

Synthetic Peptides in the Form of Dendrimers Become Resistant to Protease Activity*

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In previous papers, we observed that dendrimers of peptide mimotopes of the nicotinic receptor ligand site are strong antidotes against the lethality of the nicotinic receptor ligand α -bungarotoxin. Although their *in vitro* activity is identical to that of dendrimers, the corresponding monomeric peptide mimotopes are not effective *in vivo*. Because the higher *in vivo* efficiency of dendrimers could not in this case be related to polyvalent interaction, the stability to blood protease activity of monomeric versus tetrabranch dendrimeric mimotope peptides was compared here by incubating three different mimotopes with human plasma and serum. Unmodified peptides and cleaved sequences were followed by high pressure liquid chromatography and mass spectrometry. Tetrabranch peptides were shown to be much more stable in plasma and also in serum. To assess the notable stability of multimeric peptides, different bioactive neuropeptides, including enkephalins, neurotensin and nociceptin, were synthesized in monomeric and tetrabranch forms and incubated with human plasma and serum and with rat brain membrane extracts. All the tetrabranch neuropeptides fully retained biological activity and generally showed much greater stability to blood and brain protease activity. Some tetrabranch peptides were also resistant to trypsin and chymotrypsin. Our findings provide new insights into the possible therapeutic use of bioactive peptides.

Hundreds of peptides with potential therapeutic activities have been identified. These include naturally occurring peptide hormones and neurotransmitters, which influence and control series of vital functions, such as cell proliferation, tissue development, metabolism, immune defense, perception of pain, reproduction, behavior, and blood pressure. Selective agonists or antagonists of these natural peptides are extremely useful for the investigation of peptidergic systems and are also potential therapeutic agents (1). Moreover, several peptide fragments or mimotopes derived from potential therapeutic proteins show promising biological activity (2).

However, the use of peptides as therapeutic drugs has largely been limited by their short half-life *in vivo*. Because peptides are mainly broken down by proteases and peptidases,

peptide delivery is the bottleneck in the development of new peptide drugs. To increase peptide half-life, many strategies involving different levels of chemical modification are possible (3, 4). The introduction of D-amino acids, or pseudo amino acids, and peptide cyclization are the most common strategies to increase peptide stability. However, these modifications may profoundly alter peptide activity. Alternatively, peptidomimetic molecules can be developed by the synthesis of conformationally restricted compounds, in which the peptide is locally or globally constrained in order to reproduce the active conformation. The resulting structures are mostly non-peptide molecules, more resistant to degrading enzymes.

In general, peptide molecules have the advantage of good specificity and high selectivity and the disadvantage of poor metabolic stability and limited distribution. Although non-peptide molecules have better stability and pharmacokinetics, they may have very different specificity and affinity to natural peptides. The design and synthesis of new peptidomimetic compounds therefore involve careful testing of their selectivity, specificity, and affinity for target biomolecules.

In this study we report results suggesting that synthesis of bioactive peptides in multiple antigen peptide (MAP)¹ dendrimeric form can result in increased half-life, due to acquired resistance to protease and peptidase activity. MAPs have a peptidyl core of radially branched lysine residues onto which peptides can be added using standard solid-phase chemistry (5). MAPs can be more efficient than monomeric peptides in diagnostic applications. Moreover, they also proved to be good immunogens against a number of pathogens (6). The *in vitro* and *in vivo* efficiency of dendrimeric peptides like MAPs is generally ascribed to their multimeric nature, which enables polyvalent interactions.

We reported previously that the tetrameric MAP form of the peptide mimotope p6.7 and its high affinity analogues pDD and pDDD, which compete with the nicotinic receptor (nAChR) to bind the snake neurotoxin α -bungarotoxin (α -bgt), efficiently neutralize toxin lethality in mice, unlike the corresponding monomeric sequences. This happens despite the nearly identical affinity constant (K_A) and half-maximal inhibition concentration (IC_{50}) of MAPs and monomeric peptides, and their similar *in vitro* activity (7, 8). Because in the particular case of peptide ligands of a monomeric soluble toxin, MAPs have no polyvalent interaction-associated advantage with respect to

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¹ The abbreviations used are: MAP, multiple antigen peptide; nAChR, nicotinic acetylcholine receptor; α -bgt, α -bungarotoxin; M-enk, Met-enkephalin; NT, neurotensin; NC, nociceptin; MS, mass spectrometry; BSA, bovine serum albumin; PBS, phosphate-buffered saline; IAA, iodoacetamide; HPLC, high pressure liquid chromatography; MALDI-TOF, matrix-assisted laser desorption/ionization/time-of-flight; L-enk, Leu-enkephalin; Fmoc, N-(9-fluorenyl)methoxycarbonyl.

monomeric sequences (which was confirmed by results of *in vitro* competition experiments), we concluded that the notable *in vivo* efficiency of MAPs seemed to be related to a pharmacokinetic advantage, such as different clearance, or resistance to peptidase and protease activity.

In order to verify this possibility, we examined here the stability of the monomeric and MAP mimotopes in human plasma and serum. Moreover, we synthesized several tetrameric MAPs, reproducing sequences of different bioactive peptides, and we compared their resistance to proteases to that of the corresponding monomeric peptides.

EXPERIMENTAL PROCEDURES

Peptide Synthesis—Solid-phase synthesis was carried out on a MultiSynTech Syro automated multiple peptide synthesizer (Witten, Germany), employing *N*-(9-fluorenyl)methoxycarbonyl (Fmoc) chemistry with 2-(1*H*-benzotriazole-1-yl)-1,1,3,3-tetramethyluronium hexafluorophosphate/*N,N*-diisopropylethylamine activation. Side chain protecting groups were trityl for His, 2,2,4,6,7-pentamethylidihydro-benzofuran-5-sulfonyl for Arg, *tert*-butyl ether for Ser and Tyr, *tert*-butyl ester for Asp and Glu, and *tert*-butyloxycarbonyl for Trp. Monomeric peptides HRYYESLEPWYP (p6.7), FRYYESLEPWDD (pDD), FRYYESLEPWDDD (pDDD), YGGFM (Met-enkephalin, M-enk), YGGFL (Leu-enkephalin, L-enk), ELYENKRRPYIL (neurotensin, NT), RRPYIL (NT 8–13), and FGGFTGARKSARKLANQ (nociceptin, NC) were synthesized on a 4-(2',4'-dimethoxyphenyl)-Fmoc-aminomethyl-phenoxycetamido-norleucyl 4-methylbenzhydrylamine (RINK-amide MBHA) resin. MAP analogues were synthesized on Fmoc-4-Lys²-Lys-β-Ala Wang resin. Peptides were cleaved from the resins and deprotected by treatment with trifluoroacetic acid containing water and triisopropylsilane (95:2.5:2.5). After precipitation with diethyl ether, all peptides were purified by HPLC using a C18 Vydac column and characterized by Ettan™ MALDI-TOF mass spectrometry (MS) (Amersham Biosciences).

Peptide Processing in Serum and Plasma—150 μl of a 1 mg/ml solution of peptides was incubated at 37 °C with 40 μl of human serum or plasma. Samples withdrawn after 2 or 24 h were precipitated with 200 μl of methanol and centrifuged for 1 min at 10,000 × *g*, and the crude solution was analyzed. HPLC was performed on all samples using a C18 column. The crude solution was diluted in 0.1% trifluoroacetic acid, and the UV lamp was set at 280 nm for all peptides and at 220 nm for NC. Controls for peptide retention time in the mixture were obtained by adding the same concentration of peptides to supernatants of plasma or serum treated with methanol and centrifuged as above, and running the mixture immediately.

MS analysis was performed on an Ettan™ MALDI-TOF mass spectrometer on crude solutions and was repeated on HPLC-eluted peaks when these were detectable. When HPLC peaks corresponding to uncleaved peptides were no longer detectable after incubation with plasma or serum, the HPLC eluent was collected at the appropriate retention time, allowing a window of ±2 min, and analyzed by MS in order to assess the absence of uncleaved peptides.

Preparation of Rat Brain Membranes—Rats were decapitated after cervical dislocation. The brain was removed and rapidly transferred to ice-cold Tris-HCl buffer (Tris-HCl, 50 mM, pH 7.4). The brain was homogenized on ice using a Potter-Eurostar (IKA Labortechnik) without antifoaming agents. The homogenate was centrifuged at 13,500 × *g* for 10 min at 4 °C. The pellet was resuspended in Tris-HCl buffer and incubated for 30 min at 37 °C to remove endogenous peptides. The homogenate was centrifuged again at 13,500 × *g* for 10 min at 4 °C, and the pellet was resuspended in Tris-HCl buffer to reach a total concentration of 15 mg/ml. Membranes were prepared fresh upon use.

Peptide Processing in Rat Brain Membrane Extracts—50 μl of a 1 mg/ml solution of peptides was incubated at 37 °C with 50 μl of 15 mg/ml rat brain membranes. Samples withdrawn after 2 or 24 h were precipitated with 150 μl of methanol and centrifuged for 1 min at 10,000 × *g*, and the crude solutions were analyzed. All samples were diluted in 0.1% trifluoroacetic acid and analyzed by HPLC. MS analysis was performed on crude solutions or on HPLC-eluted material, as described above.

Inhibition of α-bgt Binding to nAChR—Microtiter plates (Falcon 3912, BD Biosciences) were coated overnight at 4 °C with affinity-purified nAChR from *Torpedo* electric organs (7) (5 μg/ml in 0.05 M carbonate buffer, pH 9.6) and then blocked with 3% BSA in phosphate-buffered saline (PBS), pH 7.4, for 1 h at room temperature. MAP and monomeric peptides, previously incubated in PBS or in human plasma or serum for 2 or 24 h at 37 °C, were diluted in PBS at concentrations

ranging from 100 μg/ml to 100 pg/ml and incubated with 10⁵ cpm (0.45 nM) ¹²⁵I-α-bungarotoxin (Amersham Biosciences) for 1 h at room temperature. After washing with PBS, α-bgt binding to nAChR was detected by a γ-counter. The IC₅₀ value of the peptides was evaluated by non-linear regression analysis of curves using GraphPad Prism 3.02 software.

Inhibition of Ligand Binding to Rat Brain Membranes—Monomeric and multimeric enkephalins, diluted at concentration ranging from 10⁻⁵ to 5 × 10⁻⁸ M in ice-cold 50 mM Tris-HCl buffer pH 7.4, were incubated with 15 mg/ml rat brain membranes in the presence of 0.5 nM [³H]naloxone. After 1 h at 25 °C, the reaction was stopped by adding 4 ml of ice-cold Tris-HCl. The membranes were filtered through Whatman GF/B filters, pretreated with 0.1% polyethyleneimine, and rinsed twice with 4 ml of the same ice-cold buffer using a Brandel Cell Harvester. [³H]Naloxone binding was detected by a β-counter. Nonspecific binding was determined in the presence of 10⁻⁶ M naloxone.

For experiments performed in the absence of protease activity, all samples were diluted in 50 mM Tris-HCl, 10 mM EDTA, 10 mM EGTA, 10 mM aminobenzamide, 10 mM IAA, and 10 mg/liter soybean trypsin inhibitor, pH 7.4. The IC₅₀ of the peptides was evaluated by non-linear regression analysis of curves using GraphPad Prism 3.02 software.

Monomeric and multimeric nociceptin, diluted at concentrations ranging from 10⁻⁵ to 10⁻¹² M, were incubated with 15 mg/ml rat brain membranes in the presence of 1 nM [³H]nociceptin (PerkinElmer Life Sciences). Nonspecific binding was determined in the presence of 10⁻⁵ M [*N*-Phe¹]nociceptin fragment 1–13-amide (Sigma). All experiments were performed in ice-cold 50 mM Tris-HCl, 2 mg/ml BSA, 10 mM EDTA, 10 mM EGTA, 10 mM aminobenzamide, 10 mM IAA, 10 mg/liter soybean trypsin inhibitor, pH 7.4.

Inhibition of Ligand Binding to HT 29 Cell Membranes—HT 29 cell line (Istituto Zooprofilattico Sperimentale, Brescia, Italy) were cultured in McCoy's 5A medium supplemented with 10% fetal bovine serum, 200 μg/ml glutamine, 60 μg/ml penicillin, and 100 μg/ml streptomycin. For membrane preparation, cells were homogenized with 10 volumes (w/v) of ice-cold 50 mM Tris-HCl, 10 mM EDTA, 10 mM EGTA, 10 mM aminobenzamide, 10 mM IAA, 10 mg/liter soybean trypsin inhibitor, pH 7.4. After centrifugation for 40 min at 40,000 × *g* at 4 °C, the pellet was washed, centrifuged again under the same conditions, resuspended in the same buffer, and sonicated to avoid membrane aggregation. HT 29 cell membranes were prepared fresh upon use.

Monomeric and multimeric neurotensin were incubated with 0.5 mg/ml HT 29 cell membranes diluted in ice-cold Tris-HCl, 2 mg/ml BSA, 10 mM EDTA, 10 mM EGTA, 10 mM aminobenzamide, 10 mM IAA, 10 mg/liter soybean trypsin inhibitor, pH 7.4, at concentrations ranging from 10⁻⁵ to 10⁻¹⁰ M in the presence of 1 nM [³H]neurotensin (PerkinElmer Life Sciences). Nonspecific binding was determined in the presence of 10⁻⁵ M neurotensin (Sigma).

RESULTS

Stability of Dendrimeric Mimotopes to Blood Proteases—We previously produced synthetic peptides mimicking the snake neurotoxin-binding site of the nicotinic receptor, and we found that the *in vivo* efficiency of these mimotopes reflects their affinity and IC₅₀ value only when they are synthesized in MAP form (7, 8). The efficacy of the tetrameric peptides *in vivo* could not be ascribed to any kinetic or thermodynamic effect and therefore seemed related to differences in pharmacokinetic behavior.

To verify this possibility, we compared the resistance of monomeric and tetrabrached peptide mimotopes in human plasma and serum. Monomeric and tetrabrached peptides were incubated with human plasma or serum for 2 or 24 h, and the mixture was analyzed by HPLC and mass spectrometry to follow the presence of uncleaved monomeric and MAP peptides.

As a general rule, peptides that were cleaved in plasma in 2 h were also cleaved in serum in the same time. Those resistant in serum after 24 h were also resistant in plasma (Table I).

Monomeric peptide mimotopes (p6.7, pDD, and pDDD) were completely degraded within 2 h in serum. Fragments derived from proteolytic cleavage were detected by MS in the HPLC-eluted material, after 2 h of incubation in plasma. Conversely, tetrabrached forms of the same peptides were still detected after 24 h in plasma and serum (Fig. 1). Incubation for 2 h in

TABLE I
Proteolytic stability of monomeric and MAP peptides

Monomeric and MAP peptides were incubated 2 or 24 h at 37 °C in human plasma or serum and in rat brain membrane extracts. The presence of monomeric or MAP peptides was followed by HPLC and MS.

Peptide	Plasma		Serum		Rat brain membranes ^a		Chymotrypsin		Trypsin	
	2 h	24 h	2 h	24 h	2 h	24 h	2 h	24 h	2 h	24 h
p6.7	— ^b	—	—	—	ND	ND	+	—	+	—
MAP p6.7	+ ^b	+	+	+	ND	ND	+	—	+	+
pDD	+	—	—	—	ND	ND	—	—	—	—
MAP pDD	+	+	+	+	ND	ND	—	—	+	+
pDDD	+	—	—	—	ND	ND	—	—	—	—
MAP pDDD	+	+	+	+	ND	ND	+	+	+	+
M-enk	+	—	—	—	+	—	—	—	+	+
MAP M-enk	+	+	+	—	+	—	+	—	+	+
L-enk	+	—	—	—	+	—	—	—	+	+
MAP L-enk	+	+	+	—	+	+	+	—	+	+
NT	+	—	+	—	+	—	—	—	+	—
MAP NT	+	+	+	+	+	+	+	—	+	+
NT-(8–13)	+	—	+	—	+	—	—	—	+	—
MAP NT-(8–13)	+	+	+	+	+	+	+	+	+	+
NC	+	—	—	—	+	—	—	—	+	+
MAP NC	+	+	+	+	+	+	+	+	+	+

^a Incubation with rat brain membrane extracts was only done for neuropeptides.

^b The absence (—) or presence (+) of peptides after the incubation times refers to the unproteolyzed peptide sequence as detected by HPLC and MS.

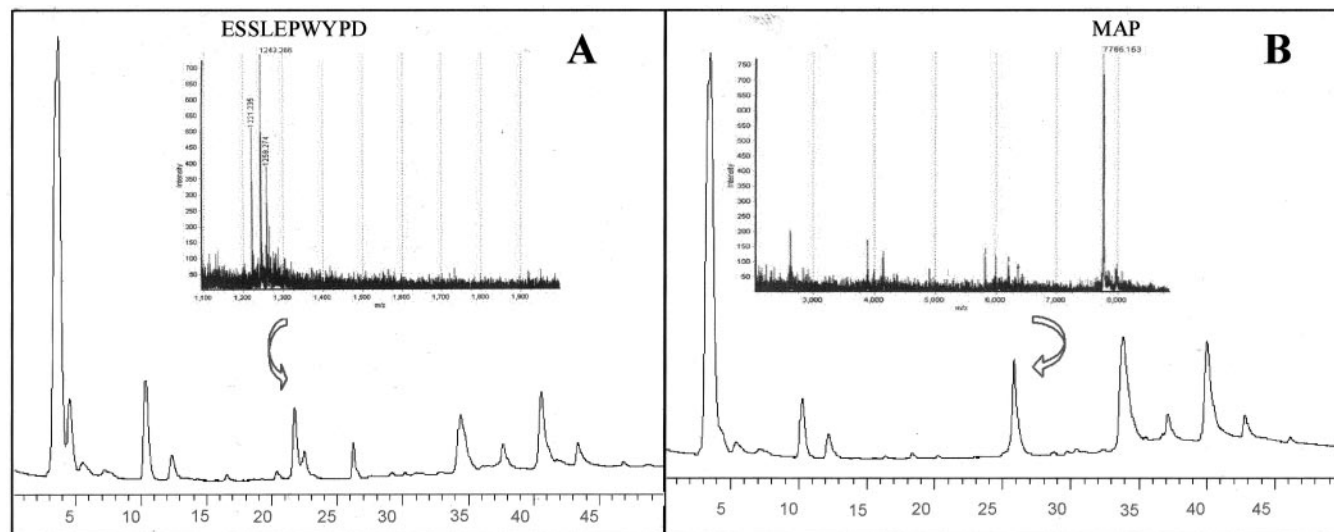


FIG. 1. HPLC and MALDI-TOF of p6.7 and MAP p6.7 after incubation in plasma or serum. A, analysis of p6.7 after incubation in plasma for 2 h. HPLC-eluted material was collected from 20 to 24 min (22 min is the retention time of uncleaved p6.7) (bottom) and analyzed by MS (top). Instead of the peptide, fragments of proteolytic cleavage were found. B, analysis of MAP p6.7 after incubation in serum for 24 h. HPLC-eluted material was collected from 24 to 28 min (26 min is the retention time of uncleaved MAP p6.7) (bottom) and analyzed by MS (top). The uncleaved sequence was still present.

human plasma or serum increased the IC₅₀ value of monomeric peptides by at least 3 log, whereas MAP peptides retained their inhibition power after 24 h (Fig. 2).

Stability of Dendrimeric Neuropeptides to Blood and Brain Proteases—In order to verify whether the enhanced stability in plasma and serum could be considered a general behavior of tetrabranch peptides, we synthesized the monomeric and the tetrameric MAP form of different natural bioactive peptides in order to compare their resistance to protease activity. We chose Met-enkephalin, Leu-enkephalin, neurotensin, and nociceptin because they have very interesting pharmacological properties but very short half-lives *in vivo*. Moreover, this choice allowed testing peptides of different length and sequence, going from 5 residues of enkephalins to 17 residues of nociceptin.

Enkephalins—Enkephalins are endogenous ligands of opioid receptors. Analogues of enkephalins are very well known μ/δ -receptor agonists with analgesic activity. Opioid receptor ligands made available so far are mainly non-peptide (morphine

and its derivatives) or pseudo-peptide molecules including pseudo or D-amino acids (9).

In this work, we synthesized and tested the natural peptide sequence of Met- and Leu-enkephalins. The MAP forms of M-enk and L-enk were analyzed for their ability to bind opioid receptors by testing inhibition of [³H]naloxone binding to rat brain cell membranes. MAPs of M- and L-enk have an IC₅₀ that is one-tenth that of their monomeric counterparts (Fig. 3). In the presence of protease inhibitors, the IC₅₀ value of tetrameric and monomeric enkephalins was nearly identical (3.7×10^{-7} and 5.7×10^{-7} M for M-enk and MAP M-enk, respectively, and 2.5×10^{-7} and 3.0×10^{-7} M for L-enk and MAP L-enk, respectively). When MAP and monomeric enkephalins resistance was tested in plasma and serum by HPLC and MS, the latter were still present after 2 h in plasma and were found to be cleaved within 2 h in serum, whereas their tetrabranch analogues were still present after 24 h of incubation in plasma (Table I).

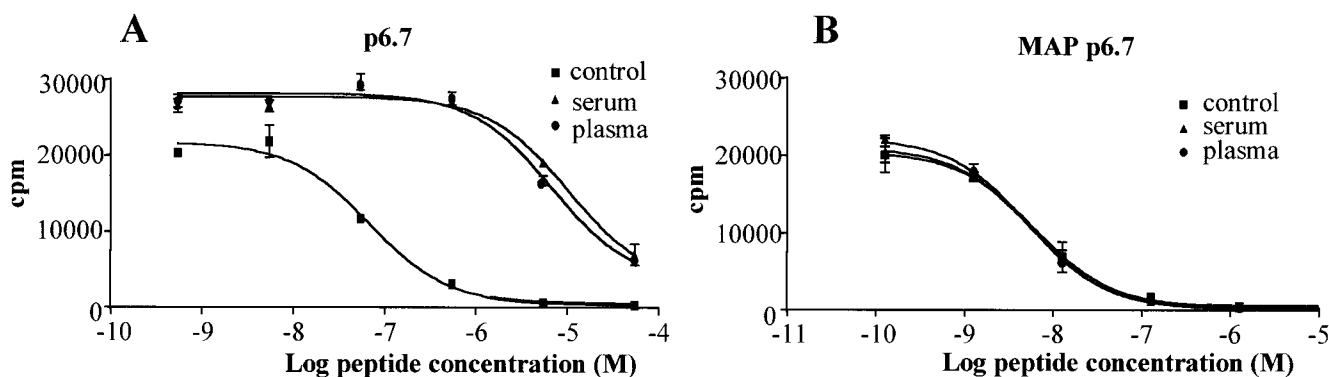


FIG. 2. Residual activity of monomeric and MAP mimotope peptides after incubation in serum or plasma. MAP and monomeric peptides, previously incubated in PBS (control), or in human plasma or serum for 2 or 24 h at 37 °C, were diluted in PBS and incubated with 10^5 cpm of ^{125}I - α -bungarotoxin for 1 h at room temperature. A, after 2 h of incubation with human plasma or serum, the ability to inhibit nAChR- α -bgt binding of the monomeric peptide p6.7 decreased to about 3 log with respect to controls. B, MAP p6.7 maintained the same IC_{50} value after incubation in plasma or serum for 24 h. Similar results were obtained with monomeric and MAP pDD and pDDD (not shown).

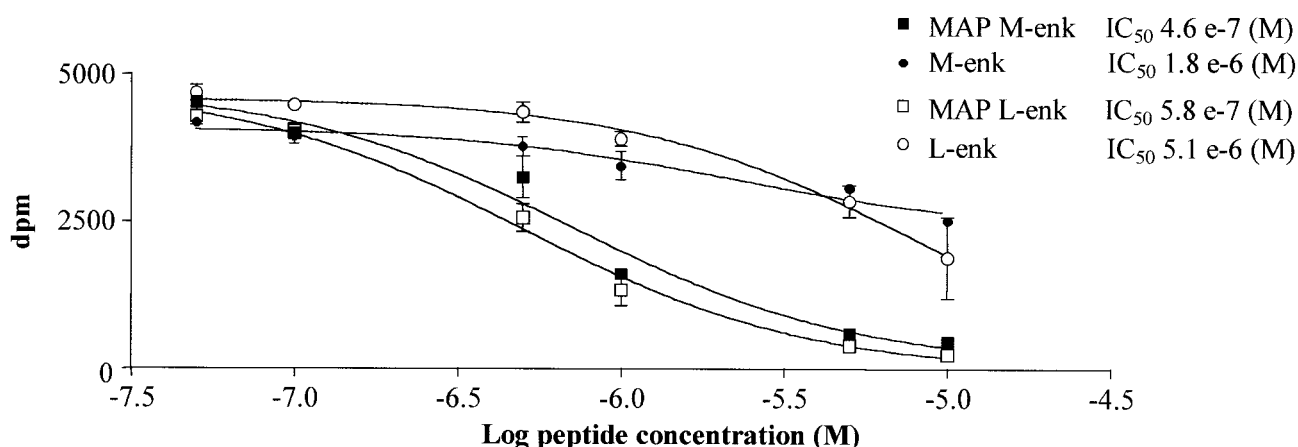


FIG. 3. Inhibition of ^{3}H naloxone binding to rat brain membranes by monomeric and tetrameric Met- and Leu-enkephalins. MAPs of Met- and Leu-enkephalin showed an IC_{50} less than one-tenth that of the monomeric peptides.

Resistance to protease activity was also tested by incubating tetrameric and monomeric peptides with rat brain membrane extracts. Monomeric Met- and Leu-enkephalins were found to be degraded within 24 h, whereas their multimeric analogues were still detectable.

Neurotensin—NT is a 13-amino acid peptide originally isolated from calf hypothalamus (10). It has the dual function of neurotransmitter or neuromodulator in the nervous system and local hormone in the periphery. NT is a neuromodulator of dopamine transmission and anterior pituitary hormone secretion and exerts potent hypothermic and analgesic effects in the brain (11). In the periphery, NT is a paracrine and endocrine modulator of the digestive tract and cardiovascular system. NT receptors are overexpressed in a number of neuroendocrine human tumors (12). Interest in NT is related to the possibility of developing antipsychotic (11) and anticancer (12, 13) drugs.

We synthesized monomeric and tetrabranch neurotensin and its short analogue NT-(8–13) (14). Bioactivity of monomeric and MAP peptides was checked by testing the inhibition of nM ^{3}H neurotensin-specific binding to membranes prepared from the human colon adenocarcinoma cell line HT 29 (15). Experiments were performed in the presence of protease inhibitors. MAP NT efficiently inhibited nM ^{3}H neurotensin-specific binding, and its IC_{50} value was analogous to that of monomeric NT (Fig. 4). NT-(8–13) was more effective than whole NT. MAP NT-(8–13) had an IC_{50} of 1.9×10^{-10} (M), which is about 1 log lower than that of monomeric NT-(8–13) and more than 2 log lower than NT and MAP NT.

The resistance to protease activity in human plasma and

serum was tested for the four peptides. We found that the MAP form was more resistant than the monomeric form. MAP forms of NT and NT-(8–13) were still detectable in serum after 24 h, when the monomeric forms had disappeared (Table I).

When the tetrameric and monomeric forms of NT and NT-(8–13) were incubated in rat brain membrane extracts, monomeric peptides were present after 2 h, but many by-products appeared at 24 h, whereas the MAP analogues resisted beyond 24 h.

Nociceptin—NC (16, 17) is an endogenous heptadecapeptide that displays nanomolar affinity for the opioid receptor-like ORL1, a G protein-coupled receptor. NC modulates nociceptive and locomotor behavior in mice (18).

We synthesized the monomeric NC and tetrameric MAP NC and tested their behavior upon plasma and serum treatment. Monomeric NC was degraded within 2 h in serum and plasma, whereas MAP NC was still detectable after 24 h in plasma.

NC and MAP NC were incubated with rat brain membrane extracts, and while the monomeric peptide disappeared within 24 h, MAP NC was still recoverable.

Bioactivity of monomeric and MAP NC was assessed by testing the inhibition of ^{3}H nociceptin-specific binding to rat brain membranes in the presence of protease inhibitors. MAP NC inhibited the binding of the labeled peptide with an IC_{50} of 1.9×10^{-9} M, whereas the IC_{50} of the monomer was 8.6×10^{-9} (M) (Fig. 5).

Stability of Dendrimeric Peptides to Trypsin and Chymotrypsin—The increased stability of dendrimeric peptides was also confirmed with purified proteases. Tetrabranch mimotopes

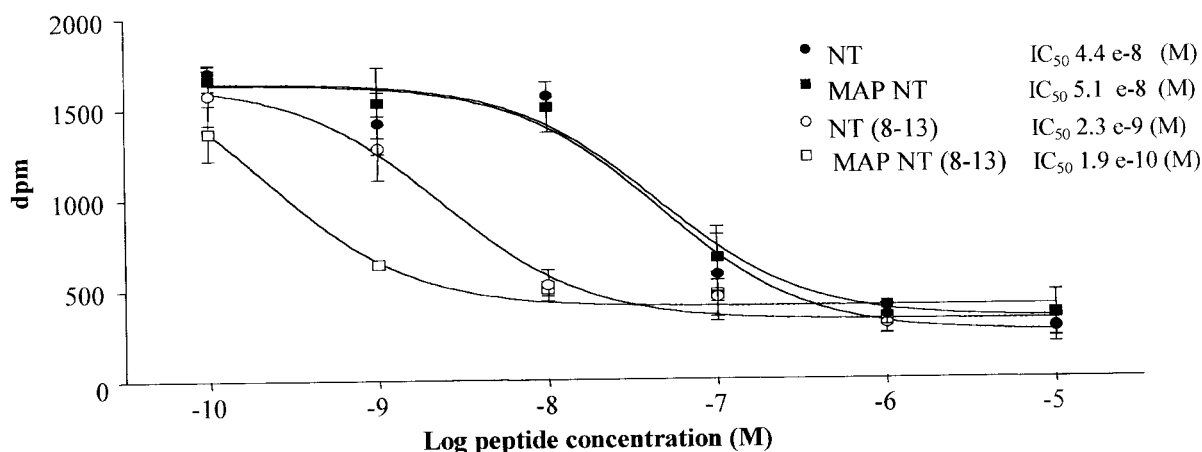


FIG. 4. Inhibition of [³H]neurotensin binding to HT 29 cell membranes by monomeric and tetrameric NT and NT-(8-13). MAPs fully retain the activity of corresponding monomeric sequences. NT-(8-13) and MAP NT-(8-13) are more efficient than peptides reproducing the entire NT sequence.

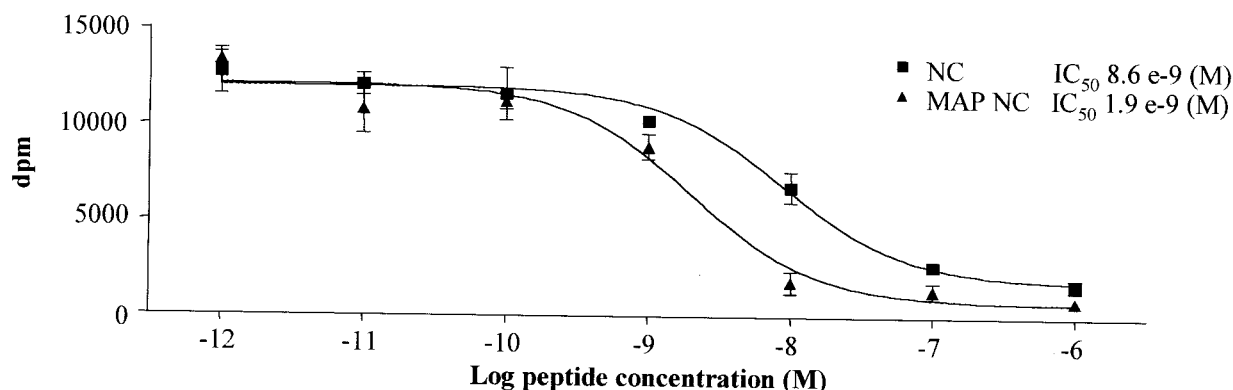


FIG. 5. Inhibition of [³H]nociceptin binding to rat brain membranes by monomeric and tetrameric NC. MAP NC fully retains NC activity.

MAP p6.7, MAP pDD, and MAP pDDD were not cleaved by trypsin in 24 h, whereas their corresponding monomers were proteolyzed within 2 h, except for peptide p6.7, which was still detectable. Most monomeric and MAP peptide mimotopes were degraded by chymotrypsin after 2 h of incubation. However, the MAP form of the peptide pDDD was still found after 24 h of incubation.

Monomeric M- and L-enk were cleaved by chymotrypsin but not by trypsin. Their multimeric analogues were detectable after exposure to chymotrypsin for 2 h.

NT was degraded by chymotrypsin within 2 h and by trypsin within 24 h when its tetrabranched analogue was still present. Moreover, the bioactive analogue NT-(8-13) in its tetrabranched form showed remarkable resistance to the purified enzymes; it remained intact after 24 h of incubation in trypsin and chymotrypsin, whereas its monomeric analogue only resisted for 2 h with trypsin and less with chymotrypsin.

NC is a substrate for chymotrypsin (degraded in 2 h) but not for trypsin; however, the MAP analogue MAP NC was stable in chymotrypsin even after 24 h (Table I).

DISCUSSION

In this paper we provide data indicating that synthesis of different peptide sequences, including natural endogenous peptides, in a MAP dendrimeric form allows retaining full peptide biological activity and results in a dramatic increase in peptide stability to blood and brain protease and peptidase activity.

The higher stability of MAP forms to protease activity could be a consequence of their unnatural branched peptide structure. Moreover, in the case of short peptides, the activity of

specific peptidases (like many membrane, blood, or central nervous system metallo-oligopeptidases) (19), which are selectively active on short sequences, could be prevented by the mass increase and steric hindrance of MAPs. This possibility seems to be supported by the reported results, indicating a higher general stability of MAPs to blood and brain proteases than to trypsin and chymotrypsin.

Synthetic peptides composed exclusively of natural amino acids, but having a longer half-life than natural peptides, may have major applications as therapeutic drugs. For example, the MAP enkephalins we produced and tested here consist of the same residues as natural enkephalins and have the same binding affinity *in vitro*. MAP enkephalins might have an improved activity related to greater stability and not to higher receptor affinity. This could be an advantage for pharmacological use as they may avoid triggering immune response, while having less intense morphine-like side effects related to overstimulation of receptors.

Another important possible development of the protease-resistant MAPs described here is the application of neurotensin to tumor therapy and *in vivo* imaging. The finding of new specifically targeted radiopharmaceuticals is an important challenge in cancer research, where tumor-specific small peptides may have major potential applications, provided that they are stabilized to overcome their degradation by proteases and peptidases. Currently, analogues of NT and particularly of NT-(8-13) are under investigation to target a variety of neuroendocrine human tumors overexpressing NT receptors (12-13, 20). We demonstrated here that NT and NT-(8-13), once

synthesized in a MAP form, fully retain their biological activity and become much more resistant to blood proteases. Moreover, the tetrabranched MAP of peptide NT-(8–13) has a 10-fold better IC₅₀ value than the corresponding monomeric peptide. These dendrimeric peptides are therefore candidates for the development of new specific tumor-targeted biomolecules.

Although the possible higher stability of dendrimeric peptides has been speculated previously (6), to our knowledge this is the first demonstration of MAPs resistance to blood and brain proteases. A higher stability of MAP activity following trypsin treatment, with respect to the corresponding linear peptide, was already reported for an antimicrobial peptide (21), although no chemical characterization of peptide proteolysis was reported. MAP stability was deduced from the higher residual antimicrobial activity after incubation with the enzyme.

Dendrimeric peptides were reported to have a higher *in vivo* efficiency with respect to corresponding monomeric sequences in several cases (6, 22), but in general they were used in conditions where they could bind through multivalent interactions. This effect can somehow have masked the pharmacokinetic advantage of dendrimeric peptides or can at least have rendered their different *in vivo* stability not immediately evident. In the particular case of dendrimeric mimotopes that bind a soluble monomeric ligand and that have identical *in vitro* activity with respect to corresponding monomers, it was clear that the *in vivo* efficiency of dendrimeric peptides could only be related to a different pharmacokinetic behavior (7–8). This enabled us to distinguish between kinetic and pharmacokinetic advantage of dendrimers *in vivo*.

In conclusion, our results indicate that synthesis in dendrimeric form may be a general method to increase *in vivo* stability, and consequently half-life, of bioactive peptides.

The biological activity of a peptide might be lost in the MAP derivative, and tests on this feature obviously should be carried out. In the cases analyzed here, the MAP forms retained the full activity of native peptides, both when peptides bind to membrane receptors, like enkephalins, neurotensin, and noci-

ceptin, and when they bind soluble proteins, like peptide mimotopes of a receptor-binding site. Moreover, biological activity and stability to protease activity was retained in dendrimers of peptides having different sequence and different length.

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