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<p>(21) International Application Number: PCT/US92/02731 (22) International Filing Date: 3 April 1992 (03.04.92) (30) Priority data: 681,802 5 April 1991 (05.04.91) US (60) Parent Application or Grant (63) Related by Continuation US 681,802 (CIP) Filed on 5 April 1991 (05.04.91) (71) Applicant (for all designated States except US): GENENTECH, INC. [US/US]; 460 Point San Bruno Boulevard, South San Francisco, CA 94080 (US).</p>	<p>(72) Inventors; and (75) Inventors/Applicants (for US only) : BURNIER, John, P. [US/US]; 202 Sterling Avenue, Pacifica, CA 94044 (US). GADEK, Thomas [US/US]; 2838 Chelsea Drive, Oakland, CA 94611 (US). McDOWELL, Robert [US/US]; 11 Hillview Court, San Francisco, CA 94124 (US). (74) Agents: WINTER, Daryl, B. et al.; Genentech, Inc., 460 Point San Bruno Boulevard, South San Francisco, CA 94080 (US). (81) Designated States: AT (European patent), AU, BE (European patent), CA, CH (European patent), DE (European patent), DK (European patent), ES (European patent), FR (European patent), GB (European patent), GR (European patent), IT (European patent), JP, LU (European patent), MC (European patent), NL (European patent), SE (European patent), US. Published <i>With international search report. Before the expiration of the time limit for amending the claims and to be republished in the event of the receipt of amendments.</i></p>	
<p>(54) Title: PLATELET AGGREGATION INHIBITORS HAVING HIGH SPECIFICITY FOR GP II_bIII_a</p> <p>(57) Abstract</p> <p>A peptide containing the tripeptide recognition sequences RGD or KGD in a cycle and an exocyclic group bearing a positive charge is provided. The compound is provided in therapeutic form for administration to a mammal and exhibits high specificity and potency as a platelet aggregation inhibitor without undesirable side effects.</p>		

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**PLATELET AGGREGATION INHIBITORS
HAVING HIGH SPECIFICITY FOR GP II_bIII_a**

5

Field of the Invention

The present invention relates to inhibitors having an anti-thrombotic effect. Specifically, the invention is directed to antagonists of the final common pathway of platelet aggregation that act as potent antithrombotics. The invention further relates to therapeutic applications of these inhibitors in diseases for which blocking of platelet aggregation is indicated.

Background of the Invention

Platelets are particles found in whole blood that initiate and provide the structural basis for the hemostatic plug necessary to stop bleeding. Platelets depend on adhesive interactions with extracellular proteins and other cells for proper function. The external platelet plasma membrane surface is covered with a variety of membrane bound glycoproteins, many of which have adhesive functions. Perhaps the most abundant platelet membrane adhesive proteins belong to the integrin superfamily which include the glycoproteins; GP II_bIII_a, GP I_aII_a, GP I_cII_a, GP I_bIX, and the fibronectin and vitronectin receptors. Each integrin receptor is an αβ heterodimer displaying characteristic affinity and specificity toward various extracellular matrix proteins such as; von Willebrand factor (vWF), collagen, entactin, tenascin, fibronectin (Fn), vitronectin (Vn), and laminin, as well as fibrinogen (Fg) and thrombospondin (see Kieffer *et al.*, *Ann. Rev. cell Biol.* 6:329-357(1990) and Ruoslahti, *J. Clin. Invest.* 87:1- 5 (1991)). The most abundant integrin found on normal platelet surfaces is GP II_bIII_a comprising about 50,000 molecules per platelet, representing about 2% of the total platelet protein. GP II_bIII_a is a non-covalent, calcium ion dependent heterodimer complex (Jennings, *et al.*, *J. Biol. Chem.* 257: 10458 (1982)) and restricted in distribution to platelets and other cells of the megakaryocytic lineage (Kieffer *et al.*, *supra*). On activated platelets, GP II_bIII_a binds a number of adhesive proteins with varying affinities; fibrinogen, fibronectin, von Willebrand factor, vitronectin and thrombospondin (Plow *et al.* *Biochemistry of Platelets*, Phillips and Shuman eds., p. 225-256, Orlando: Academic Press (1986)). It is believed the most important interactions mediating platelet aggregation involve GP II_bIII_a binding with the trinodular fibrinogen and, to a lesser extent, with the filamentous von Willebrand factor (Kieffer *et al.*, *supra* and Albeda *et al.*, *The FASEB Journal*, 4:2868-2880 (1990)).

GP II_bIII_a binding to its natural ligands can be inhibited to varying degrees by peptides and proteins containing the amino acid recognition sequences; Arg-Gly-Asp (RGD) (Ruoslahti, *supra* and EPO 0368486, assigned to Merck & Co.), Lys-Gly-Asp (KGD), and the fibrinogen γ-chain carboxy-terminal dodecapeptide HHLGGAKQAGDV and analogues thereof (Timmons *et al.*, *Biochemistry*, 28:2919-2922 (1989)).

It is known that, for the RGD recognition sequence, the conformation of the RGD sequence is important for receptor recognition. Pierschbacher *et al.* (*J. Biol. Chem.* 292:17294-17298 (1987)) found that by cyclizing the peptide Gly-Pen-Gly-Arg-Gly-Asp-Ser-Pro-Cys-Ala through a Pen-Cys disulfide bridge the peptide became 10 times more effective at inhibiting the vitronectin-vitronectin receptor interaction but ineffective at inhibiting the fibronectin-fibronectin receptor interaction when compared to the corresponding linear peptide. Similarly, Kirchofer *et al.* (*J. Biol. Chem.* 265:18525-18530 (1990)) observed that the disulfide bridged cyclic peptide, cyclic-2,10-GPenGHRGDLRCA preferentially inhibits the binding of GP IIb/IIIa to fibrinogen but does not inhibit the binding of other RGD-dependent integrins, $\alpha_v\beta_3$ and $\alpha_5\beta_1$, to their respective ligands to the same extent. These authors also observed that a smaller "non-disulfide bridged" cyclic peptide, cyclic-1,7-VRGDSPDG, preferentially inhibited $\alpha_v\beta_3$ (vitronectin receptor) binding to vitronectin.

These results demonstrate that the conformation of the recognition sequence is important to specificity, but do not provide guidance regarding what the proper conformation(s) are and what structural features produce those conformation(s). Many examples of proteins having Arg-Gly-Asp and Lys-Gly-Asp recognition sequences have been reported, each of which, due to primary, secondary and tertiary structural constraints, produces a limited number of conformations about the recognition sequence in a given protein. However, whether these conformations bind to specific integrin receptors or any receptor at all can not be deduced from knowledge of the primary structure and the mere fact that these sequences are constrained. A representative list of some of these recognition sequence containing proteins includes: Maes *et al.*, *Fed. Eur. Biochem. Soc.*, 241(1,2): 41-45 (1988); Moos *et al.*, *Nature*, 334: 701-703 (1988); Rauvala *et al.*, *J. Cell Biol.*, 107: 2293-2305 (1988); Drickamer *et al.*, *J. Biol. Chem.*, 261: 6878-6887 (1986); Bond and Strydom *et al.*, *Biochemistry*, 28: 6110-6113 (1989); Ratner *et al.*, *Nature*, 313: 277-284 (1985); Davies *et al.*, *Biochem. Soc. Trans.*, 18: 1326-1328 (1990); and Neurath *et al.*, *Mol. Immun.*, 27: 539-549 (1990).

Similarly, a number of synthetic peptides, including cyclic disulfides, have been disclosed as inhibitors of fibrinogen binding to platelets all of which contain the Arg-Gly-Asp recognition sequence. See U.S. Patent 4,683,291; WO89/05150; EPO 0 319 506 A2; EPO 0 341 915 A2; Plow *et al.*, *Proc. Natl. Acad. Sci. USA* 82: 8057-8061(1985); Ruggeri *et al.*, *Proc. Natl. Acad. Sci. USA* 83, 5708-5712(1986); Haverstick *et al.*, *Blood* 66, 946-952(1985); Plow *et al.*, *Blood* 70, 110-115(1987); F. El F. Ali, *et al.*, *Proc. Eleventh Amer. Peptide Symp.* 94-96(1990); and Pierschbacher *et al.*, *supra* (1987). None of these publications define structural features producing recognition sequence conformations that are specific for various integrin receptors.

Several synthetic cyclic peptides containing linkages other than disulfides, specifically the thioether linkage, have been reported. Gero *et al.*, *Biochem. Biophys. Res. Comm.* 120: 840-845(1984) describe a pseudohexapeptide analog of somatostatin where the

group [CH₂-S] is substituted for a peptide bond. Similarly, Edwards *et al.*, *Biochem. Biophys. Res. Comm.* 136: 730-736(1986) compare the biological activity of linear and cyclic enkephalin pseudopeptide analogs containing the thiomethylene ether linkage. Other enkephalin related pseudopeptides and macrocycles containing the [CH₂-S] substitution for peptides have been described (Spatola *et al.*, *Biopolymers* 25: 229-244(1986) and Spatola *et al.*, *Tetrahedron* 44: 821-833(1988)). No information is provided in these publications defining conformation(s) these linkages might induce in a cyclic peptide containing those linkages.

The interaction of GP II_bIII_a with fibrinogen is stimulated by certain factors released or exposed when a blood vessel is injured. Multiple factors, including a variety of physiologic stimuli and soluble mediators, initiate platelet activation via several pathways. These pathways have a common final step which is the activation of the GP II_bIII_a receptor on the platelet surface and its subsequent binding to fibrinogen followed by aggregation and thrombus formation. By virtue of these interactions GP II_bIII_a is a component of the platelet aggregation system (Pytela *et al.*, *Science* 231: 1559(1986)). Thus, inhibition of the interaction of GP II_bIII_a with Arg-Gly-Asp containing ligands such as fibrinogen is a useful means of modulating thrombus formation. An inhibitor which prevents this binding interaction would antagonize platelet aggregation following platelet activation by any stimulus and therefore would have important antithrombotic properties.

Many common human disorders are characteristically associated with a hyperthrombotic state leading to intravascular thrombi and emboli. These are a major cause of medical morbidity, leading to infarction, stroke and phlebitis, and of mortality from stroke and pulmonary and cardiac emboli. Patients with atherosclerosis are predisposed to arterial thromboembolic phenomena for a variety of reasons. Atherosclerotic plaques form niduses for platelet plugs and thrombi that lead to vascular narrowing and occlusion, resulting in myocardial and cerebral ischemic disease. This may happen spontaneously or following procedures such as angioplasty or endarterectomy. Thrombi that break off and are released into the circulation cause infarction of different organs, especially the brain, extremities, heart and kidneys.

In addition to being involved in arterial thrombosis, platelets may also play a role in venous thrombosis. A large percentage of such patients have no antecedent risk factors and develop venous thrombophlebitis and subsequent pulmonary emboli without a known cause. Other patients who form venous thrombi have underlying diseases known to predispose to these syndromes. Some of these patients may have genetic or acquired deficiencies of factors that normally prevent hypercoagulability, such as antithrombin-3. Others have mechanical obstructions to venous flow, such as tumor masses, that lead to low flow states and thrombosis. Patients with malignancy have a high incidence of thrombotic phenomena for unclear reasons. Antithrombotic therapy in this situation with currently available agents is dangerous and often ineffective.

Patients whose blood flows over artificial surfaces, such as prosthetic synthetic cardiac valves or through extracorporeal perfusion devices, are also at risk for the development of platelet plugs, thrombi and emboli. It is standard practice that patients with artificial cardiac valves be chronically anti-coagulated. However, in all instances, 5 platelet activation and emboli formation may still occur despite adequate anticoagulation treatment.

Thus, a large category of patients, including those with atherosclerosis, coronary artery disease, artificial heart valves, cancer, and a history of stroke, phlebitis, or pulmonary emboli, are candidates for limited or chronic antithrombotic therapy. The number 10 of available therapeutic agents is limited and these, for the most part, act by inhibiting or reducing levels of circulating clotting factors. These agents are frequently not effective against the patient's underlying hematologic problem, which often concerns an increased propensity for platelet aggregation and adhesion. They also cause the patient to be susceptible to abnormal bleeding. Available antiplatelet agents, such as aspirin, inhibit 15 only part of the platelet activation process and are therefore often inadequate for therapy and also cause the patient to be susceptible to abnormal bleeding.

An agent which effectively inhibits the final common pathway of platelet activation, namely fibrinogen binding to the GP II_bIII_a receptor, should accordingly be useful in a large group of disorders characterized by a hyperthrombotic state as described above. 20 The present invention contemplates such agents which are new compositions, namely cyclic polypeptides consisting in part of natural amino acids and in part of unnatural amino acids. These new compositions interfere with the interaction of Arg-Gly-Asp containing peptides and proteins, particularly fibrinogen, with the GP II_bIII_a complex thereby preventing platelet aggregation. Platelet aggregation has been identified as an early step in the 25 formation of platelet plugs, emboli and thrombi in the circulatory system which in turn have been shown to play an active role in cardiovascular complications and disease. Inhibition of fibrinogen binding to the GP II_bIII_a complex has been shown to be an effective antithrombotic treatment in animals (H. K. Gold, *et al.*, *Circulation* 77: 670-677(1988); T. Yasuda, *et al.*, *J. Clin. Invest.* 81: 1284-1291(1988); B. S. Coller, *et al.*, *Blood* 68: 783-786(1986)).

30 None of the foregoing references disclose a compound capable of potent platelet aggregation inhibition activity and low inhibitory activity for the adhesive interaction of vitronectin-vitronectin receptor, fibronectin-fibronectin receptor, and GP II_bIII_a receptor with ligands other than fibrinogen. Furthermore, none of these references disclose potent platelet aggregation inhibitors that do not produce untoward side effects such as increased cutaneous 35 bleeding time or decreased peripheral blood flow.

Accordingly, it is an object of this invention to produce compounds having potent platelet aggregation inhibition activity. It is another object of the invention to produce such compounds that are stable to degradation. It is a further object to produce potent platelet aggregation inhibitors that are specific and do not strongly inhibit RGD sensitive other

integrin interactions including the Vn-VnR, Fn-FnR, and GP IIbIIIa-vWF interactions. It is still a further object to produce potent platelet aggregation inhibitors that do not significantly increase cutaneous bleeding time or diminish other hemodynamic factors. These and other objects of this invention will be apparent from consideration of the invention as a whole.

Summary of the Invention

The objects of this invention are accomplished by providing a peptide comprising:

(a) a cyclic moiety containing a sequence selected from

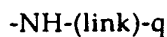
Xaa₁-Arg-Gly-Asp-Xaa₂, and

Xaa₁-Lys-Gly-Asp-Xaa₂

where Xaa₁ represents from 1 to 20 α -amino acids or α -amino acid analogues, and Xaa₂ represents an α -amino acid or α -amino acid analogue bonded to Xaa₁ through a linkage selected from the group amide, thioether, disulfide, amine, ether, sulfoxide, and sulfone, and

(b) a positively charged nitrogen containing exocyclic moiety stably bonded to Xaa₂ through a carbonyl group of Xaa₂.

Preferably the peptide contains a positively charged exocyclic moiety represented by the formula:



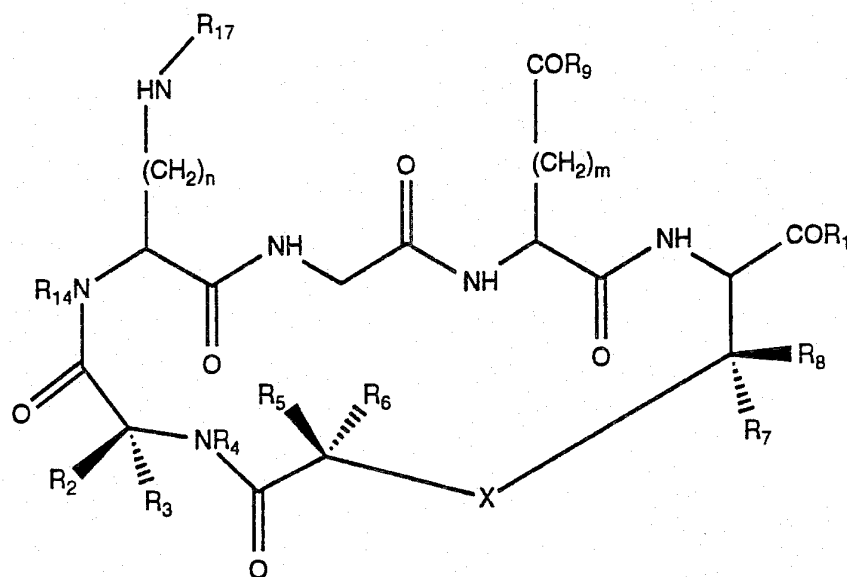
where (link) represents a linking group selected from C₃-C₁₀-alkyl either branched, linear, or cyclic, C₃-C₁₀-alkenyl, C₆-C₁₄-aryl substituted with 2 or more C₁-C₈-alkyl groups, C₅-C₁₄-heterocycle, saturated or unsaturated, containing from 1-4 heteroatoms selected from N, O, and S, C₁-C₆-alkyl substituted C₅-C₁₄, saturated or unsaturated heterocycle, containing from 1-4 heteroatoms selected from N, O, and S, and optionally, (link) may be substituted with substituents selected from COR, CONR'R", halo (F, Cl, Br, I), nitro, C₁-C₆-alkyl, phenyl, benzyl, and C₃-C₆-cycloalkyl, where R is selected from one or more of the groups hydroxy, C₁-C₈-alkoxy, C₃-C₁₂-alkenoxy, C₆-C₁₂-aryloxy, di-C₁-C₈-alkylamino-C₁-C₈-alkoxy, acylamino-C₁-C₈-alkoxy selected from the group acetylamoethoxy, nicotinoylaminoethoxy, succinamidoethoxy, and pivaloyloxyethoxy, C₆-C₁₂-aryl-C₁-C₈-alkoxy where the aryl group is unsubstituted or substituted with one or more of the groups nitro, halo (F, Cl, Br, I), C₁-C₄-alkoxy, and amino, hydroxy-C₂-C₈-alkoxy, dihydroxy-C₃-C₈-alkoxy, and where R' and R" are independently selected from hydrogen, C₁-C₁₀-alkyl either branched, linear, or cyclic, C₃-C₁₀-alkenyl, C₆-C₁₄-aryl, C₁-C₆-alkyl-C₆-C₁₀-aryl, saturated or unsaturated heterocycle having from 5-14 atoms in the cycles and from 1-4 heteroatoms selected from N, O, and S, optionally, R' and R" taken together may for trimethylene, tetramethylene, pentamethylene, and 3-oxopentamethylene, q represents a group selected from amino, amidino, and guanido, where any hydrogen bonded to any nitrogen or carbon of the

amino, amidino, or guanido group is optionally substituted with a lower C₁-C₆-alkyl group.

Typically, the positively charged exocyclic moiety is a positively charged amino acid residue selected from α -amino acids or α -amino acid analogues. These include either D or L, His, Lys, Arg or Orn, where the α -carboxyl group is derivitized with an amino or lower alkyl substituted amino group.

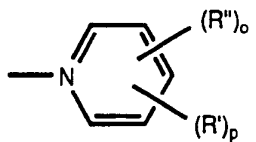
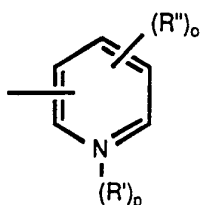
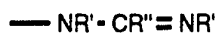
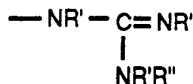
Also typically, Xaa₂ will be an amino acid residue selected from α -aminoadipic, α -amino-3-azaadipic, Cys, homo-Cys, Pen, Pas, Asp, Glu, Orn, Lys, Ser, Thr and Tyr. Most preferably, Xaa₂ will be selected from aminoadipic, Cys and Pen. Xaa₁ preferably is a single amino acid selected from Gly, D-Ala, D-Val, D-Leu, D-Ile, D-Phe, D-Tyr and D-Pro and the linkage bonding Xaa₁ with Xaa₂ will be a thioether or sulfoxide.

The preferred compound of this invention is represented by Formula I:



Formula I

where R₁ is selected from NHR₁₅Q, and NR₁₅R₁₆Q, where R₁₅ and R₁₆ are independently selected from C₃-C₁₀-alkyl either linear, branched or cyclic, C₃-C₁₀-alkenyl, C₆-C₁₄-aryl, C₁-C₆-alkyl-C₆-C₁₀-aryl, saturated or unsaturated heterocycle or C₁-C₆-alkyl substituted heterocycle containing from 5 to 14 atoms in the cycle and from 1 to 4 heteroatoms selected from N, O and S, NR₁₅ R₁₆ taken together may form a heterocycle or C₂-C₆-alkyl substituted heterocycle where R₁₅ and R₁₆ taken together are trimethylene, tetramethylene, pentamethylene or 3oxopentamethylene, and where each R₁₅ or R₁₆ may optionally be substituted with one or more substituents selected from COR₉, CONR'R", halo (F,Cl,Br,I), nitro, C₁-C₆-alkyl, phenyl, benzyl and C₃-C₆-cycloalkyl, and Q is a group bonded to R₁₅, R₁₆ or substituents bonded thereto, Q being selected from:



where R' and R'' are independently selected from hydrogen, C₁-C₁₀-alkyl either
 5 branched, linear, or cyclic, C₃-C₁₀-alkenyl, C₆-C₁₄-aryl, C₁-C₆-alkyl-C₆-C₁₀-aryl,
 saturated or unsaturated heterocycle having from 5-14 atoms in the cycles and from 1-
 4 heteroatoms selected from N, O and S, and where R' and R'' taken together are
 trimethylene, tetramethylene, pentamethylene, and 3-oxopentamethylene, and
 where R''' is C₁-C₁₀-alkyl, phenyl and benzyl, and o+ p is an integer selected from 0,
 10 1, 2, 3, and 4.

The term "functional group" when applied to amino acids and amino acid
 analogues refers to α -carboxyl, α -amino, and side-chain groups selected from the
 substituents of R₁, R₂ and R₃ as defined above. Typically, these substituents will be
 groups such as amino, amidino, guanidino, mercapto, hydroxy, carboxy, halo and
 15 aldehyde. These functional groups are capable of forming a bond with a compatible
 functional group of a second molecule.

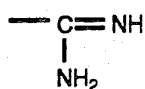
R₂, R₃, R₅, R₆, R₇, R₈ are the same or different and are selected from hydrogen, C₆-
 C₁₂-aryl where the aryl group is unsubstituted or substituted, C₁-C₁₂-alkyl, either
 substituted or unsubstituted, branched or straight chain, aromatic heterocycle where the
 20 heterocycle contains 5-10 ring atoms and one or two O, N or S heteroatoms; R₂ and R₃, R₅ and
 R₆, or R₇ and R₈ may optionally and independently be joined together to form a carbocyclic or
 heterocyclic ring of from four to seven atoms where the heteroatoms are selected from O, S or

NR₁₂, where R₁₂ is selected from hydrogen, C₁-C₈-alkyl, C₃-C₈-alkenyl, C₆-C₁₂-aryl, C₆-C₁₂-aryl-C₁-C₈-alkyl, C₁-C₈-alkanoyl and C₆-C₁₂-aroyl, and; R₄ is selected from hydrogen, C₁-C₈-alkyl, C₃-C₁₀-cycloalkyl, C₆-C₁₂-aryl, and C₆-C₁₂-aryl-C₁-C₈-alkyl; R₂ or R₃ may be optionally joined with R₄ to form a piperidine, pyrrolidine or thiazolidine ring;

5 R₉ is selected from hydroxy, C₁-C₈-alkoxy, C₃-C₁₂-alkenoxy, C₆-C₁₂-aryloxy, di-C₁-C₈-alkylamino-C₁-C₈-alkoxy, acylamino-C₁-C₈-alkoxy selected from the group acetylaminoethoxy, nicotinoylaminoethoxy, and succinamidoethoxy, pivaloyloxyethoxy, C₆-C₁₂ aryl-C₁-C₈-alkoxy where the aryl group is unsubstituted or substituted with one or more of the groups nitro, halo (F, Cl, Br, I), C₁-C₄-alkoxy, and amino, hydroxy-C₂-C₈-

10 alkoxy, dihydroxy-C₃-C₈-alkoxy, and NR₁₀R₁₁, where R₁₀ and R₁₁ are the same or different and are hydrogen, C₁-C₈-alkyl, C₃-C₈-alkenyl, C₆-C₁₂-aryl, where the aryl group is unsubstituted or substituted with one or more of the groups nitro, halo (F, Cl, Br, I), C₁-C₄-alkoxy, and amino, C₆-C₁₂-aryl-C₁-C₈-alkyl, where the aryl group is unsubstituted or substituted by one or more of the groups nitro, halo (F, Cl, Br, I), C₁-C₄-alkoxy, and amino;

15 R₁₄ is selected from hydrogen, C₁-C₈-alkyl, C₃-C₈-alkenyl, C₆-C₁₂-aryl, and C₆-C₁₂-aryl-C₁-C₈-alkyl; R₁₇ is selected from hydrogen, and



X is selected from O or S, S bearing one or two O atoms, NR₁₃ where R₁₃ is hydrogen, C₁-C₈-alkyl, C₃-C₈-alkenyl, C₆-C₁₄-aryl, C₆-C₁₄-aryl-C₁-C₈-alkyl, C₁-C₈-alkanoyl, C₆-C₁₄-aroyl, C₆-C₁₄-aryl-C₁-C₈-alkanoyl and (CH₂)_k, where k is an integer from 0 to 5, n is an integer from 1 to 6, and m is an integer from 0 to 4.

20

As used herein and unless specified otherwise, alkyl, alkenyl and alkynyl denote straight and branched hydrocarbon chains having single, double and triple bonds, respectively; C₆-C₁₂ aryl groups denote unsubstituted aromatic ring or fused rings such as, for example, phenyl or naphthyl; hetero denotes the heteroatoms O, N or S; aromatic heterocyclic groups have 5-10 ring atoms and contain up to four heteroatoms; halogen or halo denote F, Cl Br or I atoms; and alkoxy denotes an alkyl group attached to O.

25

Examples of C₁-C₈ alkyl or C₂-C₈ alkenyl groups include methyl, ethyl, propyl, isopropyl, butyl, t-butyl, pentyl, isopentyl, hexyl, vinyl, allyl, butenyl and the like; examples of C₃-C₁₀-cycloalkyl groups include cyclopropyl, cyclopentyl, cyclohexyl, and the like; aromatic heterocyclic groups include but are not limited to pyridyl, thienyl, furyl, indolyl, benzthienyl, imidazolyl, thiazolyl, quinolinyl and isoquinolinyl.

30

The present invention also includes a method for reducing platelet aggregation in a mammal. This method involves administering a therapeutically effective amount of the compounds of the present invention alone or in combination with a pharmacologically acceptable carrier. This general method may also be applied to treat a mammal having an increased propensity for thrombus formation.

35

Additionally, the present invention is directed to compositions of matter for reducing platelet aggregation in a mammal; treating a mammal having an increased propensity for thrombus formation; or inhibiting binding of a ligand to GP II_bIII_a in a mammal; wherein each of these compositions contains as an active ingredient one or more of the cyclic peptides defined in Formula I.

Detailed Description of the Invention

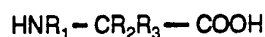
Definitions

Unless otherwise specified, the following definitions apply to the terms used throughout this specification.

10 By the term " α -amino acid" as used herein is meant naturally occurring α -amino acids encoded by mRNA and incorporated into polypeptides as they are synthesized on the ribosome. Except for glycine, these α -amino acids are chiral and of the L-stereoisomeric form.

The term "unnatural" α -amino acids is used to denote the D-isomeric form of the naturally occurring L- α -amino acids.

15 By the term " α -amino acid analogues" as used herein is meant those α -amino acids, other than the naturally occurring α -amino acids, represented by the formula:



20 Where R₁ is selected from hydrogen and C₁-C₆-alkyl. R₂ and R₃ are the same or different and are selected from: C₁-C₁₂-alkyl either linear branched or cyclic; C₂-C₁₂-alkenyl branched linear or cyclic; C₂-C₁₂-alkynyl; C₆-C₁₄-aryl, C₁-C₈-alkyl-C₆-C₁₄-aryl, C₁-C₁₃-heterocycle either saturated or unsaturated, the cycle(s) containing from 1-4 heteroatoms selected from N, O, and S; C₁-C₆-alkyl-C₁-C₁₃-heterocycle either saturated or unsaturated, the cycle(s) containing from 1-4 heteroatoms selected from N, O, and S; R₁ and
 25 R₂ or R₂ and R₃ taken together may form a carbocyclic or heterocyclic ring containing from 5 to 14 atoms of which from 1-4 heteroatoms are selected from N, O, or S. R₁, R₂ or R₃ may optionally be substituted with one or more substituents selected from:

10

halo(F, Cl, Br, I)

— OR'

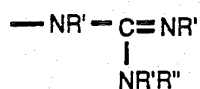
— SR'

— COOR'

— CONR'R"

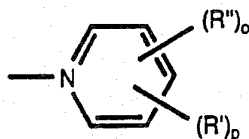
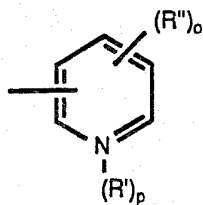
— NR'R"

— NR'R'R'''



— NR'—CR''=NR'

— NR'—CR''=NR''



5 where R' and R'' are independently selected from: hydrogen; C₁-C₁₀-alkyl either branched, linear, or cyclic; C₃-C₁₀-alkenyl; C₆-C₁₄-aryl; C₁-C₆-alkyl-C₆-C₁₀-aryl; saturated or unsaturated heterocycle having from 5-14 ring atoms and from 1-4 heteroatoms selected from N, O, and S; and where R' and R'' taken together may form a bridging functionality to produce a heterocycle, e.g., trimethylene, tetramethylene, pentamethylene, and 3-oxopentamethylene. R''' is C₁-C₁₀-alkyl, phenyl or benzyl and o+ p is an integer selected form 0, 1, 2, 3, and 4.

10 The following one-letter and three-letter abbreviations for α-amino acids and α-amino acid analogues are used herein as follows:

α -amino acids

	<u>One-Letter Code</u>	<u>Three-Letter Code</u>	<u>Common Name</u>
5	A	Ala	alanine
	R	Arg	arginine
	N	Asn	asparagine
	D	Asp	aspartic acid
	C	Cys	cysteine
	Q	Gln	glutamine
	E	Glu	glutamic acid
	G	Gly	glycine
	H	His	histidine
	I	Ile	isoleucine
	L	Leu	leucine
	K	Lys	lysine
	M	Met	methionine
	F	Phe	phenylalanine
	P	Pro	proline
	S	Ser	serine
	T	Thr	threonine
	W	Trp	tryptophan
	Y	Tyr	tyrosine
	V	Val	valine

 α -amino acid analogues

	<u>One-Letter Code</u>	<u>Three-Letter Code</u>	<u>Common Name</u>
10		Aad	α -aminoadipic acid
		Aib	α -aminoisobutyric acid
		Cha	Cyclohexylalanine
		Naa	naphthylalanine
		Nle	norleucine
	O	Om	ornithine
		Pas	6, 6-cyclopentamethylene-2-aminosuberic acid
		Pen	penicillamine
		Pmt	<i>p</i> -methyltyrosine
		SuA	succinylalanine

When capitalized, the one-letter code as used herein refers to the L or natural form of the amino acid. The corresponding lower case letter refers to the unnatural or D form of the α -amino acid or analogue.

As used herein, the term "cyclic moiety" refers to a peptide having an intramolecular bond between two adjacent amino acids within a peptide, that would not be adjacent in the absence of that bond. The intramolecular bond includes, but is not limited to; backbone to backbone, side-chain to backbone and side-chain to side-chain cyclizations.

The cyclic moiety includes the peptide bonds linking one amino acid or amino acid analogue residue with another as well as the side-chains of those amino acids or amino acid analogues. The cyclic moiety may also contain other intermolecular bonds including amide, thioether, disulfide, ether, amine, sulfoxide, sulfone and carbon-carbon bonds. The cyclic moiety may also contain groups other than amino acid and amino acid analogue residues such as acetyl or phenylacetyl groups used to form the cycle.

The term "exocyclic moiety" as used herein refers to a group having a bond connecting the group to a cycle, where breaking the bond would not destroy the cycle. By way of illustration, the exocyclic moiety may form a peptide bond by linking with an α -amino or α -carboxy of the cyclic moiety. Alternatively, the exocyclic moiety may be linked through an amide bond with the γ or δ carboxyl, or δ or ϵ amino side-chain groups of the cyclic moiety amino acid residues. Other side-chain linkages between the cyclic and exocyclic moieties may include ester and thioester, ether and thioether, aldimine, disulfide, sulfoxide, sulfone, and the like. Preferably, the exocyclic moiety will be bonded through an α , β , γ , δ , ϵ , or higher carboxyl group of an α -amino acid or α -amino acid analogue of the cyclic moiety. Most preferably, the exocyclic moiety will be bonded to the cyclic moiety through an α -carboxyl group.

As used herein, the term "stably bonded" when referring to bonding of the positively charged exocyclic moiety to the ring or cycle containing the recognition sequence means stable to hydrolysis in blood or serum. Thus, for example, amide or peptide bonds are considered stable while ester bonds, such as those employed in "prodrugs", are not considered stable.

As used herein, the term "peptide bond" refers to an amide bond between the α -carboxyl group of one α -amino acid or α -amino acid analogue and the α -amino group of another α -amino acid or α -amino acid analogue.

As used herein, the term "amide bond" is used to indicate any carboxy-amino bond other than α -carboxy- α -amino bond between α -amino acids and analogues.

Preferred Embodiments

The instant invention is a result of the unexpected discovery, that by bonding a positively charged exocyclic moiety to a cyclic peptide containing the tripeptide recognition sequences RGD or KGD, a potent platelet aggregation inhibitor is produced that is highly specific for the platelet GP IIb/IIIa receptor but which does not exhibit many of the *in vivo* side effects observed with other potent GP IIb/IIIa inhibitors.

The exocyclic moiety is typically positively charged at physiological pH (ca 7.4) and contains a nitrogen atom preferably as a substituted (alkyl) or unsubstituted amino, amidino or guanido functional group. Preferably, the exocyclic moiety will be a positively charged amino acid residue such as Lys, Arg or Orn, where the α -carboxyl is in the amide or uncharged form, and the α -amino group is linked to a carboxyl group of the cyclic moiety by a peptide or amide bond. Alternatively, the positive charge may be provided by linking a non-peptidyl group such as an alkylene diamine, e.g., ethylenediamine, propylenediamine, butylenediamine, octamethylenediamine and the like, with a free α -carboxyl group of the cyclic moiety.

10 The positively charged exocyclic moiety is bonded to the amino acid residue immediately following the RGD or KGD recognition sequence. Thus, the amino acid residue following the recognition sequence acts as a "branch point" both connecting the recognition sequence to the positively charged exocyclic moiety and participating in the formation of the cycle.

15 In one embodiment, the amino acid residue following the recognition sequence will (a) form a peptide or amide bond through its α -carboxyl group with the positively charged exocyclic group and (b) form a cycle through a functional group on the side-chain of that amino acid with either the α -amino group of the Lys or Arg of the recognition sequence or to a functional group of an amino acid or analogue on the amino side of the recognition sequence. In this embodiment, the cyclic moiety will contain, in addition to peptide bonds, a side-chain bridging bond typically selected from amide, ether, thioether, amine, sulfoxide, sulfone and disulfide.

20 In an alternative embodiment, the amino acid residue following the recognition sequence will (a) bond the exocyclic moiety through a side-chain functional group of that amino acid and (b) form the cycle through the α -carboxyl group of that amino acid residue. In this embodiment the cycle may contain all peptide bonds (e.g., backbone-backbone bridge) or may form a cycle through side-chain-side-chain or side-chain-backbone bridges between other amino acid residues.

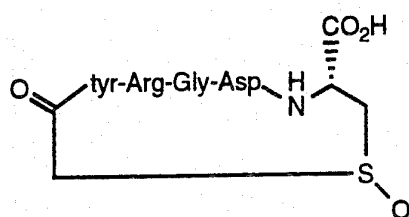
25 To be suitable for use as an antithrombotic agent, a cyclic peptide should be a potent inhibitor of platelet aggregation, that is, it must possess in IC_{50} in a human platelet aggregation inhibition assay of no more than at least about 3 μM . Preferably, the cyclic peptide will inhibit platelet aggregation with an $IC_{50} < 1 \mu M$ and most preferably the IC_{50} will range from about 50 to 500 nM.

30 In addition to being potent, the cyclic peptide should be specific, that is, not strongly inhibit the interaction of other integrin receptors for their natural ligands, or the interaction between GP II_bIII_a and ligands other than fibrinogen. Thus, the preferred cyclic peptide should not strongly inhibit the interaction between GP II_bIII_a and von Willebrand factor (vWF). Thus, in a vWF-GP II_bIII_a ELISA, as described herein, the preferred cyclic peptide should give an $IC_{50} > 1$ nM, more preferably $IC_{50} > 10$ nM, and most preferably the IC_{50}

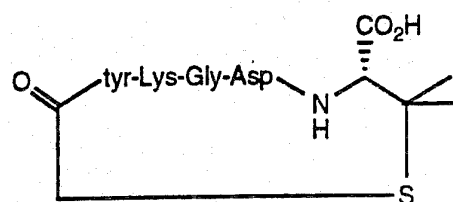
should range from at least about 15 nM to about 75 nM. Also preferably, the cyclic peptide should not strongly inhibit the interaction between Vitronectin (Vn) and the Vitronectin receptor $\alpha v \beta 3$ (VnR). Thus, in a Vn-VnR ELISA assay, as described herein, the preferred cyclic peptide should possess an $IC_{50} > 10$ nM, more preferably $IC_{50} > 50$ nM and most preferably $IC_{50} \geq 20$ μ M. Similarly, a preferred cyclic peptide should not strongly inhibit the fibronectin-fibronectin receptor interaction.

Finally, in addition to being small, potent and specific, the preferred cyclic peptide platelet aggregation inhibitor should not produce substantial untoward *in vivo* side effects such as an increased cutaneous bleeding time, reduction in platelet count, or decreased peripheral blood flow in a mammal treated with the inhibitor. Thus, for example, the time it takes for bleeding to stop, on an incision made in a mammal treated with a platelet aggregation inhibitor, dosed to inhibit about 90-100% of platelet aggregation, should not be more than about twice that of the mammal prior to treatment. Similarly, peripheral blood flow and other hemodynamic factors should not be adversely affected by treatment with doses sufficient to achieve 90-100% inhibition of platelet aggregation.

The compounds of this invention, containing the recognition sequences RGD or KGD in a cycle and bearing an exocyclic positively charged group bonded to the amino acid immediately following the Asp residue, have been found to be potent, specific and not to exhibit any untoward *in vivo* side effects at appropriate doses. The following discussion will be more readily understood by referring to Example 12 where the structures of the various compounds and the assay results are tabulated. The numerical designations used to identify the compounds refer to the example number/compound number as provided in the Examples section. The small cyclic peptides (7/46) and (7/47) shown below containing the RGD and KGD recognition sequences respectively, are potent inhibitors of platelet aggregation. Peptide (7/46) and peptide (7/47) have IC_{50} 's for platelet aggregation inhibition of 150 nM and 480 nM, respectively.



(7/46)

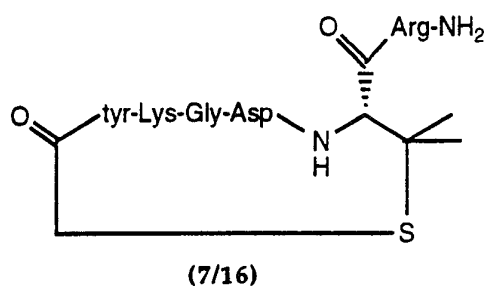
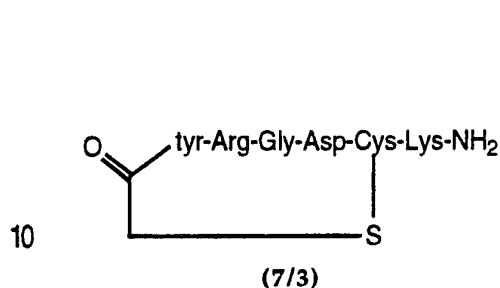
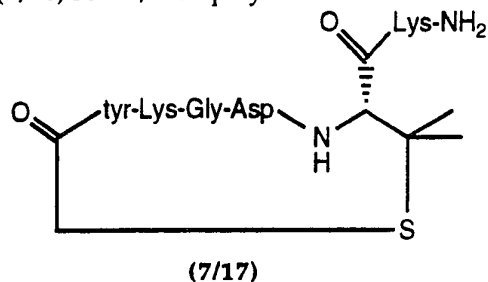
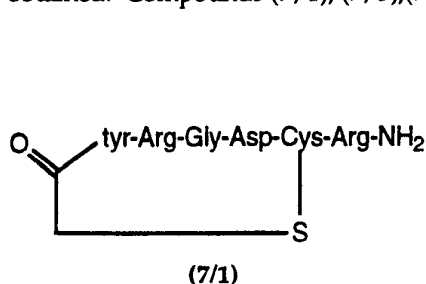


(7/47)

Compound (7/46) however is not a specific inhibitor of the fibrinogen-GP II_bIII_a interaction; rather, it potently inhibits both the GP II_bIII_a-vWF interaction ($IC_{50}=0.18$ nM) and the Vn-VnR interaction ($IC_{50}=5$ nM). Compound (7/47), while not inhibiting the Vn-VnR interaction ($IC_{50}>20$ μ M), is a fairly potent inhibitor of the GP II_bIII_a-vWF interaction ($IC_{50}=1.0$ nM). More importantly, however, both of these compounds exhibit an undesired side effect, namely, prolongation of bleeding time *in vivo* in both the rabbit and dog model. For example, in rabbit, a dose of compound (7/46 or 7/47) sufficient to achieve about 97%

inhibition of platelet aggregation, increased cutaneous bleeding time 8.0 and 8.6 times, respectively when compared to the undosed animal.

By adding an exocyclic positively charged moiety to the cyclic structures of either compound (7/46) or (7/47), more potent and more specific platelet aggregation inhibitors that do not substantially increase cutaneous bleeding time or decrease peripheral blood flow are obtained. Compounds (7/1), (7/3), (7/17), and (7/16) below, exemplify this result.

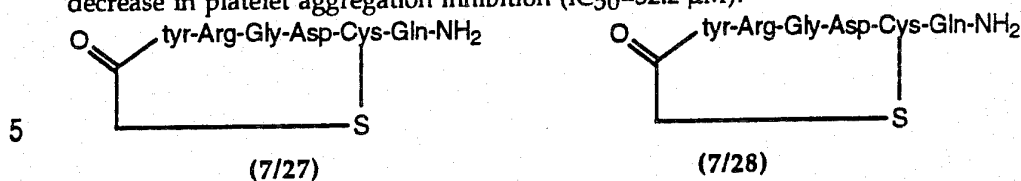


Bonding the arginylamide to the free carboxy residue of a compound of the (7/46) type produces compound (7/1) which exhibits high potency ($IC_{50}=80$ nM in the platelet aggregation assay), high specificity ($IC_{50}=13.8$ in the GPII_bIII_a-vWF ELISA), and no substantial increase in cutaneous bleeding time (1.17 times; dosed to 100% inhibition of platelet aggregation) or decrease in peripheral blood flow. Similarly, adding the positively charged lysylamide residue to a compound of the (7/46) type produces compound (7/3) which is highly potent ($IC_{50}=320$ nM in the platelet aggregation assay), highly specific ($IC_{50}=15.9$ nM in the GPII_bIII_a-vWF ELISA), and which does not substantially increase cutaneous bleeding time (1.95 times; dosed to 95% inhibition of platelet aggregation).

Compounds of the (7/47) type, that is containing the Lys-Gly-Asp sequence in the cycle, also exhibit the dramatically improved pharmacological properties upon adding an exocyclic positively charged group. Both compounds (7/17) and (7/16) containing an exocyclic lysylamide and arginylamide residue display high potency in the platelet aggregation assay (i.e. IC_{50} of 90 nM and 500 nM respectively), the highest specificity yet observed in the Vn-VnR, Fn-FnR, and GP II_bIII_a ELISA's, without substantial increased cutaneous bleeding time or decreased peripheral blood flow.

Various controls were tested in the same assays to ascertain whether the observed effect was due to the positively charged exocyclic moiety. Compound (7/27), which contains an uncharged exocyclic hydrogen-bond donor/receptor (Gln-NH₂), did not increase cutaneous blood time; however, it did not significantly improve potency. It is contemplated, however,

compound (7/27) may be modified, for example, by forming the sulfoxide or by substituting Pen for Cys to increase platelet aggregation inhibition potency. Compound (7/28), which contains a negatively charged exocyclic moiety (5-aminovaleric acid), exhibited a pronounced decrease in platelet aggregation inhibition ($IC_{50}=52.2 \mu M$).



Other positively charged non-peptidyl exocyclic groups also conferred high potency and specificity on cyclic peptides containing the tripeptide recognition sequences while not increasing cutaneous bleeding time. Exocyclic alkylene diamine groups as short as ethylenediamine produced the same effects. Similarly, inserting the amino acid residue Tyr between the positively charged exocyclic Arg and the bridging amino acid residue following the recognition sequence did not substantially decrease potency.

10

Methods for Making Platelet Aggregation Inhibitors

Polypeptides of this invention can be made by chemical synthesis or by employing recombinant technology. These methods are known to those of ordinary skill in the protein or peptide synthesis art. Chemical synthesis, especially solid phase synthesis, is preferred for short (eg. less than 50 residues) polypeptides or those containing unnatural or unusual amino acids such as; D-Tyr, Ornithine, aminoadipic acid, and the like. Recombinant procedures are preferred for longer polypeptides or for mutant or variant peptides containing the KGD or RGD sequence.

15

20

When recombinant procedures are selected, a synthetic gene may be constructed *de novo* or a natural gene may be mutanigized by, for example, cassette mutagenesis. Set forth below are exemplary general recombinant and chemical procedures.

General Recombinant Procedures

From a purified peptide or protein and its amino acid sequence, a KGD or RGD-containing peptide or protein may be produced using standard recombinant DNA techniques. These techniques contemplate, in simplified form, taking the gene, either natural or synthetic, for the peptide or protein; inserting it into an appropriate vector; inserting the vector into an appropriate host cell; culturing the host cell to cause expression of the gene; and purifying the peptide or protein produced thereby.

25

30

Somewhat more particularly, the DNA sequence encoding a KGD or RGD-containing peptide or protein is cloned and manipulated so that it may be expressed in a convenient host. DNA encoding parent polypeptides can be obtained from a genomic library, from cDNA derived from mRNA from cells expressing the peptide or protein, or by synthetically constructing the DNA sequence (Sambrook, J., Fritsch, E.F., and Maniatis, T., (1989), *Molecular Cloning* (2d ed.), Cold Spring Harbor Laboratory Press, N.Y.).

35

The parent DNA is then ligated into an appropriate plasmid or vector which is used to transform a host cell. In general, plasmid vectors containing replication and control

sequences which are derived from species compatible with the host cell are used in connection with those hosts. The vector ordinarily carries a replication site, as well as sequences which encode peptides or proteins that are capable of providing phenotypic selection in transformed cells.

5 For example, E. coli may be transformed using pBR322, a plasmid derived from an E. coli species (Mandel, M. *et al.* (1970) *J. Mol. Biol.* 53, 154). Plasmid pBR322 contains genes for ampicillin and tetracycline resistance, and thus provides easy means for selection. Other vectors include different features such as different promoters, which are often important in expression. For example, plasmids pKK223-3, pDR720, and pPL-lambda represent
10 expression vectors with the tac, trp, or PL promoters that are currently available (Pharmacia Biotechnology).

A preferred vector is pB0475. This vector contains origins of replication for phage and E. coli which allow it to be shuttled between such hosts, thereby facilitating both mutagenesis and expression (Cunningham, B., *et al.* (1989), *Science* 243, 1330-1336; Wells, J. and Cunningham, B., co-pending application U.S.S.N. 07/428,066 filed 26 October 1989.
15 Other preferred vectors are pR1T5 and pR1T2T (Pharmacia Biotechnology). These vectors contain appropriate promoters followed by the Z domain of peptide or protein A, allowing genes inserted into the vectors to be expressed as fusion peptides or proteins. Further discussion of these vectors may be found below.

20 Other preferred vectors can be constructed using standard techniques by combining the relevant traits of the vectors described above. Relevant traits include the promoter, the ribosome binding site, the target gene or gene fusion (the Z domain of peptide or protein A and the target gene and its linker), the antibiotic resistance markers, and the appropriate origins of replication.

25 The host cell may be prokaryotic or eukaryotic. Prokaryotes are preferred for cloning and expressing DNA sequences to produce parent polypeptides, segment substituted polypeptides, residue-substituted polypeptides and polypeptide variants. For example, E. coli K12 strain 294 (ATCC No. 31446) may be used as E. coli B, E. coli X1776 (ATCC No. 31537), and E. coli c600 and c600hfl, E. coli W3110 (F-, gamma-, prototrophic / ATCC No.
30 27325), bacilli such as *Bacillus subtilis*, and other enterobacteriaceae such as *Salmonella typhimurium* or *Serratia marcesans*, and various pseudomonas species. The preferred prokaryote is E. coli W3110 (ATCC 27325). When expressed by prokaryotes the polypeptides typically contain an N-terminal methionine or a formyl methionine and are not glycosylated. In the case of fusion peptides or proteins, the N-terminal methionine or
35 formyl methionine resides on the amino terminus of the fusion peptide or protein or the signal sequence of the fusion peptide or protein. These examples are, of course, intended to be illustrative rather than limiting.

In addition to prokaryotes, eukaryotic organisms, such as yeast cultures, or cells derived from multicellular organisms may be used. In principle, any such cell culture is

workable. However, interest has been greatest in vertebrate cells, and propagation of vertebrate cells in culture (tissue culture) has become a reproducible procedure (*Tissue Culture*, Academic Press, Kruse and Patterson, editors (1973)). Examples of such useful host cell lines are VERO and HeLa cells, Chinese Hamster Ovary (CHO) cell lines, W138, 293
5 (ATCC # CRL 1573), BHK, COS-7 and MDCK cell lines.

Gene Fusions

A variation on the above procedures contemplates the use of gene fusions, wherein the gene encoding the desired peptide or protein is associated, in the vector, with a gene
10 encoding another peptide or protein or a fragment of another peptide or protein. This results in the desired peptide or protein - here, a KGD or RGD-containing peptide or protein being produced by the host cell as a fusion with another peptide or protein. The "other" peptide or protein is often a protein or peptide which can be secreted by the cell, making it possible to isolate and purify the desired peptide or protein from the culture medium and eliminating the necessity of destroying the host cells which arises when the desired peptide or protein
15 remains inside the cell. Alternatively, the fusion peptide or protein can be expressed intracellularly. It is useful to use fusion peptides or proteins that are highly expressed.

The use of gene fusions, though not essential, can facilitate the expression of heterologous peptides or proteins in *E. coli* as well as the subsequent purification of those gene products (Harris, T. J. R. (1983) in *Genetic Engineering*, Williamson, R., Ed., Academic, London, Vol. 4, p. 127; Uhlen, M., Moks, To. (1989) *Methods Enzymol.* (in press)). Protein A
20 fusions are often used because the binding of protein A, or more specifically the Z domain of protein A, to IgG provides an "affinity handle" for the purification of the fused peptide or protein. It has also been shown that many heterologous peptides or proteins are degraded when expressed directly in *E. coli*, but are stable when expressed as fusions (Marston, F. A.
25 O., (1986) *Biochem J.* 240, 1).

A KGD or RGD-containing peptide or protein expressed as a fusion peptide or protein may be properly folded or may require folding or crosslinking to obtain the native structure. The properly folded fusion peptide or protein may be active and useful as a GP II_b III_a
30 antagonist and inhibitor of platelet aggregation. More preferred would be the correctly folded "native" peptide or protein that is obtained from the fusion peptide or protein by methods known in the art. Fusion peptides or proteins can be cleaved using chemicals, such as cyanogen bromide, which cleaves at a methionine, or hydroxylamine, which cleaves between an asn and gly. Using standard recombinant DNA methodology, the nucleotide base pairs encoding these amino acids may be inserted just prior to the 5' end of the KGD-
35 containing peptide or protein gene.

Alternatively, one can employ proteolytic cleavage of fusion peptides or proteins, which has been recently reviewed (Carter, P. (1990) in *Protein Purification: From Molecular Mechanisms to Large-Scale Processes*, Ladisch, M. R., Willson, R. C., Painton, C. C., and Builder, S. E., eds., American Chemical Society Symposium Series No. 427, Ch 13, 181-193).

Proteases such as Factor Xa, thrombin, subtilisin and mutants, and a number of other have been successfully used to cleave fusion peptides or proteins. Typically, a peptide linker that is amenable to cleavage by the protease used is inserted between the "other" peptide or protein (e.g., the Z domain of peptide or protein A) and the KGD or RGD-containing peptide or protein of interest. Using recombinant DNA methodology, the nucleotide base pairs encoding the linker are inserted between the genes or gene fragments coding for the other peptides or proteins. Proteolytic cleavage of the partially purified fusion peptide or protein containing the correct linker can then be carried out on either the native fusion peptide or protein, or the reduced or denatured fusion peptide or protein.

The peptide or protein may or may not be properly folded when expressed as a fusion peptide or protein. Also, the specific peptide linker containing the cleavage site may or may not be accessible to the protease. These factors determine whether the fusion peptide or protein must be denatured and refolded, and if so, whether these procedures are employed before or after cleavage.

When denaturing and refolding are needed, typically the peptide or protein is treated with a chaotrope, such as guanidine HCl, and is then treated with a redox buffer, containing, for example, reduced and oxidized dithiothreitol or glutathione at the appropriate ratios, pH, and temperature, such that the peptide or protein of interest is refolded to its native structure.

General Chemical Synthetic Procedures

When peptides are not prepared using recombinant DNA technology, they are preferably prepared using solid-phase synthesis, such as that generally described by Merrifield, *J. Am. Chem. Soc.* (1963) 85, 2149, although other equivalent chemical syntheses known in the art are employable. Solid-phase synthesis is initiated from the C-terminus of the peptide by coupling a protected α -amino acid to a suitable resin. Such a starting material can be prepared by attaching an α -amino-protected amino acid by an ester linkage to a chloromethylated resin or a hydroxymethyl resin, or by an amide bond to a BHA resin or MBHA resin. The preparation of the hydroxymethyl resin is described by Bodansky *et al.*, *Chem. Ind.* (London) (1966) 38,1597-1598. Chloromethylated resins are commercially available from BioRad Laboratories, Richmond, CA and from Lab. Systems, Inc. The preparation of such a resin is described by Stewart *et al.*, "Solid Phase Peptide Synthesis" (Freeman & Co., San Francisco 1969), Chapter 1, pp. 1-6. BHA and MBHA resin supports are commercially available and are generally used only when the desired polypeptide being synthesized has an unsubstituted amide at the C-terminus.

The amino acids are coupled to the peptide chain using techniques well known in the art for the formation of peptide bonds. One method involves converting the amino acid to a derivative that will render the carboxyl group more susceptible to reaction with the free N-terminal amino group of the peptide fragment. For example, the amino acid can be converted to a mixed anhydride by reaction of a protected amino acid with ethylchloroformate, phenyl

chloroformate, sec-butyl chloroformate, isobutyl chloroformate, pivaloyl chloride or like acid chlorides. Alternatively, the amino acid can be converted to an active ester such as a 2,4,5-trichlorophenyl ester, a pentachlorophenyl ester, a pentafluorophenyl ester, a p-nitrophenyl ester, a N-hydroxysuccinimide ester, or an ester formed from 1-

5 hydroxybenzotriazole.

Another coupling method involves use of a suitable coupling agent such as N,N'-dicyclohexylcarbodiimide or N,N'-diisopropylcarbodiimide. Other appropriate coupling agents, apparent to those skilled in the art, are disclosed in E. Gross & J. Meienhofer, *The Peptides: Analysis, Structure, Biology*, Vol. I: Major Methods of Peptide Bond Formation

10 (Academic Press, New York, 1979).

It will be recognized that the α -amino group of each amino acid employed in the peptide synthesis must be protected during the coupling reaction to prevent side reactions involving their active α -amino function. It will also be recognized that certain amino acids contain reactive side-chain functional groups (e.g. sulfhydryl, amino, carboxyl, and

15 hydroxyl) and that such functional groups must also be protected with suitable protecting groups to prevent a chemical reaction from occurring at that site during both the initial and subsequent coupling steps. Suitable protecting groups, known in the art, are described in E. Gross & J. Meienhofer, *The Peptides: Analysis, Structure, Biology*, Vol.3: Protection of Functional Groups in Peptide Synthesis (Academic Press, New York, 1981).

In the selection of a particular side-chain protecting group to be used in synthesizing the peptides, the following general rules are followed. An α -amino protecting group (a) must render the α -amino function inert under the conditions employed in the coupling reaction, (b) must be readily removable after the coupling reaction under conditions that will not remove side-chain protecting groups and will not alter the structure of the peptide fragment, and (c)

25 must eliminate the possibility of racemization upon activation immediately prior to coupling. A side-chain protecting group (a) must render the side chain functional group inert under the conditions employed in the coupling reaction, (b) must be stable under the conditions employed in removing the α -amino protecting group, and (c) must be readily removable upon completion of the desired amino acid peptide under reaction conditions that will not alter the

30 structure of the peptide chain.

It will be apparent to those skilled in the art that the protecting groups known to be useful for peptide synthesis will vary in reactivity with the agents employed for their removal. For example, certain protecting groups such as triphenylmethyl and 2-(*p*-biphenyl)isopropylloxycarbonyl are very labile and can be cleaved under mild acid

35 conditions. Other protecting groups, such as *t*-butyloxycarbonyl (BOC), *t*-amyloxycarbonyl, adamantyl-oxycarbonyl, and *p*-methoxybenzyloxycarbonyl are less labile and require moderately strong acids, such as trifluoroacetic, hydrochloric, or boron trifluoride in acetic acid, for their removal. Still other protecting groups, such as benzyloxycarbonyl (CBZ or Z), halobenzyloxycarbonyl, *p*-nitrobenzyloxycarbonyl, cycloalkyloxycarbonyl, and

isopropylloxycarbonyl, are even less labile and require stronger acids, such as hydrogen fluoride, hydrogen bromide, or boron trifluoroacetate in trifluoroacetic acid, for their removal. Among the classes of useful amino acid protecting groups are included:

- 5 (1) for an α -amino group, (a) aromatic urethane-type protecting groups, such as fluorenylmethyloxycarbonyl (Fmoc) CBZ, and substituted CBZ, such as, e.g., p-chlorobenzylloxycarbonyl, p-6-nitrobenzylloxycarbonyl, p-bromobenzylloxycarbonyl, and p-methoxybenzylloxycarbonyl, o-chlorobenzylloxycarbonyl, 2,4-dichlorobenzylloxycarbonyl, 2,6-dichlorobenzylloxycarbonyl, and the like; (b) aliphatic urethane-type protecting groups, such as BOC, t-amylloxycarbonyl, 10 isopropylloxycarbonyl, 2-(p-biphenyl)-isopropylloxycarbonyl, allyloxycarbonyl and the like; (c) cycloalkyl urethane-type protecting groups, such as cyclopentylloxycarbonyl, adamantylloxycarbonyl, and cyclohexylloxycarbonyl; and d) allyloxycarbonyl. The preferred α -amino protecting groups are BOC or Fmoc;
- 15 (2) for the side chain amino group present in Lys, protection maybe by any of the groups mentioned above in (1) such as BOC, p-chlorobenzylloxycarbonyl, etc.;
- (3) for the guanidino group of Arg, protection may be by nitro, tosyl, CBZ, adamantylloxycarbonyl, 2,2,5,7,8-pentamethylchroman-6-sulfonyl or 2,3,6-trimethyl-4-methoxyphenylsulfonyl, or BOC;
- (4) for the hydroxyl group of Ser, Thr, or Tyr, protection maybe, for example, by C1-C4 alkyl, such as t-butyl; benzyl (BZL); substituted BZL, such as p-methoxybenzyl, p-nitrobenzyl, p-chlorobenzyl, o-chlorobenzyl, and 2,6-dichlorobenzyl;
- 20 (5) for the carboxyl group of Asp or Glu, protection may be, for example, by esterification using groups such as BZL, t-butyl, cyclohexyl, cyclopentyl, and the like;
- 25 (6) for the imidazole nitrogen of His, the tosyl moiety is suitably employed;
- (7) for the phenolic hydroxyl group of Tyr, a protecting group such as tetrahydropyranyl, tert-butyl, trityl, BZL, chlorobenzyl, 4-bromobenzyl, and 2,6-dichlorobenzyl are suitably employed. The preferred protecting group is 2,6-dichlorobenzyl;
- 30 (8) for the side chain amino group of Asn or Gln, xanthyl (Xan) is preferably employed;
- (9) for Met, the amino acid is preferably left unprotected;
- (10) for the thio group of Cys, p-methoxybenzyl is typically employed.

The C-terminal amino acid, e.g., Lys, is protected at the N-amino position by an
35 appropriately selected protecting group, in the case of Lys, BOC. The BOC-Lys-OH can be first coupled to the benzylhydramine or chloromethylated resin according to the procedure set forth in Horiki *et al.*, *Chemistry Letters*, (1978)165-168 or using isopropylcarbodiimide at about 25°C for 2 hours with stirring. Following the coupling of the BOC-protected amino acid to the resin support, the α -amino protecting group is removed, as by using trifluoroacetic acid

(TFA) in methylene chloride or TFA alone. The deprotection is carried out at a temperature between about 0°C and room temperature. Other standard cleaving reagents, such as HCl in dioxane, and conditions for removal of specific α -amino protecting groups are described in Schroder & Lubke, *supra*, (Chapter I, pp. 72-75).

5 After removal of the α -amino protecting group, the remaining α -amino and side-chain protected amino acids are coupled step within the desired order. As an alternative to adding each amino acid separately in the synthesis, some may be coupled to one another prior to addition to the solid-phase synthesizer. The selection of an appropriate coupling reagent is within the skill of the art. Particularly suitable as a coupling reagent is N,N'-
10 dicyclohexylcarbodiimide or diisopropylcarbodiimide.

 Each protected amino acid or amino acid sequence is introduced into the solid-phase reactor in excess, and the coupling is suitably carried out in a medium of dimethylformamide (DMF) or CH₂Cl₂ or mixtures thereof. If incomplete coupling occurs, the coupling procedure is
15 repeated before removal of the N-amino protecting group prior to the coupling of the next amino acid. The success of the coupling reaction at each stage of the synthesis may be monitored. A preferred method of monitoring the synthesis is by the ninhydrin reaction, as described by Kaiser *et al.*, *Anal. Biochem*, 34: 595 (1970). The coupling reactions can be performed automatically using well known methods, for example, a Biosearch 9500 Peptide
20 Synthesizer.

 Upon completion of the desired peptide sequence, the protected peptide must be cleaved from the resin support, and all protecting groups must be removed. The cleavage reaction and removal of the protecting groups is suitably accomplished simultaneously or
25 stepwise. When the resin support is a chloro-methylated polystyrene resin, the bond anchoring the peptide to the resin is an ester linkage formed between the free carboxyl group of the C-terminal residue and one of the many chloromethyl groups present on the resin matrix. It will be appreciated that the anchoring bond can be cleaved by reagents that are known to be capable of breaking an ester linkage and of penetrating the resin matrix. One especially convenient method is by treatment with liquid anhydrous hydrogen fluoride. This reagent not only will cleave the peptide from the resin but also will remove all protecting
30 groups. Hence, use of this reagent will directly afford the fully deprotected peptide. When the chloromethylated resin is used hydrogen fluoride treatment results in the formation of the free peptide acids. When the benzhydrylamine resin is used, hydrogen fluoride treatment results directly in the free peptide amines. Reaction with hydrogen fluoride in the presence of anisole and dimethylsulfide at 0°C for one hour will simultaneously remove the
35 side-chain protecting groups and release the peptide from the resin.

 When it is desired to cleave the peptide without removing protecting groups, the protected peptide-resin can undergo methanolysis to yield the protected peptide in which the C-terminal carboxyl group is methylated. The methyl ester is then hydrolyzed under mild alkaline conditions to give the free C-terminal carboxyl group. The protecting groups on

the peptide chain then are removed by treatment with a strong acid, such as liquid hydrogen fluoride. A particularly useful technique for methanolysis is that of Moore *et al.*, *Peptides, Proc. Fifth Amer. Pept. Symp.*, M. Goodman and J. Meienhofer, Eds., (John Wiley, N.Y., 1977), p. 518-521, in which the protected peptide-resin is treated with methanol and potassium cyanide in the presence of crown ether.

Another method for cleaving the protected peptide from the resin when the chloromethylated resin is employed is by ammonolysis or by treatment with hydrazine. If desired, the resulting C-terminal amide or hydrazide can be hydrolyzed to the free C-terminal carboxyl moiety, and the protecting groups can be removed conventionally.

It will also be recognized that the protecting group present on the N-terminal α -amino group may be removed preferentially either before or after the protected peptide is cleaved from the support.

Purification of the polypeptides of the invention is typically achieved using conventional procedures such as preparative HPLC (including reversed phase HPLC) or other known chromatographic techniques such as gel permeation, ion exchange, partition chromatography, affinity chromatography (including monoclonal antibody columns) or countercurrent distribution.

Polypeptide chains are polymerized by crosslinking monomer chains with polyfunctional crosslinking agents, including compound 1, either directly or indirectly through multifunctional polymers. Ordinarily, two substantially identical polypeptides are crosslinked at their C or N termini using a bifunctional crosslinking agent. The agent is used to crosslink the terminal amino and/or carboxyl groups. Generally, both terminal carboxyl groups or both terminal amino groups are crosslinked to one another, although by selection of the appropriate crosslinking agent the alpha amino of one polypeptide is crosslinked to the terminal carboxyl group of the other polypeptide. Preferably, the polypeptides are substituted at their C-termini with cysteine. Under conditions well known in the art a disulfide bond can be formed between the terminal cysteines, thereby crosslinking the polypeptide chains. For example, disulfide bridges are conveniently formed by metal-catalyzed oxidation of the free cysteines or by nucleophilic substitution of a suitably modified cysteine residue. Selection of the crosslinking agent will depend upon the identities of their active side chains of the amino acids present in the polypeptides. For example, disulfide crosslinking would not be preferred if cysteine was present in the polypeptide at additional sites other than the C-terminus. Also within the scope hereof are peptides crosslinked with methylene bridges.

Suitable crosslinking sites on the peptides, aside from the N-terminal amino and C-terminal carboxyl groups, include epsilon amino groups found on lysine residues, as well as amino, imino, carboxyl, sulfhydryl and hydroxyl groups located on the side chains of internal residues of the peptides or residues introduced into flanking sequences. Crosslinking through externally added crosslinking agents is suitably achieved, e.g., using any of a number of

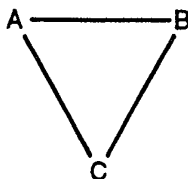
reagents familiar to those skilled in the art, for example, via carbodiimide treatment of the polypeptide. Other examples of suitable multifunctional (ordinarily bifunctional) crosslinking agents include 1,1-bis(diazoacetyl)-2-phenylethane; glutaraldehyde; N-hydroxysuccinimide esters (Bragg and Hou, *Arch. Biochem. Biophys.* (1975) 167, 311-321; Anjaneyula and Staros, *Int. J. Pep. Pro. Res.* (1987) 30, 117-124), such as esters with 4-azidosalicylic acid; homobifunctional imidoesters including disuccinimidyl esters such as 3,3'-dithiobis (succinimidyl-propionate) and dimethyladipimidate dihydrochloride (Zahn, *Agnew. Chem.* (1955) 67, 561-572; Golden and Harrison, *Biochemistry* (1982) 21, 3862-3866); bifunctional maleimides such as bis-N-maleimido-1,8-octane; disuccinimidyl suberate (Novick *et al.*, *J. Biol. Chem.* (1987) 262, 8483-8487), bis(sulfosuccinimidyl) suberate (Lee and Conrad, *J. Immunol.* (1985) 134, 518-525); heterobifunctional crosslinking reagents (Lomants and Fairbanks, *Arch. Biochem. Biophys.* (1976) 167, 311-321; Anjaneyula and Staros, *supra*; Partis *et al.*, *J. Pro.Chem.* (1983) 2, 263-277; Weltman *et al.*, *BioTechniques*, (1983) 1, 148-152; Yoshtake *et al.*, *J. Biochem.* (1982) 92, 1423-1424), including those with an N-hydroxysuccinimide moiety at one end and a maleimido group on the other end; succinimidyl 4-(N-maleimidomethyl) cyclohexane-1-carboxylate (SMCC) (Mahan *et al. Anal. Biochem.* (1987) 162, 163-170); sulfo-SMCC (Hashida *et al.*, *J. Applied Biochem.* (1984) 6, 56-63); *m*-maleimidobenzoyl-N-hydroxysuccinimide ester (MBS); sulfo-MBS; succinimidyl 4-(*p*-maleimidophenyl) butyrate (SMPB); sulfo-SMPB; N-succinimidyl(4-iodoacetyl)aminobenzoate (SIAB); sulfo-SIAB; 1-ethyl-3-(3-dimethylaminopropyl)carbodiimide hydrochloride (EDC); and N-hydroxysulfosuccinimide. Crosslinking agents such as methyl-3-[(*p*-azido-phenyl)dithio] propioimidate yield photoactivatable intermediates which are capable of forming crosslinks in the presence of light. If necessary, sensitive residues such as the side chains of the diargininyl group are protected during crosslinking and the protecting groups removed thereafter.

Polymers capable of multiple crosslinking serve as indirect crosslinking agents. For example, cyanogen bromide activated carbohydrates and the systems described in U.S. patents 3,959,080; 3,969,287; 3,691,016; 4,195,128; 4,247,642; 4,229,537; 4,055,635 and 4,330,440 are suitably modified for crosslinking the peptides herein. Crosslinking to amino groups of the peptides is accomplished by known chemistries based upon cyanuric chloride, carbonyl diimidazole, aldehyde reactive groups (PEG alkoxide plus diethyl acetal of bromoacetaldehyde; PEG plus DMSO and acetic anhydride, or PEG chloride plus the phenoxide of 4-hydroxybenzaldehyde). Also useful are succinimidyl active esters, activated dithiocarbonate PEG, and 2,4,5-trichlorophenyl-chloroformate- or *p*-nitrophenyl-chloroformate-activated PEG. Carboxyl groups are derivatized by coupling PEG-amine using carbodiimide. Ordinarily, however, the crosslinking agent is not a multifunctional polymer but instead is a small molecule being less than about 500 in MW.

The peptides of this invention also may be conformationally stabilized by cyclization. The peptides ordinarily are cyclized by covalently bonding the N and C-

terminal domains of one peptide to the corresponding domain of another peptide of this invention so as to form cyclooligomers containing two or more iterated peptide sequences, each internal peptide having substantially the same sequence. Further, cyclized peptides (whether cyclooligomers or cyclomonomers) are crosslinked to form 1-3 cyclic structures having from 2 to 6 peptides comprised therein. The peptides preferably are not covalently bonded through α -amino and main chain carboxyl groups (head to tail), but rather are cross-linked through the side chains of residues located in the N and C-terminal domains. The linking sites thus generally will be between the side chains of the residues.

The cyclic structures of the present invention will have the general formula:



wherein A and B represent the peptides of this invention and are the same or different. A and B are single peptides or head-to-tail polymers of two or more of such peptides. C represents one or more bonds or crosslinking moieties.

Many suitable methods per se are known for preparing mono- or poly-cyclized peptides as contemplated herein. Lys/Asp cyclization has been accomplished using N- α -Boc-amino acids on solid-phase support with Fmoc/9-fluorenylmethyl (OFm) side-chain protection for Lys/Asp; the process is completed by piperidine treatment followed by cyclization.

Glu and Lys side chains also have been crosslinked in preparing cyclic or bicyclic peptides: the peptide is synthesized by solid phase chemistry on a p-methylbenzhydrylamine resin. The peptide is cleaved from the resin and deprotected. The cyclic peptide is formed using diphenylphosphorylazide in diluted methylformamide. For an alternative procedure, see Schiller *et al.*, *Peptide Protein Res.* (1985) 25, 171-177. See also U.S. Patent 4,547,489.

Disulfide crosslinked or cyclized peptides are generated by conventional methods. The method of Pelton *et al.*, (*J. Med. Chem.* (1986) 29, 2370-2375) is suitable, except that a greater proportion of cyclooligomers are produced by conducting their action in more concentrated solutions than the dilute reaction mixture described by Pelton *et al.*, for the production of cyclomonomers. The same chemistry is useful for synthesis of dimers or cyclooligomers or cyclomonomers. Also useful are thiomethylene bridges (*Tetrahedron Letters* (1984) 25, 2067-2068). See also Cody *et al.*, *J. Med. Chem.* (1985) 28, 583.

The preferred thioether sulfoxide, and sulfone cyclization method employed in this invention is described in WO 91/01331, published 7 February 1991.

The desired cyclic or polymeric peptides are purified by gel filtration followed by reversed-phase high pressure liquid chromatography or other conventional procedures. The

peptides are sterile filtered and formulated into conventional pharmacologically acceptable vehicles.

The compounds described in this invention may be isolated as the free acid or base or converted to salts of various inorganic and organic acids and bases. Such salts are within the scope of this invention. Examples of such salts include ammonium, metal salts like sodium, potassium, calcium and magnesium; salts with organic bases like dicyclohexylamine, N-methyl-D-glucamine and the like; and salts with amino acids like arginine or lysine. Salts with inorganic and organic acids may be likewise prepared, for example, using hydrochloric, hydrobromic, sulfuric, phosphoric, trifluoroacetic, methanesulfonic, malic, maleic, fumaric and the like. Non-toxic and physiologically compatible salts are particularly useful although other less desirable salts may have use in the processes of isolation and purification.

A number of methods are useful for the preparation of the salts described above and are known to those skilled in the art. For example, reaction of the free acid or free base form of a compound of Formula I with one or more molar equivalents of the desired acid or base in a solvent or solvent mixture in which the salt is insoluble; or in a solvent like water after which the solvent is removed by evaporation, distillation or freeze drying. Alternatively, the free acid or base form of the product may be passed over an ion exchange resin to form the desired salt or one salt form of the product may be converted to another using the same general process.

Use of Inhibitors

As previously indicated, many common human disorders are characteristically associated with a hypercoagulable state leading to intravascular thrombi and emboli. These are a major cause of medical morbidity, leading to phlebitis, infarction, and stroke, and of mortality, from stroke and pulmonary and cardiac emboli. A large percentage of such patients have no antecedent risk factors, and develop venous thrombophlebitis and subsequent pulmonary emboli without a known cause. Other patients who form venous thrombi have underlying diseases known to predispose to these syndromes.

Some of these patients may have genetic or acquired deficiencies of factors that normally prevent hypercoagulability, such as anti-thrombin 3. Others have mechanical obstructions to venous flow, such as tumor masses, that lead to low flow states and thrombosis. Patients with malignancy have a high incidence of thrombotic phenomena, for unclear reasons. Antithrombotic therapy in this situation with currently available agents is dangerous and often ineffective.

Alternatively, the platelet aggregation inhibitors of this invention may be used in a pharmaceutical composition, optionally with a thrombolytic agent or anticoagulant to treat a mammal usually having an increased propensity for thrombus formation. Representative thrombolytic agents include but are not limited to; tissue plasminogen activator (t-PA), streptokinase, acylated plasminogen/streptokinase activator complex (APSAC), urokinase,

Pro-urokinase (suc-PA), and the like. Representative anticoagulants include but are not limited to heparin, dicumarol, warfin, hirudin, and the like (see e.g., Colman, *et al.*, *Hemostasis and Thrombosis*, 2nd Edition, J.B. Lippincott Co., Philadelphia (1987)).

5 Patients with atherosclerosis are predisposed to arterial thromboembolic phenomena for a variety of reasons. Atherosclerotic plaques form niduses from platelet plugs and thrombi that lead to vascular narrowing and occlusion, resulting in myocardial and cerebral ischemic disease. Thrombi that break off and are released into the circulation can cause infarction of different organs, especially the brain, extremities, heart and kidneys. After myocardial infarctions, clots can form in weak, poorly functioning cardiac chambers and be released into the circulation to cause emboli. All such patients with atrial fibrillation are
10 felt to be at great risk for stroke and require antithrombotic therapy.

In addition, thrombolytic therapy for acute myocardial infarction has become an established procedure for patients (Collen, D. and Stump, D. (1988) *Ann Rev Med* 39, 405-423). However, currently available thrombolytic agents are not effective in all patients which is
15 manifest by reocclusion, resistance to reperfusion, prolonged times to achieve normal coronary flow and the like. Since platelet mediated thrombosis is a major mechanism involved in the efficacy of thrombolytic therapy, agents which can be used to affect platelet aggregation in adjunctive therapy to treat acute myocardial infarction would have significant beneficial effects. Suitable thrombolytic agents include: tissue plasminogen activator, streptokinase,
20 urokinase, prourokinase, and modifications thereof.

Patients whose blood flows over artificial surfaces, such as prosthetic synthetic cardiac valves or through extracorporeal perfusion devices, are also at risk for the development of platelet plugs, thrombi, and emboli. It is standard practice that patients with artificial cardiac valves be chronically anti-coagulated.

25 Thus, a large category of patients, including those with cancer, atherosclerosis, coronary artery disease, artificial heart valves, and a history of stroke, phlebitis, or pulmonary emboli, are candidates for limited or chronic antithrombotic therapy. However, this therapy is often ineffective or morbid in its own right. This is partially because the number of available therapeutic agents is limited and these, for the most part, act by
30 reducing levels of circulating clotting factors. These agents are, therefore, not necessarily aimed at the patient's underlying hematologic problem, which often concerns an increased propensity for platelet aggregation and adhesion. They also cause the patient to be very susceptible to abnormal bleeding. Available antiplatelet agents, such as aspirin, inhibit the cyclooxygenase-induced activation of platelets only and are often inadequate for therapy.

35 In the management of thromboembolic disorders the compounds of this invention may be utilized in compositions such as tablets, capsules or elixers for oral administration; suppositories for rectal administration; sterile solutions or suspensions for injectable administration, atomized droplets for pulmonary or nasal administration, and the like. Animals in need of treatment using compounds of this invention can be administered dosages

that will provide optimal efficacy. The dose and method of administration will vary from animal to animal and be dependent upon such factors as weight, diet, concurrent medication and other factors which those skilled in the medical arts will recognize.

Dosage Formulations

5 Dosage formulations of the cyclic polypeptides of the present invention are prepared for storage or administration by mixing the the cyclic polypeptide having the desired degree of purity with physiologically acceptable carriers, excipients, or stabilizers. Such materials are non-toxic to the recipients at the dosages and concentrations employed, and include buffers such as phosphate, citrate, acetate and other organic acid salts; antioxidants such as ascorbic
10 acid; low molecular weight (less than about ten residues) peptides such as polyarginine, proteins, such as serum albumin, gelatin, or immunoglobulins; hydrophilic polymers such as polyvinylpyrrolidone; amino acids such as glycine, glutamic acid, aspartic acid, or arginine; monosaccharides, disaccharides, and other carbohydrates including cellulose or its derivatives, glucose, mannose, or dextrans; chelating agents such as EDTA; sugar alcohols such
15 as mannitol or sobitol; counterions such as sodium and/or nonionic surfactants such as Tween, Pluronic or polyethyleneglycol.

Dosage formulations of the cyclic polypeptides of the present invention to be used for therapeutic administration must be sterile. Sterility is readily accomplished by filtration through sterile filtration membranes such as 0.2 micron membranes. Cyclic polypeptide
20 formulations ordinarily will be stored in lyophilized form or as an aqueous solution. The pH of the cyclic polypeptide preparations typically will be between 3 and 11, more preferably from 5 to 9 and most preferably from 7 to 8. It will be understood that use of certain of the foregoing excipients, carriers, or stabilizers will result in the formation of cyclic polypeptide salts. While the preferred route of administration is by hypodermic injection needle, other
25 methods of administration are also anticipated such as suppositories, aerosols, oral dosage formulations and topical formulations such as ointments, drops and dermal patches.

Therapeutic cyclic polypeptide formulations generally are placed into a container having a sterile access port, for example, an intravenous solution bag or vial having a stopper pierceable by hypodermic injection needle.

30 Therapeutically effective dosages may be determined by either *in vitro* or *in vivo* methods. One method of evaluating therapeutically effective dosages consists of taking the cyclic polypeptide cyclo-S-acetyl-dTyr-Lys-Gly-Asp-Cys-Lys-NH₂ and determining a 50% inhibitory concentration (IC₅₀) of inhibiting fibrinogen binding to the GP II_b III_a platelet receptor. Similarly, in a platelet aggregation assay using the same cyclic peptide, the IC₅₀
35 is measured. Based upon such *in vitro* assay techniques, a therapeutically effective dosage range may be determined. For each particular cyclic polypeptide of the present invention, individual determinations may be made to determine the optimal dosage required. The range of therapeutically effective dosages will naturally be influenced by the route of administration. For injection by hypodermic needle it may be assumed the dosage is

delivered into the body's fluids. For other routes of administration, the absorption efficiency must be individually determined for each cyclic polypeptide by methods well known in pharmacology.

The range of therapeutic dosages is from about 0.001 nM to 1.0 mM, more preferably
5 from 0.1 nM to 100 μ M, and most preferably from 1.0 nM to 50 μ M.

Typical formulation of compounds of Formula I as pharmaceutical compositions are discussed below.

About 0.5 to 500 mg of a compound or mixture of compounds of Formula I, as the free
10 acid or base form or as a pharmaceutically acceptable salt, is compounded with a physiologically acceptable vehicle, carrier, excipient, binder, preservative, stabilizer, flavor, etc., as called for by accepted pharmaceutical practice. The amount of active ingredient in these compositions is such that a suitable dosage in the range indicated is obtained.

Typical adjuvants which may be incorporated into tablets, capsules and the like are
15 a binder such as acacia, corn starch or gelatin; an excipient such as microcrystalline cellulose; a disintegrating agent like corn starch or alginic acid; a lubricant such as magnesium stearate; a sweetening agent such as sucrose or lactose; a flavoring agent such as peppermint, wintergreen or cherry. When the dosage form is a capsule, in addition to the above materials it may also contain a liquid carrier such as a fatty oil. Other materials of various types may
20 be used as coatings or as modifiers of the physical form of the dosage unit. A syrup or elixer may contain the active compound, a sweetener such as sucrose, preservatives like propyl paraben, a coloring agent and a flavoring agent such as cherry. Sterile compositions for injection can be formulated according to conventional pharmaceutical practice. For example, dissolution or suspension of the active compound in a vehicle such as water or naturally
25 occurring vegetable oil like sesame, peanut, or cottonseed oil or a synthetic fatty vehicle like ethyl oleate or the like may be desired. Buffers, preservatives, antioxidants and the like can be incorporated according to accepted pharmaceutical practice.

The present invention has of necessity been discussed herein by reference to certain
30 specific methods and materials. It is to be understood that the discussion of these specific methods and materials in no way constitutes any limitation on the scope of the present invention, which extends to any and all alternative materials and methods suitable for accomplishing the objectives of the present invention.

EXAMPLES

35 In the following Examples, common α -amino acids may be described by the standard three letter amino acid code when referring to intermediates and final products. When the three-letter code begins with a lower-case letter, it is understood the amino acid is unnatural or the D-isomeric form. Standard abbreviations are listed in The Merck Index, 10th Edition, pp Misc-2 - Misc-3. Modified or unusual α -amino acids such as norleucine (Nle) and ornithine
40 (Orn) are designated as described in U.S. Patent and Trademark Office Official Gazette

1114TMOG, May 15, 1990. If the product or intermediate name is preceded by "cyclo-S-" this shall be taken to mean that the peptide has been cyclized through a sulfur atom.

EXAMPLE 1

Bleeding Time and Ex-Vivo Platelet Aggregation Rabbit Model

5 A. Animal Preparation and Blood Sampling

Unanesthetized male New Zealand White rabbits (2.5-3.5 kg) are placed in a standard rabbit restrainer. Ears are shaved and a 20G teflon catheter with flowswitch (Viggo) is placed in the medial artery, flushed with saline and locked with 1ml of heparinized saline (10 m/ml). A 22G catheter (Abbott) fitted with an injection cap (Medex) is placed in the marginal vein of the same ear. Saline or a GP II_bIII_a receptor antagonist, at a concentration of 1 to 3 mg/ml, is infused via the venous catheter. At time 0, 41% of the dose is given as a bolus over 2 minutes. The remainder is continuously infused over the following 60 minutes. Blood samples (3.2 ml) are collected into syringes without needles via the arterial catheter at -10, -5, 10, 45, and 60 minutes. The first 0.5 ml is discarded and the following 2.7 ml is collected directly into a syringe containing 0.3 ml of 3.8% sodium citrate. The sample is divided into 1.5 ml aliquots and centrifuged at room temperature for 5 seconds at 12,000 G. The resulting platelet rich plasma (PRP) is used to measure *ex-vivo* platelet aggregation (XPA). At -10 and 60 minutes an additional 1ml sample is drawn for an automated blood count (Baker instruments). Catheters are flushed and locked after every sample.

20 B. Ex-Vivo Platelet Aggregation (XPA)

300 ml of PRP is placed in a disposable glass cuvette with a stir bar. The cuvette is placed in the temperature regulated light path of a light transmittance aggregometer (Chrono-log) and equilibrated to a 37 °C. Baseline transmittance is recorded for 30 seconds, after which 10 ml of ADP (1mM) is added and the change in transmittance recorded. The maximum change from baseline (dT) is noted for each sample. The extent of inhibition of XPA that is produced by an inhibitor is calculated for each animal as follows: Mean dTs are calculated for the pre and post infusion values, and then, percent inhibition is calculated as $(1-dt \text{ (post)} / ddt \text{ (pre)}) \times 100$.

C. Cutaneous Bleeding Times (CBT)

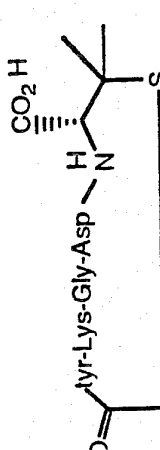
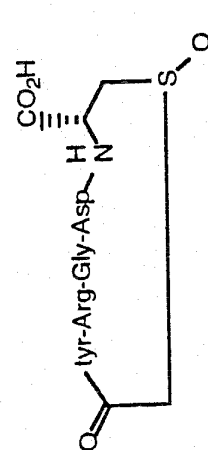
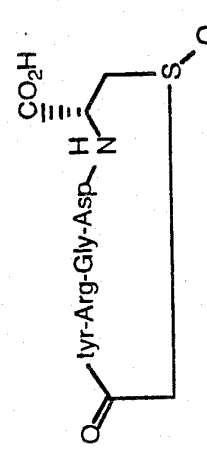
30 CBT is measured at -10, -5, 10, and 45 minutes on the opposite ear, using an automated incision-making instrument (Surgicutt®, ITD). An incision (5mm x 1mm deep) is made on the dorsal surface of the ear at sites not supplied by major blood vessels. Blood is blotted away with absorbant paper placed near the incision site, every 2 to 15 seconds, to a maximum of 15 minutes. Cessation of bleeding is defined as no blood forming at the incision site for 15 seconds. The range of duplicate CBT in 40 normal rabbits was 0.88 to 3.38 minutes.

D. Peripheral Blood Flow (PBF)

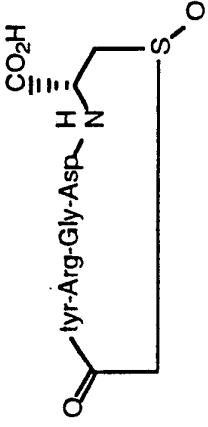
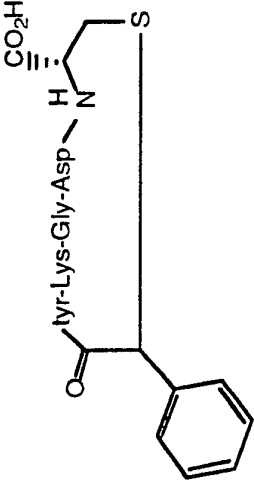


For the experiments summarized in Table 1, PBF was monitored by observation of the condition of the blood vessels in the rabbits' ears, prior to and during the infusion. Normal flow is defined as ears that appear pink to red, with no visible constriction of the major blood

vessels. Decreased flow applies to ears that have constricted vessels resulting in cold, pallid ears for up to 40 minutes following the start of the inhibitor infusion.

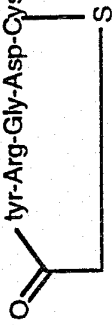

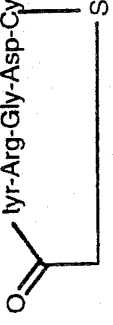
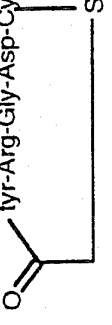
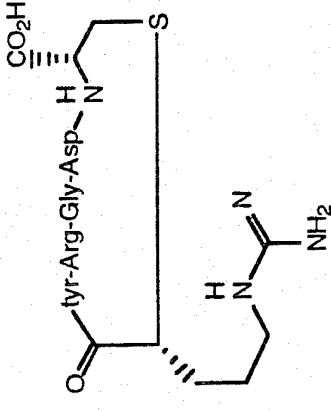
TABLE 1
Rabbit Cutaneous Bleeding Time, Peripheral Blood Flow v. Ex Vivo Platelet Aggregation and Inhibitor Dose

<u>Compound</u> ¹	<u>Dose(mg/kg)</u> ²	<u>CBT(post/pre)</u> ³	<u>XPA(%Inhib)</u> ⁴	<u>PBF</u> ⁵	<u>Number of Samples Tested</u>
Control	Saline	1.09	11	NORMAL	2
	1.5 + 0.036	8.57	96	NORMAL	1
	0.013 + 0.0003	2.74	55	NORMAL	1
	.375 + 0.009	5.67	95	NORMAL	1

Rabbit Cutaneous Bleeding Time, Peripheral Blood Flow v. Ex Vivo Platelet Aggregation and Inhibitor Dose

<u>Compound</u> ¹	<u>Dose(mg/kg)</u> ²	<u>CBT(post/pre)</u> ³	<u>XPA(%Inhib)</u> ⁴	<u>PBF</u> ⁵	<u>Number of Samples Tested</u>
	0.5 + 0.012	8.77	99	NORMAL	2
	1.5 + 0.036	4.47	94	NORMAL	2
	0.5 + 0.012	1.71	100	NORMAL	1
	1.5 + 0.036	6.72	95	NORMAL	2

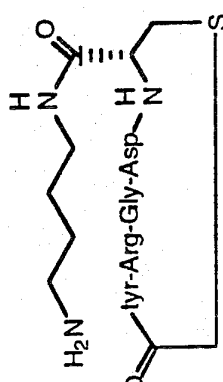
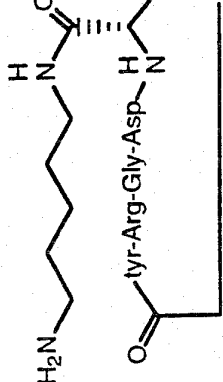
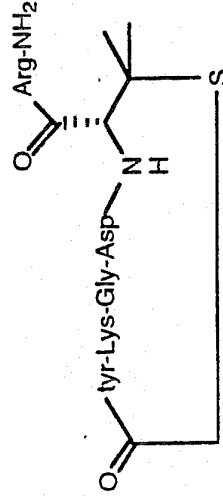
Rabbit Cutaneous Bleeding Time, Peripheral Blood Flow v. Ex Vivo Platelet Aggregation and Inhibitor Dose

<u>Compound</u> ¹	<u>Dose(mg/kg)</u> ²	<u>CBT(post/pre)</u> ³	<u>XPA(%Inhib)</u> ⁴	<u>PBF5</u>	<u>Number of Samples Tested</u>
 Tyr-Arg-Gly-Asp-Cys-Lys-NH ₂	1.5 + 0.036	1.95	97	NORMAL	2
 Tyr-Arg-Gly-Asp-Cys-Orn-NH ₂	1.5 + 0.036	1.26	100	NORMAL	2
 Tyr-Arg-Gly-Asp-Cys-orn-NH ₂	1.5 + 0.036	1.40	53	NORMAL	1
 Tyr-Arg-Gly-Asp-Cys-Tyr-Arg-NH ₂	1.5 + 0.036	4.10	98	NORMAL	2
 Tyr-Arg-Gly-Asp-Cys-Lys-NH ₂	1.5 + 0.036	1.46	55	NORMAL	1

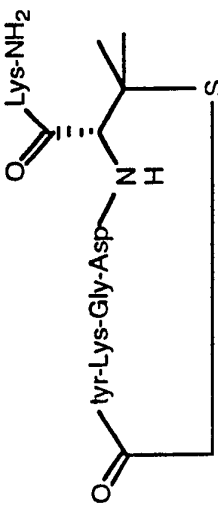
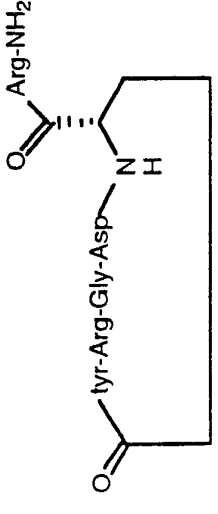
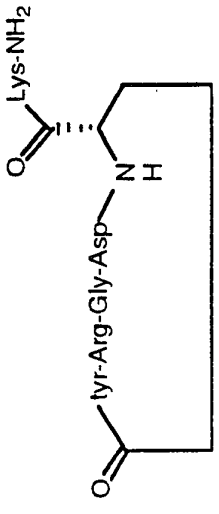
Rabbit Cutaneous Bleeding Time, Peripheral Blood Flow v. Ex Vivo Platelet Aggregation and Inhibitor Dose

<u>Compound</u> ¹	<u>Dose</u> (mg/kg) ²	<u>CBI</u> (post/pre) ³	<u>XPA</u> (%Inhib) ⁴	<u>PBF</u> ⁵	<u>Number of Samples Tested</u>
	1.5 + 0.036	1.00	1	NORMAL	1
	1.5 + 0.036	8.17	100	NORMAL	1
	1.5 + 0.036	1.42	35	NORMAL	2

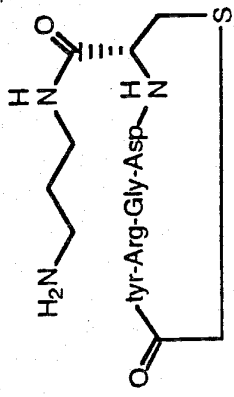
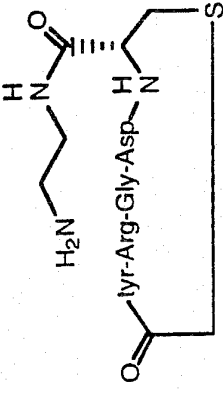
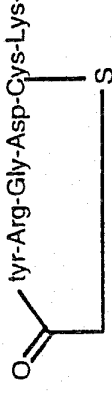
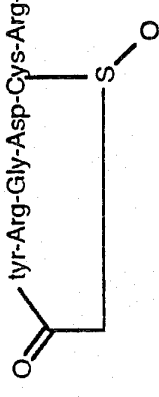
Rabbit Cutaneous Bleeding Time, Peripheral Blood Flow v. Ex Vivo Platelet Aggregation and Inhibitor Dose

<u>Compound</u> ¹	<u>Dose(mg/kg)</u> ²	<u>CBI(post/pre)</u> ³	<u>XPA(%Inhib)</u> ⁴	<u>PBF</u> ⁵	<u>Number of Samples Tested</u>
	1.5 + 0.036	0.94	96	NORMAL	1
	1.5 + 0.036	1.13	98	NORMAL	1
	1.5 + 0.036	3.29	100	NORMAL	1

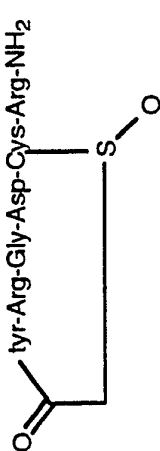
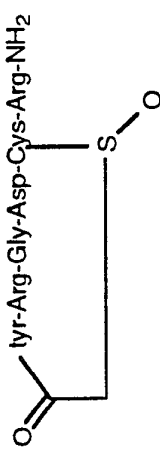
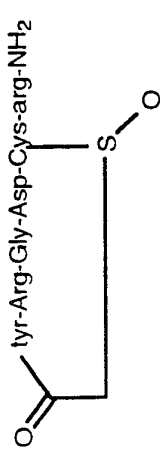
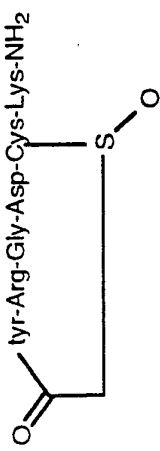
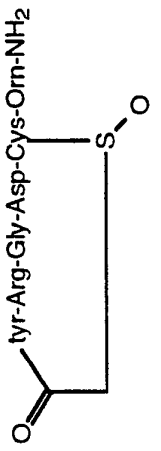
Rabbit Cutaneous Bleeding Time, Peripheral Blood Flow v. Ex Vivo Platelet Aggregation and Inhibitor Dose

<u>Compound¹</u>	<u>Dose(mg/kg)²</u>	<u>CBT(post/pre)³</u>	<u>XPA(%Inhib)⁴</u>	<u>PBF⁵</u>	<u>Number of Samples Tested</u>
	1.5 + 0.036	1.13	100	NORMAL	1
	1.5 + 0.036	7.98	98	NORMAL	1
	1.5 + 0.036	10.00	99	NORMAL	1

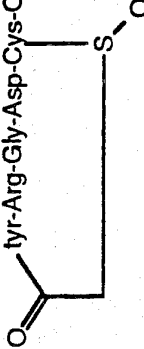
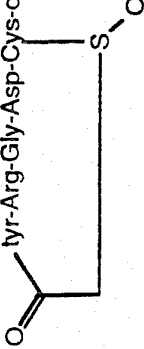
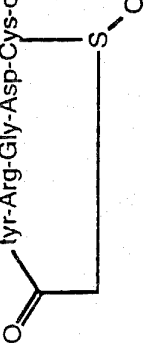
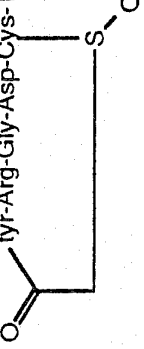
Rabbit Cutaneous Bleeding Time, Peripheral Blood Flow v. Ex Vivo Platelet Aggregation and Inhibitor Dose

Compound ¹	Dose(mg/kg) ²	CBT(post/pre) ³	XPA(%Inhib) ⁴	PBF ⁵	Number of Samples Tested
	1.5 + 0.036	1.27	99	NORMAL	1
	1.5 + 0.036	0.81	88	NORMAL	1
	1.5 + 0.036	6.14	100	NORMAL	1
	0.5 + 0.012	3.90	100	NORMAL	1

Rabbit Cutaneous Bleeding Time, Peripheral Blood Flow v. Ex Vivo Platelet Aggregation and Inhibitor Dose

Compound ¹	Dose(mg/kg) ²	CBT(post/pre) ³	XPA(%Inhib) ⁴	PBF ⁵	Number of Samples Tested
	1.5 + 0.036	16.67	95	NORMAL	1
	1.5 + 0.036	1.17	99	NORMAL	1
	1.5 + 0.036	0.93	64	NORMAL	1
	1.5 + 0.036	0.77	91	NORMAL	1
3-bromoyl tyrosine					
	1.5 + 0.036	2.63	99	NORMAL	1

Rabbit Cutaneous Bleeding Time, Peripheral Blood Flow v. Ex Vivo Platelet Aggregation and Inhibitor Dose

<u>Compound¹</u>	<u>Dose(mg/kg)²</u>	<u>CBT(post/pre)³</u>	<u>XPA(%Inhib)⁴</u>	<u>PBF⁵</u>	<u>Number of Samples Tested</u>
 Tyr-Arg-Gly-Asp-Cys-Orn-NH ₂	1.5 + 0.036	1.09	71	NORMAL	1
 Tyr-Arg-Gly-Asp-Cys-orn-NH ₂	1.5 + 0.036	1.50	98	NORMAL	1
 Tyr-Arg-Gly-Asp-Cys-orn-NH ₂	1.5 + 0.036	1.14	41	NORMAL	1
 Tyr-Arg-Gly-Asp-Cys-Tyr-Arg-NH ₂	1.5 + 0.036	1.18	52	NORMAL	1

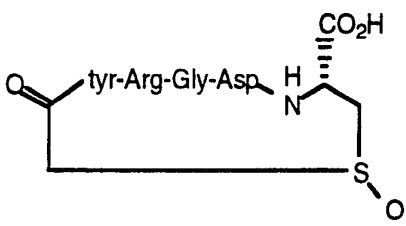
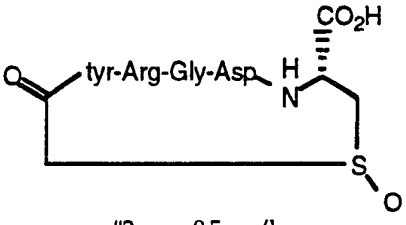
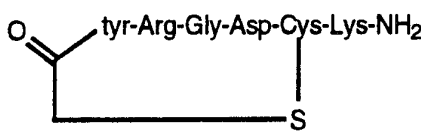
- 1 - Structure using three-letter code for amino acids; capital letters indicate L-configuration, lower case indicates D-configuration.
- 2 - Dose= Bolus dose in mg/kg; the additional amount indicated (also in mg/kg) was given by continuous infusion over 60 minutes.
- 3 - CBT=Cutaneous bleeding time in a ratio of an average of the post infusion bleeding times divided by the average of the pre bleeding times.
- 4 - XPA=Ex vivo platelet aggregation given in % inhibition of platelet aggregation.
- 5 - PBF=Peripheral blood flow in arbitrary units and is measured visually.

In an alternative series of experiments, (Table 2) PBF was measured quantitatively with a laser doppler flow probe (Perimed). The probe was positioned securely over the vascular bed of one ear and flow monitored continuously. Each inhibitor was infused and CBT measured in the opposing ear. No arterial catheter was placed for blood sampling, consequently XPA was not measured in these animals. However, the doses used were shown in previous experiments to effectively inhibit XPA.

TABLE 2

Rabbit Peripheral Blood Flow measured by dopler laser

10

<u>Compound</u> ¹	<u>Time (Min)</u> ²	<u>CBT(min)</u> ³	<u>PBF(units)</u> ⁴
 <p>#1 0.5mg/kg</p>	0	1.2	211
	10	15.0	220
	45		
 <p>#2 0.5mg/kg</p>	0	2.5	395
	10	15.0	380
	45	15.0	356
 <p>1.5mg/kg</p>	0	1.4	304
	10	1.8	283
	45	2.0	338

1 Structure using three-letter code for amino acids; capital letters indicate L-configuration, lower case indicates D-configuration.

15 2 Time of sampling.

3 CBT=Cutaneous bleeding time in a ratio of an average of the post infusion bleeding times divided by the average of the pre bleeding times.

4 PBF=Peripheral blood flow in arbitrary units and is measured using a laser doppler
flow probe.

E. Results

5 CBT, XPA, and observed PBF are summarized in Table 1. The ratio of the post to pre
treatment CBT was calculated for each animal by dividing the mean of the 2 post-treatment
samples by the mean of the pre samples. In two saline control rabbits the mean \pm sd ratio of
post treatment CBT (n=10) to a mean pre-treatment CBT was 1.12 ± 0.19 .

The experiments measuring PBF by laser doppler probe are summarized in Table 2. As
a positive control, epinephrine was infused intravenously (1 mg over 2 minutes.) at 60 minutes.
10 The resulting vasoconstriction reduced flow to near 0 flow units within 5 minutes.

The doses listed in the Tables refer to the bolus portion only. There were no
significant changes in any of the blood indices measured.

EXAMPLE 2

Fibrinogen-GP II_bIII_a Receptor ELISA Binding Assay

15 The method used is essentially that described in Nachman and Leung (J. Clin. Invest.
, 69: 263-269 [1982]). The GP II_bIII_a is essentially purified as described in Fitzgerald *et al.*,
(Anal. Biochem. 151: 169-177 [1985]).

A. GP II_bIII_a Purification

Outdated human platelets are washed 3 times with 10 mM tris-HCl, 150 mM NaCl
20 (TBS), 1 mM EDTA, pH 7.5, and centrifuged at 2000 x g to pellet cells. Cells are lysed in 5
pellet volumes of TBS, 1% Triton X-100, 1 mM Ca₂Cl₂, and followed by centrifugation at
30,000 x g. The supernatant fraction is collected and the supernatant is loaded onto a
concanavalin-A column, previously equilibrated in TBS, 1 mM Ca₂Cl₂, 0.1% Triton, 0.05%
NaN₃ and eluted with 0.2 M α -methylmannoside. Fractions are pooled and loaded onto a
25 heparin-agarose column. The flowthrough is collected and concentrated on an Amicon YM 30
filter to a volume of approximately 5-10 ml. The concentrate is then applied to an S-300
column (500 ml) and 6 ml fractions are collected. The GP II_bIII_a containing fractions are
collected, pooled, and stored at -80° C.

B. Purification of Low Solubility Fraction of Fibrinogen

30 The purification of fibrinogen is conducted essentially as described by Lipinska *et al.*,
(J. Lab. Clin. Med. 507, [1974]). Briefly, a 0.3 % w/v solution of human fibrinogen (Kabi
#5302) is dissolved in 150 mM NaCl. Saturated (NH₄)₂SO₄ is added dropwise with stirring
to the fibrinogen solution to obtain about 16% saturation. The precipitate is spun down in
appropriate size bottles at 2000 x g. The supernatant is decanted and the precipitate
35 resuspended in 150 mM NaCl (approximately 50% of the original volume). NH₄SO₄ is
again added dropwise to obtain 16% saturation. The suspension is spun down and the
precipitate is resuspended in Tris-saline in a minimal volume (approximately 5% of the
original volume). Any remaining insoluble material is spun down at 2000 rpm in a Sorval
type centrifuge and the fibrinogen supernatant is decanted and dialyzed overnight at 4° C

against Tris-saline. Characterization of the fibrinogen is by the Bradford protein assay, SDS-PAGE, and/or Western blotting using well known standard procedures.

C. ELISA Assay

Briefly, 96 well plates are coated (Type Nunc 1 Maxisorp™) with 10 µg/ml purified
5 fibrinogen (100 µl/well), and allowed to stand overnight at 4°C. The plates are washed
three times with PBS Tween (0.137 M NaCl, 0.003 M KCl, 0.008 M Na₂HPO₄, 0.001 M
KH₂PO₄, pH 7.4 at room temperature, 0.05% Tween-20) and blocked for 1 to 2 hours at room
temperature with 200 µl/well TNCNT (which is 0.5% BSA, 20mM Tris, pH 7.5 at room
temperature, 120mM NaCl, 0.2% NaN₃, 2mM CaCl₂, 0.05% Tween 20, 0.5% BSA [Calbiochem
10 RIA grade or better]) on a plate shaker. The plates are again washed three times with
PBS/Tween and then 50 µl of sample in TNCNT is added. The mixture is incubated for 15
minutes at room temperature on a plate shaker. The stock solution of purified GP IIbIIIa
receptor from human platelets, (0.4 - 1.0 mg/ml GP IIbIIIa in 0.1% Triton X-100, 1mM CaCl₂,
20 mM Tris, 150 mM NaCl, 0.05% NaN₃ in 0.3 M N-acetyl glucosamine pH 7.5, stored at -
15 70°C), is reconstituted to about 40 µg/ml in TNCNT. Fifty µl of this diluted GP IIbIIIa is
then added to each well and incubated on a plate shaker at room temperature. After one
hour, the plates are washed four times with PBS/Tween and 100 µl of a polyclonal or
monoclonal antibody specific for GP IIIa such as AP3 (1µg/ml) (See e.g. Newman *et al.*, *Blood*,
65: 227-232 [1985]) in ELISA buffer (PBS, 0.5% BSA, 0.05% Tween 20, 0.01% Thimerosal) is
20 added. After a one hour incubation at room temperature on a plate shaker, the samples are
washed 4 times with PBS/Tween. One hundred µl of GAM-HRP (horse radish peroxidase
conjugate of goat anti-mouse IgG [Pel-Freeze Cat. 715305-1] dissolved in ELISA buffer)
previously diluted to 1:10,000 is then added and the samples are incubated 1 hour at room
temperature on a plate shaker. The samples are then washed 4 times with PBS/Tween and
25 100 µl OPD/H₂O₂ substrate is added (OPD/H₂O₂ substrate: 10 mg o-phenylenediamine in
15 ml phosphate/citrate buffer, at room temperature and covered with foil; just before use,
6.25 µl of 30% H₂O₂ is added to give a final solution of 0.67 mg OPD/ml in 0.0125% H₂O₂).
(The phosphate/citrate buffer consists of 16 mM Citric Acid, 50 mM Na₂ HPO₄, pH 5.0). The
color develops within about 3 to 20 minutes and the reaction is stopped with 100 µl 1 M
30 H₂SO₄. The optical density at 492 nm vs 405 nm is recorded and IC₅₀ values are determined.

EXAMPLE 3

Human Vitronectin-Vitronectin Receptor ($\alpha_v\beta_3$) ELISA Assay

A. Human Vitronectin Purification

Human vitronectin (Vn) is isolated from human plasma and purified by affinity
35 chromatography by the method of Yatohgo *et al.*, (*Cell Structure and Function* 13: 281-292
[1988]).

B. Human Vitronectin receptor ($\alpha_v\beta_3$) Purification

Human vitronectin receptor (VnR) is purified from human placenta by the method of
Pytela *et al.*, (*Methods Enzymol.*, 144: 475 [1987]). Alternatively the $\alpha_v\beta_3$ receptor can be

purified from some cell lines (e.g., human embryonic kidney 293 cells) transfected with DNA sequences for both the α_v and β_3 subunits. The subunits are purified by employing octylglucoside extraction followed by Con-A, Heparin-Sepharose, and S-300 Chromatography.

5 C. Monoclonal Antibodies

Anti-GP II_bIII_a monoclonal antibodies specific for human GP III_a are prepared by the method of Newman *et al.* (*Blood*, 65: 227-232 [1985]), or a similar procedure. This mouse Mab is specific for the β_3 subunit of the vitronectin receptor.

Rabbit Fab 2 anti-mouse Fc fragment horse radish peroxidase conjugate (anti-MuFc HRP) is obtained from PelFreeze (cat. no. 715305-1).

10 D. ELISA Assay

Maxisorp microtiter plates are coated with 2 $\mu\text{g}/\text{ml}$ human vitronectin dissolved in PBS (50 ml/well) and stored overnight at 4°C. The plates are washed two times with PBS-0.05% Tween-20 (wash buffer) and blocked by incubating with about 150 $\mu\text{l}/\text{well}$ of assay buffer (1%, BSA [RIA grade or better] in 50 mM Tris-HCl, 100 mM NaCl, 1mM MgCl₂, CaCl₂, MnCl₂ pH 7.4) for 60 minutes. Dilutions of standards are prepared and putative inhibitors (Table 3) are dissolved in assay buffer. The blocked plates are emptied and 25 $\mu\text{l}/\text{well}$ of inhibitor or standard solution is added to each well. Twenty-five μl of a 30 $\mu\text{g}/\text{ml}$ solution of purified $\alpha_v\beta_3$ in assay buffer is pipetted into the coated plate. The final concentration of receptor in the assay well is about 15 $\mu\text{g}/\text{ml}$. The plate is incubated on a shaker for 60 minutes. Meanwhile, for each microtiter plate, 6 ml buffer solution containing 1.5 $\mu\text{g}/\text{ml}$ of mouse monoclonal antibody specific for β_3 is prepared. To this solution is added μl of the secondary antibody, which is anti-mouse-Fc-HRP antibody conjugate. For example, for one plate, prepare 6 ml of a 1.5 $\mu\text{g}/\text{ml}$ mouse Mab solution to which is added 1 μl of anti-mouse-Fc-HRP antibody stock, (this represents a 1:6000 dilution of the antibody - HRP conjugate). This mixture is allowed to incubate during the receptor-inhibitor incubation. The assay plates are washed 4 times with PBS-Tween and 50 $\mu\text{l}/\text{well}$ of the antibody mixture is then pipetted into the plate for a 60 minute incubation. The plate is washed 4 times and the color reaction is developed with 50 $\mu\text{l}/\text{well}$ of 0.67 mg/ml *o*-phenyldiamine in PBS containing 0.012% H₂O₂. Alternatively, 16 mM citric acid, 50 mM Na₂PO₄ at pH 5.0 can be used as a substrate buffer. The reaction is stopped with 50 $\mu\text{l}/\text{well}$ 1 M H₂SO₄. The plates are read at 492-405 nm and the data analyzed by four-parameter fit.

EXAMPLE 4

GP II_bIII_a-von Willebrand factor (vWF) ELISA Assay

35 A. ELISA Assay

Microtiter plates are coated with 1.0 $\mu\text{g}/\text{ml}$ GP II_bIII_a, prepared by the method of Fitzgerald *et al.*, (*Anal. Biochem.* 151: 169-177 [1985]) and allowed to incubate overnight in coat buffer. The plates are then washed three times in wash buffer (0.05% Tween 20 in PBS) and 150 μl of assay buffer is added and allowed to incubate for 1-2 hours at room temperature

on plate shaker. The plates are washed three times and 50 μ l of 2x inhibitor in assay buffer (Assay buffer: 0.5% BSA/50mM Tris, 100mM NaCl, 1.0mM CaCl₂, 1.0mM MgCl₂, 1.0mM MnCl₂; coat buffer is the same but without BSA) is added. Fifty μ l of 4.0 μ g/ml vWF (prepared as described by Ledford *et al.*, *Thrombosis and Haemostasis*, 64(4): 569-575 [1990]) in assay buffer is then added and allowed to incubate for one hour at room temperature on plate-shaker. The plates are washed three times and the antibody mixture is added (1:5000 of mouse anti-vWF and 1:5000 of rabbit-anti-mouse-Fc-HRP, both commercially available) in assay buffer and incubated for 1 hour at room temperature on plate-shaker. Plates are again washed three times and 100 μ l of substrate solution (10 mg OPD, 6.5 μ l H₂O₂, 15 ml phosphate citrate buffer) is added and incubated at room temperature. The color change of OPD/H₂O₂ reagent is read at 492 nm with a 405 nm reference wavelength on the filter photometer.

EXAMPLE 5

In Vitro Human Platelet Aggregation Assay

Platelet aggregation assays are performed in human platelet rich plasma (PRP). Fifty milliliters of whole human blood (9 parts) is drawn on 3.6% sodium citrate (1 part) from a donor who has not taken aspirin or related medications for at least two weeks. The blood is centrifuged at 160 x g for 10 minutes at 22°C and allowed to stand for 5 minutes after which the PRP is decanted. Platelet poor plasma (PPP) is isolated from the remaining blood after centrifugation at 2000 x g for 25 minutes. The platelet count of the PRP is diluted to about 300,000 platelets per microliter with PPP.

A 225 μ l aliquot of PRP plus 25 μ l of either a dilution of the test inhibitor sample or a control (PBS) is incubated for 5 minutes in a Chrono-log Whole Blood Aggregometer at 25°C. An aggregating agent (collagen, 1 μ g/ml; U46619, 100 ng/ml; or ADP, 17 μ M) is added and the transmission is recorded.

EXAMPLE 6

Human Fibronectin-Fibronectin Receptor ELISA Assay

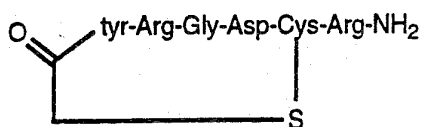
A. Fibronectin-Fibronectin Receptor Purification

Fibronectin receptor is purified according to the procedures of Pytela *et al.*, (*Methods Enzymol.* 144: 475 [1985]). Briefly, Fibronectin receptor is purified employing an Arg-Gly-Asp affinity chromatography from (100mM) octylglucoside (OG) extracted human placenta. The OG extract is filtered over sepharose 6B GRGDSPK column. The column is washed three times with three column volumes of Tris-buffered saline (TBS) TBS, 1mM CaCl₂, and 25mM OG. The receptor is eluted with TBS, 20mM EDTA, and 25mM Octyl thioglucoside, and is stored in 1mM CaCl₂, 1mM MgCl₂, at -80°C.

B. ELISA Assay

Microtiter plates are coated with 110 μ l human fibronectin (at 2 μ g/ml) in TBS. Plates are washed three times with TBS containing 0.05% Tween 20. Test inhibitors are then added after washing in 10 microliter aliquots. The fibronectin receptor is added in 2-fold

serial dilutions with TBS containing 20 mM octyl glucoside and 2 mM MnCl₂. Plates are incubated three hours at room temperature, and washed with 200 μ l TBS-Tween 20 buffer. 100 μ l of affinity-purified rabbit anti-human fibronectin receptor antibody is added to the wells and the plates are incubated two hours, washed twice with TBS-Tween and once with distilled water. Affinity-purified goat anti-rabbit IgG conjugated to horseradish peroxidase (100 μ l) is added to each well followed by incubation for 60 minutes at room temperature. Plates are again washed twice with TBS-Tween and distilled water. 100 μ l of substrate mixture (10 mg *o*-phenylenediamine in 25 ml 0.1 M citrate-phosphate buffer, pH 5.0 6 μ l 30% H₂O₂) is added to the plates and allowed to develop. The development process is stopped by adding 50 μ l of 4N H₂SO₄ to each well and read at 492/405 nM.

EXAMPLE 7cyclo-S-acetyl-dTyr-Arg-Gly-Asp-Cys-Arg-NH₂

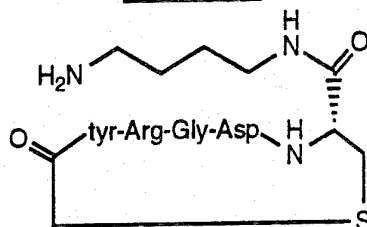
The title compound is prepared in protected form by standard solid phase peptide synthesis on 2% cross-linked *p*-methylbenzhydryl amine polystyrene resin (Merrifield resin). The peptide is prepared by sequential addition of suitably protected amino acid residues from the resin bound arginine to the *d*-tyrosine residue. After the amino protecting group of the *d*-tyrosine residue has been removed, bromoacetic acid is coupled with diisopropylcarbodiimide. Treatment of the resin bound intermediate with liquid hydrogen fluoride induces concomitant cleavage of the protecting groups from the title compound as well as cleavage of the peptide from the resin. The crude peptide is dissolved in deionized water (1mg/ml) and the pH of the solution is adjusted to 8.0 - 8.5 with ammonium hydroxide to effect cyclization. After stirring for 4 hr at ambient temperature the reaction solution is acidified to pH 3.0 - 3.5 with trifluoroacetic acid and then lyophilized. The resulting crude cyclic product is purified by reverse phase high performance liquid chromatography (HPLC) using a 4.6 mm x 250 mm column containing 10 micron, 300 Angstrom pore size C-18 packing. The elution of the column is with an acetonitrile/0.1% aqueous trifluoroacetic acid gradient going from 0% - 40% acetonitrile linearly over 80 minutes. The title compound elutes at 19 minutes. FAB mass spectrum: calc. 807.2; obs. 808.3 (M+1).

Except for compound 1, the following compounds were prepared by analogous procedures.

<u>Compound No.</u>	<u>Sequence</u>
1	cyclo-S-acetyl-dTyr-Arg-Gly-Asp-Cys-Arg-NH ₂
2	cyclo-S-acetyl-dTyr--Arg-Gly-Asp-Cys-dArg-NH ₂
3	cyclo-S-acetyl-dTyr--Arg-Gly-Asp-Cys-Lys-NH ₂
4	cyclo-S-acetyl-dTyr--Arg-Gly-Asp-Cys-dLys-NH ₂
5	cyclo-S-acetyl-dTyr--Arg-Gly-Asp-Cys-Orn-NH ₂

	6	cyclo-S-acetyl-dTyr--Arg-Gly-Asp-Cys-dOrn-NH ₂	
	7	cyclo-S-acetyl-dTyr--Arg-Gly-Asp-Cys-Tyr-Arg-NH ₂	
	16	cyclo-S-acetyl-dTyr-Lys-Gly-Asp-Pen-Arg-NH ₂	
	17	cyclo-S-acetyl-dTyr-Lys-Gly-Asp-Pen-Lys-NH ₂	
5	24	cyclo-S-acetyl-dTyr-Arg-Gly-Asp-Cys-Arg-OH	
	25	cyclo-S-acetyl-dTyr-Arg-Gly-Asp-Cys-Lys-OH	
	26	cyclo-S-acetyl-dTyr-Lys-Gly-Asp-Pen-Arg-OH	
	27	cyclo-S-acetyl-dTyr-Arg-Gly-Asp-Cys-Gln-NH ₂	
	28	cyclo-S-(1-5)-dTyr-Arg-Gly-Asp-Cys-X (X=aminovaleric acid)	
10	29	cyclo-S-(1-5)-dTyr-Arg-Gly-Asp-Cys-X (X=aminovaleric amide)	
	30	cyclo-S-acetyl-Tyr-Arg-Gly-Asp-Cys-Arg-NH ₂	
	31	cyclo-S-acetyl-dTyr-Arg-Gly-Asp-Cys-Arg-NH ₂	Sulfoxide (isomer 1)
	32	cyclo-S-acetyl-dTyr-Arg-Gly-Asp-Cys-Arg-NH ₂	Sulfoxide (isomer 2)
	33	cyclo-S-acetyl-dTyr-Arg-Gly-Asp-Cys-dArg-NH ₂	Sulfoxide (isomer 1)
15	34	cyclo-S-acetyl-dTyr-Arg-Gly-Asp-Cys-dArg-NH ₂	Sulfoxide (isomer 2)
	35	cyclo-S-acetyl-dTyr-Arg-Gly-Asp-Cys-Lys-NH ₂	Sulfoxide (isomer 1)
	36	cyclo-S-acetyl-dTyr-Arg-Gly-Asp-Cys-Lys-NH ₂	Sulfoxide (isomer 2)
	37	cyclo-S-acetyl-dTyr-Arg-Gly-Asp-Cys-dLys-NH ₂	Sulfoxide (isomer 1)
	38	cyclo-S-acetyl-dTyr-Arg-Gly-Asp-Cys-dLys-NH ₂	Sulfoxide (isomer 2)
20	39	cyclo-S-acetyl-dTyr-Arg-Gly-Asp-Cys-Orn-NH ₂	Sulfoxide (isomer 1)
	40	cyclo-S-acetyl-dTyr-Arg-Gly-Asp-Cys-Orn-NH ₂	Sulfoxide (isomer 2)
	41	cyclo-S-acetyl-dTyr-Arg-Gly-Asp-Cys-dOrn-NH ₂	Sulfoxide (isomer 1)
	42	cyclo-S-acetyl-dTyr-Arg-Gly-Asp-Cys-dOrn-NH ₂	Sulfoxide (isomer 2)
	43	cyclo-S-acetyl-dTyr-Arg-Gly-Asp-Cys-Tyr-Arg-NH ₂	Sulfoxide (isomer 1)
25	44	cyclo-S-acetyl-dTyr-Arg-Gly-Asp-Cys-Tyr-Arg-NH ₂	Sulfoxide (isomer 2)
	45	cyclo-S-phenylacetyl-dTyr-Lys-Gly-Asp-Cys-OH	
	46	cyclo-S-acetyl-dTyr-Arg-Gly-Asp-Cys-OH	
	47	cyclo-S-acetyl-dTyr-Arg-Gly-Asp-Pen-OH	
	48	Cyclo-S-acetyl-dVal-Arg-Gly-Asp-Cys-Lys-NH ₂	sulfoxide (isomer 1)
30	49	Cyclo-S-acetyl-dVal-Arg-Gly-Asp-Cys-Lys-NH ₂	sulfoxide (isomer 2)
	50	Cyclo-S-acetyl-dVal-Arg-Gly-Asp-Cys-Arg-NH ₂	sulfoxide (isomer 1)
	51	Cyclo-S-acetyl-dVal-Arg-Gly-Asp-Cys-Arg-NH ₂	sulfoxide (isomer 2)
	52	Cyclo-S-acetyl-dVal-Lys-Gly-Asp-Cys-Lys-NH ₂	sulfoxide (isomer 1)
	53	Cyclo-S-acetyl-dVal-Lys-Gly-Asp-Cys-Lys-NH ₂	sulfoxide (isomer 2)
35	54	Cyclo-S-acetyl-dVal-Arg-Gly-Asp-Cys-NH-X (X=butylamine)	
		sulfoxide (isomer 1)	
	55	Cyclo-S-acetyl-dVal-Arg-Gly-Asp-Cys-NH-X (X=butylamine)	
		sulfoxide (isomer 2)	
	56	Cyclo-S-acetyl-dVal-Arg-Gly-Asp-Cys-NH-X (X=butylguanidine)	

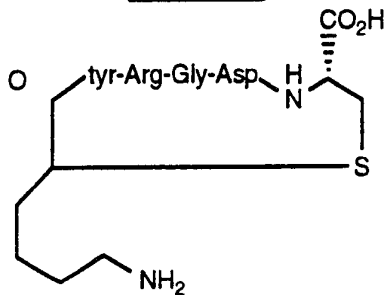
		sulfoxide (isomer 1)	
57	Cyclo-S-acetyl-dVal-Arg-Gly-Asp-Cys-NH-X (X=butylguanidine)		
		sulfoxide (isomer 2)	
58	Cyclo-S-acetyl-dTyr-Lys-Gly-Asp-Cys-Lys-NH ₂	sulfoxide (isomer 1)	
5	59 Cyclo-S-acetyl-dTyr-Lys-Gly-Asp-Cys-Lys-NH ₂	sulfoxide (isomer 2)	
60	Cyclo-S-(phenylacetyl)-dTyr-Lys-Gly-Asp-Cys-Lys-NH ₂	sulfoxide (isomer 1)	
61	Cyclo-S-(phenylacetyl)-dTyr-Lys-Gly-Asp-Cys-Lys-NH ₂	sulfoxide (isomer 2)	
62	Cyclo-S-acetyl-dPro-Arg-Gly-Asp-Cys-Arg-NH ₂	sulfoxide (isomer 1)	
63	Cyclo-S-acetyl-dPro-Arg-Gly-Asp-Cys-Arg-NH ₂	sulfoxide (isomer 2)	
10	64 Cyclo-S-acetyl-dPro-Arg-Gly-Asp-Cys-Lys-NH ₂	sulfoxide (isomer 1)	
65	Cyclo-S-acetyl-dPro-Arg-Gly-Asp-Cys-Lys-NH ₂	sulfoxide (isomer 2)	

EXAMPLE 8cyclo-S-acetyl-dTyr-Arg-Gly-Asp-Cys-NH-(CH₂)₄-NH₂

- 15 The title compound is prepared in protected form by standard solid phase peptide synthesis. Hydroxymethyl methyl polystyrene is treated with 20% phosgene in toluene for one hour. The resin is then washed six times with toluene. Excess 1,4 diaminobutane is added in toluene and mixed with the resin for 2 hours. The resin is washed with toluene followed by dimethylacetamide. The protected cysteine residue is coupled to the amine on the resin with
- 20 diisopropylcarbodiimide. Standard solid phase techniques are used for the addition of the subsequent amino acids. After the amino protecting group of the d-Tyrosine residue has been removed bromoacetic acid is coupled with diisopropylcarbodiimide. Treatment of the resin bound intermediate with liquid hydrogen fluoride induces concomitant cleavage of the protecting groups from the title compound as well as cleavage of the peptide from the resin.
- 25 The crude peptide is dissolved in deionized water (1mg/ml) and the pH of the solution is adjusted to 8.0 - 8.5 with ammonium hydroxide. After stirring for 4 hr at ambient temperature the reaction solution is acidified to pH 3.0 - 3.5 with trifluoroacetic acid and then lyophilized. The resulting crude cyclic product is purified by reverse phase high performance
- 30 liquid chromatography (HPLC) using a 4.6 mm x 250 mm column containing 10 micron, 300 Angstrom pore size C-18 packing. The elution of the column is with an acetonitrile/0.1% aqueous trifluoroacetic acid gradient going from 0% - 40% acetonitrile linearly over 80 minutes. The title compound elutes at 18 minutes. FAB mass spectrum: calc. 722.2; obs. 723.2(M+1).

35 Except for compound 12, the following compounds were prepared by analogous procedures:

<u>Compound No.</u>	<u>Sequence</u>	
12	cyclo-S-acetyl-dTyr-Arg-Gly-Asp-Cys-NH-X	(X=butylamine)
13	cyclo-S-acetyl-dTyr-Arg-Gly-Asp-Cys-NH-X	(X=pentylamine)
14	cyclo-S-acetyl-dTyr-Arg-Gly-Asp-Cys-NH-X	(X=hexyl amine)
5 15	cyclo-S-acetyl-dTyr-Lys-Gly-Asp-Cys-NH-X	(X=butyl amine)
18	cyclo-S-acetyl-dTyr-Lys-Gly-Asp-Pen-NH-X	(X=butyl amine)
22	cyclo-S-acetyl-dTyr-Arg-Gly-Asp-Cys-NH-X	(X=propyl amine)
23	cyclo-S-acetyl-dTyr-Arg-Gly-Asp-Cys-NH-X	(X=ethyl amine)

EXAMPLE 9

10

cyclo-S-2-(6-aminocaproyl)dTyr-Arg-Gly-Asp-Cys-OH

The title compound is prepared in protected form by standard solid phase peptide synthesis on 2% cross-linked polystyrene resin (Merrifield resin). After removal of the t-butyloxycarbonyl group from the d-tyrosine residue and subsequent neutralization, the 2-bromo 6-(2-chlorobenzyloxycarbonyl) aminocaproic acid, synthesized from ε-(2-chloro-CBZ)-L-lysine, was coupled using diisopropylcarbodiimide. Treatment of the resin bound intermediate with liquid hydrogen fluoride induces concomitant cleavage of the protecting groups from the title compound as well as cleavage of the peptide from the resin. The crude peptide is dissolved in deionized water (1mg/ml) and the pH of the solution is adjusted to 8.0 - 8.5 with ammonium hydroxide. After stirring for 4 hr at ambient temperature the reaction solution is acidified to pH 3.0 - 3.5 with trifluoroacetic acid and then lyophilized. The resulting crude cyclic product is purified by reverse phase high performance liquid chromatography (HPLC) using a 4.6 mm x 250 mm column containing 10 micron, 300 Angstrom pore size C-18 packing. The elution of the column is with an acetonitrile/0.1% aqueous trifluoroacetic acid gradient going from 0% - 40% acetonitrile linearly over 80 minutes. The title compound elutes at 14 minutes. FAB mass spectrum: calc. 723.2; obs. 724.2 (M+1).

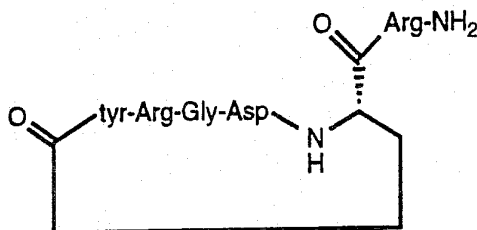
25

Except for compounds 10 and 11, the following compounds were prepared by analogous procedures:

<u>Compound No.</u>	<u>Sequence</u>	
30 8	cyclo-S-2-L-(5-guanidinovaleryl)-dTyr-Arg-Gly-Asp-Cys-OH	
9	cyclo-S-2-D-(5-guanidinovaleryl)-dTyr-Arg-Gly-Asp-Cys-OH	
10	cyclo-S-2-L-(6-aminocaproyl)-Lys-dTyr-Arg-Gly-Asp-Cys-OH	(isomer 1)
11	cyclo-S-2-L-(6-aminocaproyl)-Lys-dTyr-Arg-Gly-Asp-Cys-OH	(isomer 2)

EXAMPLE 10

50



cyclo(1-5)dTyr-Arg-Gly-Asp-L-α-aminoadipic acid-Arg-NH₂

The title compound is prepared in protected form by standard solid phase peptide synthesis on 2% cross-linked para methylbenzhydryl amine polystyrene resin (Merrifield resin). The amino adipic acid residue is incorporated as the N-Boc-δ-allyl-α-aminoadipic acid derivative. After deprotection and neutralization at the d-tyrosine residue the allyl group is removed and the peptide is cyclized with BOP. Treatment of the resin bound intermediate with liquid hydrogen fluoride induces concomitant cleavage of the protecting groups from the title compound as well as cleavage of the peptide from the resin. The resulting crude cyclic product is purified by reverse phase high performance liquid chromatography (HPLC) using a 4.6 mm x 250 mm column containing 10 micron, 300 Angstrom pore size C-18 packing. The elution of the column is with an acetonitrile/0.1% aqueous trifluoroacetic acid gradient going from 0% - 40% acetonitrile linearly over 80 minutes. The title compound elutes at 14 minutes. FAB mass spectrum: calc. 789.3; obs. 790.3 (M+1).

Except for compound 19, the following compounds were prepared by analogous procedures:

<u>Compound No.</u>	<u>Sequence</u>
19	cyclo(1-5)dTyr-Arg-Gly-Asp-L-α-aminoadipic acid-Arg-NH ₂
20	cyclo(1-5)dTyr-Arg-Gly-Asp-L-α-aminoadipic acid-Lys-NH ₂
20	21 cyclo(1-5)dTyr-Arg-Gly-Asp-L-α-aminoadipic-butylamine

EXAMPLE 11

Bleeding Time and EX-VIVO Platelet Aggregation - Dog Model

A. Animal preparation

Beagle dogs were premedicated with atropine (3 mg subcutaneously) and anesthetized with intravenous sodium pentobarbital solution (65mg/ml, to effect, induction and maintenance). The dogs were placed in dorsal recumbency on heating pads, intubated, and placed on oxygen-room air mixture (free breathing). A pulse oximeter was attached and heart rate and oxygen saturation monitored. An indwelling catheter was placed in one cephalic vein through which normal saline drip (for patency) and maintenance anesthetic doses were administered, and in one saphenous vein through which test or control compound was bolused or infused.

At each proscribed time point a 5 ml whole blood sample was withdrawn via jugular venipuncture for *ex vivo* platelet aggregation and complete blood cell count. Manual pressure was applied to the venipuncture site following blood withdrawal to minimize hematoma formation.

At one "pre" time point, 14ml of blood was withdrawn in order to conduct *in vitro* dose-response experiments.

At several timepoints bleeding times were measured. The forearm of the dog was closely shaven. A Surgicutt® automatic bleeding time device was employed to make an
 5 approximate 1cm clean incision through the skin. Blood was blotted with the edge of a Whatman's #4 filter paper every 15 seconds. Bleeding time was measured as the time from incision to the last time blood was absorbed by the filter paper.

B. Handling of blood samples

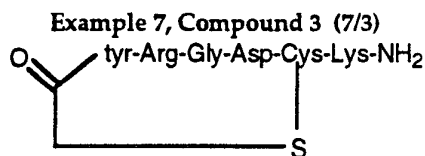
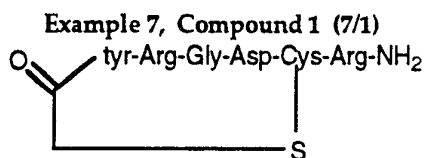
Approximately one ml of blood withdrawn at each timepoint was placed on EDTA
 10 (purple-top vacutainer) and used to measure the complete blood cell count on a Cell-Dyne 1500 automated hematology analyzer. 3.6ml of blood withdrawn at each timepoint was placed on a 3.8% sodium citrate (1 part to 9 parts blood) and centrifuged at 12,000rpm for 4 seconds in an Eppendorf microcentrifuge. The resultant platelet rich plasma (PRP) was pipetted off for use in *ex vivo* aggregation procedures.

15 C. Ex Vivo platelet aggregation

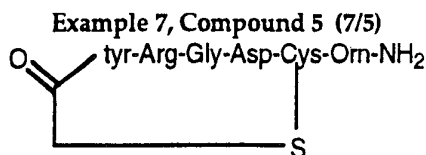
Platelet aggregation was measured in a Payton dual-channel aggregometer. The equipment was calibrated with PRP assigned as 0% light transmission and PPP as 100% light transmission. PRP was allowed to rest at room temperature for 20 minutes following separation from whole blood. 300 µl PRP was incubated at 37°C in the aggregometer for
 20 approximately 5 minutes with the chart recorder on to establish a baseline. 10µl ADP (1mM) was added to the PRP and the aggregation response (change in light transmission) recorded. An aliquot of the PRP was used to perform a platelet count on the Cell-Dyne 1500 automated hematology analyzer.

D. Results

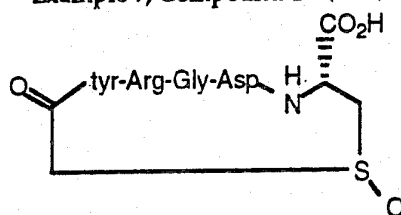
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30



Example 7, Compound 46 (7/46)

**Example 7, Compound 1**

5 Compared to 7/46, 7/1 was equipotent at inhibiting human platelet aggregation *in vitro*. The effect on platelet aggregation in the dog of 7/46, i.e., a bolus of 125 $\mu\text{g}/\text{kg}$ plus an infusion of 3 $\mu\text{g}/\text{kg}/\text{min}$ gave 100% inhibition of *ex vivo* aggregation throughout the infusion period. At this dose, 7/46 prolonged bleeding time to >30 minutes during the infusion. With 7/1, bleeding time was increased from preinfusion value of 3 minutes to a mean of 6.25 minutes during the infusion. This effect is no different to that seen with a normal saline infusion.

10 **Example 7, Compound 3**

At a dose of 300 $\mu\text{g}/\text{kg}$ bolus plus an infusion of 7.2 $\mu\text{g}/\text{kg}/\text{min}$ this peptide inhibited *ex vivo* aggregation >95% without prolongation of the bleeding time. Based on potency *in vitro*, this was a dose equivalent to 75 $\mu\text{g}/\text{kg}$ bolus plus an infusion of 2 $\mu\text{g}/\text{kg}/\text{min}$ for compound D which gave a prolongation of bleeding.

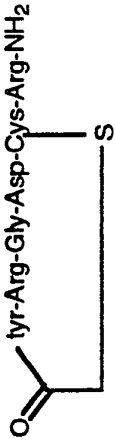

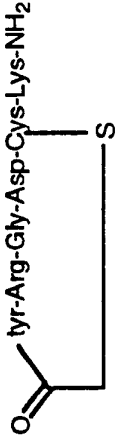
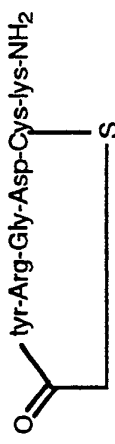
15 **Example 7, Compound 5**

At a dose of 200 $\mu\text{g}/\text{kg}$ bolus plus an infusion of 4.8 $\mu\text{g}/\text{kg}/\text{min}$, gave >90% inhibition of aggregation with minimal prolongation of bleeding.

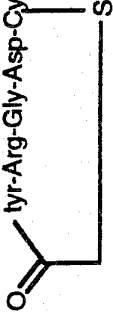
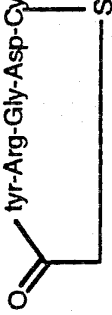
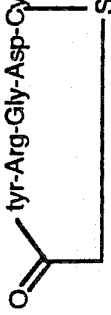
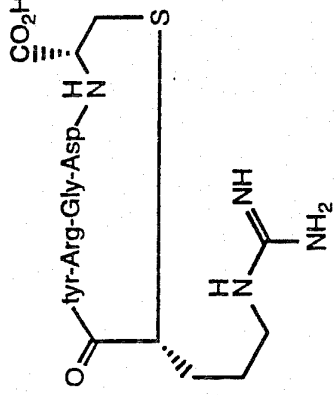
EXAMPLE 12

In Vivo and In Vitro Assay Results For Platelet Aggregation Inhibitors

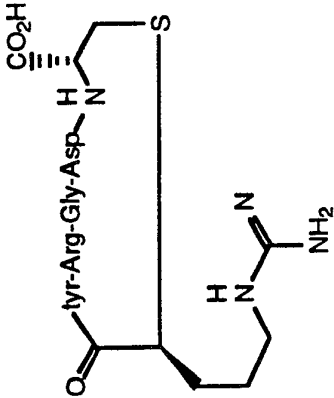
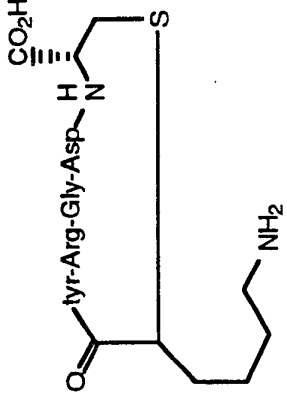
-----ELISA IC₅₀ nM-----

Compound ¹ Ex(#)/Cpd.(#)	Sequence ²	Fibrinogen ³ GPIIb/IIIa	Vitronectin ⁴ VnR	GPIIb/IIIa ⁵ vWF	Fibronectin ⁶ FnR	Human ⁷ Platelet Aggregation IC ₅₀ nM	Rabbit ⁸ BT Ratio (%P.L.A. Inhibition)
7 / 1		10	49	13.8	57	80	.71(100)
7 / 2		13.5	22	2.93	42		
7 / 3		17	60	15.9	201	320	1.95(97)
7 / 4		8.3	17	1.56	66		

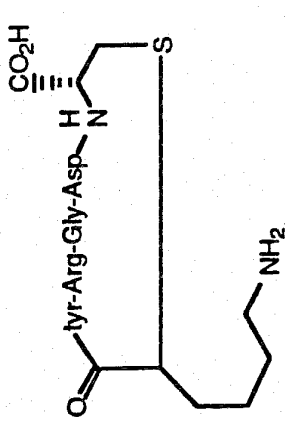
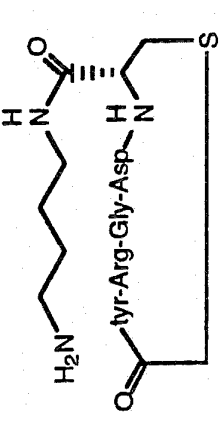
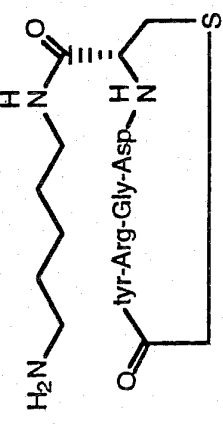
-----ELISA IC₅₀ nM-----

Compound 1 Ex(#)/Cpd.(#)	Sequence ²	Fibrinogen ³ GPIIb,IIIa	Vitronectin ⁴ VnR	GPIIb,IIIa ⁵ vWF	Fibronectin ⁶ FnR	Human ⁷ Platelet Aggregation IC ₅₀ nM	Rabbit ⁸ BT Ratio (%P.A. Inhibition)
7 / 5		11	45	6.84	63	190	1.26(100)
7 / 6		10.5	55	1.92	161	1,176	1.4(53)
7 / 7		8.9	51		27	320	4.10(98)
9 / 8		8.4	7	2.45	216	790	1.46(55)

-----ELISA IC₅₀ nM-----

Compound 1 Ex(#)/Cpd.(#)	Sequence ²	Fibrinogen ³ GPIIb/IIIa	Vitronectin ⁴ VnR	GPIIb/IIIa ⁵ vWF	Fibronectin ⁶ FnR	Human ⁷ Platelet Aggregation IC ₅₀ nM	Rabbit ⁸ BT Ratio (%P.A. Inhibition)
9 / 9		50	17	13.8	273	6,300	1.0(1)
9 / 10		5.3	9.6	0.785	2	498	8.17(100)

-----ELISA IC₅₀ nM-----

Compound ¹ Ex(#)/Cpd.(#)	Sequence ²	Fibrinogen ³ GPIIb/IIIa	Vitronectin ⁴ VnR	GPIIb/IIIa ⁵ vWF	Fibronectin ⁶ FnR	Human ⁷ Platelet Aggregation IC ₅₀ nM	Rabbit ⁸ BT Ratio (%P.A. Inhibition)
9 / 11		14.3	11.2	6.95	160	3,177	
8 / 12		10	70	5.40	920	390	0.94(96)
8 / 13		9	32	5.30	473	590	1.13(98)

-----ELISA IC₅₀ nM-----

Human⁷
Platelet
Aggregation
IC₅₀nM

Rabbit⁸
BT Ratio
(%P.A. Inhibition)

Fibrinogen³
GPII_bIII_a

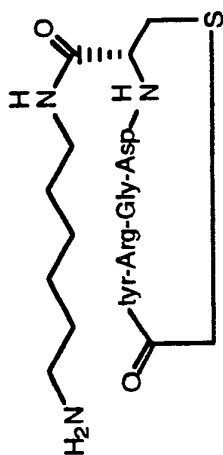
Vitronectin⁴
VnR

GPII_bIII_a
vWF

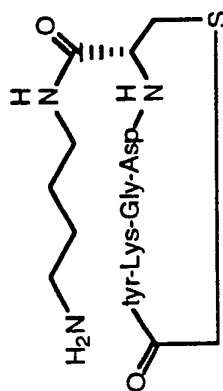
Fibronectin⁶
FnR

Compound 1
Ex(#)/Cpd.(#)

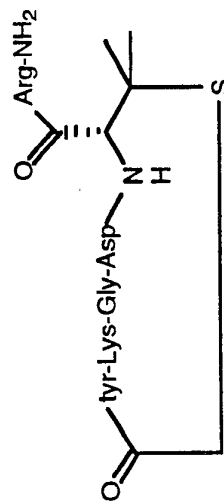
8 / 14



8 / 15



7 / 16



8.29(100)

506

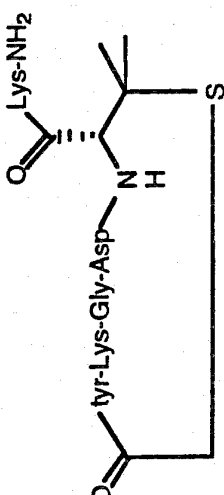
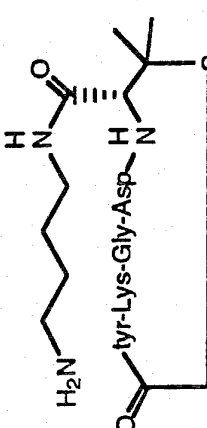
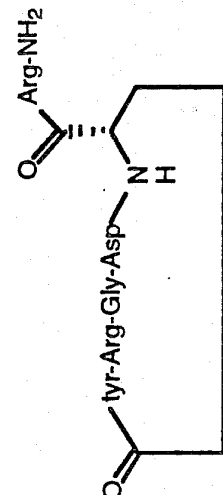
>10,000

53.5

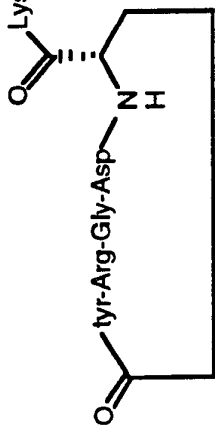
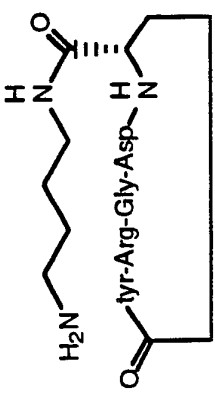
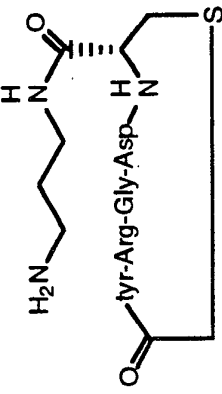
120,000

150

-----ELISA IC₅₀ nM-----

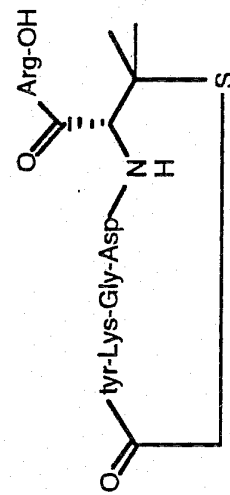
Compound ¹ Ex(#)/Cpd.(#)	Sequence ²	Fibrinogen ³ GPIIb/IIIa	Vitronectin ⁴ VnR	GPIIb/IIIa ⁵ vWF	Fibronectin ⁶ FnR	Human ⁷ Platelet Aggregation IC ₅₀ nM	Rabbit ⁸ BT Ratio (%P.A. Inhibition)
7 / 17		238	100,000	73.1	>10,000	1160	1.13(100) 1.0(100)
8 / 18							
10 / 19		8.9	32	2.90	4400	180 205	7.98(98)

-----ELISA IC₅₀ nM-----

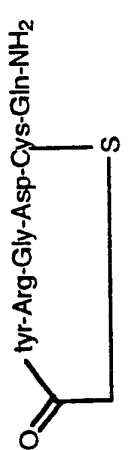
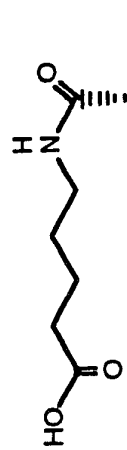
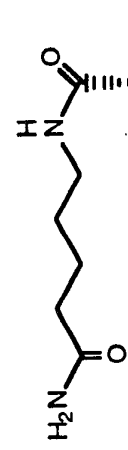
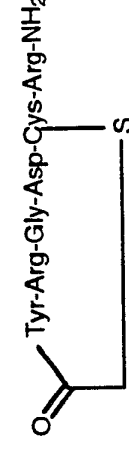
Compound 1 Ex(#)/C.pd.(#)	Sequence ²	Fibrinogen ³ GPIIb/IIIa	Vitronectin ⁴ VnR	GPIIb/IIIa ⁵ vWF	Fibronectin ⁶ FnR	Human ⁷ Platelet Aggregation IC ₅₀ nM	Rabbit ⁸ BT Ratio (%P.A. Inhibition)
10 / 20		12.4	20	2.74	2,310	183	10(99)
10 / 21		3.4				150	
8 / 22		15	36	4.73	176	387	1.27(99)

-----ELISA IC₅₀ nM-----

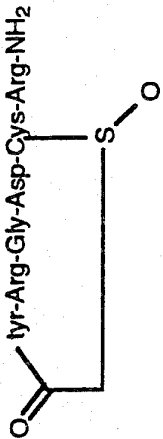
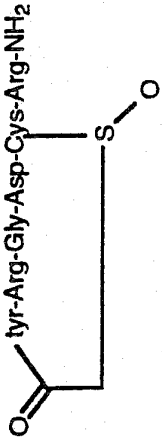
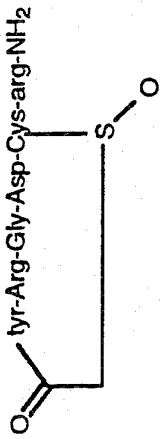
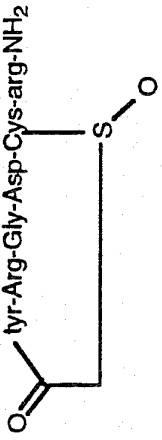
Compound 1 Ex(#) /Cpd.(#)	Sequence ²	Fibrinogen ³ GPIIb/IIIa	Vitronectin ⁴ VnR	GPIIb/IIIa ⁵ vWF	Fibronectin ⁶ FnR	Human ⁷ Platelet Aggregation IC ₅₀ nM	Rabbit ⁸ BT Ratio (%P.A. Inhibition)
8 / 23		14	35	4.67	61	470	0.81(88)
7 / 24		4.4	210	3.80	1130	240	
7 / 25		7.4	385	8.90		470	6.14(100)
7 / 26							



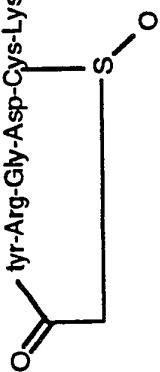
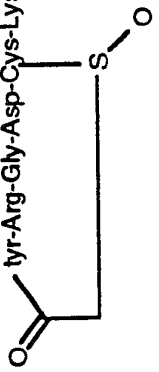
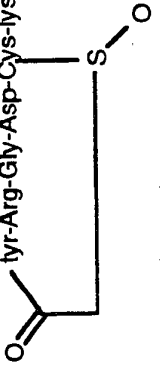
-----ELISA IC₅₀ nM-----

Compound 1 Ex(#)/Cpd.(#)	Sequence ²	Fibrinogen ³ GPIIb/IIIa	Vitronectin ⁴ VnR	GPIIb/IIIa ⁵ vWE	Fibronectin ⁶ FnR	Human ⁷ Platelet Aggregation IC ₅₀ nM	Rabbit ⁸ BT Ratio (%P.A. Inhibition)
7 / 27		3.4	327	2.20		1760	0.79(72)
7 / 28		70.6	1670	64.00		2.0	1.75(81)
7 / 29						1670	0.86(77)
7 / 30		370	1118	183.00		3780	

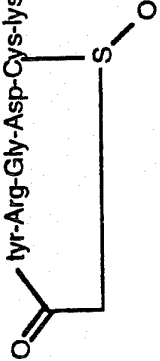
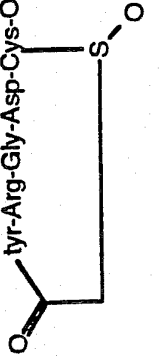
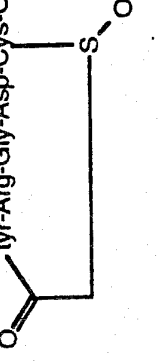
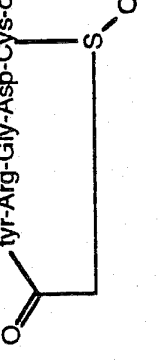
-----ELISA IC₅₀ nM-----

Compound ¹ Ex(#)/Cpd.(#)	Sequence ²	Fibrinogen ³ GPIIb/IIIa	Vitronectin ⁴ VnR	GPIIb/IIIa ⁵ vWF	Fibronectin ⁶ FnR	Human ⁷ Platelet Aggregation IC ₅₀ nM	Rabbit ⁸ BT Ratio (%P.A. Inhibition)
7 / 31		9.5	39	2.04	330	230	
7 / 32		55	82	59.3	386	750	1.17(99)
7 / 33		6.8	22	0.994	470	500	
7 / 34		48	57	33.6	409	4,900	0.94(64)

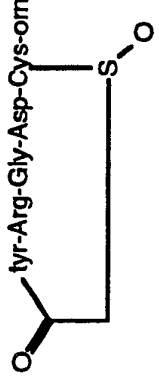
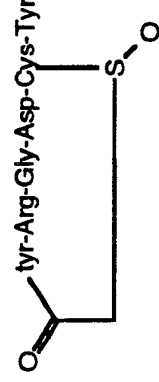
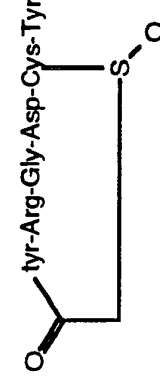
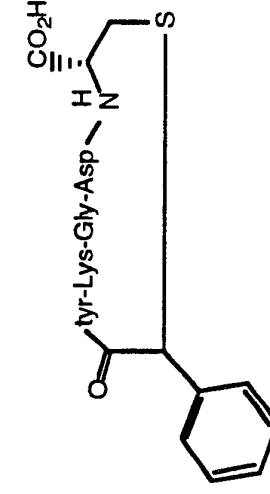
-----ELISA IC₅₀ nM-----

Compound 1 Ex(#) /Cpd.(#)	Sequence ²	Fibrinogen ³ GPIIb/IIIa	Vitronectin ⁴ VnK	GPIIb/IIIa ⁵ vWF	Fibronectin ⁶ FnK	Human ⁷ Platelet Aggregation IC ₅₀ nM	Rabbit ⁸ BT Ratio (%P.A. Inhibition)
7 / 35	 <p>tyr-Arg-Gly-Asp-Cys-Lys-NH₂</p> <p>3-bromoyl tyrosine</p>	16.5	27	6.11	1,070		
7 / 36	 <p>tyr-Arg-Gly-Asp-Cys-Lys-NH₂</p> <p>3-bromoyl tyrosine</p>	108	78	45.6	524		0.77(91)
7 / 37	 <p>tyr-Arg-Gly-Asp-Cys-lys-NH₂</p> <p>3-bromoyl tyrosine</p>	6.3	38	1.82	994		

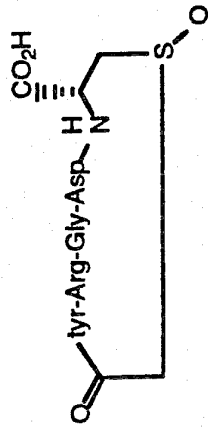
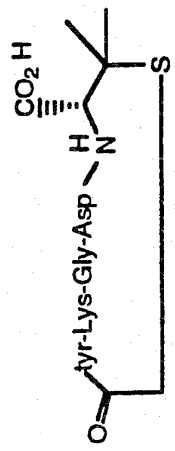
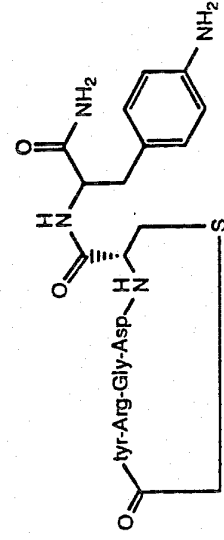
-----ELISA IC₅₀ nM-----

Compound 1 Ex(#)/Cpd.(#)	Sequence ²	Fibrinogen ³ GPIIb/IIIa	Vitronectin ⁴ VnR	GPIIb/IIIa ⁵ vWF	Fibronectin ⁶ FnR	Human ⁷ Platelet Aggregation IC ₅₀ nM	Rabbit ⁸ BT Ratio (%P.A. Inhibition)
7 / 38		69	65	47.3	615		
	3-bromyl tyrosine						
7 / 39		5.3	64		3.27	93	
7 / 40		43	300		67.7	1.950	1.09(71)
7 / 41		5.3	42		1.21	340	1.50(98)

-----ELISA IC₅₀ nM-----

Compound ¹ Ex(#)/Cpd.(#)	Sequence ²	Fibrinogen ³ GPIIb/IIIa	Vitronectin ⁴ VnR	GPIIb/IIIa ⁵ vWF	Fibronectin ⁶ FnR	Human ⁷ Platelet Aggregation IC ₅₀ nM	Rabbit ⁸ BT Ratio (%P.A. Inhibition)
7 / 42		23	153		42.1	6,680	1.14(41)
7 / 43						350	
7 / 44		94	160		60.3	8,470	1.18(52)
7 / 45		>20,000	>20,000	1	>10,000	220	4.47(94)

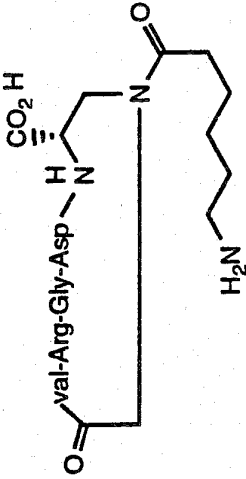
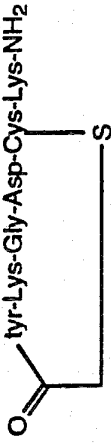
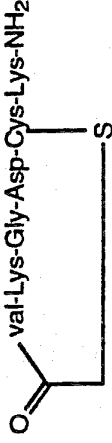
-----ELISA IC₅₀ nM-----

Compound 1 Ex(#)/Cpd.(#)	Sequence ²	Fibrinogen ³ GPIIb/IIIa	Vitronectin ⁴ VnR	GPIIb/IIIa ⁵ vWF	Fibronectin ⁶ FnR	Human ⁷ Platelet Aggregation IC ₅₀ nM	Rabbit ⁸ BT Ratio (%P.A. Inhibition)
7 / 46		3.2	5	0.18	3,390	150	8.77(99) 2.74(55)
7 / 47		4.2	>20,000	10.3	>10,000	486	8.57(96)
		5	154	1.60		2000	

-----ELISA IC₅₀ nM-----

Compound 1 Ex(#)/Cpd.(#)	Sequence ²	Fibrinogen ³ GPIIb/IIIa	Vitronectin ⁴ VnR	GPIIb/IIIa ⁵ vWF	Fibronectin ⁶ FnR	Human ⁷ Platelet Aggregation IC ₅₀ nM	Rabbit ⁸ BT Ratio (%P.A. Inhibition)
	<chem>NC(=O)C[C@H](N)C(=O)C[C@H](N)C(=O)C[C@H](N)C(=O)C[C@H](N)C(=O)N</chem>	8.6	417	6.40	280	240	
	<chem>NC(=O)C[C@H](N)C(=O)C[C@H](N)C(=O)C[C@H](N)C(=O)C[C@H](N)C(=O)N</chem>	18	1158	16.00		590	
	<chem>NC(=O)C[C@H](N)C(=O)C[C@H](N)C(=O)C[C@H](N)C(=O)C[C@H](N)C(=O)N</chem>				2410	760	
	<chem>NC(=O)C[C@H](N)C(=O)C[C@H](N)C(=O)C[C@H](N)C(=O)C[C@H](N)C(=O)N</chem>						

-----ELISA IC₅₀ nM-----

Compound 1 Ex(#)/Cpd.(#)	Sequence ²	Fibrinogen ³ GPIIb/IIIa	Vitronectin ⁴ VnR	GPIIb/IIIa ⁵ vWF	Fibronectin ⁶ FnR	Human ⁷ Platelet Aggregation IC ₅₀ nM	Rabbit ⁸ BT Ratio (%P.A. Inhibition)
		52	100000	97.00		1290	
		177	100000	231.00		7800	
		8100	100000	7548			

-----ELISA IC₅₀ nM-----

Human⁷
Platelet
Aggregation
IC₅₀nM

Rabbit⁸
BT Ratio
(%P.A. Inhibition)

Fibronectin⁶
FnR

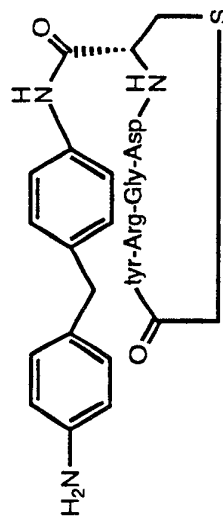
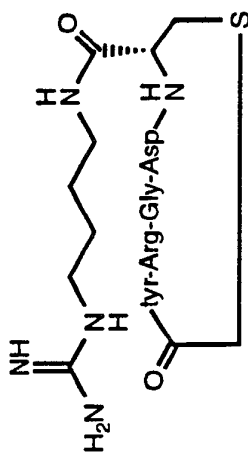
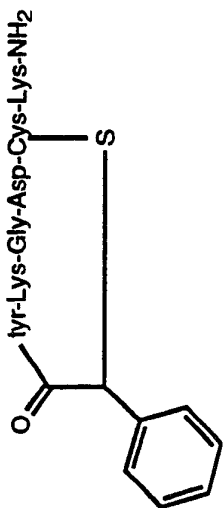
GPIIb,IIIa⁵
vWF

Vitronectin⁴
VnR

Fibrinogen³
GPIIb,IIIa

Sequence²

Compound¹
Ex(#)/Cpd.(#)



5.4

94

7.40

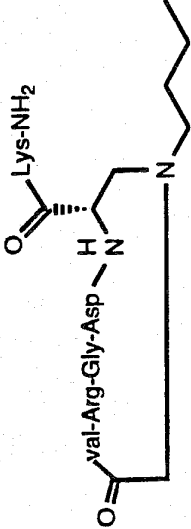
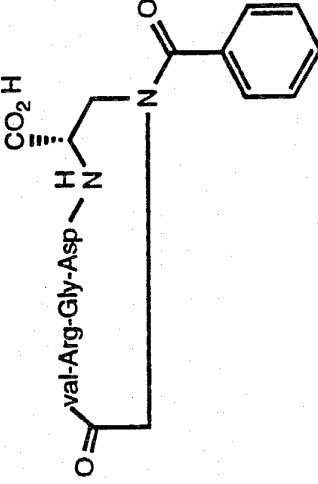
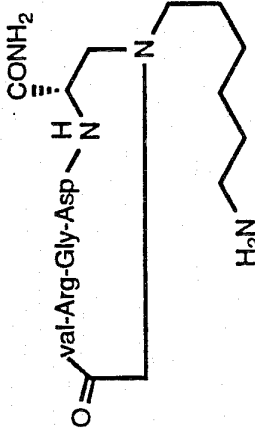
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260

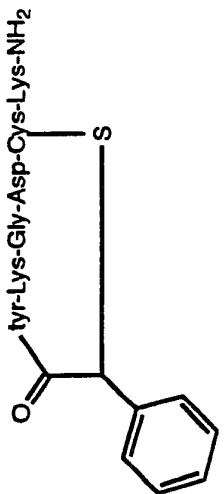
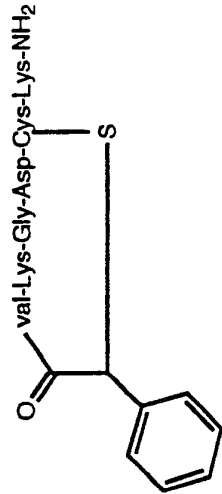
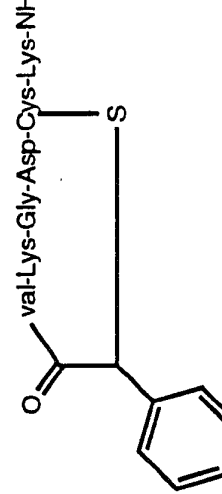
6.26

980

-----ELISA IC₅₀ nM-----

Compound 1 Ex(#)/Cpd.(#)	Sequence ²	Fibrinogen ³ GPIIb/III _a	Vitronectin ⁴ VnR	GPIIb/III _a ⁵ vWF	Fibronectin ⁶ FnR	Human ⁷ Platelet Aggregation IC ₅₀ nM	Rabbit ⁸ BT Ratio (%P.A. Inhibition)
		284	3400	424.00		3120	
		5.6	44	2.00		840	
		80	516			10900	

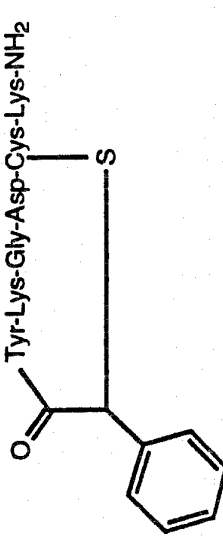
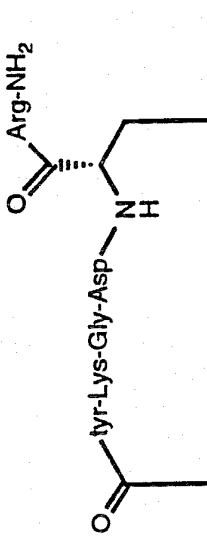
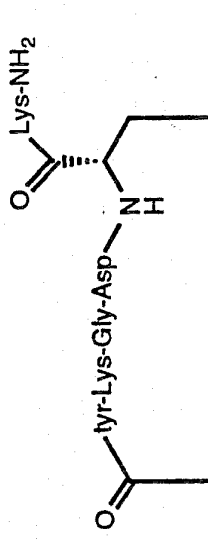
-----ELISA IC₅₀ nM-----

Compound ¹ Ex(#)/Cpd.(#)	Sequence ²	Fibrinogen ³ GPIIb/IIIa	Vitronectin ⁴ VnR	GPIIb/IIIa ⁵ vWF	Fibronectin ⁶ FnR	Human ⁷ Platelet Aggregation IC ₅₀ nM	Rabbit ⁸ BT Ratio (%P.A. Inhibition)
						980	
						12350	
						850	

120

15

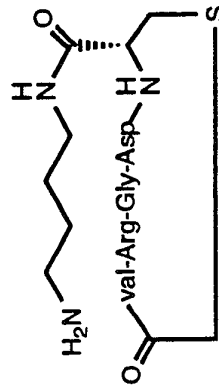
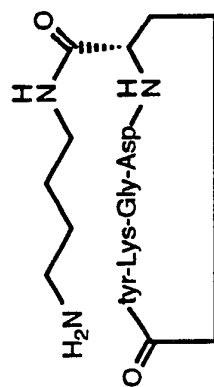
-----ELISA IC₅₀ nM-----

Compound 1 Ex(#)/Cpd.(#)	Sequence ²	Fibrinogen ³ GPIIb/IIIa	Vitronectin ⁴ VnR	GPIIb/IIIa ⁵ vWF	Fibronectin ⁶ FnR	Human ⁷ Platelet Aggregation IC ₅₀ nM	Rabbit ⁸ BT Ratio (%P.A. Inhibition)
	 <p>Tyr-Lys-Gly-Asp-Cys-Lys-NH₂</p>						
	 <p>Tyr-Lys-Gly-Asp-Arg-NH₂</p>	11.8	100000	14.70	>10000	180	
	 <p>Tyr-Lys-Gly-Asp-Lys-NH₂</p>	30	100000	22.50	>10000	730	

-----ELISA IC₅₀ nM-----

Compound 1 Ex(#)/Cpd.(#)	Sequence ²	Fibrinogen ³ GPIIb/III _a	Vitronectin ⁴ VnR	GPIIb/III _a ⁵ vWF	Fibronectin ⁶ FnR	Human ⁷ Platelet Aggregation IC ₅₀ nM	Rabbit ⁸ BT Ratio (%P.A. Inhibition)
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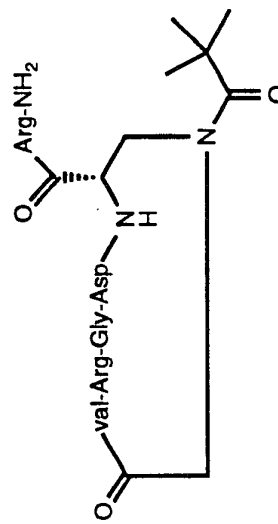
4.1



5.5

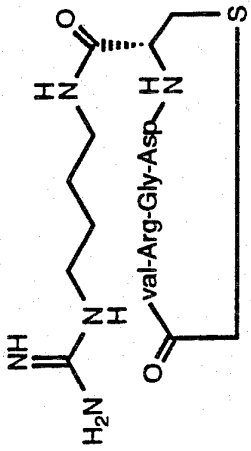
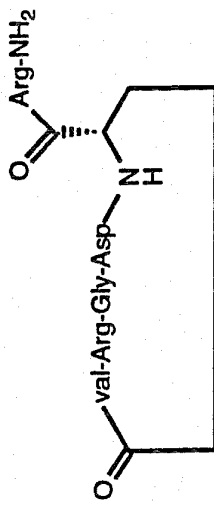
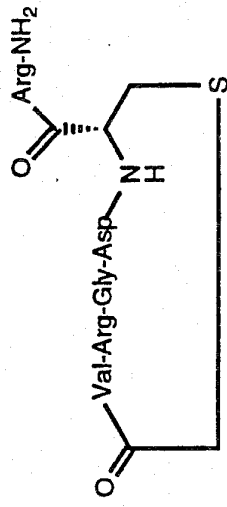
4.20

270



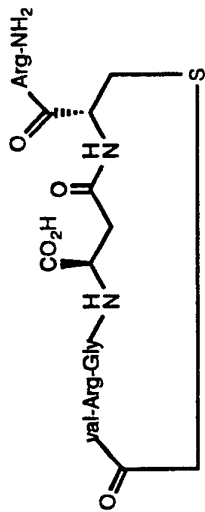
170

-----ELISA IC₅₀ nM-----

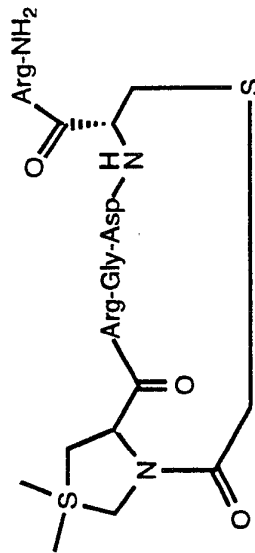
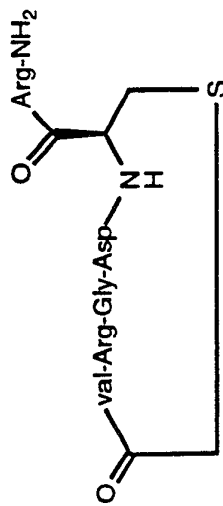
Compound ¹ Ex(#)/Cpd.(#)	Sequence ²	Fibrinogen ³ GPIIb/IIIa	Vitronectin ⁴ VnR	GPIIb/IIIa ⁵ vWF	Fibrinectin ⁶ FnR	Human ⁷ Platelet Aggregation IC ₅₀ nM	Rabbit ⁸ BT Ratio (%P.A. Inhibition)
		6.9	81	5.70			
		3.2	400	1.00	8350	70	
		130				2730	

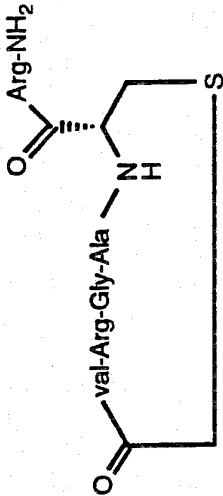
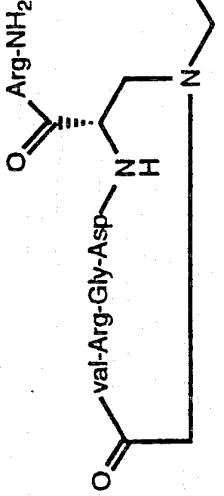
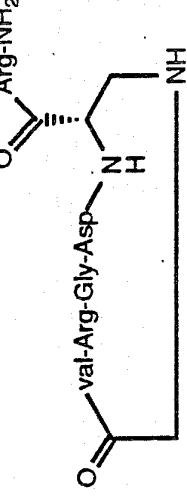
-----ELISA IC₅₀ nM-----

Compound 1 Ex(#)/Cpd.(#)	Sequence ²	Fibrinogen ³ GPIIb/IIIa	Vitronectin ⁴ VnR	GPIIb/IIIa ⁵ vWF	Fibronectin ⁶ FnR	Human ⁷ Platelet Aggregation IC ₅₀ nM	Rabbit ⁸ BT Ratio (%P.A. Inhibition)
							neg
						16560	



110

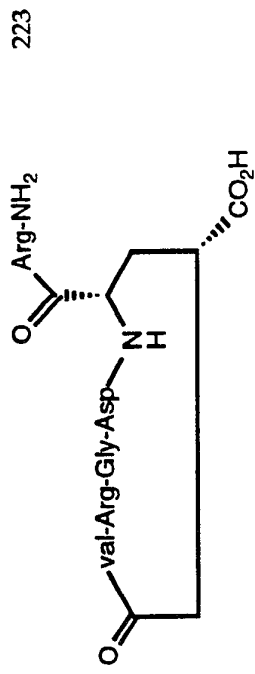


Compound ¹ Ex(#)/Cpd.(#)	Sequence ²	-----ELISA IC ₅₀ nM-----				Human ⁷ Platelet Aggregation IC ₅₀ nM	Rabbit ⁸ BT Ratio (%P.A. Inhibition)
		Fibrinogen ³ GPIIb/IIIa	Vitronectin ⁴ VnR	GPIIb/IIIa ⁵ vWF	Fibronectin ⁶ FnR		
					neg	84	
					330		
					330		

11

-----ELISA IC₅₀ nM-----

Compound 1 Ex(#)/Cpd.(#)	Sequence ²	Fibrinogen ³ GPIIb/III _a	Vitronectin ⁴ VnR	GPIIb/III _a ⁵ vWF	Fibronectin ⁶ FnR	Human ⁷ Platelet Aggregation IC ₅₀ nM	Rabbit ⁸ BT Ratio (%P.A. Inhibition)
-----------------------------	-----------------------	---	---------------------------------	--	---------------------------------	--	---



- 1 The numerical designation refers to the example and compound number in that example where the method of synthesis is described.
- 2 The three letter code used in this table refers to the naturally occurring α -amino acid residues unless the code begins with a lower case letter in which case it represents the unnatural α -amino acid. Two formats for depicting the structures of the cyclic peptides are used in this table. When the residue Cys is connected to the letter "S", it represents a thioether linkage where "S" is the sulfur from Cys.
- 3 This assay was conducted as described in Example 2.
- 4 This assay was conducted as described in Example 3.
- 5 This assay was conducted as described in Example 4.
- 6 This assay was conducted as described in Example 6.
- 7 This assay was conducted as described in Example 5.
- 8 This assay was conducted as described in Example 1. BT Ratio is the bleeding time ratio pre-administration/post-administration. %P.A. inhibition is the percent *ex vivo* platelet aggregation inhibition in the rabbit.

What is claimed is:

1. A peptide comprising:

- 5 (a) a cyclic moiety containing a sequence selected from;
Xaa₁-Arg-Gly-Asp-Xaa₂ and
Xaa₁-Lys-Gly-Asp-Xaa₂

where;

Xaa₁ represents from 1 to 20 α -amino acids or

α -amino acid analogues, and

10 Xaa₂ represents an α -amino acid or α -amino acid
analogue bonded to Xaa₁ through a bond
selected from the group;

amide,

thioether,

15 disulfide,

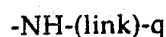
ether,

sulfoxide,

sulfone, and

- 20 (b) a positively charged nitrogen containing exocyclic moiety stably
bonded to Xaa₂ through a functional group of Xaa₂.

2. The peptide of claim 1 wherein the positively charged exocyclic
moiety is represented by the formula:



where;

(link) represents a linking group selected from;

30 C₃-C₁₀-alkyl, either branched, linear, or cyclic,

C₃-C₁₀-alkenyl,

C₃-C₁₀-alkynyl,

C₆-C₁₄-aryl substituted with 2 or more C₁-C₈-alkyl
groups,

C₁-C₁₃-heterocycle, saturated or unsaturated, containing

35 1-4 heteroatoms selected from N, O and S,

C₁-C₆-alkyl substituted C₅-C₁₄ saturated or unsaturated
heterocycle, containing from 1-4 heteroatoms
selected from N, O and S, and

optionally, (link) may be substituted with substituents
selected from;

5 COR,
 CONR'R",
 halo (F,Cl,Br,I),
 nitro,
 C₁-C₆-alkyl,
 phenyl,
 benzyl, and
10 C₃-C₆-cycloalkyl,

wherein R is selected from;

 hydroxy,
 C₁-C₈-alkoxy,
 C₃-C₁₂-alkenoxy,
15 C₆-C₁₂-aryloxy,
 di-C₁-C₈-alkylamino-C₁-C₈-alkoxy,
 acylamino-C₁-C₈-alkoxy selected from the
 group;
20 acetylaminoethoxy,
 nicotinoylaminoethoxy,
 succinamidoethoxy, and
 pivaloyloxyethoxy,
 C₆-C₁₂-aryl-C₁-C₈-alkoxy where the aryl group
 is unsubstituted or substituted with one or
25 more of the groups;
 nitro,
 halo (F, Cl, Br, I),
 C₁-C₄-alkoxy, and
 amino,
30 hydroxy-C₂-C₈-alkoxy,
 dihydroxy-C₃-C₈-alkoxy, and

wherein R' and R" are independently selected from;

 hydrogen,
 C₁-C₁₀-alkyl either branched, linear or
35 cyclic, C₃-C₁₀-alkenyl provided the
 double bond is not adjacent any
 N,
 C₆-C₁₄-aryl,
 C₁-C₆-alkyl-C₆-C₁₀-aryl

saturated or unsaturated heterocycle
having from 5-14 atoms in the
cycles and from 1-4 heteroatoms
selected from N, O and S,

5 optionally, R' and R" taken together may form
trimethylene,
tetramethylene,
pentamethylene, and
3-oxopentamethylene,

10 q represents a group selected from;

amino,
amidino, and
guanidino,
wherein any hydrogen bonded to any nitrogen or carbon
15 of the amino, amidino or guanidino group is optionally
substituted with a lower C₁-C₆-alkyl group.

3. The peptide of claim 2 wherein the positively charged exocyclic moiety
is a positively charged amino acid residue selected from α -amino acids
20 or α -amino acid analogues.

4. The peptide of claim 3 wherein the positively charged exocyclic moiety
is a D or L α -amino acid selected from;

25 His,
Lys,
Arg, and
Om,

wherein the α -carboxyl group of the D or L α -amino acid is optionally
derivitized with an amino or lower alkyl substituted amino group.

30 5. The peptide of claim 4 wherein Xaa₂ is an amino acid residue selected
from the group;

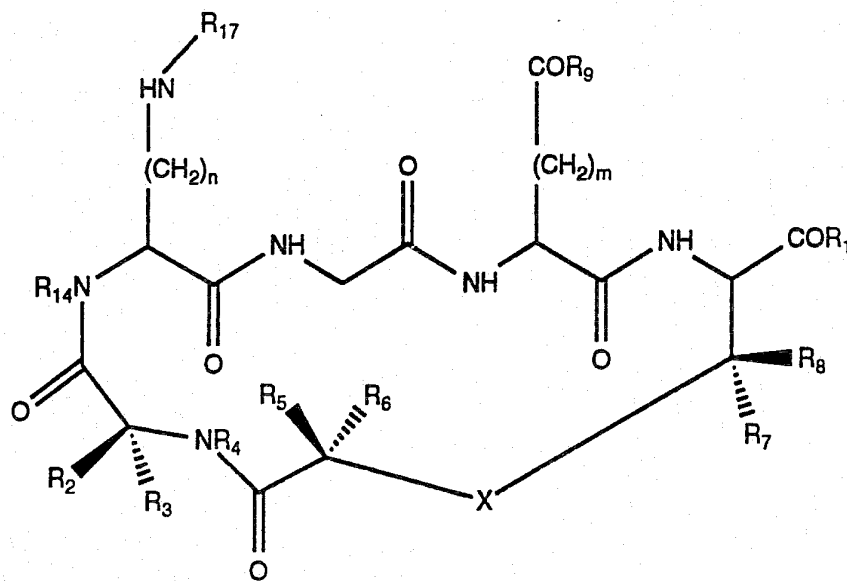
α -aminoadipic,
35 Cys,
homo-Cys,
Pen,
Pmp,
Pas,
Asp,

- 5 Glu,
 Orn,
 Lys,
 Ser,
 Thr, and
 Tyr.
6. The peptide of claim 5 where Xaa₂ is selected from Cys and Pen.
- 10 7. The peptide of claim 6 wherein Xaa₁ is one or more naturally occurring
 α-amino acids or the D stereoisomers thereof.
8. The peptide of claim 7 wherein Xaa₁ is a single amino acid selected
 from;
- 15 Gly,
 D-Ala,
 D-Val,
 D-Leu,
 D-Ile,
20 D-Phe,
 D-Tyr, and
 D-Pro.
9. The peptide of claim 8 wherein the linkage bonding Xaa₁ and Xaa₂ is
25 selected from thioether and sulfoxide.
10. The peptide of claim 9 wherein the linkage further comprises an acetyl
 group wherein the β-carbon of the acetyl group is bonded to the sulfur
 and wherein the carbonyl forms an amide with the α-amino of Xaa₁.
- 30 11. A peptide selected from the group:
- cyclo-S-acetyl-dTyr-Arg-Gly-Asp-Cys-Arg-NH₂;
 cyclo-S-acetyl-dTyr-Arg-Gly-Asp-Cys-dArg-NH₂;
35 cyclo-S-acetyl-dTyr-Arg-Gly-Asp-Cys-Lys-NH₂;
 cyclo-S-acetyl-dTyr-Arg-Gly-Asp-Cys-dLys-NH₂;
 cyclo-S-acetyl-dTyr-Arg-Gly-Asp-Cys-Orn-NH₂;
 cyclo-S-acetyl-dTyr-Arg-Gly-Asp-Cys-dOrn-NH₂;
 cyclo-S-acetyl-dTyr-Arg-Gly-Asp-Cys-Tyr-Arg-NH₂;

	cyclo-S-acetyl-dTyr-Arg-Gly-Asp-Cys-NH-X	(X=butylamine);
	cyclo-S-acetyl-dTyr-Arg-Gly-Asp-Cys-NH-X	(X=pentylamine);
	cyclo-S-acetyl-dTyr-Arg-Gly-Asp-Cys-NH-X	(X=hexylamine);
	cyclo-S-acetyl-dTyr-Lys-Gly-Asp-Cys-NH-X	(X=butylamine);
5	cyclo-S-acetyl-dTyr-Lys-Gly-Asp-Pen-NH-X	(X=butylamine);
	cyclo(1-5)dTyr-Arg-Gly-Asp-L- α -aminoadipic acid-Arg-NH ₂ ;	
	cyclo(1-5)dTyr-Arg-Gly-Asp-L- α -aminoadipic acid-Lys-NH ₂ ;	
	cyclo(1-5)dTyr-Arg-Gly-Asp-L- α -aminoadipic-butylamine;	
	cyclo-S-acetyl-dTyr-Arg-Gly-Asp-Cys-NH-X	(X=propylamine);
10	cyclo-S-acetyl-dTyr-Arg-Gly-Asp-Cys-NH-X	(X=ethylamine);
	cyclo-S-acetyl-dTyr-Lys-Gly-Asp-Pen-Arg-NH ₂ ;	
	cyclo-S-acetyl-dTyr-Lys-Gly-Asp-Pen-Lys-NH ₂ ;	
	cyclo-S-acetyl-dTyr-Arg-Gly-Asp-Cys-Arg-OH;	
	cyclo-S-acetyl-dTyr-Arg-Gly-Asp-Cys-Lys-OH;	
15	cyclo-S-acetyl-dTyr-Lys-Gly-Asp-Pen-Arg-OH;	
	cyclo-S-acetyl-dTyr-Arg-Gly-Asp-Cys-Gln-NH ₂ ;	
	cyclo-S-acetyl-Tyr-Arg-Gly-Asp-Cys-Arg-NH ₂ ;	
	cyclo-S-acetyl-dTyr-Arg-Gly-Asp-Cys-Arg-NH ₂	Sulfoxide (isomer 1);
	cyclo-S-acetyl-dTyr-Arg-Gly-Asp-Cys-Arg-NH ₂	Sulfoxide (isomer 2);
20	cyclo-S-acetyl-dTyr-Arg-Gly-Asp-Cys-dArg-NH ₂	Sulfoxide (isomer 1);
	cyclo-S-acetyl-dTyr-Arg-Gly-Asp-Cys-dArg-NH ₂	Sulfoxide (isomer 2);
	cyclo-S-acetyl-dTyr-Arg-Gly-Asp-Cys-Lys-NH ₂	Sulfoxide (isomer 1);
	cyclo-S-acetyl-dTyr-Arg-Gly-Asp-Cys-Lys-NH ₂	Sulfoxide (isomer 2);
	cyclo-S-acetyl-dTyr-Arg-Gly-Asp-Cys-dLys-NH ₂	Sulfoxide (isomer 1);
25	cyclo-S-acetyl-dTyr-Arg-Gly-Asp-Cys-dLys-NH ₂	Sulfoxide (isomer 2);
	cyclo-S-acetyl-dTyr-Arg-Gly-Asp-Cys-Orn-NH ₂	Sulfoxide (isomer 1);
	cyclo-S-acetyl-dTyr-Arg-Gly-Asp-Cys-Orn-NH ₂	Sulfoxide (isomer 2);
	cyclo-S-acetyl-dTyr-Arg-Gly-Asp-Cys-dOrn-NH ₂	Sulfoxide (isomer 1);
	cyclo-S-acetyl-dTyr-Arg-Gly-Asp-Cys-dOrn-NH ₂	Sulfoxide (isomer 2);
30	cyclo-S-acetyl-dTyr-Arg-Gly-Asp-Cys-Tyr-Arg-NH ₂	Sulfoxide (isomer 1);
	cyclo-S-acetyl-dTyr-Arg-Gly-Asp-Cys-Tyr-Arg-NH ₂	Sulfoxide (isomer 2);
	Cyclo-S-acetyl-dVal-Arg-Gly-Asp-Cys-Lys-NH ₂ ;	
	Cyclo-S-acetyl-dVal-Arg-Gly-Asp-Cys-Lys-NH ₂	sulfoxide (isomer 1);
	Cyclo-S-acetyl-dVal-Arg-Gly-Asp-Cys-Lys-NH ₂	sulfoxide (isomer 2);
35	Cyclo-S-acetyl-dVal-Arg-Gly-Asp-Cys-Arg-NH ₂ ;	
	Cyclo-S-acetyl-dVal-Arg-Gly-Asp-Cys-Arg-NH ₂	sulfoxide (isomer 1);
	Cyclo-S-acetyl-dVal-Arg-Gly-Asp-Cys-Arg-NH ₂	sulfoxide (isomer 2);
	Cyclo-S-acetyl-dVal-Lys-Gly-Asp-Cys-Lys-NH ₂ ;	
	Cyclo-S-acetyl-dVal-Lys-Gly-Asp-Cys-Lys-NH ₂	sulfoxide (isomer 1);

- Cyclo-S-acetyl-dVal-Lys-Gly-Asp-Cys-Lys-NH₂ sulfoxide (isomer 2);
 Cyclo-S-(phenyl)acetyl-dVal-Lys-Gly-Asp-Cys-Lys-NH₂;
 Cyclo-S-(phenyl)acetyl-dVal-Lys-Gly-Asp-Cys-Lys-NH₂ sulfoxide (isomer 1);
 Cyclo-S-(phenyl)acetyl-dVal-Lys-Gly-Asp-Cys-Lys-NH₂ sulfoxide (isomer 2);
 5 Cyclo-S-acetyl-dVal-Arg-Gly-Asp-Cys-NH-X (X=butylamine);
 Cyclo-S-acetyl-dVal-Arg-Gly-Asp-Cys-NH-X (X=butylamine)
 sulfoxide (isomer 1);
 Cyclo-S-acetyl-dVal-Arg-Gly-Asp-Cys-NH-X (X=butylamine)
 sulfoxide (isomer 2);
 10 Cyclo-(1-5)-dVal-Arg-Gly-Asp-L- α -aminoadipic acid-Arg-NH₂;
 Cyclo-S-acetyl-dVal-Arg-Gly-Asp-Cys-NH-X (X=butylguanidine);
 Cyclo-S-acetyl-dVal-Arg-Gly-Asp-Cys-NH-X (X=butylguanidine)
 sulfoxide (isomer 1);
 Cyclo-S-acetyl-dVal-Arg-Gly-Asp-Cys-NH-X (X=butylguanidine)
 15 sulfoxide (isomer 2);
 Cyclo-S-acetyl-dTyr-Lys-Gly-Asp-Cys-Lys-NH₂;
 Cyclo-S-acetyl-dTyr-Lys-Gly-Asp-Cys-Lys-NH₂ sulfoxide (isomer 1);
 Cyclo-S-acetyl-dTyr-Lys-Gly-Asp-Cys-Lys-NH₂ sulfoxide (isomer 2);
 Cyclo-S-(phenyl)acetyl-dTyr-Lys-Gly-Asp-Cys-Lys-NH₂;
 20 Cyclo-S-(phenyl)acetyl-dTyr-Lys-Gly-Asp-Cys-Lys-NH₂ sulfoxide (isomer 1);
 Cyclo-S-(phenyl)acetyl-dTyr-Lys-Gly-Asp-Cys-Lys-NH₂ sulfoxide (isomer 2);
 Cyclo-(1-5)-dTyr-Lys-Gly-Asp-L- α -aminoadipic acid-Arg-NH₂;
 Cyclo-(1-5)-dTyr-Lys-Gly-Asp-L- α -aminoadipic acid-Lys-NH₂;
 Cyclo-(1-5)-dTyr-Lys-Gly-Asp-L- α -aminoadipic acid-NH-X (X=butylamine);
 25 Cyclo-S-acetyl-dPro-Arg-Gly-Asp-Cys-Arg-NH₂ ;
 Cyclo-S-acetyl-dPro-Arg-Gly-Asp-Cys-Arg-NH₂ sulfoxide (isomer 1);
 Cyclo-S-acetyl-dPro-Arg-Gly-Asp-Cys-Arg-NH₂ sulfoxide (isomer 2);
 Cyclo-S-acetyl-dPro-Arg-Gly-Asp-Cys-Lys-NH₂;
 Cyclo-S-acetyl-dPro-Arg-Gly-Asp-Cys-Lys-NH₂ sulfoxide (isomer 1);
 30 Cyclo-S-acetyl-dPro-Arg-Gly-Asp-Cys-Lys-NH₂ sulfoxide (isomer 2);
 Cyclo-(1-5)-dVal-Arg-Gly-Asp-L- α -amino-3-azaadipic acid-Arg-NH₂;
 Cyclo-(1-5)-dVal-Arg-Gly-Asp-L- α -amino-3-(N-pivaloyl)azaadipic acid-Arg-NH₂;
 Cyclo-(1-5)-dVal-Arg-Gly-Asp-L- α -amino-3-(N-neopentyl)azaadipic acid-Arg-NH₂;

12. A compound of the formula:



5 wherein

R₁ is selected from;

NHR₁₅Q, and

NR₁₅R₁₆Q,

wherein R₁₅ and R₁₆ are independently selected from;

10

C₃-C₁₀-alkyl either linear, branched or cyclic,

C₃-C₁₀-alkenyl,

C₃-C₁₀-alkynyl,

C₆-C₁₄-aryl,

C₁-C₆-alkyl-C₆-C₁₀-aryl,

15

saturated or unsaturated heterocycle or C₁-C₆-

alkyl substituted heterocycle

containing from 5 to 14 atoms in the

cycle and from 1 to 4 heteroatoms

selected from N, O and S,

20

NR₁₅ R₁₆ taken together may form a heterocycle

or C₂-C₆-alkyl substituted heterocycle

wherein R₁₅ and R₁₆ taken together are

trimethylene, tetramethylene,

pentamethylene or 3-

25

oxopentamethylene,

and wherein each R₁₅ or R₁₆ may optionally be

substituted with one or more substituents selected

from;

85

COR₉,

CONR'R",

halo (F,Cl,Br,I),

nitro,

5

C₁-C₆-alkyl,

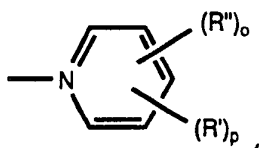
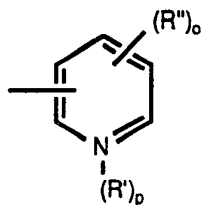
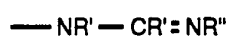
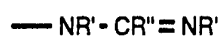
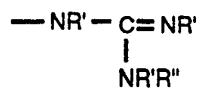
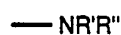
phenyl,

benzyl, and

C₃-C₆-cycloalkyl, and

Q is a group bonded to R₁₅, R₁₆ or substituents bonded
thereto, Q being selected from;

10



wherein;

15

R' and R" are independently selected from;

hydrogen,

C₁-C₁₀-alkyl either branched, linear or
cyclic,C₃-C₁₀-alkynyl,

20

C₆-C₁₄-aryl,C₁-C₆-alkyl-C₆-C₁₀-aryl

saturated or unsaturated heterocycle
 having from 5-14 atoms in the
 cycles and from 1-4 heteroatoms
 selected from N, O and S,

5 and wherein R' and R" taken together are

trimethylene,
 tetramethylene,
 pentamethylene, and
 3-oxopentamethylene,

10 R''' is C₁-C₁₀-alkyl, phenyl and benzyl, and
 o+ p is an integer selected from 0, 1, 2, 3, and 4;

R₂, R₃, R₅, R₆, R₇, R₈ are the same or different and are selected from;

hydrogen,

15 C₆-C₁₂-aryl where the aryl group is unsubstituted or substituted by one or
 more of the groups; nitro, hydroxy, halo (F, Cl, Br, I), C₁-C₈-alkyl, halo-
 C₁-C₈-alkyl, C₁-C₈-alkoxy, amino, phenyloxy, phenyl, acetamido,
 benzamido, di-C₁-C₈-alkylamino, C₁-C₈-alkylamino, C₆-C₁₂-aroyl, C₁-
 C₈-alkanoyl and hydroxy-C₁-C₈-alkyl,

20 C₁-C₁₂-alkyl either substituted or unsubstituted, branched or straight chain
 where the substituents are selected from;

halo (F, Cl, Br, I),

C₁-C₈-alkoxy,

25 C₆-C₁₂-aryloxy where the aryl group is unsubstituted or substituted by
 one or more of the groups; nitro, hydroxy, halo (F, Cl, Br, I), C₁-
 C₈-alkyl, C₁-C₈-alkoxy, amino, phenyloxy, acetamido,
 benzamido, di-C₁-C₈-alkylamino, C₁-C₈-alkylamino, C₆-C₁₂
 aroyl and C₁-C₈-alkanoyl,

isothioureido,

C₃-C₇-cycloalkyl,

30 ureido,

amino,

C₁-C₈-alkylamino,

di-C₁-C₈-alkylamino,

hydroxy,

35 amino-C₂-C₈-alkylthio,

amino-C₂-C₈-alkoxy,

acetamido, and

benzamido wherein the phenyl ring is unsubstituted or substituted by
 one or more of the groups; nitro, hydroxy, halo (F, Cl, Br, I), C₁-

- C₈-alkyl, C₁-C₈-alkoxy, amino, phenyloxy, acetamido,
 benzamido, di-C₁-C₈-alkylamino, C₁-C₈-alkylamino, C₆-C₁₂
 aroyl, and C₁-C₈-alkanoyl,
 C₆-C₁₂-arylamino wherein the aryl group is unsubstituted or
 5 substituted by one or more of the groups; nitro, hydroxy, halo,
 C₁-C₈-alkyl, C₁-C₈-alkoxy, amino, phenyloxy, acetamido,
 benzamido, di-C₁-C₈-alkylamino, C₁-C₈-alkylamino, C₆-C₁₂-
 aroyl, and C₁-C₈-alkanoyl,
 guanidino,
 10 phthalimido,
 mercapto,
 C₁-C₈-alkylthio,
 C₆-C₁₂-arylthio,
 carboxy,
 15 carboxamide,
 carbo-C₁-C₈-alkoxy, and
 C₆-C₁₂-aryl wherein the aryl group is unsubstituted or substituted by
 one or more of the groups nitro, hydroxy, halo, C₁-C₈-alkyl, C₁-
 C₈-alkoxy, amino, phenyloxy, acetamido, benzamido, di-C₁-C₈
 20 alkylamino, C₁-C₈-alkylamino, hydroxy-C₁-C₈-alkyl, C₆-C₁₂
 aroyl and C₁-C₈-alkanoyl, and
 aromatic heterocycle wherein the heterocycle contains 5-10 ring
 atoms and one or two O, N or S heteroatoms;
 R₂ and R₃, R₅ and R₆, or R₇ and R₈ may optionally and independently be joined
 25 together to form a carbocyclic or heterocyclic ring of from four to seven atoms
 where the heteroatoms are selected from O, S or NR₁₂ where R₁₂ is selected
 from;
 hydrogen,
 C₁-C₈-alkyl,
 30 C₃-C₈-alkenyl,
 C₆-C₁₂-aryl,
 C₆-C₁₂-aryl-C₁-C₈-alkyl,
 C₁-C₈-alkanoyl, and
 C₆-C₁₂-aroyl,
 35 R₄ is selected from;
 hydrogen,
 C₁-C₈-alkyl,
 C₃-C₁₀-cycloalkyl,
 C₆-C₁₂-aryl, and

C₆-C₁₂-aryl-C₁-C₈-alkyl;

R₂ or R₃ may be optionally joined with R₄ to form a piperidine, pyrrolidine or thiazolidine ring;

R₉ is selected from;

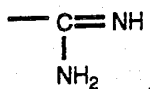
- 5 hydroxy,
 C₁-C₈-alkoxy,
 C₃-C₁₂-alkenoxy,
 C₆-C₁₂-aryloxy,
 di-C₁-C₈-alkylamino-C₁-C₈-alkoxy,
 10 acylamino-C₁-C₈-alkoxy selected from the group acetylaminoethoxy,
 nicotinoylaminoethoxy and succinamidoethoxy,
 pivaloyloxyethoxy,
 C₆-C₁₂ aryl-C₁-C₈-alkoxy where the aryl group is unsubstituted or substituted
 with one or more of the groups; nitro, halo (F, Cl, Br, I), C₁-C₄-alkoxy,
 15 and amino,
 hydroxy-C₂-C₈-alkoxy,
 dihydroxy-C₃-C₈-alkoxy, and
 NR₁₀R₁₁ wherein R₁₀ and R₁₁ are the same or different and are hydrogen,
 C₁-C₈-alkyl, C₃-C₈-alkenyl, C₆-C₁₂-aryl where the aryl group is
 20 unsubstituted or substituted with one or more of the groups
 nitro, halo (F, Cl, Br, I), C₁-C₄-alkoxy, and amino, C₆-C₁₂-aryl-
 C₁-C₈-alkyl where the aryl group is unsubstituted or substituted
 by one or more of the groups nitro, halo (F, Cl, Br, I), C₁-C₄-
 alkoxy, and amino;

- 25 R₁₄ is selected from;
 hydrogen, C₁-C₈-alkyl, C₃-C₈-alkenyl, C₆-C₁₂-aryl and C₆-C₁₂-aryl-C₁-C₈-
 alkyl;

R₁₇ is selected from;

hydrogen, and

30



X is selected from;

O or S,

35

S bearing one or two O atoms,

NR₁₃ wherein R₁₃ is;

hydrogen,

C₁-C₈-alkyl,

C₃-C₈-alkenyl,
C₆-C₁₄-aryl,
C₆-C₁₄-aryl-C₁-C₈-alkyl,
C₁-C₈-alkanoyl,
C₆-C₁₄-aroyl,
C₆-C₁₄-aryl-C₁-C₈-alkanoyl, and

(CH₂)_k where k is an integer from 0 to 5;

n is an integer from 1 to 6;

m is an integer from 0 to 4; and

pharmaceutically acceptable salts thereof.

13. A pharmaceutical composition comprising a pharmaceutically acceptable excipient and the compound of claim 1.

14. A method for inhibiting platelet aggregation which method comprises administering a platelet aggregation inhibiting amount of the compound of claim 1.

15. A method for reducing platelet aggregation in a mammal, comprising administering a pharmaceutically effective amount of the composition of matter as defined by claim 1 to the mammal.

16. The method of claim 15, further comprising administering said composition of matter to the mammal in admixture with a pharmaceutically acceptable carrier.

17. A method for treating a mammal who has an increased propensity for thrombus formation, comprising administering a pharmaceutically effective amount of the composition of matter as defined by claim 1 to the mammal.

18. A method for treating a mammal who has an increased propensity for thrombus formation, comprising administering a pharmaceutically effective amount of the composition of matter as defined by claim 1 in combination with a thrombolytic agent.

19. A method for treating a mammal who has an increased propensity for thrombus formation, comprising administering a pharmaceutically effective amount of the composition of matter as defined by claim 1 in combination with an anticoagulant.

20. A method for treating a mammal who has an increased propensity for thrombus formation, comprising administering a pharmaceutically effective amount of the composition of matter as defined by claim 12.
- 5 21. The cyclic peptide of claim 1 wherein the cycle contains 17 or 18 atoms in the ring.

INTERNATIONAL SEARCH REPORT

International Application No **PCT/US 92/02731**

I. CLASSIFICATION OF SUBJECT MATTER (if several classification symbols apply, indicate all) ⁶		
According to International Patent Classification (IPC) or to both National Classification and IPC		
Int. Cl. 5 C 07 K 7/10	C 07 K 7/02 A 61 K 37/02	C 07 K 7/06 C 07 K 7/08
II. FIELDS SEARCHED		
Minimum Documentation Searched ⁷		
Classification System	Classification Symbols	
Int. Cl. 5	C 07 K A 61 K	
Documentation Searched other than Minimum Documentation to the Extent that such Documents are Included in the Fields Searched ⁸		
III. DOCUMENTS CONSIDERED TO BE RELEVANT ⁹		
Category ¹⁰	Citation of Document, ¹¹ with indication, where appropriate, of the relevant passages ¹²	Relevant to Claim No. ¹³
A	EP, A, 0341915 (SMITHKLINE BECKMAN) 15 November 1989, see the whole disclosure -----	1-21
P, X	WO, A, 9115515 (LA JOLLA CANCER RESEARCH FOUNDATION) 17 October 1991, see the whole document, especially compounds cited in table 1; page 5, lines 29-32; page 27, lines 17-23; page 54, claim 17 -----	1-21
<div style="display: flex; justify-content: space-between;"> <div style="width: 45%;"> <p>¹⁰ Special categories of cited documents :</p> <p>"A" document defining the general state of the art which is not considered to be of particular relevance</p> <p>"E" earlier document but published on or after the international filing date</p> <p>"L" document which may throw doubts on priority claim(s) or which is cited to establish the publication date of another citation or other special reason (as specified)</p> <p>"O" document referring to an oral disclosure, use, exhibition or other means</p> <p>"P" document published prior to the international filing date but later than the priority date claimed</p> </div> <div style="width: 45%;"> <p>"T" later document published after the international filing date or priority date and not in conflict with the application but cited to understand the principle or theory underlying the invention</p> <p>"X" document of particular relevance; the claimed invention cannot be considered novel or cannot be considered to involve an inventive step</p> <p>"Y" document of particular relevance; the claimed invention cannot be considered to involve an inventive step when the document is combined with one or more other such documents, such combination being obvious to a person skilled in the art.</p> <p>"&" document member of the same patent family</p> </div> </div>		
IV. CERTIFICATION		
Date of the Actual Completion of the International Search		Date of Mailing of this International Search Report
27-07-1992		20. 08. 92
International Searching Authority		Signature of Authorized Officer
EUROPEAN PATENT OFFICE		MASTURZO

INTERNATIONAL SEARCH REPORT

International application No.

PCT/US 92/02731

Box I Observations where certain claims were found unsearchable (Continuation of item 1 of first sheet)

This international search report has not been established in respect of certain claims under Article 17(2)(a) for the following reasons:

1. Claims Nos.: 14-20
because they relate to subject matter not required to be searched by this Authority, namely:
Although claims 14-20 refer to a method of treatment of the human body, the search was carried out and based on the alleged effects of the compounds.
2. Claims Nos.:
because they relate to parts of the international application that do not comply with the prescribed requirements to such an extent that no meaningful international search can be carried out, specifically:
3. Claims Nos.:
because they are dependent claims and are not drafted in accordance with the second and third sentences of Rule 6.4(a).

Box II Observations where unity of invention is lacking (Continuation of item 2 of first sheet)

This International Searching Authority found multiple inventions in this international application, as follows:

1. As all required additional search fees were timely paid by the applicant, this international search report covers all searchable claims.
2. As all searchable claims could be searched without effort justifying an additional fee, this Authority did not invite payment of any additional fee.
3. As only some of the required additional search fees were timely paid by the applicant, this international search report covers only those claims for which fees were paid, specifically claims Nos.:
4. No required additional search fees were timely paid by the applicant. Consequently, this international search report is restricted to the invention first mentioned in the claims; it is covered by claims Nos.:

Remark on Protest

- The additional search fees were accompanied by the applicant's protest.
- No protest accompanied the payment of additional search fees.

ANNEX TO THE INTERNATIONAL SEARCH REPORT
ON INTERNATIONAL PATENT APPLICATION NO.

US 9202731

SA 58760

This annex lists the patent family members relating to the patent documents cited in the above-mentioned international search report. The members are as contained in the European Patent Office EDP file on 04/08/92
The European Patent Office is in no way liable for these particulars which are merely given for the purpose of information.

Patent document cited in search report	Publication date	Patent family member(s)	Publication date
EP-A- 0341915	15-11-89	AU-A- 3458889	09-11-89
		JP-A- 2062892	02-03-90
WO-A- 9115515	17-10-91	AU-A- 7763191	30-10-91