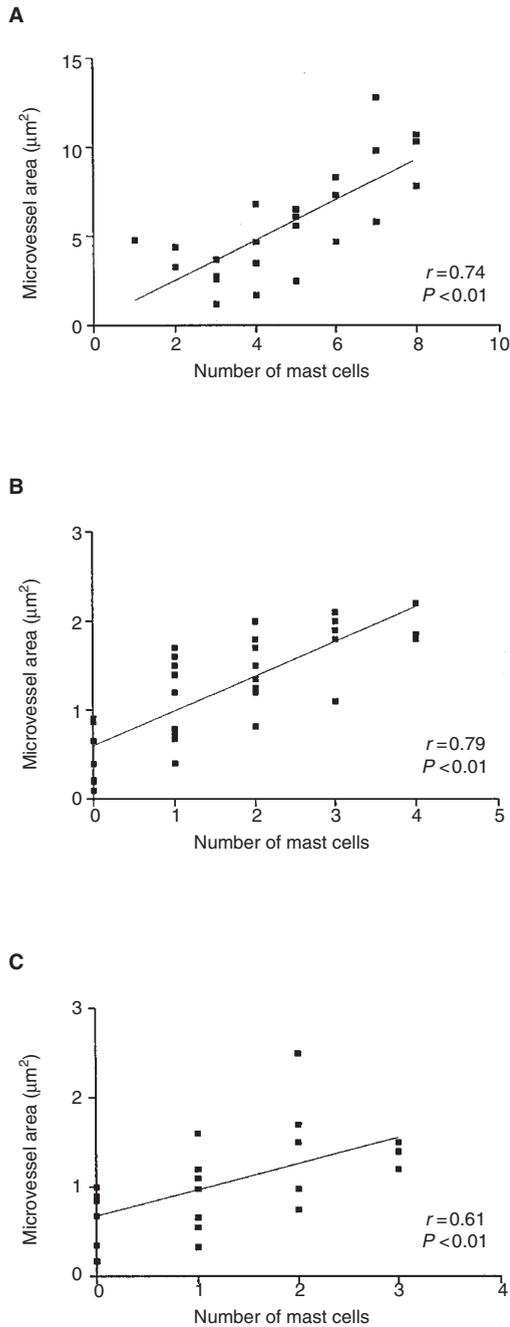
The study was approved by the local ethics committee and all patients gave their informed consent.

### Measurement of bone marrow angiogenesis

All blood vessels were displayed in 6- $\mu\text{m}$  sections of 4% paraformaldehyde-fixed paraffin-embedded biopsies by staining endothelial cells with the anti-factor VIII murine monoclonal antibody M616 (IgG1; Dako, Glostrup, Denmark) and a three-layer biotin-avidin-peroxidase system described previously (Vacca et al, 1994). The very few megakaryocytes also stained by factor VIII

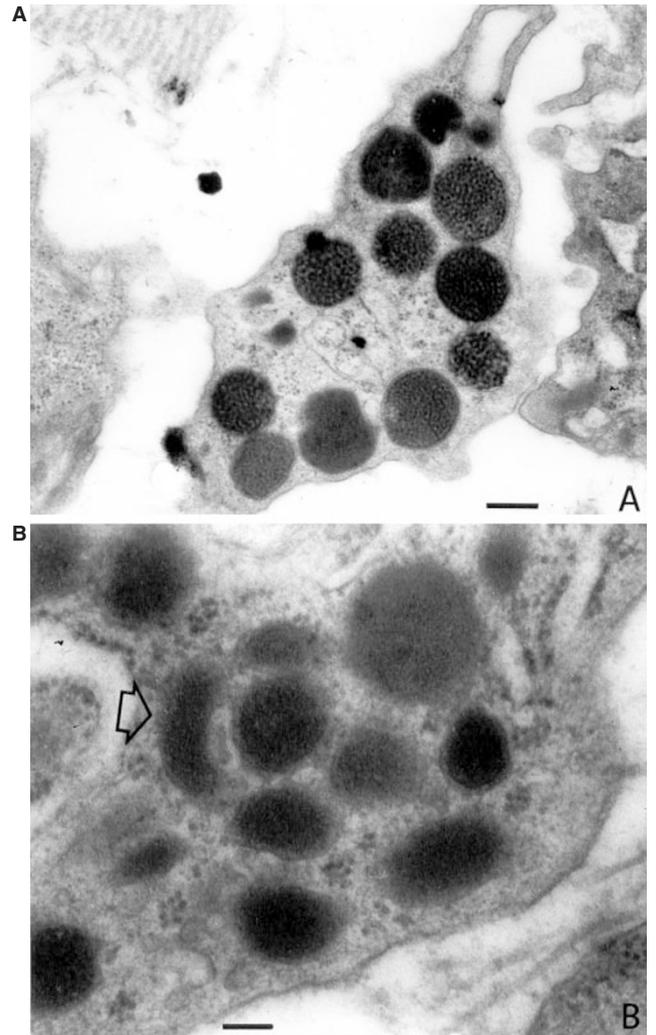


**Figure 1** Adjacent sections of bone marrow biopsies stained with factor VIII for microvessels (A, C, E) and with toluidine blue for mast cells (B, D, F) from patients with: (A) and (B) active MM (relapse); (C) and (D) non-active MM (plateau); and (E) and (F) MGUS. Note the higher density of microvessels and mast cells (some are arrowheaded) in the active MM patient. Bar = 10  $\mu\text{m}$



**Figure 2** Mast cell counts in comparison with the microvessel area in the bone marrow of patients with (A) active and (B) non-active MM and (C) with MGUS. Significance of the regression analysis was calculated by the Pearson's (*r*) test

were easily distinguishable by their morphology. Angiogenesis was measured as microvessel area without knowledge of final diagnosis. Briefly, six to eight 250× fields covering the whole of each of two sections per biopsy were examined with a superimposed 484-point square reticulum (125 µm²) to identify microvessels (capillaries and small venules) as endothelial cells either single or clustered in nests or tubes, and clearly separated from one another, and either without or with a lumen (not exceeding 10 µm). A planimetric point count method (Elias and Hyde, 1983) with slight modifications for the



**Figure 3** Ultrastructural findings of bone marrow biopsies from patients with active MM. In (A), a mast cell with typical electron-dense round granules and in (B), at higher magnification, a cytoplasmic granule with a semilunar aspect (arrow), among other typical round granules, is recognizable. Bar, (A), 0.08 µm; (B) 0.02 µm

computed image analysis (Leica Quantimet 500, Wetzlar, Germany) was applied to measure the microvessel area within the cellular area (reticulum area minus connective tissue, fat, bone lamellae, necrosis and haemorrhage areas) (Vacca et al, 1994). Values are expressed as means ± 1 standard deviation (s.d.) per group of patients.

**MC counts**

MCs were highlighted in two sections adjacent to that stained for microvessels with 0.5% aqueous solution of toluidine blue (Merk, Darmstadt, Germany). Cells were counted in six to eight 250× fields inside the reticulum and calculated as means ± 1 s.d. for each group of patients.

**Electron microscopy**

Small pieces (approximately 1 mm³) of tissue were fixed in 3% glutaraldehyde in 0.1 M phosphate-buffered saline (PBS) for 3 h,

washed in the same buffer for 12 h, post-fixed in 1% osmium tetroxide, dehydrated in graded ethanols and embedded in Epon 812. Ultrathin sections were cut with a diamond knife on a LKB V ultratome, stained with uranyl acetate followed by lead citrate, and examined in a 9A Zeiss electron microscope.

## Statistics

The significance of changes in the microvessel area and MC counts in the groups was determined with the parametric (Fisher's test) and non-parametric (Kruskal-Wallis test) analysis of variance, followed by Duncan (*t*), Bonferroni (*t*), and Wilcoxon tests to compare groups two by two. Correlations between microvessel area and MC counts in the groups were assessed with the Pearson's (*r*) coefficient and simple regression analysis. Data were computed with the Statistical Analysis Software (SAS, SAS Institute, Cary, NC, USA).

## RESULTS

Table 2 shows the microvessel area normalized to the total cellular area and the MC counts on bone marrow adjacent sections of patients with active MM, non-active MM and MGUS. The area was significantly larger in patients with active MM than in those with non-active MM and with MGUS, between which variations were negligible. In parallel, the MC counts were significantly higher in active MM than in the other groups. The differences in microvessels and MC are also shown in Figure 1. The within-group comparison showed that both parameters were always significantly correlated (Figure 2). At the ultrastructural level, typical MCs with cytoplasmic matrix filled by numerous electron dense secretory granules (Figure 3A), and MCs with semilunar aspect of granules (Figure 3B) were recognizable. The latter imply slow, chronic release of mediators in response to a moderate, progressive, degranulatory stimulus (Kops et al, 1984; Ribatti et al, 1988).

## DISCUSSION

In the current study, we showed that bone marrow angiogenesis (evaluated as microvessel area) and MC counts were highly correlated in patients with non-active and active MM and in those with MGUS, and that both parameters increased simultaneously in active MM. As the progression from in situ to invasive and metastatic solid tumours is accompanied and facilitated by the switch from the prevascular to the vascular phase (Hanahan and Folkman, 1996), our findings suggest that the active MM represent the 'vascular phase' of plasma cell tumours, whereas the non-active MM and MGUS represent the 'prevascular phase'. Bone marrow angiogenesis may, therefore, favour the progression from MGUS or non-active MM to active MM. As in solid tumours, where angiogenesis could be stimulated either directly or indirectly, after the tumour cells have recruited inflammatory cells stimulating them to secrete their own angiogenic factors (Hanahan and Folkman, 1996), the switch in MGUS and non-active MM may be induced by tumour plasma cells via secretion of angiogenic factors, namely interleukin 1 (IL-1) (Cozzolino et al, 1989), IL-6 (Schwab et al, 1991), tumour necrosis factor alpha (TNF- $\alpha$ ) (Lichtenstein et al, 1989), macrophage colony-stimulating factor (M-CSF) (Nakamura et al, 1989), transforming growth factor beta (TGF- $\beta$ ) (Klein, 1995), and by inflammatory cells, including MCs, via secretion of their angiogenic factors.

MCs are strikingly associated with angiogenesis, as found in chronic inflammatory diseases, namely rheumatoid arthritis and psoriasis, and in tumours, namely haemangiomas, carcinomas and lymphomas (Meininger and Zetter, 1992; Norrby and Woolley, 1993; Qu et al, 1995; Ribatti et al, 1998). In tumours, MCs are recruited and activated via several factors secreted by tumour cells: the *c-kit* receptor, or stem cells factor (Poole and Zetter, 1983; Norrby and Woolley, 1993), as well as the basic fibroblast growth factor (FGF-2), vascular endothelial growth factor (VEGF-A) and platelet-derived endothelial cell growth factor (Gruber et al, 1995). MCs contain, in secretory granules, heparin that in vitro stimulates endothelial cell proliferation and migration (Thorton et al, 1983; Alessandri et al, 1984), whereas in vivo it has been shown to have variable effects on angiogenesis; it may, thus, stimulate (Ribatti et al, 1987; Norrby, 1993), inhibit (Jakobson and Hahnenberger, 1991; Norrby, 1993; Wilks et al, 1991) or have no effect (Castellot et al, 1982; Taylor and Folkman, 1982). Histamine and tryptase, other MC-derived factors, also stimulate angiogenesis (Sorbo et al, 1994; Blair et al, 1997). In addition, MCs produce a variety of multifunctional cytokines and growth factors, such as IL-6 and IL-8 (Motro et al, 1990; Norrby, 1996), TNF- $\alpha$  (Beil et al, 1994), granulocyte-macrophage colony-stimulating factor (GM-CSF) (Bussolino et al, 1991), TGF- $\beta$  (Roberts et al, 1986), FGF-2 (Qu et al, 1995) and VEGF-A (Grutzkan et al, 1996), which may contribute to angiogenesis in active MM.

As concerns the ultrastructural features of MCs, the semilunar, or partial degranulating, aspect of their secretory granules, unlike IgE-mediated massive degranulation which occurs during the immediate hypersensitivity reactions, is typical of a slow degranulation, taking place in delayed hypersensitivity reactions and in chronic inflammatory processes (Kops et al, 1984; Ribatti et al, 1988). In tumours, such as MM, the semilunar aspect of MC secretory granules might correspond to a slow but progressive release of angiogenic factors, in consequence of a chronic and progressive stimulation of MC degranulation.

Tentatively, our data suggest that an increasing number of MCs may be recruited and activated by more malignant plasma cells in active MM, and that angiogenesis in this disease phase may be mediated, at least in part, by angiogenic factors contained in their secretory granules.

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