

Somatostatin receptor subtype expression in cells of the rat immune system during adjuvant arthritis

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Abstract

Somatostatin is a neuropeptide that is widely distributed throughout the body. It acts as a neurohormone and a neurotransmitter and may also have an immunomodulatory role. The genes for five subtypes of somatostatin receptors (sst) have been cloned, suggesting that the diverse effects of the peptide might be mediated by different receptors.

We are interested in studying the role of sst in inflammation, using an animal model. Because of the up-regulation of sst expression in inflamed joints in human rheumatoid arthritis, we chose rat adjuvant arthritis as an experimental model. In order to determine which of the sst subtypes might be important in immune modulation, subtype expression in leukocytes isolated from different lymphoid tissues of the rat was studied. Also, the expression levels of the most abundantly expressed sst mRNAs in leukocytes from spleen and blood were compared in rats with adjuvant arthritis and controls, using a semi-quantitative approach. Furthermore, the effect of systemic administration of a long-acting somatostatin analogue, octreotide, which binds selectively to sst subtypes 2 and 5 (sst₂ and sst₅), on the incidence and the severity of rat adjuvant arthritis, was studied.

The main sst expressed in cells of the rat immune system, both resting and activated, were found to be sst₃ and sst₄. This contrasts with the human and murine situations, in which sst₂ appears to be the main subtype expressed in the immune system. No quantitative differences in sst subtype mRNA levels in leukocytes from spleen and blood were found between rats with adjuvant arthritis and controls. Finally, no effect of systemic administration of octreotide on either the incidence or severity of adjuvant arthritis in Lewis rats was found. As octreotide binds selectively to sst₂ and sst₅, the absence of an immunomodulatory effect of this analogue in rat adjuvant arthritis corroborates our finding that these sst subtypes are not expressed in cells of the rat immune system. In conclusion, cells of the rat immune system appear to express a spectrum of sst (sst₃ and sst₄) different from that found in human granulomatous and autoimmune disease (mainly sst₂). Therefore, the rat adjuvant arthritis model appears to be suitable only for studying the immunomodulatory effects of somatostatin analogues which have a high affinity for sst₃ and sst₄, but not for studying the immunomodulatory effects of octreotide, which has a high affinity only for sst₂ and sst₅.

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Introduction

Somatostatin is a neuropeptide that is widely distributed throughout the body. It was first identified as a growth hormone release-inhibiting factor synthesised in the hypothalamus. Outside the central nervous system, the peptide is present in a variety of endocrine and non-endocrine tissues, and it appears to have many functions. In the central nervous system it can act as a neurotransmitter and neurohormone, while in peripheral tissues it regulates endocrine and exocrine secretion and acts as a

modulator of motor activity in the gastrointestinal tract (Reichlin 1983, Brazeau 1986). There is also increasing evidence that somatostatin can have immunomodulatory actions (for a review see Van Hagen *et al.* (1994a)). Specific receptors (sst) for somatostatin have been identified in lymphoid tissues and on the surface of various white blood cells and cell lines by classic ligand-binding studies (Van Hagen *et al.* 1994a). In addition it was found, in a number of granulomatous (i.e. sarcoidosis) and autoimmune diseases (i.e. Graves' ophthalmopathy and rheumatoid arthritis), that sites of active inflammation

could be visualised after the administration of an isotope-coupled somatostatin analogue (Van Hagen *et al.* 1994b,c).

Since we are interested in studying the role of sst in such inflammatory diseases, we decided to use an animal model. Because of the up-regulation of sst expression in inflamed joints in human rheumatoid arthritis (Reubi *et al.* 1994, Van Hagen *et al.* 1994c), we specifically chose rat adjuvant arthritis as an experimental model (for a review, see Wauben *et al.* (1994)). The low numbers of leukocytes that can be obtained from the joints of rats with adjuvant arthritis prohibit analysis of the sst subtypes expressed by these cells. However, as adjuvant arthritis is considered to be a systemic disease in which the causative agent disseminates throughout the tissues of the body (Van Arman 1976, Vernon-Roberts *et al.* 1976, Kleinau *et al.* 1995), an experimental approach was set up to study the sst subtype expression in leukocytes isolated from lymphoid organs and peripheral blood of rats with adjuvant arthritis and controls. We also made a comparison of the expression levels of the most abundantly expressed sst mRNAs in leukocytes from spleen and blood of rats with adjuvant arthritis and controls, using a semi-quantitative approach. Furthermore, we studied the effect of systemic administration of a long-acting somatostatin analogue, octreotide, which binds specifically to sst subtypes 2 and 5 (sst₂ and sst₅), on the incidence and the severity of rat adjuvant arthritis, as somatostatin has been shown to have beneficial effects in human rheumatoid arthritis (Coari *et al.* 1995).

Materials and Methods

Animals

Female Lewis rats (Harlan Sprague-Dawley, Bicester, UK), 7–8 weeks of age, were used. The animals were kept under clean conventional conditions with free access to pelleted food and water in the animal facilities of the Department of Immunology. The experimental protocol was approved by the Animal Welfare Committee of Erasmus University.

Induction of adjuvant arthritis

In order to induce adjuvant arthritis, rats were injected i.d. in the base of the tail with 1 mg of a suspension of heat-killed *Mycobacterium tuberculosis* strain H37 RA (Difco Laboratories, Detroit, MI, USA) in incomplete Freund's adjuvant (IFA) (Difco), prepared by grinding in a mortar, as described by Wauben *et al.* (1994). This preparation will be further referred to as complete Freund's adjuvant (CFA). Control rats were injected with IFA only.

Preparation of cell suspensions

In order to study the sst subtype expression in rats in an inflammatory versus a non-inflammatory state, adjuvant

arthritis was induced in ten rats, as described above. Ten control rats were injected with IFA only. The rats were killed by decapitation 3 weeks after injection, at peak inflammation. Cell suspensions were prepared as described below for analysis by reverse transcription polymerase chain reaction (RT-PCR).

Peripheral blood Peripheral blood was collected in polypropylene tubes containing 100 IE heparin (Leo Pharmaceutical Products BV, Weesp, The Netherlands). Peripheral blood mononuclear cells (PBMC) were purified by density centrifugation on Ficoll-Paque, density = 1.077 kg/l (Pharmacia, Uppsala, Sweden). PBMC from rats within one experimental group were pooled.

Lymphoid organ cell suspensions Spleen, thymus and peripheral lymph nodes (popliteal, inguinal and axillary nodes) were removed from the animals and cell suspensions were prepared in RPMI 1640 medium (Gibco BRL, Grand Island, NY, USA) supplemented with 10% fetal calf serum (FCS) by pressing the organs through nylon gauze (mesh size 100 µm).

Erythrocytes were depleted from the spleen cell suspension by hypotonic lysis in a 10 mM KHCO₃ buffer (pH 7.4) supplemented with 155 mM NH₄Cl and 100 mM EDTA (3 ml per spleen). After 3 min at 4 °C, the cells were washed twice in RPMI 1640 medium with 10% FCS. Half of the total lymph node suspension was used to prepare a non-adherent cell fraction by passage over nylon wool (Leuko-Pak, Fenwall Laboratories, Deerfield, IL, USA).

The different cell populations were pooled per experimental group. The cells were washed in ice-cold PBS, pelleted and stored as dry pellets at –80 °C.

RNA isolation and cDNA reaction

PolyA⁺ mRNA was isolated from the cells using paramagnetic oligo(dT) beads (Dynabeads Oligo(dT)₂₅, Dynal AS, Oslo, Norway) according to the manufacturer's instructions. The polyA⁺ mRNA was eluted in 30 µl diethyl pyrocarbonate-treated water. All mRNA preparations were treated with deoxyribonuclease I (Amplification Grade, Gibco) for 15 min at room temperature in the buffer provided with the enzyme. The reaction was stopped by the addition of 2.5 mM EDTA followed by heating at 65 °C for 10 min.

mRNA was extracted from 10⁷ cells in the case of mixed cell populations (spleen, thymus, total lymph node cells) and 2 × 10⁶ cells in the case of more purified cell populations (PBMC and lymph node non-adherent cells).

Complementary DNA (cDNA) synthesis was performed starting with one-third of the isolated mRNA. Random hexamer primers were used to ensure that all RNA was represented equally in the cDNA pool (Tan & Weis 1992). The reaction mixture contained 50 mM

Tris-HCl buffer, pH 8.3, 10 mM MgCl₂, 50 mM KCl, 1 mM dithiothreitol, 1 mM EDTA, 10 µg/ml BSA, 1 mM salmon sperm HCl (Sigma Chemical Co., St Louis, MO, USA), 1 mM of each of the four deoxynucleotide triphosphates (DNA Polymerisation Mix, Pharmacia), 20 U of RNA Guard (Pharmacia), 0.02 optical density units (dN)₆ (Pharmacia), 0.2 µg oligo(dT)₁₂₋₁₈ (Boehringer Mannheim, Mannheim, Germany) and 5 U reverse transcriptase (from avian myeloblastoma virus; Promega, Madison, WI, USA) in a total reaction volume of 20 µl. The reaction mixture was incubated at 41 °C for 1 h and the reaction was stopped by heating at 75 °C for 10 min. Each cDNA reaction was carried out in duplicate.

Controls

To ascertain that no genomic DNA was present in the polyA⁺ preparations (which would give false-positive signals upon amplification, as *sst* genes do not contain introns), the cDNA reactions were also performed once without reverse transcriptase (negative control samples). These negative control samples never yielded positive signals on amplification.

cDNA from the GH3 rat pituitary cell line was used as a positive control for *sst*₁ mRNA expression and cDNA from the transplantable prolactin-secreting rat pituitary tumour 7315b as a positive control for *sst*₂ mRNA expression (Visser-Wisselaar *et al.* 1997). Rat genomic DNA, isolated by lysis of rat splenocytes in PCR buffer, was used as a positive control in the PCR reactions for *sst*₃₋₅.

The PCR reactions were optimised to detect at least 10⁴ copies of the sequence of interest (corresponding to 0.1 µg genomic DNA) in ethidium bromide-stained gels. If, in a particular series of PCR reactions, 10⁴ copies of genomic DNA after amplification did not give a clear signal in ethidium bromide-stained gel, the series was discarded.

PCR

One-tenth of the synthesised cDNA was used for each PCR reaction. The reaction mixture consisted of 10 mM Tris-HCl buffer, pH 9.0, 50 mM KCl, 1.5 mM MgCl₂, 0.01% (w/v) gelatine, 0.1% (v/v) Triton X-100, 0.05 mM of each of the four deoxynucleotide triphosphates (DNA Polymerisation Mix; Pharmacia), 5 pmol each of the forward and reverse primers specific for one of the five rat *sst* subtypes or for the β-actin gene, and 1 U AmpliTaq DNA polymerase (Perkin-Elmer-Cetus, Norwalk, CT, USA). PCR reactions for all primer pairs were performed with each of the duplicate cDNA samples and positive and negative control samples.

The primers specific for the human *sst*₁ and *sst*₂ as described by Wulfsen *et al.* (1993) and for *sst*₃₋₅ as described by Kubota *et al.* (1994), were modified by us to match the rat sequences (Meyerhof *et al.* 1991, 1992,

Bruno *et al.* 1992, Kluxen *et al.* 1992, Li *et al.* 1992, Yasuda *et al.* 1992). The rat β-actin primers were as described by Vidal *et al.* (1994). *sst*₁ forward 5'-ATGGTG GCCCTCAAGGCCGG-3', reverse 5'-GGCAGTGGC GTAGTAGTCAA-3' (product size 318 bp); *sst*₂ forward 5'-TCCTCTGGGATCCGAGTGGG-3', reverse 5'-TT GTCCTGCTTACTGTCGCT-3' (product size 332 bp); *sst*₃ forward 5'-TGCCAGTGGGTACAGGCACC-3', reverse 5'-CTGGAGGGCCAGACCCTGGC-3' (product size 328 bp); *sst*₄ forward 5'-TGCGGGCTGGCTGG CAACAA-3', reverse 5'-GTAGTCCAGGGGCTCTT CCT-3' (product size 311 bp); *sst*₅ forward 5'-AGCCT TCATCACCTACACGT-3', reverse 5'-GGCCAGGT TGACGATGTTGA-3' (product size 226 bp); β-actin forward 5'-TCATGCCATCCTGCGTCTGGACCT-3', reverse 5'-CCGGACTCATCGTACTCCTGCTTG-3' (product size 582 bp).

The reactions were carried out in a DNA thermal cycler (Perkin-Elmer-Cetus). After an initial denaturation for 5 min at 94 °C, the samples were subjected to 35 cycles (β-actin) or 40 cycles (*sst* subtypes) of denaturation at 94 °C for 1 min, annealing at 59 °C (β-actin and *sst*₁₋₄) or 55 °C (*sst*₅) for 2 min and extension at 72 °C for 1 min. These numbers of cycles had previously been shown to lie within the linear range of the reaction. After a final extension phase of 10 min at 72 °C, 40 µl aliquots of the amplified products were analysed on 2% agarose gels stained with ethidium bromide. Weak bands were confirmed by hybridising Southern blots of the PCR products with subtype-specific oligonucleotide probes.

For semi-quantitative comparison of the amount of mRNA, a limiting dilution RT-PCR method was used (Kincy-Cain & Bost 1997). Briefly, 2-fold serial dilutions of the cDNA preparations were normalised to give approximately equal β-actin signals, before performing PCR for the *sst* subtypes. The linearity of the dilution series was assessed by scanning the photographs of the ethidium bromide-stained gels and determining the intensity of the bands (Koning *et al.* 1995). For the most accurate quantification, care was taken to include a dilution of which the β-actin signal was no longer visible when analysed on ethidium bromide-stained gel. Comparisons were based on this dilution and the two preceding it.

Receptor saturation curves

The radioligands used in the binding studies were [¹²⁵I-Tyr³]-octreotide and [¹²⁵I-Tyr⁰]-somatostatin-28. [Tyr³]-octreotide was iodinated and HPLC purified as described previously (Bakker *et al.* 1990). [¹²⁵I-Tyr⁰]-somatostatin-28 was purchased from ARC-Biotrend (Cologne, Germany). Reaction conditions were the same as described by Reubi (1985). Briefly, cell membrane preparations (corresponding to 15–30 µg protein) were incubated at room temperature for 60 min with increasing

concentrations of radioligand in a total volume of 100 μ l Hepes buffer (10 mM Hepes, 5 mM MgCl_2 and 0.2 g/l bacitracin, pH 7.6) containing 0.2% (w/v) BSA. After the incubation, 1 ml ice-cold Hepes buffer was added to the assay mixture and membrane-bound radioactivity was separated from unbound activity by centrifugation (2 min 10 000 *g* in an Eppendorf centrifuge). The remaining pellet was washed twice with ice-cold Hepes buffer and radioactivity remaining in the final pellet was counted in a gamma counter. Specific binding was taken to be the total bound activity minus the activity bound in the presence of 1 μ M unlabelled octreotide (Sandostatin; Novartis Pharma, Basel, Switzerland) or somatostatin-28 (Bachem, Bubendorf, Switzerland) respectively. As a positive control for binding, cell membranes from the mouse pituitary cell line AtT20 were included in each test.

Systemic administration of octreotide and monitoring of arthritis

The effect of systemic administration of octreotide, a long-acting somatostatin analogue, on adjuvant arthritis in the rat was studied. Sandostatin-LAR, a slow-release formulation of octreotide, was used. This formulation gives a continuous release of octreotide for a period of 6 weeks after s.c. injection.

One day prior to the induction of arthritis, ten rats were injected s.c. in the dorsal neck region with a suspension of Sandostatin-LAR (a gift from Novartis Pharma) in standard injection vehicle (buffered sodium carboxyl-methylcellulose), corresponding to 10 (experiments 1 and 2) or 30 mg/kg octreotide (experiment 3). Ten control rats were injected with standard injection vehicle only. The next day adjuvant arthritis was induced in all the rats as described above. Disease course was monitored twice weekly for 6 weeks by measuring the diameter of all four paws using industrial callipers. The animals were weighed once a week and their general appearance and behaviour were monitored to assess their condition. Blood was collected by orbital puncture on days 21 and 42 of the experiment. The rats were killed by CO_2 asphyxiation on day 42.

Results

With a view to using rat adjuvant arthritis as an animal model in which to study the role of sst in inflammation, we determined the expression pattern of the five known sst subtypes in the major lymphoid organs of the rat, using RT-PCR.

In order to study mainly leukocytes, stromal components were removed from the lymphoid organs under study by passing them through nylon gauze. Other authors have studied only whole organs of the rat (Bruno *et al.* 1993, Raulf *et al.* 1994).

Table 1 Sst subtype mRNAs in cells of the rat immune system: summary of RT-PCR study

	sst ₁	sst ₂	sst ₃	sst ₄	sst ₅
Source of leukocytes					
Spleen	±	—	+	+	—
Peripheral blood	—	—	+	+	—
Thymus	—	—	+	±	—
Lymph nodes	—	—	+	±	—
Lymph nodes (non-adherent cells)	—	—	+	±	—

+, visible on ethidium bromide-stained agarose gel; ±, not visible, but visualised by Southern blotting.

The results of the RT-PCR studies are shown in Table 1. The major sst subtypes expressed in rat leukocytes were found to be sst₃ and sst₄. A very weak sst₁ signal was observed in spleen cells in some, but not all, experiments. No expression of sst₂ or sst₅ was observed in any of the cell populations, even after a repeat amplification. A ladder pattern of amplification products was consistently observed in the sst₄ PCR of thymocytes, but hybridisation of a Southern blot of the PCR gel with an sst₄-specific probe yielded only a single band of the expected size. The sst₄-specific band for the total lymph node and lymph node non-adherent cell preparations was barely visible in the ethidium bromide-stained gel, but could be visualised by Southern blot hybridisation with an sst₄-specific probe. The pattern of expression of sst in rat leukocytes was confirmed in splenocytes from other strains of rats (females: Wistar, Brown Norway, Wag/Rij, Sprague-Dawley, Buffalo and Fisher F344). Again only sst₃ and sst₄ were expressed, with occasionally a very weak sst₁ signal (not shown).

Receptor saturation curves

Receptor saturation experiments were carried out on membrane preparations of splenocytes and thymocytes of control Lewis rats using the radioligands [¹²⁵I-Tyr⁰]-somatostatin-28 and [¹²⁵I-Tyr³]-octreotide. *K_d* values and receptor densities were determined by Scatchard analysis of the saturation curves.

No membrane binding of [¹²⁵I-Tyr³]-octreotide was found (not shown), but [¹²⁵I-Tyr⁰]-somatostatin-28 bound with a *K_d* of 0.5 and 0.3 nM for splenocytes and thymocytes respectively. As a control for binding, sst-positive AtT20 mouse pituitary tumour cells were used (*K_d*=0.9 nM). The calculated numbers of binding sites were 755, 492 and 2679 fmol/mg protein for splenocytes, thymocytes and AtT20 cells respectively. Figure 1 shows the Scatchard plots for the binding of [¹²⁵I-Tyr⁰]-somatostatin-28.

Semi-quantitative comparison of sst₃ expression

Limiting dilution RT-PCR was performed to semi-quantitatively compare sst₃ mRNA levels in splenocytes

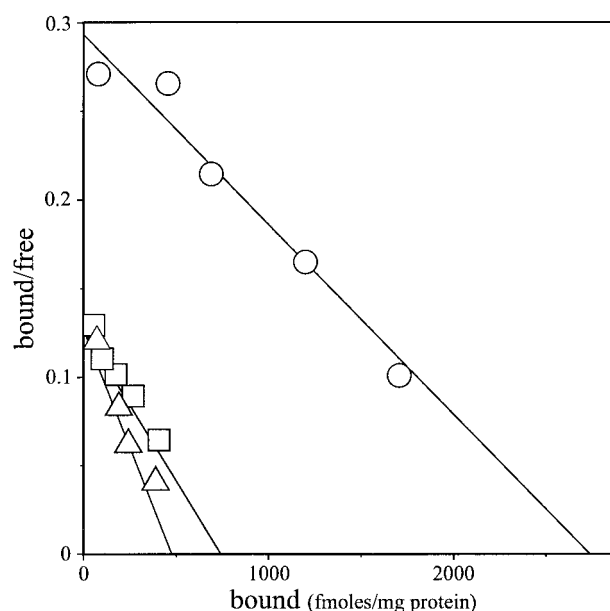


Figure 1 Sst expression at the protein level in rat splenocytes and thymocytes. Scatchard analysis of [125 I-Tyr 0]-somatostatin-28 binding to Lewis rat splenocytes and thymocytes and the sst-positive mouse pituitary tumour cell line, AtT20. \circ AtT20 cells ($K_d=0.9$ nM, $n=2679$ fmol/mg), \square Lewis splenocytes ($K_d=0.5$ nM, $n=755$ fmol/mg), \triangle Lewis thymocytes ($K_d=0.3$ nM, $n=492$ fmol/mg).

from CFA-injected rats with adjuvant arthritis (further referred to as CFA-injected rats) and controls. Two-fold serial dilutions of cDNA were titrated to give approximately equal β -actin signals and were then subjected to PCR for sst $_3$. The results are shown in Fig. 2. No substantial differences in the amount of sst $_3$ mRNA in CFA-injected versus control rats could be demonstrated.

Differential expression of sst $_4$ mRNA in leukocyte subpopulations

Sst $_4$ was expressed at levels readily detectable on ethidium bromide-stained gels only in splenocytes and peripheral blood, whereas expression in the other lymphoid cell preparations was only detectable after Southern blotting. This was true for both CFA-injected rats and control rats. Peripheral blood and spleen, as well as peripheral lymph nodes harbour large numbers of activated cells during systemic inflammation. However, spleen and peripheral blood contain a relatively larger proportion of monocytes in addition to their lymphocytes. The difference observed in sst $_4$ expression levels between peripheral blood and splenocytes on the one hand and lymph nodes on the other hand, suggested that sst $_4$ might be expressed by monocytes and cells of the monocyte lineage such as dendritic cells. A cell population enriched for dendritic cells, isolated from the spleen of a normal Lewis rat, was found to express sst $_4$

mRNA (not shown). We therefore set out to prepare, from peripheral blood, a cell population enriched for monocytes, using density centrifugation over Percoll. Sst $_3$ and sst $_4$ were expressed in the unseparated population and also in both populations obtained after density centrifugation (not shown). The cellular composition of the different populations is summarised in Table 2.

Limiting dilution RT-PCR was carried out as described above to semi-quantitatively compare the sst $_4$ mRNA expression in the different populations. The results are shown in Fig. 3. It was shown that the interphase cell population had a relatively higher sst $_4$ expression than either the pellet population or total PBMCs. This was true for both CFA-injected and control rats.

The effect of systemic administration of octreotide on adjuvant arthritis

We studied the effect of systemic administration of octreotide on the incidence and the severity of rat adjuvant arthritis, as somatostatin has been shown to have beneficial effects in human rheumatoid arthritis (Coari *et al.* 1995). Sandostatin-LAR was used in three separate experiments (see Table 3). The data show that neither a low dose (10 mg/kg) nor a higher dose (30 mg/kg) of Sandostatin-LAR had an effect on incidence and severity of the arthritis.

Arthritis was confirmed histologically. Marginal erosion of the bone and hyperplastic synovium extending into the joint cavity was observed, with an infiltrate consisting mainly of lymphocytes, polymorphonuclear cells and histiocytes.

In the first experiment the octreotide levels in the serum were determined on days 21 and 42. The mean concentrations \pm s.e.m. were 2.353 ± 0.641 and 1.524 ± 0.744 ng/ml respectively for the octreotide-treated group, and undetectable (<0.1 mg/ml) for the control group. Nine rats out of ten still had detectable serum levels of octreotide on day 42. These data show that the octreotide was released from the s.c. depot continuously throughout the course of the experiment.

Discussion

Somatostatin is a ubiquitous neuropeptide with diverse physiological functions. The recent cloning of a family of sst and the demonstration that these receptors differ in their affinities for synthetic somatostatin analogues (Shimon *et al.* 1997) suggest that the diverse actions of somatostatin might be mediated by different receptor subtypes.

We are interested in the immunomodulatory effects of somatostatin and somatostatin analogues. Because of the up-regulation of sst expression in inflamed joints in human rheumatoid arthritis, we chose rat adjuvant arthritis as an

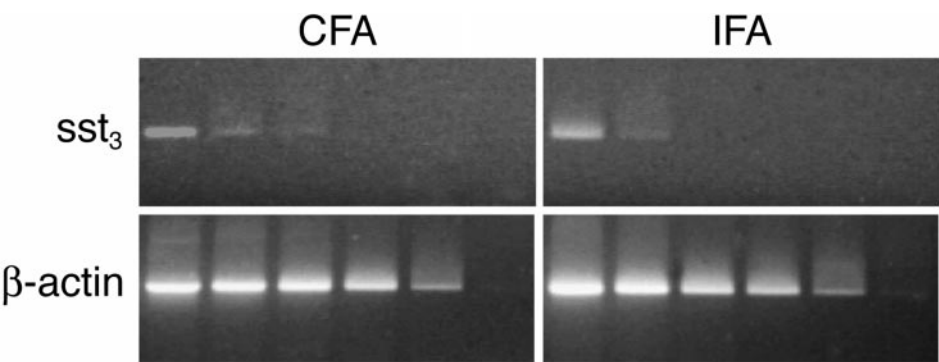


Figure 2 Sst₃ mRNA expression levels in rats with adjuvant arthritis and controls. For semi-quantitative comparison of sst₃ mRNA expression in splenocytes of Lewis rats with adjuvant arthritis (CFA-injected) versus controls (IFA-injected), serial 2-fold dilutions of cDNA were subjected to PCR for β -actin and sst₃.

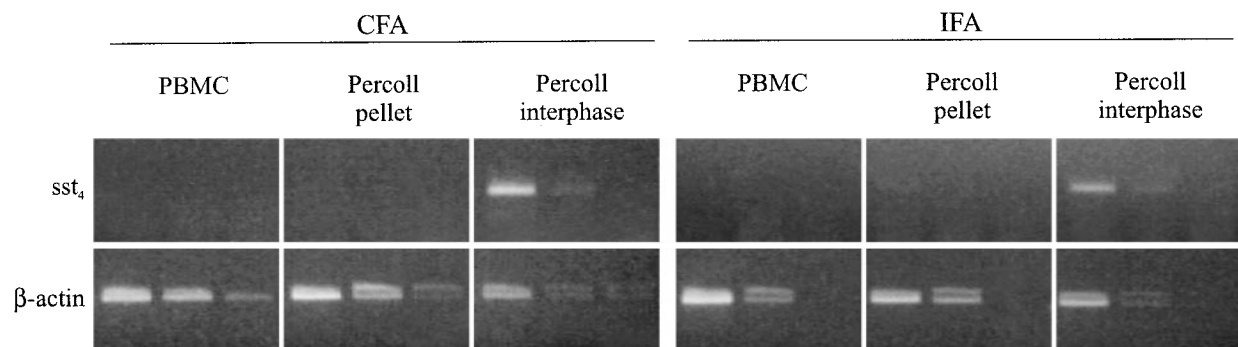


Figure 3 Increased expression of sst₄ mRNA in a population of rat peripheral blood cells enriched for monocytes. Subpopulations of PBMC from Lewis rats with adjuvant arthritis (CFA injected) and controls (IFA injected) were prepared by density centrifugation over Percoll. For semi-quantitative comparison of sst₄ mRNA expression levels, serial 2-fold dilutions of cDNA were subjected to PCR for β -actin and sst₄. The cDNA samples in the β -actin panel of the figure are diluted 8-fold as compared with the corresponding samples in the sst₄ panel.

Table 2 Cellular composition (%) by nuclear morphology of the peripheral blood cell populations before and after Percoll density centrifugation

	CFA			IFA		
	PMN	Lymphocyte	Monocyte	PMN	Lymphocyte	Monocyte
PBMC	7	91	2	12	80	8
Percoll pellet	9	85	5	2	90	8
Percoll interphase	4	78	18	4	80	16

PMN, polymorphonuclear.

experimental model. In order to determine which of the sst subtypes might be important in immune modulation, we studied the sst subtype expression in lymphoid tissues and cells of the rat, using RT-PCR.

The main sst expressed in cells of the rat immune system, both resting and activated, were found to be sst₃ and sst₄. This contrasts with the situation in humans and mice, in which sst₂ appears to be the main subtype expressed in the immune system (Elliott *et al.* 1994, Van Hagen *et al.* 1994a and unpublished observations).

Furthermore, our results indicate that sst₄ is expressed by distinct sub-populations of rat leukocytes. A peripheral

blood cell fraction enriched for monocytes showed an increased sst₄ expression. However, we were not able to obtain a highly purified rat monocyte population using density centrifugation. Flow cytometric analysis suggests that, in contrast to what is seen in humans, rat PBMC exhibit a wide range of sizes and densities, and cannot be identified as a separate population in forward-sideward-scatter plots (Scriba *et al.* 1996 and unpublished observations). The low-density cell population also contains activated lymphocytes (lymphoblasts), leaving open the possibility that sst₄ mRNA is up-regulated upon activation. However, we found no quantitative differences in

Table 3 Effect of systemic administration of octreotide (as Sandostatin-LAR) on the incidence and severity of adjuvant arthritis in Lewis rats

	Incidence			Severity*		
	Sandostatin-LAR	Controls	P-value**	Sandostatin-LAR	Controls	P-value**
Octreotide dose (mg/kg)						
10 (experiment 1)	7/9	10/10	0.21	38.1 ± 10.9	59.2 ± 14.1	0.29
10 (experiment 2)	8/10	7/10	0.50	58.3 ± 15.0	57.4 ± 12.4	0.97
30 (experiment 3)	8/9	10/10	0.47	105.0 ± 18.3	116.3 ± 10.2	0.81

*Severity expressed as mean maximal increase in total paw diameter (in 0.1 mm) ± S.E.M.

**Statistics: incidence was compared using a Fisher exact test. Severity was compared using a Mann-Whitney test (two-sided).

sst subtype mRNA levels between CFA-injected rats with adjuvant arthritis and control rats. This observation again seems in contrast with the situation in the human immune system in which binding sites for somatostatin seem to be preferentially expressed on activated leukocytes (Van Hagen *et al.* 1994a).

Our results on the sst subtype expression in rat splenocytes contrast with the expression patterns found by others in intact rat spleen. Raulf *et al.* (1994) detected sst₃ mRNA but not sst₄ mRNA in rat spleen and lymph nodes. These authors also found a weak sst₅ signal in the spleen. However, this expression might be due to the presence of stromal cells, e.g. nerves, which are largely eliminated in our procedure of making cell suspensions. Using a highly sensitive nuclease protection assay, Bruno *et al.* (1993) found mRNA for all five sst subtypes in rat spleen. It was clear from their results also that sst₃ was the main subtype in spleen, whereas the other four subtypes showed much lower expression. Our results on the sst subtype expression in rat thymocytes contrast with the findings of Sedqi *et al.* (1996), who showed transcripts for sst₂ in resting and activated thymocytes and sst₁ in activated thymocytes, but who did not detect mRNA for sst₃. A contributing factor to the major discrepancy in sst₂ expression reported in the two papers could be the fact that Sedqi *et al.* analysed cells from male rats, whereas we used female rats on account of their susceptibility to adjuvant arthritis. It is known that the expression of sst₂ can be influenced by oestrogens, e.g. the transplantable rat prolactin-secreting pituitary tumour 7315b expresses sst₂ when grown *in vitro*, but loses this expression when grown *in vivo* under the influence of circulating oestrogens (Visser-Wisselaar *et al.* 1997). However, the difference in expression of sst₃ as determined in the study of Sedqi *et al.* and in this study needs further investigation.

At the protein level, no binding of the sst₂- and sst₅-selective radioligand [¹²⁵I-Tyr³]-octreotide was found. This result was in accordance with the lack of sst₂ and sst₅ expression, as found by RT-PCR. Other studies have generally found a good correlation between the presence of octreotide binding sites and the presence of sst₂ and/or sst₅ mRNA (Reubi *et al.* 1996). The radioligand [¹²⁵I-Tyr⁰]-somatostatin-28 was bound, demonstrating the presence of

other subtypes of sst. We therefore conclude that the mRNA for sst₃ and sst₄, which was detected by RT-PCR, is functional, and can give rise to membrane expression of sst proteins in rat leukocytes.

Finally, no therapeutic effect of systemic administration of octreotide, in the form of Sandostatin-LAR, was found on the incidence and severity of adjuvant arthritis in Lewis rats. This may be due to the absence in the immune system of the rat of sst₂ and sst₅, which have a high affinity for this octapeptide somatostatin analogue. The absence of octreotide binding sites *in vitro* is correlated with the absence of *in vivo* biochemical response to octreotide, at least in the case of hormone-secreting tumours (Reubi *et al.* 1990). Using a different somatostatin analogue, BIM23014 (1 µg/h; the doses we used correspond to 2.5 and 7.5 µg octreotide/h), Rees *et al.* (1989) did find a statistically significant decrease in arthritis severity, although a different measure of severity was used. The effect cannot be explained by differences in binding profiles of octreotide and BIM23014 (Shimon *et al.* 1997) on the distinct sst subtypes (although to our knowledge no comparisons of the binding profiles of these compounds on the rat sst subtypes are available). Alternatively, the somatostatin analogues might not have acted directly on the immune cells in these studies, but might have had a more generalised effect through neuroendocrine mechanisms. However, no differences were found in the efficacies of octreotide and BIM23014 in inhibiting the release of several hormones (Hofland *et al.* 1997, Shimon *et al.* 1997).

In human rheumatoid arthritis the inflamed joints may be visualised by *in vivo* scintigraphy using the sst₂- and sst₅-selective somatostatin analogue [¹¹¹In-DTPA-D-Phe¹]-octreotide (Van Hagen *et al.* 1994c). In rat adjuvant arthritis the affected joints could not be visualised using this analogue (Breeman *et al.* 1996), indicating that the cells of the inflammatory infiltrate in the joint also lack the sst subtypes (sst₂ and sst₅) which have a high affinity for octreotide. We recently showed the presence of sst_{2A} in endothelial cells and cells of the monocyte/macrophage lineage in human rheumatoid synovium (ten Bokum *et al.*, unpublished observations).

In conclusion, cells of the rat immune system appear to express a spectrum of sst (sst₃ and sst₄) different from that

found in human granulomatous and autoimmune diseases (Van Hagen *et al.* 1994b,c). Therefore, the rat adjuvant arthritis model appears to be suitable only for studying the immunomodulatory effects of somatostatin analogues which have a high affinity for sst₃ and sst₄, but not for studying the immunomodulatory effects of octreotide, which has a high affinity only for sst₂ and sst₅.

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