The burden of the rheumatic diseases in the general adult population of Greece: the ESORDIG study

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The burden of the rheumatic diseases in the general adult population of Greece: the ESORDIG study

Abstract
OBJECTIVE: To estimate the burden of rheumatic diseases in terms of disability and health-care utilization in the Greek general adult population. METHODS: The study was conducted on the total adult population of seven communities (8547 subjects), as well as on 2100 out of 5686 randomly selected subjects in an additional two communities. Rheumatologists visited the participants at their homes to assess the prevalence of six morbidity indicators concerning disability and health-care utilization associated with rheumatic diseases or other major disease groups. RESULTS: The participation rate in the study was 82.1%. The prevalence of chronic health problems, long-term disability, short-term disability, physician office visits and prescription or non-prescription drug use due to rheumatic diseases in the total target adult population was 14.3, 4.3, 2.9, 2.8, 7.2 and 2.0%, respectively. Compared with all other major disease groups, rheumatic diseases were the most common cause of chronic health problems (38.7%), long-term disability (47.2%), short-term disability (26.2%) and physician office visits (20.5%), while they ranked second for the use of prescription (24.0%) or non-prescription drugs (17.7%). Rheumatic diseases were the main cause of morbidity in five out of six indicators in subjects aged < or =65 yr. Logistic regression analysis revealed an association of female gender, age > or =45 yr and obesity with almost all morbidity indicators related to rheumatic diseases. CONCLUSION: These findings suggest that rheumatic diseases constitute a major public health problem and should be considered in planning undergraduate and postgraduate medical education, research and health-care services.

Keywords
burden, adult, rheumatic, population, greece, esordig, study, diseases, general

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Systemic levels of interleukin-6 and matrix metalloproteinase-9 in patients with multiple myeloma may be useful as prognostic indexes of bone disease

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Abstract

Multiple myeloma is characterized by accelerated production of the proteolytic enzyme matrix metalloproteinase (MMP)-9. We hypothesized that myeloma-produced MMP-9 may influence the rate of bone turnover in a paracrine manner. Thus, we examined the correlations of MMP-9 levels, disease severity, and bone turnover rate as evaluated by markers of bone formation and resorption.

Thirty-seven newly diagnosed multiple myeloma patients (nine of Durie-Salmon stage I, 12 of stage II and 16 of stage III) and 12 age-matched controls were studied. Serum MMP-9 levels were significantly higher at stage II compared to stage I (188.78 ± 91.27 vs. 59.25 ± 33.09 ng/mL, p < 0.004). Additionally, free urine pyridinolines (F-Pyd), free urine deoxy-pyridinolines (F-Dpd) and urine N-telopeptide fragment (NTx) were elevated, their levels correlating with disease stage (p < 0.001, p < 0.03, p < 0.001, respectively), as were bone marrow infiltration and serum interleukin-6 (IL-6) levels (p < 0.001, p < 0.01, respectively). MMP-9 levels were lower in patients compared with controls (p < 0.001), whereas IL-6 and bone resorption marker levels were higher in patients than in controls (p < 0.001 in all cases). Significant correlation was found between infiltration, MMP-9, free urine pyd, free urine dpd and NTx for each stage of the disease (p < 0.03, p < 0.003, p < 0.002, p < 0.003 and p < 0.001, respectively). Levels of MMP-9 and of IL-6 in multiple myeloma correlate well with bone turnover rate and may be useful in disease evaluation.

Keywords: matrix metalloproteinases; multiple myeloma; N-telopeptide.

Introduction

Multiple myeloma (MM) is a malignant monoclonal proliferation of bone marrow plasma cells, with debilitating skeletal involvement (1, 2). Skeletal morbidity in MM correlates well with the degree of plasma cell infiltration (3, 4). The progression of the disease involves the production by the tumor cells of a specific class of extracellular matrix-degrading metalloenzymes, including matrix metalloproteinase (MMP)-9, a 92-kDa type IV collagenase (5, 6). MMP-9 degrades collagen IV, the major constituent of basement membrane, and this appears an essential step for cellular invasion and tumor progression (7–9). The pathogenesis of myeloma-induced bone disease is associated with accelerated bone turnover, leading to an imbalance between bone formation and resorption (10). MMP-9 is predominantly expressed in cells of macrophage lineage, such as osteoclasts that use it to hydrolyze bone collagen type I (11, 12). It is postulated that myeloma cell-derived MMP-9 – produced in the vicinity of bones – may enhance organic bone matrix absorption in a direct, paracrine fashion (7, 13). Malignant plasma cell infiltration has been shown to be directly responsible for the increased resorptive activity and abnormal remodeling, either by locally produced autocrine or paracrine osteoclast-activating factors or by direct cell to cell contact (1). Furthermore, the production of MMP-9 by myeloma cells could be accelerated locally by cytokines, chemokines and growth factors produced by bone cells, aggravating, thus, the invasive behavior of the malignant cells, by the establishment of a local self-accelerating cycle (14). Indeed, bone marrow endothelial cells were found to enhance MMP-9 production in MM, indicating that endothelial cells accelerate MM cell invasion via MMP-9 secretion (7, 13, 15). Bone formation and resorption markers represent sensitive indices of the rate of bone turnover (16–18). The aim of the present work was to examine the levels of MMP-9 in patients with MM and age-matched controls, and correlate them to the severity of disease, as well as to the rate of bone turnover. For this purpose, we measured MMP-9 levels and bone resorption biochemical indi-
ces in our patients, and correlated them with the stage of their disease.

Materials and methods

Patients

The study population included 37 newly diagnosed patients with MM: nine in stage I, 12 in stage II and 16 in stage III according to Durie-Salmon classification (19). Serum samples from 12 persons, age- and sex-matched healthy volunteers, were used as controls. Informed consent for the study was obtained from all subjects.

The median age of the patient group was 63 years (range 39–84 years). There were 17 males (age 39–81 years) and 20 females (age 39–81 years). Monoclonal immunoglobulins IgG, IgA and light chain disease were found in 22, 11 and 39–84 years). There were 17 males (age 39–81 years) and 20 females (age 39–81 years). Monoclonal immunoglobulins IgG, IgA and light chain disease were found in 22, 11 and 4 females (age 39–84 years). There were 17 males (age 39–81 years) and 20 females (age 39–81 years). Monoclonal immunoglobulins IgG, IgA and light chain disease were found in 22, 11 and 4 females, respectively. No patient had received chemotherapy or radiotherapy prior to initial sampling. None of the patients had evidence of thyroid or parathyroid hormone-related abnormalities or had received any therapy known to affect bone metabolism. According to standard X-ray evaluation of the patients, bone involvement was graded into four scores: no lesions (score 0), one bone involved or diffuse osteoporosis (score 1), more than one but less than four bone lesions (score 2) and more than four bone lesions or bone fracture present (score 3). Using these definitions, 11 patients had score 0, four had score 1, six had score 2, and 16 patients had score 3.

Methods

Serum samples were collected from patients and controls, aliquoted into separate vials and stored at −70°C after collection, and assayed at the end of the study, in order to avoid inter-assay variability.

Urinary N-telopeptide (NTx) cross-links of type I collagen were measured by a competitive-inhibition enzyme-linked immunosorbent assay (ELISA) (Osteomark Ostex International, Seattle, WA, USA), using a monoclonal antibody for NTx labeled with horseradish peroxidase. Results were collected for 24-h urinary concentration of creatinine and expressed as nanomoles of bone collagen equivalent per millimole of creatinine (nmol BCE/mM creat). Free urinary pyridinolines (F-Pyd) were measured by a competitive enzyme immunoassay (METRA PYD EIA assay, Metra biosystems, Mountain View, CA, USA) using monoclonal antibodies. Free urine deoxy-pyridinolines (F-Dpd) alone were measured by METRA DPD EIA assay (Metra biosystems). Results for F-Pyd and F-Dpd alone were corrected for 24-h urinary concentration of creatinine and expressed as nanomoles per millimole of creatinine.

Measurements of IL-6 and MMP-9 in serum samples collected from patients and controls were performed by solid-phase sandwich ELISA with commercially available test kits, Quantikine® Human MMP-9 (total) and IL-6 (R&D Systems Inc., Minneapolis, MN, USA).

Bone marrow cellularity and percentage of bone marrow infiltration by myeloma cells was estimated by bone marrow biopsies that were taken during local anesthesia from a point below the iliac crest.

Statistical analysis

Results are expressed as mean ± SD. The non-parametric Kruskal-Wallis test and one-way analysis of variance (ANOVA) were assessed to test the existence of differences between different stages. The Student-Newman-Keuls test was used for pairwise comparison of subgroups. Statistical comparisons between the MM group and the control group were made using the non-parametric Mann-Whitney test. Correlations between the various measured parameters were calculated by Spearman’s rank correlation coefficient. p-Values < 0.05 were considered to be statistically significant.

Results

The mean concentration of IL-6 in the entire group of patients was 6.69 ± 5.51 pg/mL and was significantly higher in comparison to that found in the control group (p < 0.001) (Table 1). The mean values for infiltration and IL-6 in the group of MM patients were significantly higher with increasing stage of disease (p < 0.001, p < 0.01, respectively) (Table 1). Subsequently, we evaluated the measured parameters of disease activity, infiltration and IL-6 according to the grade of bone disease. Bone marrow infiltration was significantly higher in grade 3 than in grade 0 (51.88 ± 21.28 vs. 28 ± 16.28, p = 0.02). In contrast, mean serum levels of IL-6 were higher with advancing bone disease, but without statistical significance.

The mean concentrations of NTx, F-Pyd and F-Dpd were significantly higher in MM patients in comparison to those found in the group of controls (p < 0.001

Table 1 Values of plasma cell infiltration, IL-6, MMP-9, and of biochemical markers of bone resorption in 37 untreated MM patients according to disease stage, and in controls. With the exception of MMP-9, values of all measured parameters were significantly higher in MM patients than in controls. Values for infiltration, IL-6, and urinary N-telopeptide (NTx) were significantly higher with advancing stage of the disease.

<table>
<thead>
<tr>
<th>Multiple myeloma group</th>
<th>Stage I (Mean ± SD)</th>
<th>Stage II (Mean ± SD)</th>
<th>Stage III (Mean ± SD)</th>
<th>Controls</th>
</tr>
</thead>
<tbody>
<tr>
<td>Infiltration</td>
<td>22.56 ± 16.09</td>
<td>33.58 ± 19.78</td>
<td>56.25 ± 16.78*</td>
<td></td>
</tr>
<tr>
<td>IL-6, pg/mL</td>
<td>3.06 ± 1.33</td>
<td>5.71 ± 3.48</td>
<td>9.47 ± 6.79**</td>
<td>0.98 ± 0.64*</td>
</tr>
<tr>
<td>MMP-9, ng/mL</td>
<td>59.25 ± 33.09</td>
<td>188.78 ± 91.27***</td>
<td>131.2 ± 96.32</td>
<td>289.8 ± 110.46*</td>
</tr>
<tr>
<td>NTx</td>
<td>86 ± 67.49</td>
<td>159 ± 61.23</td>
<td>222.4 ± 47.68*</td>
<td>32.67 ± 3.11*</td>
</tr>
<tr>
<td>F-Pyd</td>
<td>26.6 ± 6.5</td>
<td>54.3 ± 19.6**</td>
<td>58.4 ± 23.8*</td>
<td>20.83 ± 2.62*</td>
</tr>
<tr>
<td>F-Dpd</td>
<td>5.7 ± 1.6</td>
<td>12.3 ± 5.3***</td>
<td>11.9 ± 5.3***</td>
<td>4.17 ± 0.72*</td>
</tr>
</tbody>
</table>

*Stage III vs. I and II p < 0.001, **Stage III vs. I p < 0.002, ***Stage II vs. I p < 0.004, *control group vs. multiple myeloma group p < 0.001.
Correlation between serum MMP-9 levels and urinary N-telopeptide (NTx) values in 37 newly diagnosed multiple myeloma patients. A significant positive correlation was found between serum MMP-9 and NTx values (Spearman’s correlation coefficient, $r_s = 0.340$, $p < 0.04$).

There was a positive correlation between serum MMP-9 and F-Pyd values, which did not reach statistical significance (Spearman’s correlation coefficient, $r_s = 0.299$, $p = 0.07$).

There was a positive correlation between serum MMP-9 and N-Tx levels in all cases) (Table 1). In the group of MM patients, the mean concentrations of NTx, F-Pyd and F-Dpd were significantly higher as the stage of disease advanced ($p < 0.001$, $p < 0.001$, $p < 0.003$, respectively).

The biochemical parameters were analyzed according to the severity of bone disease at diagnosis, as estimated by radiographs of the whole skeleton (grade 0 to 3). Patients with progressive bone disease had elevated levels of F-Pyd and F-Dpd in comparison to patients without osteolytic lesions. More specifically, the mean concentrations of F-Pyd and F-Dpd were higher in grade 3 compared to grade 0 ($61.81 \pm 22.17$ vs. $30.55 \pm 15.91$, $p < 0.002$, and $13.13 \pm 5.46$ vs. $6.27 \pm 2.72$, $p < 0.003$, respectively). In contrast, the mean concentrations of F-Pyd and F-Dpd in grades 0, 1 and 2 were increasing gradually with advancing bone disease, but differences did not reach statistical significance. Moreover, mean concentrations of NTx were higher in grade 3 compared to grades 0, 1 and 2 ($p < 0.001$ in all cases).

We further evaluated the levels of MMP-9 in healthy controls and in the group of patients. The mean serum concentration of MMP-9 ($132.37 \pm 95.14$ ng/mL) was lower in the group of patients compared to the control group ($289.80 \pm 110.46$ ng/mL, $p < 0.001$) (Table 1). Furthermore, MMP-9 mean serum concentration was significantly higher in stage II, in comparison to stage I ($p < 0.004$). Stratification of MMP-9 serum levels according to the grade of bone disease revealed higher levels in grade 2 vs. grades 0 and 1 ($p < 0.003$, $p < 0.02$, respectively). Of note, a positive correlation was found between infiltration and NTx levels ($r = 0.340$, $p < 0.03$) as well as between serum IL-6 and NTx levels ($r = 0.340$, $p < 0.04$). A significant correlation was found between serum MMP-9 and NTx values ($r = 0.340$, $p < 0.04$) (Figure 1). In addition, a trend was detected between MMP-9 and IL-6, and F-Pyd and F-Dpd, which approached statistical significance ($r = 0.299$, $p = 0.07$ and $r = 0.228$, $p = 0.08$, respectively) (Figures 2 and 3).

We and others have shown that there is an enhancement of osteoclastic activity in MM patients (3, 4, 20). This process is mediated by factors produced locally by malignant and non-malignant cells in the microenvironment of bones. We have found that MMP-9...
may play an important role in worsening bone disease in patients with MM.

In the present study, serum levels of MMP-9 in stages II and III MM patients were higher than in stage I patients, but serum MMP-9 levels in the whole patient group were lower compared to the age-matched control group. The explanation for this discrepancy remains unclear. In MM patients, reduced MMP-9 levels have been detected in tumors with elevated levels of syndecan-1 (21), however, decreased MMP-9 does not result from association with syndecan-1 expression in malignant cells (22). Lower MMP-9 levels in myeloma patients may be due to a reduction in MMP-9 production from normal sources in MM, which leads to a net decrease in serum MMP-9 (21). It has been postulated that the predominant sources of serum MMP-9 activity in normal people are the circulating lymphocytes, neutrophils and monocytes in which MMP-9 is thought to play an important physiological role (23). As such cells may be crowded by malignant plasma cells in MM, future relevant studies of MMP-9 levels from the bone lesions may help clarify this issue. From this point of view, mast cells may also be important as they have been reported to produce MMP-9 (24), and mast cell chymase activates pro-MMP-9 (25). Mast cell release of MMP-9 and other lytic enzymes has been associated with tumor progression and aggressiveness (26).

The progression of the disease in MM patients is accompanied by an accelerated angiogenesis and production of proteolytic enzymes (6, 27). Furthermore, the growth of myeloma cells is regulated by a cytokine network in which IL-6 represents a major growth player. We found higher serum MMP-9 levels in patients with bone disease scores 2 and 3, in comparison to scores 0 and 1. This observation suggests that the excessive MMP-9 activity in bone marrow plays a functional role in pathological conditions associated with MM, such as tumor progression and osteolysis.

We found that the levels of serum MMP-9 correlated significantly to the estimated rate of bone turnover. Indeed, we report a positive correlation between urinary levels of NTx, MMP-9 ($r = 0.328, p < 0.04$) and IL-6 ($r = 0.340, p < 0.04$). Based on this finding, we and others postulate that the accelerated bone turnover is a result of locally produced factors (including MMP-9) in the microenvironment of the bone, which either accelerate bone turnover (3) or enhance their biological effect via MMP-9. In fact, our results show a trend for direct association of MMP-9 values with F-Pyd and F-Dpd suggesting the possible role of MMP-9, produced locally by myeloma cells, in the promotion of osteolytic bone disease. Published reports suggest that MMP-9 plays an important role in bone resorption (7, 16), and that MMP-9 production by plasma cells accelerates invasiveness and angiogenesis (28). The latter is related to the proliferative activity of malignant plasma cells, according to our previous results (29).

In the present study, serum levels of IL-6 correlated well with urinary levels of NTx, and also, in contrast with MMP-9, with disease stage, being significantly higher in MM patients than in controls, confirming previous results (29). Measurement, thus, of IL-6 may provide a useful marker for monitoring disease activity in MM. Interestingly, serum IL-6 levels have been reported to correlate well with bone involvement in systemic mastocytosis patients (30).

We have found that the mean concentrations of NTx, F-Pyd and F-Dpd were significantly higher in the MM patients compared to age-matched healthy controls ($p < 0.001$). Moreover, in the group of MM patients the mean concentrations of NTx, F-Pyd and F-Dpd were increasing significantly as the disease progressed ($p < 0.001, p < 0.001, p < 0.003$, respectively).

Finally, biochemical parameters were found to correlate to the severity of bone disease at diagnosis, as estimated by X-rays. Thus, patients with progressive bone disease had elevated levels of F-Pyd and F-Dpd compared to patients without osteolytic lesions. Specifically, mean concentrations of F-Pyd and F-Dpd were higher in grade 3 than in grade 0 (61.81±22.17 vs. $30.55 \pm 15.91$, $p < 0.002$, and $13.13 \pm 5.46$ vs. $6.27 \pm 2.72$, $p < 0.003$, respectively).

In conclusion, we have found that levels of MMP-9 and of IL-6 in the circulation of patients with MM correlate with both the radiological disease score and the rate of bone turnover as judged by biochemical markers of bone resorption and formation. Based on these findings, we propose measuring the levels of MMP-9 in the blood of patients with MM, since this determination may add information regarding the severity of their disease and the degree of its effect on bone well being.

References


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