Increased circulating nerve growth factor is directly correlated with disease activity in juvenile chronic arthritis

F Falcini, M Matucci Cerinic, A Lombardi, S Generini, A Pignore, P Tirassa, M Ermini, L Lepore, G Partsch, L Aloé

Abstract

Objective—To determine the circulating serum concentrations of nerve growth factor (NGF) and compare them with indices of disease activity in juvenile chronic arthritis.

Methods—NGF concentrations were evaluated with a two site immunoenzymatic assay (ELISA), in 17 children with systemic, 39 with polyarticular, and 24 with pauciarticular onset juvenile chronic arthritis. Each subset was divided according to different variables, appropriate to each subset, reflecting active and inactive disease.

Results—NGF concentrations were significantly higher in children with systemic [254 (SD 256.1) pg ml⁻¹; P < 0.001], polyarticular [165.2 (300.8) pg ml⁻¹; P < 0.05], and pauciarticular [106.8 (111.8) pg ml⁻¹; P < 0.005] onset juvenile chronic arthritis than in controls. In all subsets, NGF concentrations were higher in the active than in the inactive phase of the disease. A significant direct correlation between NGF concentrations and erythrocyte sedimentation rate was found both in the systemic and in the polyarticular onset juvenile chronic arthritis.

Conclusions—The increase in NGF concentrations in all juvenile chronic arthritis subsets and the correlation with disease activity suggest that NGF may take an active part in joint inflammation.


Nerve growth factor (NGF) is a neurotrophic protein responsible for the development and differentiation of sympathetic, sensory, and cholinergic neurones.1 In the peripheral nervous system, NGF is involved in the modulation of the function of the nerve terminals, controlling their development, survival, and production and release of neurotransmitters.1 NGF is produced by different cells, such as keratinocytes, smooth muscle cells, fibroblasts, and lymphocytes.1 NGF has been shown to increase the chemotaxis of polymorphonuclear cells, the B cell production of immunoglobulins, and the T cell dependent production of autoantibodies.1 NGF interacts specifically with nociceptive sensory neurones, by increasing the content of neuropeptides, in particular substance P. The release of neuropeptides from nerve terminals elicits a local flare called neurogenic inflammation1 mainly linked to the release of substance P which is involved in the activation of synoviocytes and cartilage breakdown in arthritis (for review see Matucci-Cerinic, 1993). The presence of NGF is essential in the production and in particular the release of neuropeptides induced by different stimuli.3 This evidence suggests that the presence of NGF is associated with the maintenance of neurogenic inflammation.4

Previous studies showed that NGF is increased in diseases with an immune pathogenesis, such as multiple sclerosis,7 systemic lupus erythematosus (SLE),8 systemic sclerosis (unpublished data), and, in particular, Kawasaki's disease (F Falcini et al, unpublished data). In adult rheumatoid arthritis, high NGF concentrations have been reported in synovial fluid and synovial membrane9 and in serum.9

Increased NGF concentrations have also been found in the joint tissues of rats with adjuvant arthritis,10 and in arthritic transgenic mice carrying and expressing the human tumour necrosis factor α (TNF-α) gene.11 Juvenile chronic arthritis is a chronic inflammatory disease characterised by articular and systemic involvement in which three types of onset are recognised: systemic, polyarticular, and pauciarticular.13 As far as we know, there are no published studies concerning the role of the peripheral nervous system and neurogenic inflammation in juvenile chronic arthritis, and no studies evaluating NGF, either in the synovium and synovial fluid or in circulation, in this condition are available. High concentrations of NGF were found in the synovial fluid of one child affected with juvenile chronic arthritis.10 To date, NGF concentrations have not been investigated in a large population of children with juvenile chronic arthritis and this prompted us to investigate serum NGF and to evaluate its possible role in the genesis of inflammation by comparing it with clinical and haematological indices of disease activity.

Methods

Patients

Eighty children (54 girls and 26 boys), aged from 15 months to 14 years (mean age 7 years 8 months), all fulfilling EULAR criteria for the diagnosis of juvenile chronic arthritis,14 were selected and entered the study. Seventeen had systemic (age 5 to 12 years), 39 polyarticular (age 3 to 4 years), and 24 pauciarticular onset (age 11 months to 4 years). The duration of the
disease ranged from 8 months to 12 years (mean duration 8 years, 9 months). Each subset was then divided according to the clinical findings (fever, rash, joint inflammation) and haematological data, erythrocyte sedimentation rate (ESR), and C reactive protein into two groups, one with active and the other with inactive disease. No patient was in remission.

Systemic juvenile chronic arthritis was defined active in presence of all the following variables: spiking fever \( > 38.5^\circ \text{C} \); typical rash; joint swelling with reduced range of motion in one or more joints; Hb \( < 10 \text{ g d}^1 \); 5. ESR \( > 25 \text{ mm h}^1 \).

Polyarticular arthritis was defined active in the presence of: (1) ESR \( > 25 \text{ mm h}^1 \); (2) swelling with reduced range of motion in more than five joints. Pauciarticular arthritis was defined active in presence of joint swelling with reduced range of motion in one to four joints.

The basic treatment in children with systemic juvenile chronic arthritis consisted of steroids and non-steroidal anti-inflammatory drugs (NSAID), while in polyarticular subset, methotrexate represented the second line drug used in association with NSAID. All patients in the pauciarticular onset group were given only NSAID. Forty healthy children (25 girls, 15 boys; mean age 6 years, 7 months) matched for sex and age were chosen as controls.

Disease activity was scored following a previous scoring system modified by us\(^{15}\): 0 = no activity; systemic subset 1 = fever \( < 38^\circ \text{C} \), mild joint swelling, ESR 25-40 mm h\(^1\), Hb 9-10 g d\(^1\); 2 = fever \( < 39^\circ \text{C} \), moderate joint swelling, ESR 40-60 mm h\(^1\), Hb 8-9 g d\(^1\); 3 = fever \( < 39^\circ \text{C} \), severe joint swelling, ESR > 60 mm h\(^1\), Hb 6-8 g d\(^1\); polyarticular subset 1 = mild joint swelling, ESR 25-40 mm h\(^1\); 2 = moderate joint swelling, ESR 40-60 mm h\(^1\); 3 = severe joint swelling, ESR > 60 mm h\(^1\); pauciarticular subset 1 = mild, 2 = severe joint swelling.

**PROCEDURES**

After consent had been obtained from the parents, blood was drawn during routine haematological analysis in the morning between 8 am and 9 am and quickly centrifuged. Sera were then stored at \(-70^\circ \text{C} \) until used for NGF determination. NGF was prepared from mouse salivary glands following the procedure of Bocchini and Angeletti\(^{16}\) and a monoclonal anti-NGF antibody, which recognises rodent and human NGF, was supplied to our laboratory. The NGF content in human sera was measured by a two site immunoenzymatic assay (ELISA). Briefly, 96-well polystyrene microtitre immunoplates (Nunc) were coated with affinity purified polyclonal goat anti-NGF antibody. Parallel wells were coated with affinity purified polyclonal goat IgG for evaluation of the non-specific signal. After 5 h at \(20^\circ \text{C} \), the plates were washed (in subsequent steps) with phosphate buffered saline (PBS) / 0.0005%. Tween 20 / 1% fetal calf serum (FCS) to block non-specific binding sites. After washing, the NGF standard solutions, ranging from 0 to 1 ng ml\(^1\), were distributed in each plate. The samples were ultrasonicated at 4°C in the sample buffer and centrifuged at 6,700 g for 30 min at 4°C. A 100 \( \mu \text{l} \) aliquot of supernatant was added to each well and the plates incubated overnight at 4°C. The plates were then washed and 100 \( \mu \text{l} \) of monoclonal antibody against NGF, diluted 1:100 in PBS, were added to each well. After incubation for 4 h at \(20^\circ \text{C} \), the plates were washed and incubated with biotinylated rat immunoglobulins (1:8,000, Zymed). The subsequent incubation with peroxidase conjugated streptavidin (Zymed) and addition of orthophenyldiamine resulted in a colorimetric reaction the optical density of which was measured at 490 nm using a Dynatech MR 5000 microplate reader. Specificity for NGF was also assessed using a recombinant human NGF.\(^{7,12}\) Recombinant brain derived neurotrophic factor is not recognised in the ELISA at concentrations up to 20 ng ml\(^1\).

Serum NGF was correlated with disease activity and haematological variables (that is, ESR and C reactive protein).

**STATISTICAL ANALYSIS**

Evaluation of the differences in NGF levels observed between the three groups of juvenile chronic arthritis and controls was performed with the unpaired Student \( t \) test. The significance of the differences was also confirmed by the parameter-free Mann-Whitney \( U \) test. The correlation between NGF and ESR, C reactive protein, and IgG, in both active and inactive disease, was tested by Pearson’s correlation coefficient.

**Results**

Of the 80 children studied, 53 were in the active phase of the disease and 27 in the inactive phase. The mean NGF value in healthy children was 6 (SD 4.5) pg ml\(^1\). The results show significantly raised concentrations of NGF in children with systemic juvenile chronic arthritis [254 (256.1) pg ml\(^1\); \( P < 0.001 \)] than in controls, in both the active phase [259.3 (280.4) pg ml\(^1\); \( P < 0.001 \)] and the inactive phase [91.6 (87.2) pg ml\(^1\); \( P < 0.001 \)] of the disease. NGF concentrations were raised in all patients with active disease and—although at a lesser extent—also in the inactive phase of the disease. In the polyarticular subset, NGF values were significantly increased in the whole group [165.2 (300.8) pg ml\(^1\); \( P < 0.05 \)], and were always higher in the former subset than in the latter [33.3 (16.2) pg ml\(^1\)]. The mean concentration of NGF in all children with pauciarticular arthritis increased [106.8 (111.8) pg ml\(^1\); \( P < 0.005 \)], and was always higher in the former subset than in the latter [33.3 (16.2) pg ml\(^1\)]. In each subset, a statistical correlation was performed between serum NGF and ESR, C reactive protein, and the activity scores. In systemic onset, a very significant correlation was found with ESR (\( r = 0.9; P < 0.001 \)) and C reactive protein (\( r = 0.8; P < 0.001 \)). In polyart-
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Figure 1. Nerve growth factor (NGF) concentrations (pg ml⁻¹) detected for each subgroup of juvenile chronic arthritis (JCA) in different grades of disease activity.

Table: NGF concentrations detected for each subgroup of juvenile chronic arthritis (JCA) in different grades of disease activity.

![Table: NGF concentrations detected for each subgroup of juvenile chronic arthritis (JCA) in different grades of disease activity.]

**Discussion**

Our data show a striking rise in circulating NGF in juvenile chronic arthritis. The highest values are detected in systemic onset juvenile chronic arthritis, but in the other two subsets the values are higher than in controls. NGF concentrations are increased in the active phase of each subset and remain raised in the inactive phase indicating also that if disease activity is decreasing the serum NGF is kept at a higher level by mechanism still to be determined. NGF concentrations are also high in the pauciarticular subset of juvenile chronic arthritis, where the inflammatory process is limited to one or few joints and systemic involvement is usually lacking.

The source of the increased circulating concentrations of NGF is still a matter of debate (fig 2). The local increase may be derived from synoviocyte proliferation or lymphocytes or peripheral nervous terminals activation in inflamed joints, or a combination of all these. Some pathogenic events (for example, local production of TNF-α and interleukin 1) may trigger the significant increase of NGF in synovial fluid and in the systemic circulation. Cytokines such as interleukin 1, interleukin 6, and TNF-α play an important role in the pathogenesis of juvenile chronic arthritis.15 17 18 These cytokines have been shown to stimulate NGF synthesis by the peripheral nervous system during chronic inflammatory processes.19 NGF is produced by peripheral nerve endings,20 fibroblasts,21 and lymphocytes22 (see fig 2), and the interleukin 1 and TNF-α dependent stimulation of these cells might be related—as potential alternative sources—to the striking increase in the protein in the serum of systemic onset patients. This hypothesis may be supported by the evidence that in systemic subsets of juvenile chronic arthritis, where constitutional findings such as fever are prominent and joint involvement is less relevant, we detected the highest circulating NGF concentrations. In this case, the production of NGF by lymphocytes may also have a particular role in the disease pathogenesis. B and T lymphocytes possess the Trk protein family on their membrane,22 serving as signal transducing receptor unit for NGF. They may favour the activity of NGF as an autocrine protein, thus enhancing an amplificatory loop involving macrophages and lymphocytes—fundamental cells for the development and maintenance of juvenile chronic arthritis.13

On the other hand, local production (that is, by synoviocytes) might account for the presence of high NGF values in polyarticular subset where fever is rarely present, indicating mild systemic involvement, and in pauciarticular onset, where systemic findings are absent.

Another point that should be stressed is the possible participation of NGF in pain transmission. It has been shown that cleaved NGF alters pain threshold in injured target regions of NGF responsive neurones.23 This property might suggest that increased NGF concentrations may enhance the sensitivity to pain of the peripheral terminals of primary afferent nociceptors involved in joint inflammation in juvenile chronic arthritis. NGF is retrogradely transported by sensory nerves as well as by sympathetic nerves and has been shown to increase neuropeptide content in adult sensory neurones.24 NGF induces the synthesis and the release from sensory endings of substance P, which is
mainly related to neurogenic inflammation.\textsuperscript{3,4} The presence of NGF is considered to be the main switch to activate the release of neuropeptides from C fibres and generate neurogenic inflammation.\textsuperscript{4} In this perspective, the increased concentrations of NGF, either in the circulation or in synovial fluid, might amplify the inflammatory joint process by facilitating the development of neurogenic inflammation.

It is interesting to stress that NGF concentrations are directly correlated to ESR and in particular to the progressive increase in disease activity. This kind of behaviour is similar to that observed in multiple sclerosis, where NGF increases during acute attacks and diminishes during remission.\textsuperscript{7} In SLE, increased NGF concentrations have been found concomitantly with disease relapses.\textsuperscript{8}

Our data clearly show that NGF concentrations are increased in all juvenile chronic arthritis subsets and are correlated with disease activity. This evidence seems to strengthen the hypothesis that NGF is involved—primarily or secondarily—in inflammation, though it is still unknown whether it plays an active role or is just the result of a defence mechanism in the pathogenesis of juvenile chronic arthritis. Further investigations are needed to determine the source of circulating NGF, to discover whether NGF is a marker of disease activity, and to identify its real significance in the evolution of the disease.

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