INTRODUCTION

Bone destruction is a hallmark of myeloma and affects 80% of patients. Myeloma cells promote bone destruction by activating osteoclasts. In investigating the underlying mechanism, we found that C-reactive protein (CRP), a protein secreted in increased amounts by hepatocytes in response to myeloma-derived cytokines, activated myeloma cells to promote osteoclastogenesis and bone destruction in vivo. In mice bearing human bone grafts and injected with multiple myeloma cells, CRP bound to surface CD32 (also known as FcγRII) on myeloma cells, which activated a pathway mediated by the kinase p38 MAPK and the transcription factor Twist that enhanced the cells’ secretion of osteolytic cytokines. Furthermore, analysis of clinical samples from newly diagnosed myeloma patients revealed a positive correlation between the amount of serum CRP and the number of osteolytic bone lesions. These findings establish a mechanism by which myeloma cells are activated to promote bone destruction and suggest that CRP may be targeted to prevent or treat myeloma-associated bone disease in patients.

RESULTS

CRP promotes lytic bone lesions in myeloma-bearing mice

To determine the effect of CRP on bone destruction under physiological conditions, we injected human recombinant CRP, serum amyloid P component (SAP; an inactive analog of CRP), or phosphate-buffered saline (PBS) (vehicle) into tumor-free severe combined immunodeficient (SCID) or SCID-hu mice in which the SCID mice were implanted with human bone chips. None caused bone destruction in the implanted human bones in the mice (Fig. 1A, left).

We then examined the effect of human CRP on bones in SCID mice xenografted with human myeloma cells. CRP, SAP, or PBS was
injected into SCID-hu mice bearing primary human myeloma cells or SCID mice bearing myeloma cell lines. Lytic bone lesions were detected in the implanted human bone chips of SCID-hu mice 8 weeks after tumor inoculation (Fig. 1A, right), whereas in mice injected with CRP but not SAP or PBS, lytic bone lesions were detected at week 4 after tumor injection. CRP not only accelerated the appearance of bone lesions but also caused more bone destruction, evident by the greater number of lytic bone lesions (Fig. 1A, right), larger lytic area (Fig. 1B), and lower trabecular bone volume (Fig. 1C) compared with controls. Similarly, CRP also resulted in more bone lesions in murine femurs of SCID mice bearing human myeloma cells (Fig. 1, D to F). The earlier appearance and larger number of bone lesions in CRP-injected myeloma-bearing mice were not due to higher tumor burden because no difference in tumor burden, detected as circulating human immunoglobulin G (IgG) secreted by myeloma cells, was detected in myeloma-bearing SCID-hu (fig. S1A) or SCID (fig. S1B) mice injected without CRP, indicating that CRP affected not myeloma growth but rather bone destruction in vivo. We injected 20 μg of CRP per mouse twice a week continuously for 8 weeks, which maintained CRP concentrations of 3 to 8 μg/ml in mouse serum (fig. S1C and D). Mean values of serum CRP in myeloma patients reported for different studies range from 3.8 to 12.0 mg/liter, with measured values ranging from 1.5 to 120 mg/liter (9, 11, 16). Therefore, the CRP concentrations in the mice were comparable to those seen in myeloma patients.

To confirm the results, we generated human myeloma cell lines (ARP-1 and MM.1S) that stably produced (fig. S2A) and secreted (fig. S2B) human CRP protein. After myeloma was established in SCID mice, these cells also caused more lytic bone lesions in the femurs of SCID mice bearing CRP-expressing or vector-control myeloma cells (Fig. 1, G and H). Injection of a neutralizing antibody against human CRP diminished the ability of CRP-producing, but not control, myeloma cells to cause lytic bone lesion in the mice (Fig. 1I). Because these CRP-expressing myeloma cells secreted much lower amounts of CRP than those observed in myeloma patients, it is possible that this autocrine CRP, although at low concentrations but constantly secreted, may accumulate on myeloma cells to effectively bind with the surface receptors to activate signaling in myeloma cells. Thus, these results indicate that human
CRP is capable of promoting myeloma cell–mediated lytic bone disease in vivo.

**CRP enhances OC differentiation and bone resorption activity**

Myeloma manifests as an osteolytic bone disease mainly due to enhanced OC activity (17). To determine whether CRP affected osteoclastogenesis, primary myeloma-bearing mice were sacrificed 8 weeks after tumor injection, and sections of murine femurs or implanted human bone chips were stained for Tartrate-resistant acid phosphatase (TRAP) to identify OCs. CRP injection resulted in significantly more TRAP+ OCs localized on the trabecular bone surface of the bone-tumor interface (Fig. 2, A and B) and more erosion on trabecular bones (Fig. 2C) in the implanted human bones of myeloma-bearing SCID-hu mice compared with controls. Similarly, more OCs (Fig. 2D) or bone erosion (Fig. 2E) were found in the femurs of myeloma-bearing, CRP-injected SCID mice than those injected with PBS and in mice bearing CRP-producing myeloma cells than those injected with control myeloma cells (Fig. 2F). Injection of CRP-neutralizing antibodies but not control IgG abrogated the enhanced bone destruction produced by CRP-producing myeloma cells (Fig. 2G).

Because the in vivo findings suggested that CRP affected osteoclastogenesis in myeloma-bearing mice, we investigated the role of CRP in OC differentiation in vitro. We observed that adding CRP (5 μg/ml, equal to the average serum levels of CRP in myeloma patients) to cultured OC precursor cells did not induce the formation of mature (multinuclear, TRAP+) OCs (Fig. 3A) or increase the levels of TRAP5b (Fig. 3B), which is an active component of TRAP and is secreted only by mature OCs; the TRAP5b level reflects OC-mediated bone resorption activity (18). However, coculture of OC precursors with myeloma cells, without addition of RANKL in the medium, induced formation of mature OCs and addition of CRP (5 μg/ml) to the coculture further and significantly enhanced the generation of mature OCs (Fig. 3, A and B). In line with these results, the expression of OC-associated genes, such as TRAP and those encoding calcitonin receptor (CALCR) and cathepsin K (CTSK) (19), was highly enhanced in OCs cocultured with myeloma cells in the presence of CRP (Fig. 3C), as compared with controls. Coculturing OC precursors

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**Fig. 2. CRP enhances osteoclastogenesis in myeloma-bearing mice.** (A) Representative images of Tartrate-resistant acid phosphatase (TRAP)+ osteoclasts (OCs) in human trabecular bones from SCID-hu mice or 8 weeks after inoculation with patient myeloma cells (Pt 1) and treatment with CRP or PBS. Arrows indicate OCs in bone sections. No MM, tumor-free mice. Scale bars, 10 μm. (B and C) Histomorphometric quantitative analysis of (B) the number of OCs on the bone surface (Oc. S/BS) and (C) OC-mediated erosion of bone surface, expressed as a percentage of total bone surface (ES/BS), in the implanted human bones of SCID-hu mice that were either tumor-free (−) or bearing primary myeloma cells from one of five patients (Pt MM) and treated with CRP (+) or PBS (−). (D and E) Histomorphometric analysis of (D) Oc. S/BS and (E) ES/BS in the femurs of SCID mice (n = 10 per group) bearing myeloma cells and treated with CRP or PBS. (F) Oc. S/BS in the femurs of SCID mice bearing CRP-expressing (MM-CRP) or vector-control (MM-Vector) cells. (G) Oc. S/BS in the femurs of SCID mice bearing MM-CRP or MM-Vector cells treated with antibodies against CRP (αCRP) or control (IgG). Tumor-free mice (no MM) treated with agents or antibodies served as controls for baseline levels. Data from three independent experiments are shown. *P < 0.05, **P < 0.01 by Student’s t test.

**Fig. 3. CRP enhances myeloma-induced OC formation and activity in vitro.** (A and B) The number of multinuclear TRAP+ cells (A) and the amount of secreted TRAP5b (B) in cultures of OC precursor cells cultured alone or with either primary myeloma cells isolated from five myeloma patients (Pt MM) or with human myeloma cell lines (MM) in medium without receptor activator of nuclear factor κB (NF-κB) ligand (RANKL) in the presence of CRP (5 μg/ml) or PBS. (C) Real-time polymerase chain reaction (PCR) analysis of the expression of CALCR, CTSK, and TRAP in mature OCs cocultured with OC precursors and ARP-1 or MM.1S cells with or without CRP (5 μg/ml). (D) TRAP staining for the number of multinuclear TRAP+ cells in cultures of OC precursors with vector- or CRP-expressing ARP-1 or MM.1S cells (MM-Vector or MM-CRP) without RANKL. (E) As in (D), in the presence of antibodies against CRP (αCRP; 5 μg/ml) or control (IgG). *P < 0.05, **P < 0.01 by Student’s t test. Data from four independent experiments are shown.
with CRP-producing human myeloma cells also enhanced mature OC formation (Fig. 3D). Once again, neutralizing CRP in the coculture abrogated CRP-induced, myeloma-mediated OC formation (Fig. 3E). Together, these results indicate that CRP enhances myeloma cell-mediated osteoclastogenesis in vitro.

**CRP stimulates the production of OC activators by myeloma cells**

Because our results showed that addition of CRP to the transwell coculture of OC precursors and myeloma cells induced OC differentiation (Fig. 3), we hypothesized that CRP mediates this effect by up-regulating myeloma cell secretion of cytokines that activate OC differentiation. Microarray analysis of gene profiling identified more than 500 genes with twofold increased expression in CRP-treated myeloma cells compared to PBS-treated cells, including 11 that encode cytokines known to activate osteoclastogenesis: *IL-1*, *MCP-1*, *IL-11*, *IL-6*, *HGF*, *IL-17*, *RANKL*, *M-CSF*, *BAFF*, *DcR3*, and *MIP-1α* (fig. S3A). Real-time polymerase chain reaction (PCR) validated that among these cytokines, *RANKL*, *MCP-1*, and *MIP-1α* were highly expressed in CRP-treated myeloma cells, including patient-derived primary myeloma cells and myeloma cell lines ARP-1 and MM.1S, as compared with control myeloma cells (fig. S3B). Not surprisingly, CRP-expressing ARP-1 and MM.1S cells also expressed a higher level of these cytokines as compared with vector-control myeloma cells (fig. S3C).

To determine the involvement of these cytokines in CRP-enhanced OC differentiation, we added neutralizing antibodies against these cytokines to the coculture of OC precursors and myeloma cells in the presence of CRP. The neutralizing antibodies significantly reduced the number of TRAP+ multinuclear OCs (fig. S4A) and level of TRAP5b (fig. S4B). These results indicate that CRP enhances OC differentiation by up-regulating myeloma cell production of OC activators such as RANKL, MCP-1, and MIP-1α.

**CRP binds to CD32 on myeloma cells, promotes cytokine production, and induces OC differentiation**

Because CRP has been shown to bind to FcγRs such as CD32 on human cells including myeloma cells (7, 15), we hypothesized that CD32 expressed on myeloma cells may be involved in CRP-induced myeloma-mediated OC differentiation. We generated stable myeloma cell lines (ARP-1 and MM.1S) with significantly reduced expression of CD32 [herein called CD32-knockdown (CD32-KD); Fig. 4A]. CD32-KD myeloma cells were less able to promote OC differentiation in response to CRP (Fig. 4, B and C). Moreover, CD32-KD myeloma cells caused less bone disease in mice injected with CRP (Fig. 4, D and E), although similar numbers or percentages of CD32-KD and control MM cells were found in the bone marrow of mice (Fig. 4F). As expected, CD32-KD myeloma cells had lower CD32 expression in vivo compared with control cells (Fig. 4G). We then added CRP to cultures of CD32-KD or control ARP-1 or MM.1S cells. The CD32-KD myeloma cells...
CRP regulates the expression of osteolytic cytokines in myeloma cells through p38 MAPK–Twist signaling

Next, we sought to elucidate the molecular mechanism underlying CRP–CD32–mediated signaling pathways that regulate the production of osteolytic cytokines in myeloma cells. First, we analyzed the expression of transcriptional factors in myeloma cell lines by microarray and identified 31 genes that were at least twofold increased in expression of transcriptional factors in myeloma cell lines by microarray and identified 31 genes that were at least twofold increased in response to CRP (Fig. 6A). We used the software Genomatix (20) to analyze the transcription factor binding sites on the promoter regions of the osteolytic cytokine genes and predicted several potential Twist binding sites. To confirm the results of microarray analysis, we performed a chromatin immunoprecipitation (ChIP) assay to examine the ability of Twist to bind to the promoters of two relevant cytokine (RANKL and MCP-1) genes. We found that Twist was enriched on RANKL and MCP-1 promoters in ARP-1 and MM.1S promoters (Fig. 6B). To investigate the importance of Twist in CRP-induced cytokine production by myeloma cells, we used short hairpin RNA (shRNA) to knock down Twist (Twist-KD) abundance in myeloma cells (Fig. 6C). CRP induction of myeloma cell expression of the osteolytic cytokines MCP-1 and RANKL was significantly reduced in Twist-KD myeloma cells as compared to control cells (Fig. 6C). We also cloned the Twist-containing transcription start site of the RANKL and MCP-1 promoters into a luciferase reporter vector. Adding CRP increased the luciferase activity of Luc-RANKL and Luc-MCP-1 in ARP-1 and MM.1S, and truncation of the Twist binding sites in the MCP-1 or RANKL gene promoters significantly reduced the activity (Fig. 6, D and E), suggesting that the potential Twist binding sites for RANKL locate at −1500 to −800 base pairs (bp) of its start codon and those for MCP-1 locate at −350 to −150 bp of its start codon. Next, we performed site-directed mutagenesis on the two putative binding sites for RANKL and three putative binding sites for MCP-1 within each region. Not surprisingly, mutating a single binding site for either RANKL or MCP-1 blocked the transcriptional activation induced by CRP (Fig. 6, D and E). Furthermore, CRP–treated Twist-KD myeloma cells were less able to induce the expression of OC genes in cocultured OC progenitor cells (Fig. 6F), indicating that Twist is crucial to myeloma cell–mediated OC differentiation in response to CRP.

To further examine the effect of CRP on Twist expression, we treated ARP-1 or MM.1S cells with various concentrations (1 to 10 μg/ml) of Twist binding sites. To confirm the results of microarray analysis, we performed a chromatin immunoprecipitation (ChIP) assay to examine the ability of Twist to bind to the promoters of two relevant cytokine (RANKL and MCP-1) genes. We found that Twist was enriched on RANKL and MCP-1 promoters in ARP-1 and MM.1S promoters (Fig. 6B). To investigate the importance of Twist in CRP-induced cytokine production by myeloma cells, we used short hairpin RNA (shRNA) to knock down Twist (Twist-KD) abundance in myeloma cells (Fig. 6C). CRP induction of myeloma cell expression of the osteolytic cytokines MCP-1 and RANKL was significantly reduced in Twist-KD myeloma cells as compared to control cells (Fig. 6C). We also cloned the Twist-containing transcription start site of the RANKL and MCP-1 promoters into a luciferase reporter vector. Adding CRP increased the luciferase activity of Luc-RANKL and Luc-MCP-1 in ARP-1 and MM.1S, and truncation of the Twist binding sites in the MCP-1 or RANKL gene promoters significantly reduced the activity (Fig. 6, D and E), suggesting that the potential Twist binding sites for RANKL locate at −1500 to −800 base pairs (bp) of its start codon and those for MCP-1 locate at −350 to −150 bp of its start codon. Next, we performed site-directed mutagenesis on the two putative binding sites for RANKL and three putative binding sites for MCP-1 within each region. Not surprisingly, mutating a single binding site for either RANKL or MCP-1 blocked the transcriptional activation induced by CRP (Fig. 6, D and E). Furthermore, CRP–treated Twist-KD myeloma cells were less able to induce the expression of OC genes in cocultured OC progenitor cells (Fig. 6F), indicating that Twist is crucial to myeloma cell–mediated OC differentiation in response to CRP.

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CRP and assessed Twist expression by Western blot. CRP enhanced Twist expression in a dose-dependent manner (Fig. 7A). In contrast, Twist expression was reduced in CRP-treated CD32-KD myeloma cells (Fig. 7B). To further investigate the potential downstream targets of Twist, we added various concentrations (1 to 10 µg/ml) of CRP to cultures of myeloma cells and found that CRP enhanced phosphorylation of p38 MAPK but not extracellular signal-regulated kinase (ERK) or NF-κB in a dose-dependent manner (Fig. 7C). CRP promoted the levels of phosphorylated p38 MAPK in control and in CD32-overexpressing myeloma cells but not in CD32-KD myeloma cells (Fig. 7D), indicating that CRP activates the p38 MAPK pathway in myeloma cells through surface CD32. To examine the importance of p38 MAPK in CRP-induced cytokine production in myeloma cells, we added the p38 MAPK–specific inhibitor SB202190 to cultures of myeloma cells along with CRP. SB202190 significantly reduced the expression of osteolytic cytokine mRNAs (Fig. 7E) and reduced CRP-induced Twist protein abundance (Fig. 7F). These results showed that CRP binds to CD32 and activates p38 MAPK–Twist signaling pathways, leading to increased production of osteolytic cytokines by myeloma cells.

**DISCUSSION**

Myeloma is accompanied by bone disease in the vast majority of patients (1, 2). Myeloma cells disrupt the delicate balance between bone formation and resorption, leading to debilitating osteolytic bone lesions (3). Although it is well established that myeloma cells enhance OC differentiation and activity via secreting osteolytic cytokines (17), the mechanism underlying how myeloma cells are regulated and activated to secrete these cytokines remains elusive. Here, we showed that human CRP, which is secreted in elevated amounts by hepatocytes in serum CRP levels significantly and positively correlate with bone lesions in patients with newly diagnosed myeloma

Finally, we examined the relationship between serum CRP and development of bone disease in patients with myeloma. In newly diagnosed myeloma patients, serum CRP was quantified by enzyme-linked immunosorbent assay (ELISA), and bone disease and the number of bone lesions were detected by x-ray and magnetic resonance imaging. As shown in Fig. 8A, serum CRP level significantly and positively correlated with the number of bone lesions (n = 244; r = 0.6096; P < 0.0001). We examined bone marrow biopsies from myeloma patients with high (≥6 bone lesions) or low (<3 bone lesions) numbers of bone lesions and found that more CRP protein was present in the bone marrow of myeloma patients with more bone disease (Fig. 8B). Moreover, positive correlations were also found between the levels of serum RANKL (Fig. 8C) or MCP-1 (Fig. 8D) and CRP level. We quantified serum collagen type I C-telopeptide (CTx)–1, which reflects the bone resorption activity of OCs, and observed that CTx-1 and CRP level (Fig. 8E) also positively correlated. These results are clinically significant because they suggest that high levels of CRP may be a key causative factor in the development of the bone disease observed in 80% of patients with myeloma.
response to cytokines (IL-1 or IL-6) derived from myeloma cells in patients (6), may be responsible, at least in part, for activating myeloma cells to promote osteoclastogenesis and inducing bone destruction in vivo. CRP binds with surface CD32/FcγRII, activates p38 MAPK–Twist pathways, and stimulates the secretion of osteolytic cytokines by myeloma cells.

Our previous study showed that CRP at a much higher concentration (25 μg/ml) activated ERK and NF-κB but inhibited p38 MAPK (15). Our clinical analyses examining the relationship between the level of serum CRP and the number of osteolytic bone lesions in newly diagnosed patients support this conclusion. Thus, this study reveals a previously unknown mechanism that explains how myeloma cells are activated to promote osteoclastogenesis and induce bone destruction in human myeloma. Our study also suggests that CRP may be a therapeutic target for preventing or treating bone disease in patients with myeloma.

CRP is a sensitive systemic marker of inflammation and tissue damage (6). Its level in the serum is increased in patients with infection, inflammatory disease, necrosis, or malignancy, including myeloma, lymphoma, and carcinoma, and high levels of circulating CRP correlate with poor prognosis in patients with myeloma or lymphoma (9–13). We hypothesized that CRP may not be just a surrogate for tumor burden in myeloma but also plays a central role in myeloma growth, survival, and drug resistance. We reported that adding human CRP to myeloma cultures at levels seen in patients with myeloma or other tumors promoted tumor cell proliferation under stressed conditions and protected tumor cells from apoptosis induced by chemotherapy, IL-6 withdrawal, or serum deprivation in vitro and in vivo (15). The results of this study further showed that CRP may be an important player in myeloma-mediated bone destruction in human myeloma. As inflammation has long been hypothesized to be linked to cancer (21), our studies suggest a possible link between

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**Fig. 7. Cross-linking CD32 by CRP activates p38 MAPK–Twist in myeloma cells.**

(A) Western blot for Twist abundance in ARP-1 or MM.1S cells treated with various doses of CRP for 24 hours. (B) Real-time PCR for TWIST mRNA levels in CD32-KD or control myeloma cells cultured with or without CRP. (C) Western blot analysis for dose-dependent effects of CRP on the phosphorylation (p) of p38 mitogen-activated protein kinase (MAPK), extracellular signal–regulated kinase (ERK), and NF-κB in myeloma cells. (D) Western blot analysis for effects of CRP on the phosphorylation of p38 MAPK in CD32-overexpressing (CD32) or knockdown (CD32-KD) ARP-1 cells compared with the respective controls. (E) Real-time PCR analysis of the expression of MCP-1, MIP-1α, and RANKL in ARP-1 cells cultured with or without CRP and the p38 MAPK inhibitor SB202190 (SB20). (F) Western blot analysis for the effect of SB20 on CRP-induced Twist abundance. *P < 0.05, **P < 0.01 by Student’s t test. Data from three independent experiments are shown.

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**Fig. 8. Relationship between serum CRP level and osteolytic bone lesion number in patients with newly diagnosed myeloma.**

(A) Correlation between serum CRP level and the number of lytic bone lesions (BLs) in 244 patients with newly diagnosed myeloma. CRP was measured by ELISA, and BLs were detected by radiography (x-ray or magnetic resonance imaging). (B) Representative images of immunohistochemical staining for CRP accumulation in the bone marrow of myeloma patients (three per group) with low (BL < 3) or high (BL ≥ 6) numbers of BLs. Scale bar, 50 μm. (C to E) Correlations between the level of circulating CRP and (C) MCP-1, (D) RANKL, and (E) CTx-1 in 28 randomly selected patients with newly diagnosed myeloma, calculated by Pearson’s correlation coefficient analysis.
inflammation or its products and the pathogenesis and bone destruction in myeloma.

Twist is a basic helix-loop-helix (bHLH) transcriptional factor essential in embryological morphogenesis, and it belongs to the HLH superfamily of proteins, which are involved in a variety of regulatory processes in organisms (22). Initially identified in Drosophila where it is involved in mesodermal patterning and morphogenetic movement (23), Twist is a highly conserved protein and a master regulator of morphogenesis. Previous studies have also shown that Twist is important in bone development and is expressed in primary osteoblastic cells (24) and preosteoblasts (25). In situ analysis of murine embryonic development provided evidence that Twist abundance is reduced during endochondral and intramembranous fetal bone development (26). Twist proteins inhibit skeletal development through direct and indirect inhibition of Runx2 (27, 28), a key regulator of osteogenic differentiation. Finally, expression of Twist1 and Twist2 in mesenchymal cells may promote adipogenesis (29). Together, these observations suggest that Twist proteins are important to normal bone marrow stromal cell function. However, Twist is also essential to tumor metastasis. Ectopic expression of Twist results in loss of E-cadherin-mediated cell-cell adhesion, activation of mesenchymal markers, and induction of cell motility; it contributes to metastasis by promoting an epithelial-mesenchymal transition. In human breast cancers, high Twist expression correlates with the presence of invasive lobular carcinoma (30). The role of Twist in bone destruction in myeloma is unclear. Our results indicate that Twist mediates CRP-induced up-regulation of MCP-1, RANKL, and MIP-1α expression in myeloma cells, leading to enhanced OC-mediated bone resorption. Further studies are needed to elucidate its effect on pathogenesis of myeloma-associated bone disease.

MATERIALS AND METHODS

Primary myeloma cells and myeloma cell lines

This study was approved by the institutional review board at University of Texas MD Anderson Cancer Center (UTMDACC), and all patients provided written informed consent. Bone marrow aspirates were obtained from newly diagnosed patients with myeloma as part of their routine clinical workup. Patient CD138+ myeloma cells were obtained from newly diagnosed patients with myeloma as part of their routine clinical workup. Patient CD138+ myeloma cells were provided by the Arkansas Cancer Research Center. Other cell lines were purchased from the American Type Culture Collection. All myeloma cells were cultured in RPMI-1640 medium supplemented with 10% fetal bovine serum (FBS) (Life Technologies).

Antibodies and reagents

Neutralizing antibodies against CRP, MIP-1α, MCP-1, RANKL, and control IgG, and recombinant human CRP, M-CSF, RANKL, and SAP were purchased from R&D Systems. The p38 MAPK-specific inhibitor was purchased from Axon Medchem BV. Except where specified, antibodies for Western blotting analysis were purchased from Cell Signaling Technology.

In vitro OC formation and function assays

Human OCs were generated from peripheral blood, CD14 antibody-coated magnetic bead-purified (Miltenyi Biotec) monocytes of healthy blood donors, and cultured in α-minimum essential medium supplemented with 10% FBS with M-CSF (25 ng/ml) (R&D Systems) for 7 days to obtain the precursors of OCs as we previously reported (5). The precursors were then cocultured with myeloma cells in medium without or with a low dose of RANKL (10 ng/ml) for an additional 7 days to induce mature OC formation. To examine the effects of myeloma cells and CRP on OC differentiation, OC precursors (1 × 10⁵ cells/ml) were seeded in the wells and cocultured with myeloma cells (0.5 × 10⁶ cells/ml) in transwell inserts for 7 days, in OC medium without the addition of RANKL, in the presence or absence of CRP. Cells were fixed with 4% formaldehyde and stained for TRAP by using the Leukocyte Acid Phosphatase Kit (Sigma-Aldrich). At the end of the 2-week culture, culture supernatants were collected and used for quantifying by ELISA (CrossLaps, Biosciences Diagnostics) secreted TRAP5b.

Knockdown and overexpression in myeloma cells

The shRNAs specific for CD32 or Twist were purchased from Santa Cruz Technology according to the manufacturer’s instructions. After lentiviral infection, transduced cells were harvested to examine CD32 or Twist expression or used for experiments. The cells with CD32 or Twist knockdown were drug-selected. CRP and CD32 overexpression was achieved using the retroviral expression vector pBABEhybro, according to the manufacturer’s instructions.

Western blotting analysis

Cells were harvested and lysed with lysis buffer [50 mM tris (pH 7.5), 140 mM NaCl, 5 mM EDTA, 5 mM NaN₃, 1% Triton X-100, 1% NP-40, and 1× protease inhibitor cocktail]. Cell lysates were subjected to SDS–polyacrylamide gel electrophoresis, transferred to a polyvinylidene difluoride membrane, and immunoblotted with antibodies against CRP (R&D Systems) and phosphorylated or nonphosphorylated p38 MAPK (Cell Signaling Technology) at 4°C overnight. Antibodies were diluted according to the manufacturer’s recommendations. The membrane was stripped and reprobed with anti-β-actin antibodies (Sigma-Aldrich) to ensure equal protein loading. Secondary antibodies conjugated to horseradish peroxidase were used for detection and followed by enhanced chemiluminescence (Pierce Biotechnology) and autoradiography.

Real-time PCR

Total RNA was isolated with an RNeasy kit (Qiagen). An aliquot of 1 μg of total RNA was subjected to reverse transcription (RT) with SuperScript II (Invitrogen) RT-PCR kit, and 1 ng of the final complementary DNA was applied to real-time PCR amplification with SYBR Green using StepOnePlus real-time PCR system (Life Technologies). The primers used are listed in table S1.

Myeloma cell apoptosis

The fraction of apoptotic cells was determined by staining with fluorescein isothiocyanate–conjugated annexin V and propidium iodide and analyzed by flow cytometry (BD LSRFortessa, BD Biosciences) according to the manufacturer’s instructions.

Mouse models, radiography, histology, and bone histomorphometry

CB.17 SCID mice were purchased from Harlan. All mice were maintained in American Association of Laboratory Animal Care–accredited facilities, and the studies were approved by the Institutional Animal Care and Use Committee of the UTMDACC. Six- to 8-week-old mice were injected intravenously with (1 × 10⁶ cells per mouse) human myeloma cell lines ARP-1 and MM.1S, and myeloma-bearing mice were...
then treated with CRP (20 μg per mouse; twice a week for 6 to 8 weeks) or vehicle (PBS). Serum was collected from mice daily during the treatment and tested for myeloma-secreted M-proteins (human Ig) or their light chains by ELISA.

SCID-hu hosts were developed as described previously (31). Fetal bones (10-mm pieces, purchased from Advanced Bioscience Resources Inc.) were implanted subcutaneously and allowed to become vascularized (6 to 10 weeks). Purified myeloma cells (≥10⁶ cells per mouse) were injected into implanted human bones to establish myeloma (4 to 6 weeks) (n = 5 per group). After 7 days, mice were treated with CRP or SAP (20 μg per mouse) or equal volume of PBS twice a week for 8 weeks. Serum was obtained from the mice weekly and assayed by ELISA for the human M-component.

Mice were x-rayed weekly starting at 1 to 2 weeks after tumor injection. To measure the size of lytic bone lesions, radiographs were scanned with a Faxitron x-ray cabinet (Faxitron x-ray). The total number of osteolytic lesions or area was quantified using ImageJ software (National Institutes of Health). For histological and bone histomorphometric analyses, mice were euthanized, and mouse femurs were fixed in 10% neutral-buffered formalin for 18 hours. To detect mature OCs, a leukocyte acid phosphatase staining kit (Sigma-Aldrich) was used to stain sections with TRAP according to the manufacturer’s instructions. The number of TRAP-positive, multinuclear (>3) OCs per millimeter of bone at the bone-tumor interface was calculated using a computerized image analysis system.

**Immunohistochemistry**

Formalin-fixed, paraffin-embedded sections of SCID mouse femurs were deparaffinized with xylene and rehydrated to water through a graded alcohol series. Endogenous peroxidase activity was quenched with 3% hydrogen peroxide. Expression of CD138 and CD32 was detected using specific antibodies (R&D Systems and LifeSpan BioSciences Inc.). Signal was detected using secondary biotinylated antibodies and streptavidin/horseradish peroxidase. Chromagen 3,3-diaminobenzidine/ H2O2 (Dako) was used, and slides were counterstained with hematoxylin. All slides were observed under light microscopy, and images were captured with a SPOT RT camera (Diagnostic Instruments).

**ChIP assays**

ChIP analysis with anti-Twist was performed using a commercially available kit (Upstate Biotechnology). Immunoprecipitates and total chromatin input were reverse cross-linked, DNA was isolated, and 1-μl DNA was used for PCR with primers specific for the gene promoter region for RANKL and for MCP-1. The primer sequences were as follows: (i) RANKL: forward, CTTGGACCTCCAGAAAGACAG; reverse, ACTCTTATAAACCAGCTTGAGAG; (ii) MCP-1: forward, CTGCTAGGCTTCGTTCTTCTTATGATGCTAC; reverse, TCACTGCTGAGACCAAATGAG.

**Transfection and luciferase assay**

Cells at 50% confluence were transfected in 12-well petri dishes using Lipofectamine (Gibco-BRL). The total amount of DNA was adjusted to 1.0 μg by adding empty vector. We determined luciferase activity by using a luciferase assay system (Promega). As a reference plasmid to normalize transfection efficiency, 25 ng of pRL-CMV plasmid (Promega) was cotransfected in all experiments.

**Microarray analysis**

We performed the microarray analysis as described previously (32). At least a twofold difference between control myeloma cells and CRP-treated myeloma cells was required. Microarray data have been deposited in the Gene Expression Omnibus (GEO) database (GSE103697).

**Statistical analysis**

All data are shown as means ± SD. Student’s t test was used to compare various experimental groups; significance was set at an α level of 0.05.


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C-reactive protein promotes bone destruction in human myeloma through the CD32–p38 MAPK–Twist axis

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Liver protein instructs bone loss in myeloma

Bone loss is common in patients with multiple myeloma (MM). MM cells activate osteoclasts, cells that degrade bone. The sera of MM patients typically have increased amounts of C-reactive protein (CRP), which is secreted by the liver in response to cytokines associated with tissue inflammation and physiological stress, including those secreted by MM. Using samples from patients, as well as human cell lines and mice bearing human bone grafts and MM cells, Yang et al. found that CRP is not merely a diagnostic marker for MM but that rather it feeds back on MM cells to stimulate the expression and secretion of osteoclast-activating cytokines from MM cells, thereby driving bone loss in patients. These findings suggest that targeting the cyclical CRP signaling axis may reduce or prevent MM-associated bone loss.