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(54) **ANALOGS OF PARATHYROID HORMONE**

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(63) Continuation of application No. 09/399,499, filed on Sep. 20, 1999, now Pat. No. 6,544,949, and a continuation-in-part of application No. 09/341,217, filed as application No. PCT/US97/022498 on Dec. 8, 1997, now abandoned, which is a continuation-in-part of application No. 08/813,534, filed on Mar. 7, 1997, now Pat. No. 5,955,574, which is a continuation-in-part of application No. 08/779,768, filed on Jan. 7, 1997, now Pat. No. 5,969,095, which is a continuation-in-part of application No. 08/626,186, filed on Mar. 29, 1996, now Pat. No. 5,723,577

(60) Provisional application No. 60/001,105, filed on Jul. 13, 1995, and provisional application No. 60/003,305, filed on Sep. 6, 1995.

(51) **Int. Cl.**
A61K 38/29 (2006.01)
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(52) **U.S. Cl.** **514/12**; 530/324

(58) **Field of Classification Search** ..... None
See application file for complete search history.

(56) **References Cited**

U.S. PATENT DOCUMENTS

4,656,250	A	*	4/1987	Morita et al.	530/324
5,434,246	A	*	7/1995	Fukuda et al.	530/324
5,455,329	A	*	10/1995	Wingender et al.	530/324
5,457,047	A	*	10/1995	Wingender et al.	435/252.3
5,589,452	A	*	12/1996	Krstenansky et al.	514/12
5,599,792	A	*	2/1997	Kronis et al.	514/12
5,723,577	A	*	3/1998	Dong	530/324
5,955,574	A	*	9/1999	Dong	530/324
5,969,095	A	*	10/1999	Dong	530/324
6,544,949	B1	*	4/2003	Dong	514/12
7,410,948	B2	*	8/2008	Dong	514/12

FOREIGN PATENT DOCUMENTS

AU	64834/96	*	7/1999	
EP	0 293 158	A *	11/1988	
EP	0 293 158		11/1988	
EP	415867		10/1991	
EP	451867	A *	10/1991	
EP	0 477 885		4/1992	
EP	0 561 412		9/1993	
EP	0 561 412	A *	9/1993	
EP	0 672 057		9/1995	
EP	748817		12/1996	
EP	748817	A *	12/1996	
GB	2269176	A *	2/1994	
GB	2269176		2/1994	
HU	P9500110	*	6/1995	
HU	P9500115	*	6/1995	
HU	217 843	B *	9/1995	
JP	6-184198		7/1994	
WO	WO 90/10067		9/1990	
WO	WO 90/10067	A *	9/1990	
WO	WO 93/06846	A *	4/1993	
WO	WO 93/06846		4/1993	
WO	WO 94/01460	A *	1/1994	
WO	94/01460		1/1994	
WO	94/02510		2/1994	
WO	WO 94/02510	A *	2/1994	
WO	WO95/02610		1/1995	
WO	95/02610		1/1995	
WO	95/04752		2/1995	
WO	96/19246		6/1996	
WO	96/40775		12/1996	
WO	97/02834		1/1997	

OTHER PUBLICATIONS

Chorev et al. Modifications of Position 12 in Parathyroid Hormone . . . . Biochemistry, vol. 29, No. 6, pp. 1580–1586 (1990).*

Lynn H. Caporale et al; "Characterization of parathyroid hormone antagonists". Peptides, Chemistry and Biology; Proceedings of the Tenth American Peptides, Chemistry and Biology; Peptide Symposium; (1987); pp. 449–451.*

Horiuchi N. et al, "Similarity of synthetic peptide from human tumor to parathyroid hormone in vivo and in vitro". Science; (1987); vol. 238, pp. 1566–1568.*

Gardella T.J. et al. "Parathyroid Hormone (PTH)–PTH–Related Peptide Hybrid Peptides Reveal Functional Interactions Between The 1–14 and 15 34 Domains of the Ligand". Journal of Biological Chemistry. vol. 270, No. 12, pp. 6584–6588 (1995).*

Barden et al. "NMR Solution Structure of Human Parathyroid Hormone (1–34)". Biochemistry, vol. 32, pp. 7126–7132 (1993).*

Cohen et al. "Analogues of Parathyroid Hormone Modified at Positions 3 and 6". The Journal of Biological Chemistry. vol. 266, No. 3, pp. 1997–2004 (1991).*

Karle et al. "Structural Characteristics of alpha–Helical Peptide Molecules Containing Aib Residues". Biochemistry, vol. 29, No. 29, pp. 6747–6756 (1990).*

(Continued)

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(57) **ABSTRACT**

The present invention is directed to peptide analogues of fragment of parathyroid hormone (PTH) or parathyroid hormone-related protein (PTHrP), a method of using said analogues alone or in combination with a bisphosphonate or calcitonin to treat osteoporosis and pharmaceutical compositions comprising said analogues alone or in combination with a bisphosphonate or calcitonin.

52 Claims, No Drawings

OTHER PUBLICATIONS

Leaffer et al. "Modulation of Osteogenic Cell Ultrastructure by RS–23581; an Analog of Human Parathyroid Hormone (PTH)–Related Peptide–(1–34). . . ". Endocrinology, vol. 136, No. 8, pp. 3624–3631 (1995).*

Li et al. "A Measure of Helical Propensity for Amino Acids in Membrane Environments". Structural Biology 1, No. 6, pp. 368–373 (1994).*

McLean et al. "Minimal Peptide Length for Interaction of Amphipathic alpha–Helical Peptides with Phosphatidylcholine Liposomes". Biochemistry. vol. 30, pp. 31–37 (1991).*

Neugebauer et al. Structural Elements of Human Parathyroid Hormone and Their Possible Relation to Biological Activities. Biochemistry. vol. 31, pp. 2056–2063 (1992).*

Strickland et al, "Structure of Human Parathyroid Hormone (1–34) in the Presence of Solvents and Micelles". Biochemistry. vol. 32, pp. 6050–6057 (1993).*

Surewicz et al. "Structure–function Relationships in Human Parathyroid Hormone: The Essential Role of Amphiphilic alpha–Helix". Peptide Hormone: 556–558 (not dated).*

* cited by examiner

ANALOGS OF PARATHYROID HORMONE

Matter enclosed in heavy brackets [ ] appears in the original patent but forms no part of this reissue specification; matter printed in italics indicates the additions made by reissue.

CROSS REFERENCE TO RELATED APPLICATIONS

This application is a continuation of U.S. application Ser. No. 09/399,499 filed Sep. 20, 1999, now issued as U.S. Pat. No. 6,544,949, which is a continuation-in-part of both U.S. application Ser. No. 08/779,768 filed Jan. 7, 1997, now issued as U.S. Pat. No. 5,969,095, and Ser. No. 09/341,217, filed Nov. 22, 1999, now abandoned, which is the national phase application of International Application No. PCT/US97/22498, filed Dec. 8, 1997, which is a continuation-in-part of U.S. application Ser. No. 08/813,354, filed Mar. 7, 1997, now issued as U.S. Pat. No. 5,955,574, which is a continuation-in-part of U.S. application Ser. No. 08/779,768, filed Jan. 7, 1997, now issued as U.S. Pat. No. 5,969,095, which is a continuation-in-part of U.S. application Ser. No. 08/626,186, filed Mar. 29, 1996, now issued as U.S. Pat. No. 5,723,577, which claims the benefit of priority of U.S. application No. 60/003,305, filed Sep. 6, 1995 and U.S. application No. 60/001,105, filed Jul. 13, 1995.

BACKGROUND OF THE INVENTION

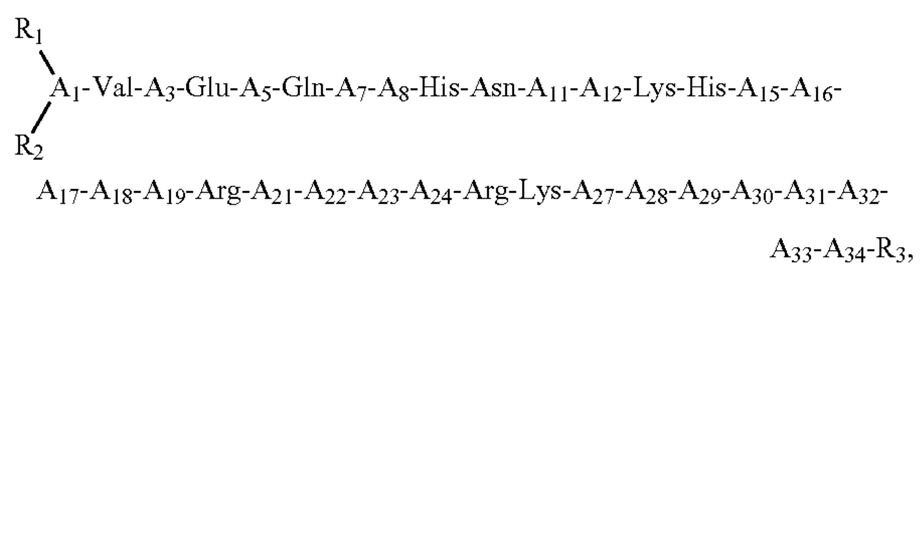
Parathyroid hormone ("PTH") is a polypeptide produced by the parathyroid glands. The mature circulating form of the hormone is comprised of 84 amino acid residues. The biological action of PTH can be reproduced by a peptide fragment of its N-terminus (e.g. amino acid residues 1 through 34). Parathyroid hormone-related protein ("PTHrP") is a 139 to 173 amino acid-protein with N-terminal homology to PTH. PTHrP shares many of the biological effects of PTH including binding to a common PTH/PTHrP receptor. Tregear, et. al, Endocrinol. 93:1349 (1983). PTH peptides from many different sources, e.g. human, bovine, rat, chicken, have been characterized. Nissenson. et al., Receptor, 3:193 (1993).

PTH has been shown to both improve bone mass and quality Dempster, et al., Endocrine Rev., 14:690 (1993); and Riggs, Amer. J. Med. 91 (Suppl 5B):37S (1991). The anabolic effect of intermittently administered PTH has been observed in osteoporitic men and women either with or without concurrent antiresorptive therapy. Slovik, et al, J. Bone Miner. Res., 1:377 (1986); Reeve, et al., Br. Med. J., 301:314 (1990); and Hesch, R-D., et al., Calcif. Tissue Int'l, 44:176 (1989).

SUMMARY OF THE INVENTION

In one aspect, the invention features a peptide of the formula (I),

[SEQ ID NO: 1]



wherein

A_1 is Ser, Ala, or Dap;

A_3 is Ser, Thr, or Aib;

A_5 is Leu, Nle, Ile, Cha, β-Nal, Trp, Pal, Acc, Phe or p-X-Phe, in which X is OH, a halogen, or CH_3;

A_7 is Leu, Nle, Ile, Cha, β-Nal, Trp, Pal, Acc, Phe, or p-X-Phe in which X is OH, a halogen, or CH_3;

A_8 is Met, Nva, Leu, Val, Ile, Cha, Acc, or Nle;

A_{11} is Leu, Nle, Ile, Cha, β-Nal, Trp, Pal, Acc, Phe or p-X-Phe in which X is OH, a halogen, or CH_3;

A_{12} is Gly, Acc, or Aib;

A_{15} is Leu, Nle, Ile, Cha, β-Nal, Trp, Pal, Acc, Phe, or p-X-Phe in which X is OH, a halogen, or CH_3;

A_{16} is Ser, Asn, Ala, or Aib;

A_{17} is Ser, Thr, or Aib;

A_{18} is Met, Nva, Leu, Val, Ile, Nle, Acc, Cha, or Aib;

A_{19} is Glu or Aib;

A_{21} is Val, Acc, Cha, or Met;

A_{22} is Acc or Glu;

A_{23} is Trp, Acc, or Cha;

A_{24} is Leu, Acc, or Cha,

A_{27} is Lys, Aib, Leu, hArg, Gin, Acc, or Cha;

A_{28} is Leu, Acc, or Cha;

A_{29} is [Gin] *Gln*, Acc, or Aib;

A_{30} is Asp or Lys;

A_{31} is Val, Leu, Nle, Acc, Cha, or deleted;

A_{32} is His or deleted;

A_{33} is Asn or deleted;

A_{34} is Phe, Tyr, Amp, Aib, or deleted; each of R_1 and R_2 is, independently, H, $C_{1\text{-}12}$ alkyl, $C_{2\text{-}12}$ alkenyl, $C_{7\text{-}20}$ phenylalkyl, $C_{11\text{-}20}$ naphthylalkyl, $C_{1\text{-}12}$ hydroxyalkyl, $C_{2\text{-}12}$ hydroxyalkenyl, $C_{7\text{-}20}$ hydroxyphenylalkyl, or $C_{11\text{-}20}$ hydroxynaphthylalkyl; or one and only one of R_1 and R_2 is COE_1 in which E_1 is $C_{1\text{-}12}$ alkyl, $C_{2\text{-}12}$ alkenyl, $C_{7\text{-}20}$ phenylalkyl, $C_{11\text{-}20}$ naphthylalkyl, $C_{1\text{-}12}$ hydroxyalkyl, $C_{2\text{-}12}$ hydroxyalkenyl, $C_{7\text{-}20}$ hydroxyphenylalkyl, or $C_{11\text{-}20}$ hydroxynaphthylalkyl; and

R_3 is OH, NH_2, $C_{1\text{-}12}$ alkoxy, or NH—Y—CH_2-Z in which Y is a $C_{1\text{-}12}$ hydrocarbon moiety and Z is H, OH, CO_2H, or $CONH_2$;

provided that at least one of A_5, A_7, A_8, A_{11}, A_{12}, A_{15}, A_{18}, A_{21}, A_{22}, A_{23}, A_{24}, A_{27}, A_{28}, A_{29}, and A_{31} is Acc; or a pharmaceutically acceptable salt thereof.

A preferred embodiment of the immediately foregoing peptide is where A_3 is Ser; A_5 is Ile or Acc; A_7 is Leu, Acc, or Cha; A_8 is Acc, Met, Nva, Leu, Val, Ile, or Nle; A_{11} is Leu, Acc, or Cha; A_{12} is Acc or Gly; A_{15} is Leu, Acc, or Cha; A_{16} is Asn or Aib; A_{17} is Ser or Aib; A_{18} is Acc, Met, or Nle; A_{21} is Val or Acc; A_{27} is Lys, hArg, Acc, or Cha; A_{31} is Val, Leu, Nle, Acc, or Cha; A_{32} is His; A_{33} is Asn; A_{34} is Phe, Tyr, Amp, or Aib; or a pharmaceutically acceptable salt thereof.

A preferred embodiment of the immediately foregoing peptide, designated Group B, is where A_5 is Ile or Ahc; A_7 is Leu, Ahc, or Cha; A_8 is Ahc, Met, or Nle; A_{11} is Leu, Ahc, or Cha; A_{12} is Ahc or Gly; A_{15} is Leu, Ahc, or Cha; A_{18} is Met or Ahc; A_{21} is Val or Ahc; A_{22} is Glu or Ahc; A_{23} is Trp, Ahc, or Cha; A_{24} is Leu, Ahc, or Cha; A_{27} is Lys, hArg, Ahc, or Cha; A_{28} is Leu, Ahc, or Cha; A_{29} is Gln, Ahc or Aib; A_{31} is Val, Leu, Nle, Ahc, or Cha; R_1 is H; R_2 is H; and R_3 is NH_2; or a pharmaceutically acceptable salt thereof.

A preferred group of peptides of Group B is where at least one of A_7, A_{11}, A_{15}, A_{23}, A_{24}, A_{27}, A_{28}, or A_{31} is Cha.

Another preferred group of peptides of Group B is where at least one of A_{16}, A_{17}, A_{19}, A_{29}, or A_{34} is Aib.

Preferred peptides of formula (I) are $[Ahc^{7,11}]$hPTH(1–34)NH_2; $[Ahc^{7,11}, Nle^{8,18}, Tyr^{34}]$hPTH(1–34)$NH_2$; $[Ahc^{11}]$hPTH(1–34)NH_2; $[Ahc^{7,11,15}]$hPTH(1–34)NH_2;

[Ahc^{7}]hPTH(1–34)NH$_{2}$; [Ahc^{23}]hPTH(1–34)NH$_{2}$; [Ahc^{24}]hPTH(1–34)NH$_{2}$; [Nle^{8,18}, Ahc^{27}]hPTH(1–34)NH$_{2}$; [Ahc^{28}]hPTH(1–34)NH$_{2}$; [Ahc^{31}]hPTH(1–34)NH$_{2}$; [Ahc^{24,28,31}]hPTH(1–34)NH$_{2}$; [Ahc^{24,28,31}, Lys^{30}]hPTH (1–34)NH$_{2}$; [Ahc^{28,31}]hPTH(1–34)NH$_{2}$; [Ahc^{5}]hPTH (1–34)NH$_{2}$; [Ahc^{24,27}, Aib^{29}, Lys^{30}]hPTH(1–34)NH$_{2}$; [Ahc^{24,27}, Aib^{29}, Lys^{30}, Leu^{31}]hPTH(1–34)NH$_{2}$; [Ahc^{5}]hPTH(1–34)NH$_{2}$; [Ahc^{12}]hPTH(1–34)NH$_{2}$; [Ahc^{27}]hPTH (1–34)NH$_{2}$; [Ahc^{29}]hPTH(1–34)NH$_{2}$; [Ahc^{24,27}]hPTH (1–34)NH$_{2}$, [Ahc^{24,27}, Aib^{29}]hPTH(1–34)NH$_{2}$; [Ahc^{24}, Aib^{29}]hPTH(1–34)NH$_{2}$; [Ahc^{27}, Aib^{29}]hPTH(1–34)NH$_{2}$; [Ahc^{18}]hPTH(1–34)NH$_{2}$; [Ahc^{8}]hPTH(1–34)NH$_{2}$; [Ahc^{18,27}, Aib^{29}]hPTH(1–34)NH$_{2}$; or [Ahc^{18,24,27}, Aib^{29}]hPTH(1–34)NH$_{2}$; [Ahc^{22}, Leu^{27}, Aib^{29}]hPTH(1–34)NH$_{2}$; [Ahc^{24}, Leu^{27}, Aib^{29}]hPTH(1–34)NH$_{2}$; [Ahc^{22}]hPTH (1–34)NH$_{2}$; and [Ahc^{22}, Aib^{29}]hPTH(1–34)NH$_{2}$; or a pharmaceutically acceptable salt thereof.

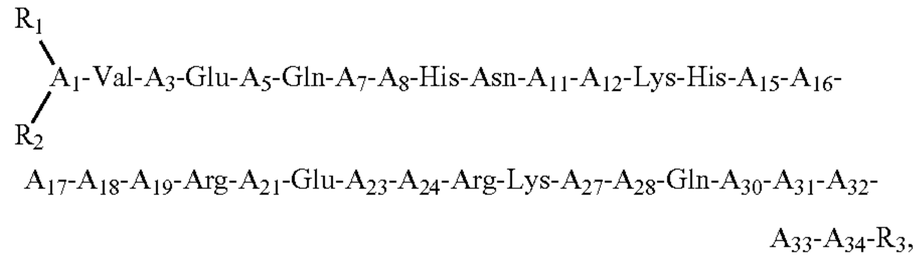
The invention also features peptides of the following formulae; [Cha^{22,23}, Glu^{25}, Lys^{26,30}, Leu^{28}, Aib^{29}]hPTHrP (1–34)NH$_{2}$; [Cha^{22,23}, Glu^{25}, Lys^{26,30}, Aib^{29}]hPTHrP(1–34) NH$_{2}$; [Glu^{22,23}, Leu^{25,28,31}, Lys^{26}, Aib^{29}, Nle^{30}]hPTHrP (1–34)NH$_{2}$; [Glu^{22,25}, Leu^{23,28,30,31}, Lys^{26}, Aib^{29}]hPTHrP (1–34)NH$_{2}$; [Glu^{22,25,29}, Leu^{23,28,30,31}, Lys^{26}]hPTHrP (1–34)NH$_{2}$; [Glu^{22,25,29}, Leu^{23,28,31}, Lys^{26}, Nle^{30}]hPTHrP (1–34)NH$_{2}$; [Ser^{1}, Ile^{5}, Met^{8}, Asn^{10}, Leu^{11,23,28,31}, His^{14}, Cha^{15}, Glu^{22,25}, Lys^{26,30}, Aib^{29}]hPTHrP(1–34)NH$_{2}$; [Glu^{22,25}, Cha^{23}, Lys^{26}, Leu^{28,31}, Aib^{29}, Nle^{30}]hPTHrP (1–34)NH$_{2}$; [Cha^{22,23}, Glu^{25}, Lys^{26,30}, Leu^{28,31}, Aib^{29}]hPTHrP(1–34)NH$_{2}$; [Cha^{22}, Leu^{23,28,31}, Glu^{25,29}, Lys^{26}, Nle^{30}]hPTHrP(1–34)NH$_{2}$; [Cha^{7,11,15}]hPTHrP(1–34)NH$_{2}$; [Cha^{7,8,15}]hPTHrP(1–34)NH$_{2}$; [Glu^{22}, Cha^{23}, Aib^{25,29}, Lys^{26,30}, Leu^{28,31}]hPTHrP(1–34)NH$_{2}$; [Glu^{22}, Cha^{23}, Aib25,29, Lys^{26}, Leu^{28}]hPTHrP(1–34)NH$_{2}$; [Glu^{22}, Leu^{23,28}, Aib^{25,29}, Lys^{26}]hPTHrP(1–34)NH$_{2}$; [Aib^{29}]hPTHrP(1–34)NH$_{2}$; [Glu^{22,25}, Cha^{23}, Lys^{26}, Leu^{28,31}, Aib^{29}, Nle^{30}]hPTHrP(1–34)NH$_{2}$; [Glu^{22,25}, Cha^{23}, Lys^{26,30}, Aib^{29}, Leu^{31}]hPTHrP(1–34)NH$_{2}$; [Glu^{22,25}, Leu^{23,28,31}, Lys^{26}, Aib^{29,30}]hPTHrP(1–34)NH$_{2}$; [Glu^{22,25}, Leu23,28,31, Lys^{26}, Aib^{29}]hPTHrP(1–34)NH$_{2}$; [Glu^{22,25}, Leu^{23,28,31}, Aib^{26,29}, Lys^{30}]hPTHrP(1–34)NH$_{2}$; [Glu^{22,25}, Cha^{23}, Lys^{26,30}, Leu^{28,31}, Aib^{29}]hPTHrP(1–34)NH$_{2}$; [Glu^{22,25}, Cha^{23}, Lys^{26,30}, Aib^{29}]hPTHrP(1–34)NH$_{2}$; [Glu^{22,25}, Cha^{23}, Lys^{26,30}, Leu^{28}, Aib^{29}]hPTHrP(1–34)NH$_{2}$; or [Leu^{27}, Aib^{29}]hPTHrP(1–34)NH$_{2}$; or a pharmaceutically acceptable salt thereof.

The following are examples of the peptides of the invention covered by the above formula: [Glu^{22,25}, Leu^{23,28}, Lys^{26,30}, Aib^{29}, Ahc^{31}]hPTHrP(1–34)NH$_{2}$; [Glu^{22,25}, Ahc^{23}, Lys^{26,30}, Leu^{28,31}, Aib^{29}]hPTHrP(1–34)NH$_{2}$, [Glu^{22,25}, Leu^{23,28,31}, Lys^{26,30}, Ahc^{27}, Aib^{29}]hPTHrP(1–34)NH$_{2}$; [Glu^{22,25,29}, Leu^{23,28,31}, Lys^{26}l Ahc^{30}]hPTHrP(1–34)NH$_{2}$; [Cha^{22}, Leu^{23,28,31}, Glu^{25}, Lys^{26,30}, Ahc^{27}, Aib^{29}]hPTHrP (1–34)NH$_{2}$; [Glu^{22,25}, Leu^{23,28,31}, Ahc^{24}, Lys^{26,30}, Aib^{29}]hPTHrP(1–34)NH$_{2}$; [Glu^{22,29}, Leu^{23,28,31}, Aib^{25}, Lys^{26,30}, Ahc^{27}]hPTHrP(1–34)NH$_{2}$; [Glu^{22}, Leu^{23,28,31}, Aib^{25,29}, Lys^{26,30}, Ahc^{27}]hPTHrP(1–34)NH$_{2}$; [Ahc^{22}, Leu23,28,31, Glu^{25}, Lys^{26,30}, Aib^{29}]hPTHrP(1–34)NH$_{2}$; [Glu^{22,25}, Leu^{23,31}, Lys^{26,30}, Ahc^{28}, Aib^{29}]hPTHrP(1–34)NH$_{2}$; [Cha^{22}, Ahc^{23}, Glu^{25}, Lys^{26,30}, Leu^{28,31}, Aib^{29}]hPTHrP (1–34)NH$_{2}$; [Ahc^{22,24,27}, Leu^{23,28,31}, Glu^{25}, Lys^{26,30}, Aib^{29}]hPTHrP(1–34)NH$_{2}$; [Cha^{22}, Leu^{23,28,31}, Ahc^{24,27}, Glu^{25}, Lys^{26,30}, Aib^{29}]hPTHrP(1–34)NH$_{2}$; [Glu^{22}, Leu^{23,28,31}, Ahc^{24,27}, Lys^{25,26}, Aib^{29}]hPTHrP(1–34)NH$_{2}$; [Ahc^{18,24,27}, Glu^{22}, Cha^{23}, Lys^{25,26}, Leu^{28}, Aib^{29}]hPTHrP(1–34)NH$_{2}$; [Glu^{22}, Cha^{23}, Ahc^{24}, Lys^{25,26}, Leu^{28}, Aib^{29}]hPTHrP(1–34) NH$_{2}$; [Glu^{22,25}, Leu^{23,28,31}, Lys^{26}, Ahc^{27}, Aib^{29}, Nle^{30}]hPTHrP(1–34)NH$_{2}$; [Ser^{1}, Ile^{5}, Cha^{7,11}, Met^{8}, Asn^{10}, His^{14}, Glu^{22,25}, Leu^{23,28,31}, Lys^{26,30}, Ahc^{27}, Aib^{29}]hPTHrP(1–34) NH$_{2}$; [Ser^{1}, Ile^{5}, Met^{8}, Asn^{10}, Leu^{11,23,28,31}, His^{14}, Cha^{15}, Glu^{22,25}, Lys^{26,30}, Ahc^{27}, Aib^{29}]hPTHrP(1–34)NH$_{2}$; [Cha^{22}, Ahc^{23}, Glu^{25}, Lys^{26,30}, Leu^{28,31}, Aib^{29}]hPTHrP(1–34)NH$_{2}$; [Glu^{22}, Ahc^{23}, Aib^{25,29}, Lys^{26,30}, Leu^{28,31}]hPTHrP(1–34) NH$_{2}$; [Glu^{22,25}, Leu^{23,28,31}, Lys^{26,30}, Ahc^{29}]hPTHrP(1–34) NH$_{2}$; [Cha^{22}, Leu^{23,28,31}, Ahc^{24}, Glu^{25}, Lys^{26,30}, Aib^{29}]hPTHrP(1–34)NH$_{2}$, [Cha^{22}, Leu^{23,28,31}, Ahc^{24,27}, Glu^{25}, Lys^{26,30}, Aib^{29}]hPTHrP(1–34)NH$_{2}$; [Glu^{22,25}, Leu^{23,28,31}, Ahc^{24,27}, Lys^{26,30}, Aib^{29}]hPTHrP(1–34)NH$_{2}$; [Ahc^{22,24,27}, Leu^{23,28,31}, Glu^{25}, Lys^{26,30}, Aib^{29}]hPTHrP(1–34)NH$_{2}$; [Cha^{22}, Leu^{23,28,31}, Aib^{25,29}, Lys^{26,30}, Ahc^{27}]hPTHrP(1–34) NH$_{2}$; [Ahc^{22,27}, Leu^{23,28,31}, Aib^{25,29}, Lys^{26,30}]hPTHrP (1–34)NH$_{2}$; [Glu^{22}, Leu^{23,28,31}, Ahc^{24,27}, Lys^{25,26,30}, Aib^{29}]hPTHrP 1–34)NH$_{2}$; [Glu^{22}, Leu^{23,28}, Ahc^{24,27}, Lys^{25,26,30}, Aib^{29}]hPTHrP(1–34)NH$_{2}$; [Glu^{22}, Cha^{23}, Ahc^{24,27}, Lys^{25,26,30}, Leu^{28}, Aib^{29}]hPTHrP(1–34)NH$_{2}$; [Glu^{22,25}, Cha^{23}, Ahc^{24,27}, Lys^{26,30}, Leu^{28,31}, Aib^{29}]hPTHrP(1–34) NH$_{2}$; [Glu^{22}, Cha^{23}, Ahc^{24,27}, Lys^{25,26,30}, Leu^{28,31}, Aib^{29}]hPTHrP(1–34)NH$_{2}$; [Glu^{22}, Leu^{23,28,31}, Ahc^{24,27}, Lys^{25,26}, Aib^{29}]hPTHrP(1–34)NH$_{2}$; [Glu^{22}, Leu^{23,28}, Ahc^{24,27}, Lys^{25,26}, Aib^{29}]hPTHrP(1–34)NH$_{2}$; [Glu^{22}, Cha^{23}, Ahc^{24,27}, Lys^{25,26}, Leu^{28,31}, Aib^{29}]hPTHrP(1–34)NH$_{2}$; [Glu^{22}, Cha^{23}, Ahc^{24,27}, Lys^{25,26}, Leu^{28}, Aib^{29}]hPTHrP (1–34)NH$_{2}$; [Glu^{22}, Leu^{23,28}, Lys^{25,26}, Ahc^{27}, Aib^{29}]hPTHrP(1–34)NH$_{2}$; [Glu^{22}, Leu^{23,28,31}, Lys^{25,26}, Ahc^{27}, Aib^{29}]hPTHrP(1–34) NH$_{2}$; [Glu^{22}, Leu^{23,28,31}, Lys^{25,26,30}, Ahc^{27}, Aib^{29}]hPTHrP (1–34)NH$_{2}$; [Glu^{22}, Leu^{23,28}, Lys^{25,26,30}, Ahc^{27}, Aib^{29}]hPTHrP(1–34)NH$_{2}$; [Glu^{22}, Cha^{23}, Ahc^{24}, Lys^{25,26}, Leu^{28}, Aib^{29}]hPTHrP(1–34)NH$_{2}$; [Glu^{22,25}, Leu^{23,28,31}, Lys^{26}, Aib^{29}, Ahc^{30}]hPTHrP(1–34)NH$_{2}$; [Aib^{22,29}, Leu^{23,28,31}, Glu^{25}, Lys^{26,30}]hPTHrP(1–34)NH$_{2}$; [Cha^{22}, Ahc^{23}, Glu^{25,29}, Lys^{26,30}, Leu^{28,31}]hPTHrP(1–34) NH$_{2}$[Cha^{22}, Leu^{23,28,31}, Ahc^{24}, Glu^{25,29}, Lys^{26,30}]hPTHrP (1–34)NH$_{2}$, [Cha^{22}, Leu^{23,28,31}, Glu^{25,29}, Lys^{26,30}, Ahc^{27}]hPTHrP(1–34) NH$_{2}$, [Cha^{22}, Leu^{23,31}, Glu^{25,29}, Lys^{26,30}, Ahc^{28}]hPTHrP (1–34)NH$_{2}$; [Cha^{22}, Leu^{23,28,31}, Glu^{25,29}, Lys^{26}, Ahc^{30}]hPTHrP(1–34)NH$_{2}$; [Cha^{22}, Leu^{23,28}, Glu^{25,29}, Lys^{26,30}, Ahc^{31}]hPTHrP(1–34)NH$_{2}$; [Glu^{22,29}, Ahc^{23}, Aib^{25}, Lys^{26,30}, Leu^{28,31}]hPTHrP(1–34)NH$_{2}$, [Ahc^{22}, Leu^{23,28,31}, Aib^{25}, Lys^{26,30}, Glu^{29}]hPTHrP(1–34) NH$_{2}$; [Glu^{22,29}, Leu^{23,28,31}, Ahc^{24}, Aib^{25}, Lys^{26,30}]hPTHrP (1–34)NH$_{2}$; [Glu^{22,29}, Leu^{23,31}, Aib^{25}, Lys^{26,30}, Ahc^{28}]hPTHrP(1–34) NH$_{2}$; [Glu^{22,29}, Leu^{23,28}, Aib^{25}, Lys^{26,30}, Ahc31]hPTHrP (1–34)NH$_{2}$; [Glu^{22,29}, Leu^{23,28,31}, Aib^{25}, Lys^{26}, Ahc^{30}]hPTHrP(1–34)NH$_{2}$; [Glu^{22,25,29}, Leu^{23,28,31}, Lys^{26}, Ahc^{27}, Aib^{30}]hPTHrP(1–34)NH$_{2}$; [Glu^{22,25,29}, Leu^{23,28,31}, Ahc^{24}, Lys^{26}, Aib^{30}]hPTHrP(1–34)NH$_{2}$; [Ahc^{22}, Leu^{23,28,31}, Glu^{25,29}, Lys^{26}, Aib^{30}]hPTHrP(1–34) NH$_{2}$; [Ahc^{22}, Leu^{23,28}, Glu^{25,29}, Lys^{26,30,31}]hPTHrP(1–34) NH$_{2}$; [Glu^{22,25,29}, Leu^{23,28}, Lys^{26,31}, Ahc^{30}]hPTHrP(1–34) NH$_{2}$; [Glu^{22,25,29}, Leu^{23,28}, Lys^{26,30,31}, Ahc^{27}]hPTHrP (1–34) NH$_{2}$; [Ahc^{22}, Cha^{23}, Glu^{25}, Lys^{26,30}, Leu^{28,31}, Aib^{29}]hPTHrP(1–34)NH$_{2}$; [Ahc^{22}, Cha^{23}, Lys^{25,26,30}, Leu^{28,31}, Aib^{29}]hPTHrP(1–34)NH$_{2}$; [Ahc^{22}, Cha^{23}, Lys^{25,26}, Leu^{28,31}, Aib^{29}]hPTHrP(1–34)NH$_{2}$, [Ahc^{22}, Leu^{23,28}, Lys^{25,26}, Aib^{29}]hPTHrP(1–34)NH$_{2}$, [Ahc^{22}, Leu^{23,28}, Arg^{25}, Lys^{26}, Aib^{29}]hPTHrP(1–34)NH$_{2}$; [Ahc^{22,24}, Leu^{23,28,31}, Glu^{25}, Lys^{26,30}, Aib^{29}]hPTHrP(1–34)NH$_{2}$; [Ahc^{22,24}, Leu$^{23,28\ 31}$, Lys^{25,26,30}, Aib^{29}]hPTHrP(1–34)NH$_{2}$; [Ahc^{22,24}, Leu^{23,28,31}, Lys^{25,26}, Aib^{29}]hPTHrP(1–34)NH$_{2}$; [Ahc^{22,24}, Leu^{23,28}, Lys^{25,26}, Aib^{29}]hPTHrP(1–34)NH$_{2}$; [Ahc^{22,24}, Leu^{23,28}, Arg^{25}, Lys^{26}, Aib^{29}]hPTHrP(1–34)NH$_{2}$; [Glu^{22}, Leu^{23,28,31}, Ahc^{24}, Lys^{25,26,30}, Aib^{29}]hPTHrP(1–34) NH$_{2}$; **⟦**[Glu^{22}, Leu^{23,28,31}, Ahc^{24}, Lys^{25,26}, Aib^{2}]hPTHrP (1–34)NH$_{2}$**⟧** *[Glu^{22}, Leu^{23,28,31}, Ahc^{24}, Lys^{25,26}, Aib^{29}] hPTHrP(1–34)NH$_{2}$*; [Glu^{22}, Leu^{23,28}, Ahc^{24}, Lys^{25,26},

Aib[29]]hPTHrP(1–34)NH_2; [Glu[22], Leu[23,28,31], Ahc[24], Arg[25], Lys[26,30], Aib[29]]hPTHrP(1–34)NH_2; [Glu[22], Leu[23,28,31], Ahc[24], Arg[25], Lys[26], Aib[29]]hPTHrP(1–34)NH_2; [Glu[22], Leu[23,28], Ahc[24], Arg[25], Lys[26], Aib[29]]hPTHrP(1–34)NH_2, [Glu[22], Ahc[23], Aib[25,29], Lys[26,30], Leu[28,31]]hPTHrP(1–34)NH_2; [Glu[22], Ahc[23], Aib[25,29], Lys[26], Leu[28]]hPTHrP(1–34)NH_2; [Glu[22], Ahc[23,31], Aib[25,29], Lys[26], Leu[28]]hPTHrP(1–34)NH_2; [Glu[22], Leu[23,28], Aib[25,29], Lys[26,30], Ahc[31]]hPTHrP(1–34)NH_2; [Glu[22], Leu[23,28], Aib[25,29], Lys[26], Ahc[31]]hPTHrP(1–34)NH_2; [Glu[22,25], Leu[23,28], Ahc[24,31], Lys[26,30], Aib[29]]hPTHrP(1–34)NH_2; or [Glu[22], Leu[23,28], Ahc[24,31], Lys[25,26], Aib[29]]hPTHrP(1–34)NH_2; [Glu[22], Leu[23,28,31], Ahc[24], Aib[25,29], Lys[26,30]]hPTHrP(1–34)NH_2; or a pharmaceutically acceptable salt thereof.

In another aspect, the invention relates to peptide variants of PTH(1–34) of the following generic formula:

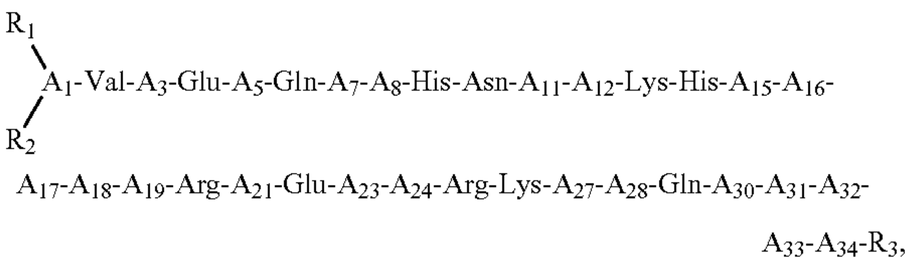
[SEQ ID NO: 2]



wherein
A_1 is Ser, Ala, or Dap;
A_3 is Ser, Thr, or Aib;
A_5 is Leu, Nle, Ile, Cha, β-Nal, Trp, Pal, Phe or p-X-Phe, in which X is OH, a halogen, or CH_3;
A_7 is Leu, Nle, Ile, Cha, β-Nal, Trp, Pal, Phe, or p-X-Phe in which X is OH, a halogen, or CH_3;
A_8 is Met, Nva, Leu, Val, Ile, Cha, or Nle;
A_{11} is Leu, Nle, Ile, Cha, β-Nal, Trp, Pal, Phe or p-X-Phe in which X is OH, a halogen, or CH_3;
A_{12} is Gly or Aib;
A_{15} is Leu, Nle, Ile, Cha, β-Nal, Trp, Pal, Phe, or p-X-Phe in which X is OH, a halogen, or CH_3;
A_{16} Is Ser, Asn, Ala, or Aib;
A_{17} is Ser, Thr, or Aib;
A_{18} is Met, Nva, Leu, Val, Ile, Nle, Cha, or Aib;
A_{19} is Glu or Aib;
A_{21} is Val, Cha, or Met;
A_{23} is Trp or Cha;
A_{24} is Leu or Cha,
A_{27} is Lys, Aib, Leu, hArg, Gln, or Cha;
A_{28} is Leu or Cha;
A_{30} is Asp or Lys;
A_{31} is Val, Nle, Cha, or deleted;
A_{32} is His or deleted;
A_{33} is Asn or deleted;
A_{34} is Phe, Tyr, Amp, Aib, or deleted;
each of R_1 and R_2 is, independently, H, C_{1-12} alkyl, C_{2-12} alkenyl, C_{7-20} phenylalkyl, C_{11-20} naphthylalkyl, C_{1-12} hydroxyalkyl, C_{2-12} hydroxyalkenyl, C_{7-20} hydroxyphenylalkyl, or C_{11-20} hydroxynaphthylalkyl; or one and only one of R_1 and R_2 is COE_1 in which E_1 is C_{1-12} alkyl, C_{2-12} alkenyl, C_{7-20} phenylalkyl, C_{11-20} naphthylalkyl, C_{1-12} hydroxyalkyl, C_{2-12} hydroxyalkenyl, C_{7-20} hydroxyphenylalkyl, or C_{11-20} hydroxynaphthylalkyl; and
R_3 is OH, NH_2, C_{1-12} alkoxy, or NH—Y—CH_2-Z in which Y is a C_{1-12} hydrocarbon moiety and Z is H, OH, CO_2H, or $CONH_2$;
provided that (i) at least one of A_5, A_7, A_8, A_{11}, A_{15}, A_{18}, A_{21}, A_{23}, A_{24}, A_{27}, A_{28}, and A_{31} is Cha, or at least one of A_3, A_{12}, A_{16}, A_{17}, A_{18}, A_{19}, and A34 is Aib; or that (ii) at least A_1 is Dap, A_7 is β-Nal, Trp, Pal, Phe, or p-X-Phe, A_{15} is β-Nal, Trp, Pal, Phe, or p-X-Phe, A_{27} is hArg, or A_{31} is Nle; or a pharmaceutically acceptable salt thereof.

In another aspect, the invention relates to peptide variants of PTH(1–34) of the following formula (II):

[SEQ ID NO: 2]



wherein
A_1 is Ser, Ala, or Dap;
A_3 is Ser, Thr, or Aib;
A_5 is Leu, Nle, Ile, Cha, β-Nal, Trp, Pal, Phe or p-X-Phe, in which X is OH, a halogen, or CH_3;
A_7 is Leu, Nle, Ile, Cha, β-Nal, Trp, Pal, Phe, or p-X-Phe in which X is OH, a halogen, or CH_3;
A_8 is Met, Nva, Leu, Val, Ile, Cha, or Nle;
A_{11} is Leu, Nle, Ile, Cha, β-Nal, Trp, Pal, Phe or p-X-Phe in which X is OH, a halogen, or CH_3;
A_{12} is Gly or Aib;
A_{15} is Leu, Nle, Ile, Cha, β-Nal, Trp, Pal, Phe, or p-X-Phe in which X is OH, a halogen, or CH_3;
A_{16} is Ser, Asn, Ala, or Aib;
A_{17} is Ser, Thr, or Aib;
A_{18} is Met, Nva, Leu, Val, Ile, Nle, Cha, or Aib;
A_{19} is Glu or Aib;
A_{21} is Val, Cha, or Met;
A_{23} is Trp or Cha;
A_{24} is Leu or Cha;
A_{27} is Lys, Aib, Leu, hArg, Gln, or Cha;
A_{28} is Leu or Cha;
A_{30} is Asp or Lys;
A_{31} is Val, Nle, Cha, or deleted;
A_{32} is His or deleted;
A_{33} is Asn or deleted;
A_{34} is Phe, Tyr, Amp, Aib, or deleted,
each of R_1 and R_2 is, independently, H, C_{1-12} alkyl, C_{2-12} alkenyl, C_{7-20} phenylalkyl, C_{11-20} naphthylalkyl, C_{1-12} hydroxyalkyl, C_{2-12} hydroxyalkenyl, C_{7-20} hydroxyphenylalkyl, or C_{11-20} hydroxynaphthylalkyl; or one and only one of R_1 and R_2 is COE_1 in which E_1 is C_{1-12} alkyl, C_{2-12} alkenyl, C_{7-20} phenylalkyl, C_{11-20} naphthylalkyl, C_{1-12} hydroxyalkyl, C_{2-12} hydroxyalkenyl, C_{7-20} hydroxy-phenylalkyl, or C_{11-20} hydroxynaphthylalkyl; and
R_3 is OH, NH_2, C_{1-12} alkoxy, or NH—Y—CH_2-Z in which Y is a C_{1-12} hydrocarbon moiety and Z is H, OH, CO_2H, or $CONH_2$;
provided that (i) at least one of A_5, A_7, A_8, A_{11}, A_{15}, A_{18}, A_{21}, A_{23}, A_{24}, A_{27}, A_{28}, and A_{31} is Cha, or at least one of A_3, A_{12}, A_{16}, A_{17}, A_{18}, A_{19}, and A_{34} is Aib, and the peptide is not [Aib[12], Tyr[34]]hPTH(1–34)NH_2 or a pharmaceutically acceptable salt thereof.

A preferred group of peptides of formula (II), designated Group (i) is where at least one of A_7, A_{11}, A_{15}, A_{23}, A_{24}, A_{27}, A_{28}, and A_{31} is Cha; or a pharmaceutically acceptable salt thereof.

A preferred group of peptides of Group (i), designated Group (ii), is where A_3 is Ser; A_5 is Ile; A_7 is Leu or Cha; A_8 is Met, Nva, Leu, Val, Ile, or Nle; A_{12} is Leu or Cha; A_{12} is

Gly; A_{15} is Leu or Cha; A_{16} is Asn or Aib; A_{17} is Ser; A_{18} is Met or Nle. A_{21} is Val; A_{27} is Lys, hArg, or Cha; A_{32} is His; A_{31} is Val, Nle, or Cha; A_{33} is Asn; A_{34} is Phe, Tyr, Amp, or Aib; R_1 is H; R_2 is H, and R_3 is NH_2; or a pharmaceutically acceptable salt thereof.

A preferred group of peptides of Group (ii), designated Group (iii), is where at least one of A_7 and A_{11} is Cha, or a pharmaceutically acceptable salt thereof.

Preferred peptides of Group (iii) are [$Cha^{7,11}$]hPTH (1–34)NH_2, [$Cha^{7,11}$, $Nle^{8,18}$, Tyr^{34}]hPTH(1–34)NH_2; [Cha^{11}]hPTH(1–34)NH_2; [$Cha^{7,11,15}$]hPTH(1–34)NH_2; and [Cha^{7}]hPTH(1–34)NH_2; or a pharmaceutically acceptable salt thereof.

Another preferred group of peptides of Group (ii), designated Group (iv), is where at least one of A_{15}, A_{23}, A_{24}, A_{27}, A_{28}, and A_{31} is Cha; or a pharmaceutically acceptable salt thereof.

Preferred peptides of Group (iv) are [Cha^{23}]hPTH(1–34) NH_2, [Cha^{24}]hPTH(1–34)NH_2, [$Nle^{8,18}$, Cha^{27}]hPTH (1–34)NH_2, [Cha^{28}]hPTH(1–34)NH_2, [Cha^{31}]hPTH(1–34) NH_2, [$Cha^{24,28,31}$]hPTH(1–34)NH_2; [$Cha^{24,28,31}$, Lys^{30}] hPTH(1–34)NH_2; [$Cha^{28,31}$]hPTH(1–34)NH_2; and [Cha^{15}] hPTH(1–34)NH_2; or a pharmaceutically acceptable salt thereof.

Another preferred group of peptides of formula (II), designated Gropu (v), is where at least one of A_3, A_{12}, A_{16}, A_{17}, A_{18}, A_{19}, and A_{34} is Aib; or a pharmaceutically acceptable salt thereof.

A preferred group of peptides of Group (v), designated Group (vi), is where A_3 is Ser or Aib; A_5 is Ile, A_7 is Leu or Cha; A_8 is Met, Nva, Leu, Val, Ile, or Nle; A_{11} is Leu or Cha; A_{15} is Leu or Cha; A_{16} is Asn or Aib; A_{18} is Met, Aib, or Nle; A_{21} is Val; A_{27} is Lys, Aib, Leu, hArg, or Cha; A_{31} is Val, Nle, or Cha; A_{32} is His; A_{33} is Asn; A_{34} is Phe, Tyr, Amp, or Aib; R_1 is H; R_2 is H, and R_3 is NH_2; or a pharmaceutically acceptable salt thereof.

A preferred group of peptides of Group (vi), designated Group (vii), is where at least one of A_3, A_{12}, A_{16}, A_{17}, A_{19}, and A_{34} is Aib; or a pharmaceutically acceptable salt thereof.

Preferred peptides of Group (vii) are [Aib^{16}]hPTH(1–34) NH_2, [Aib^{19}]hPTH(1–34)NH_2[Aib^{34}]hPTH(1–34)NH_2; [$Aib^{16,19}$]hPTH(1–34)NH_2; [Aib]hPTH(1–34)NH_2, [Aib^{17}] hPTH(1–34)NH_2; and [Aib^{12}]hPTH(1–34)NH_2; or a pharmaceutically acceptable salt thereof.

Another preferred group of peptides of formula (II), designated Group (viii), is where at least one of A_7, A_{11}, A_{15}, A_{23}, A_{24}, A_{27}, A_{28}, and A_{31} is Cha and at least one of A_3, A_{12}, A_{16}, A_{17}, A_{18}, A_{19}, and A_{34} is Aib; or a pharmaceutically acceptable salt thereof.

A preferred group of peptides of Group (viii), designated Group (ix), is where A_3 is Ser or Aib; A_5 is Ile; A_7 is Leu or Cha; A_8 is Met, Nva, Leu, Val, Ile, or Nle; A_{11} is Leu or Cha; A_{15} is Leu or Cha; A_{16} is Asn or Aib; A_{18} is Met, Aib, or Nle, A_{21} is Val; A_{27} is Lys, Aib, Leu, hArg, or Cha; A_{31} is Val, Nle, or Cha; A_{32} is His; A_{33} is Asn; A_{34} is Phe, Tyr, Amp, or Aib; R_1 is H, R_2 is H; and R_3 is NH_2, or a pharmaceutically acceptable salt thereof.

A preferred group of peptides of Group (ix), designated Group (x), is where at least one of A_7 and A_{11} is Cha and at least one of A_{16}, A_{19}, and A_{34} is Aib; or a pharmaceutically acceptable salt thereof.

Preferred peptides of Group (x) are [$Cha^{7,11}$, $Nle^{8,18}$, $Aib^{16,19}$, Tyr^{34}]hPTH(1–34)NH_2, [$Cha^{7,11}$, $Nle^{8,18,31}$, $Aib^{16,}{}_{19}$, Tyr^{34}]hPTH(1–34)NH_2, [$Cha^{7,11}$, Aib^{19}]hPTH (1–34) NH_2; [$Cha^{7,11}$, Aib^{16}]hPTH(1–34)NH_2; [$Cha^{7,11}$, $Nle^{8,18}$, Aib^{34}]hPTH(1–34)NH_2; or [$Cha^{7,11}$, Aib^{19}, Lys^{30}] hPTH (1–34)NH_2; or a pharmaceutically acceptable salt thereof.

Another preferred group of peptides of Group (ix), designated Group (xi), is where at least one of A_{24}, A_{28}, and A_{31} is Cha and at least one of A_{16} and A_{17} is Aib; or a pharmaceutically acceptable salt thereof.

Preferred peptides of Group (xi) are [Cha^{28}, $Nle^{8,18}$, $Aib^{16,19}$, Tyr^{34}]hPTH(1–34)NH_2, and [Cha^{28}, $Aib^{16,19}$]PTH (1–34)NH_2; or a pharmaceutically acceptable salt thereof.

In another aspect, the present invention is directed to a peptide of the formula (III):

[SEQ ID NO: 2]

R_1
\
A_1-Val-A_3-Glu-A_5-Gln-A_7-A_8-His-Asn-A_{11}-A_{12}-Lys-His-A_{15}-A_{16}-
/
R_2

A_{17}-A_{18}-A_{19}-Arg-A_{21}-Glu-A_{23}-A_{24}-Arg-Lys-A_{27}-A_{28}-Gln-A_{30}-A_{31}-A_{32}-A_{33}-A_{34}-R_3,

wherein

A_3 is Ser, Thr, or Aib;
A_5 is Leu, Nle, Ile, Cha, β-Nal, Trp, Pal, Phe or p-X-Phe, in which X is OH, a halogen, or CH_3;
A_7 is Leu, Nle, Ile, Cha, β-Nal, Trp, Pal, Phe, or p-X-Phe in which X is H, OH, a halogen, or CH_3;
A_8 is Met, Nva, Leu, Val, Ile, Cha, or Nle;
A_{11} is Leu, Nle, Ile, Cha, β-Nal, Trp, Pal, Phe or p-X-Phe in which X is OH, a halogen, or CH_3;
A_{12} is Gly or Aib;
A_{15} is Leu, Nle, Ile, Cha, β-Nal, Trp, Pal, Phe, or p-X-Phe in which X is OH, a halogen, or CH_3;
A_{16} is Ser, Asn, Ala, or Aib;
A_{17} is Ser, Thr, or Aib;
A_{18} is Met, Nva, Leu, Val, Ile, Nle, Cha, or Aib;
A_{19} is Glu or Aib;
A_{21} is Val, Cha, or Met;
A_{23} is Trp or Cha,
A_{24} is Leu or Cha;
A_{27} is Lys, Aib, Leu, hArg, Gln, or Cha;
A_{28} is Leu or Cha;
A_{30} is Asp or Lys;
A_{31} is Val, Nle, Cha, or deleted;
A_{32} is His or deleted;
A_{33} is Asn or deleted;
A_{34} is Phe, Tyr, Amp, Aib, or deleted;
each of R_1 and R_2 is, independently, H, $C_{1\text{-}12}$ alkyl, $C_{2\text{-}12}$ alkenyl, $C_{7\text{-}20}$ phenylalkyl, $C_{11\text{-}20}$ naphthylalkyl, $C_{1\text{-}12}$ hydroxyalkyl, $C_{2\text{-}12}$ hydroxyalkenyl, $C_{7\text{-}20}$ hydroxyphenylalkyl, or $C_{11\text{-}20}$ hydroxynaphthylalkyl; or one and only one of R_1 and R_2 is COE_1 in which E_1 is $C_{1\text{-}12}$ alkyl, $C_{2\text{-}12}$ alkenyl, $C_{7\text{-}20}$ phenylalkyl, $C_{11\text{-}20}$ naphthylalkyl, $C_{1\text{-}12}$ hydroxyalkyl, $C_{2\text{-}12}$ hydroxyalkenyl, $C_{7\text{-}20}$ hydroxy-phenylalkyl, or $C_{11\text{-}20}$ hydroxynaphthylalkyl;
R_3 is OH, NH_2, $C_{1\text{-}12}$ alkoxy, or NH—Y—CH_2-Z in which Y is a $C_{1\text{-}12}$ hydrocarbon moiety and Z is H, OH, CO_2H, or $CONH_2$;

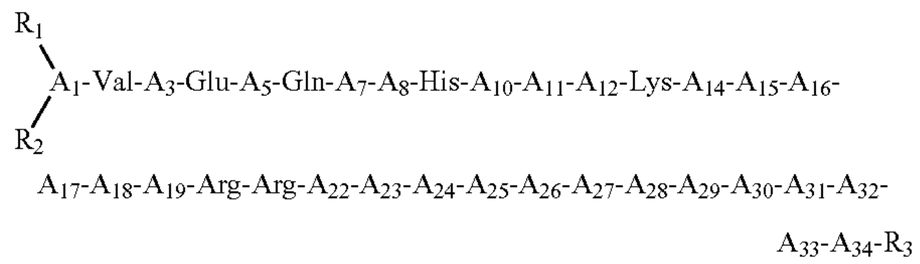
provided that at least one of A_1 is Dap, A_7 is β-Nal, Trp, Pal, Phe, or p-X-Phe; A_{15} is β-Nal, Trp, Pal, Phe, or p-X-Phe, A_{27} is hArg, or A_{31} is Nle; or a pharmaceutically acceptable salt thereof.

A preferred group of peptides of formula (III) is where A_1 is Ser, Gly, or Dap; A_3 is Ser or Aib; A_8 is Met, Nva, Leu, Val, Ile, or Nle; A_{16} is Asn or Aib; A_{18} is Met, Aib, or Nle; A_{21} is Val; A_{27} is Lys, Aib, Leu, hArg, or Cha; A_{31} is Val, Nle, or Cha; A_{32} is His; A_{33} is Asn; A_{34} is Phe, Tyr, Amp, or Aib; R_1 is H; R_2 is H; and R_3 is NH_2, or a pharmaceutically acceptable salt thereof.

Preferred peptides of the immediately foregoing peptides are [Nle^{31}]hPTH(1–34)NH_2, [hArg^{27}]hPTH(1–34)NH_2, and [Dap^{1}, Nle^{8,18}, Tyr^{34}]hPTH(1–34)NH_2; or a pharmaceutically acceptable salt thereof.

In another aspect, the present invention is directed to a peptide of the formula (IV):

[SEQ ID NO: 3]



R_1
\
A_1-Val-A_3-Glu-A_5-Gln-A_7-A_8-His-A_{10}-A_{11}-A_{12}-Lys-A_{14}-A_{15}-A_{16}-
/
R_2

A_{17}-A_{18}-A_{19}-Arg-Arg-A_{22}-A_{23}-A_{24}-A_{25}-A_{26}-A_{27}-A_{28}-A_{29}-A_{30}-A_{31}-A_{32}-

A_{33}-A_{34}-R_3

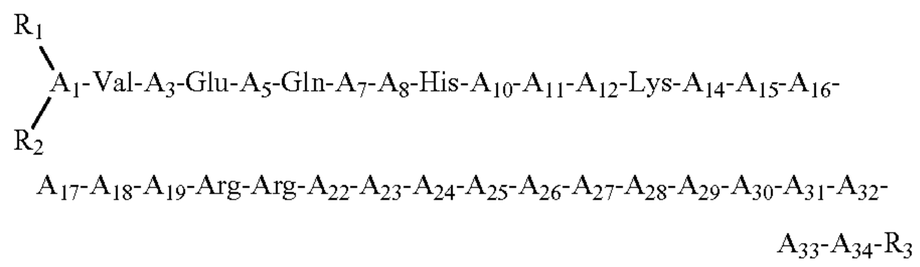
wherein
A_1 is Ala, Ser, or Dap;
A_3 is Ser or Aib;
A_5 is His, Ile, or Cha;
A_7 is Leu, Cha, Nle, β-Nal, Trp, Pal, Phe, or p-X-Phe in which X is OH, a halogen, or CH_3;
A_8 is Leu, Met, or Cha;
A_{10} is Asp or Asn;
A_{11} is Lys, Leu, Cha, Phe, or β-Nal;
A_{12} is Gly or Aib;
A_{14} is Ser or His;
A_{15} is Ile, or Cha;
A_{16} is Gln or Aib,
A_{17} is Asp or Aib;
A_{18} is Leu, Aib, or Cha;
A_{19} is Arg or Aib;
A_{22} is Phe, Glu, Aib, or Cha;
A_{23} is Phe, Leu, Lys, or Cha;
A_{24} is Leu, Lys, or Cha;
A_{25} is His, Aib, or Glu;
A_{26} is His, Aib, or Lys;
A_{27} is Leu, Lys, or Cha;
A_{28} is Ile, Leu, Lys, or Cha;
A_{29} is Ala, Glu, or Aib;
A_{30} is Glu, Cha, Aib, or Lys;
A_{31} is Ile, Leu, Cha, Lys, or deleted;
A_{32} is His or deleted;
A_{33} is Thr or deleted;
A_{34} is Ala or deleted;
each of R_1 and R_2 is, independently, H, $C_{1\text{-}12}$ alkanyl, $C_{7\text{-}20}$ phenylalkyl, $C_{11\text{-}20}$ naphthylalkyl, $C_{1\text{-}12}$, hydroxyalkyl, $C_{2\text{-}12}$ hydroxyalkenyl, $C_{7\text{-}20}$ hydroxyphenylalkyl, or $C_{11\text{-}20}$ hydroxynaphthylalkyl; or one and only one of R_1 and R_2 is COE_1 in which E_1 is $C_{1\text{-}12}$ alkyl, $C_{2\text{-}12}$ alkyl, $C_{2\text{-}12}$ alkenyl, $C_{7\text{-}20}$ phenylalkyl, $C_{11\text{-}20}$ naphthylalkyl, $C_{1\text{-}12}$ hydroxyalkyl, $C_{2\text{-}12}$ hydroxyalkenyl, $C_{7\text{-}20}$ hydroxyphenylalkyl, or $C_{11\text{-}20}$ hydroxynaphthylalkyl; and
R_3 is OH, NH_2, $C_{1\text{-}12}$ alkoxy, or NH—Y—CH_2-Z in which Y is a $C_{1\text{-}12}$ hydrocarbon moiety and Z is H, OH, CO_2H, or $CONH_2$;
provided that at least one of A_5, A_7, A_8, A_{11}, A_{15}, A_{18}, A_{22}, A_{23}, A_{24}, A_{27}, A_{28}, A_{30}, or A_{31} is Cha, or at least one of A_3, A_{12}, A_{16}, A_{17}, A_{18}, A_{19}, A_{22}, A_{25}, A_{26}, A_{29}, A_{30}, or A_{34} is Aib; or a pharmaceutically acceptable salt thereof.

A preferred group of peptides of formula (IV) is where A_{22} is Phe or Cha; A_{23} is Phe or Cha; A_{25} is His; A_{26} is His; A_{27} is Leu or Cha; A_{28} is Ile or Cha; A_{29} is Ala; A_{30} is Glu or Lys; A_{31} is Ile or Cha; A_{32} is His; A_{33} is Thr; and A_{34} is Ala; or a pharmaceutically acceptable salt thereof. Two preferred groups of peptides of the immediately foregoing group of peptides is where at least one of A_7 and A_{11} is Cha; or where at least one of A_{16} or A_{19} is Aib; or a pharmaceutically acceptable salt thereof.

Another preferred group of peptides of formula (IV), is where A_{22} is Glu, Aib, or Cha; A_{23} is Leu, Lys, or Cha; A_{25} is Aib or Glu; A_{26} is Aib or Lys; A_{28} is Leu, Lys, or Cha; A_{29} is Glu or Aib; A_{30} is Cha, Aib, or Lys, A_{31} is Leu, Cha, or Lys A_{32} is His; A_{33} is Thr; and A_{34} is Ala; or a pharmaceutically acceptable salt thereof. Two preferred groups of peptides of the immediately foregoing group of peptides is where at least one of A_7 and A_{11} is Cha; or where at least one of A_{16} or A_{19} is Aib; or a pharmaceutically acceptable salt thereof.

In another aspect, this invention is directed to a peptide of the formula (V):

[SEQ ID NO: 4]



R_1
\
A_1-Val-A_3-Glu-A_5-Gln-A_7-A_8-His-A_{10}-A_{11}-A_{12}-Lys-A_{14}-A_{15}-A_{16}-
/
R_2

A_{17}-A_{18}-A_{19}-Arg-Arg-A_{22}-A_{23}-A_{24}-A_{25}-A_{26}-A_{27}-A_{28}-A_{29}-A_{30}-A_{31}-A_{32}-

A_{33}-A_{34}-R_3

wherein
A_1 is Ala, Ser or Dap,
A_3 is Ser or Aib,
A_5 is His, Ile or Cha,
A_7 is Leu, Cha, Nle, β-Nal, Trp, Pal, Phe, or p-X-Phe in which X is OH, a halogen, or CH_3;
A_8 is Leu, Met or Cha;
A_{10} is Asp or Asn;
A_{11} is Lys, Leu, Cha, Phe or β-Nal;
A_{12} is Gly or Aib;
A_{14} is Ser or His;
A_{15} is Ile or Cha;
A_{16} is Gln or Aib;
A_{17} is Asp or Aib;
A_{18} is Leu, Aib or Cha;
A_{19} is Arg or Aib;
A_{22} is Phe, Glu, Aib, Acc or Cha;
A_{23} is Phe, Leu, Lys, Acc or Cha;
A_{24} is Leu, Lys, Acc or Cha;
A_{25} is His, Aib or Glu;
A_{26} is His, Aib or Lys,
A_{27} is Leu, Lys, Acc or Cha,
A_{28} is Ile, Leu, Lys, Acc or Cha;
A_{29} is Ala, Glu or Aib;
A_{30} is Glu, Cha, Aib, Acc or Lys;
A_{31} is Ile, Leu, Cha, Lys, Acc or deleted;
A_{32} is His or deleted;
A_{33} is Thr or deleted;
A_{34} is Ala or deleted;
each of R_1 and R_2 is, independently, H, $C_{1\text{-}12}$ alkanyl, $C_{7\text{-}20}$ phenylalkyl, $C_{11\text{-}20}$ naphthylalkyl, $C_{1\text{-}12}$, hydroxyalkyl, $C_{2\text{-}12}$ hydroxyalkenyl, $C_{7\text{-}20}$ hydroxyphenylalkyl, or $C_{11\text{-}20}$ hydroxynaphthylalkyl; or one and only one of R_1 and R_2 is COE_1 in which E_1 is $C_{1\text{-}12}$ alkyl, $C_{2\text{-}12}$ alkyl, $C_{2\text{-}12}$ alkenyl, $C_{7\text{-}20}$ phenylalkyl, $C_{11\text{-}20}$ naphthylalkyl, $C_{1\text{-}12}$ hydroxyalkyl, $C_{2\text{-}12}$ hydroxyalkenyl, $C_{7\text{-}20}$ hydroxyphenylalkyl, or $C_{11\text{-}20}$ hydroxynaphthylalkyl; and
R_3 is OH, NH_2, $C_{1\text{-}12}$ alkoxy, or NH—Y—CH_2-Z in which Y is a $C_{1\text{-}12}$ hydrocarbon moiety and Z is H, OH, CO_2H, or $CONH_2$;
provided that at least one of A_{23}, A_{24}, A_{27}, A_{28}, or A_{31} is Lys; or a pharmaceutically acceptable salt thereof.

A preferred group of peptides of formula (V) is where A_{22} is Glu, Aib, Acc, or Cha; A_{23} is Leu, Lys, Acc, or Cha; A_{25} is

Aib or Glu; A_{26} is Aib or Lys, A_{28} is Leu, Lys, Acc, or Cha; A_{29} is Glu or Aib; A_{30} is Cha, Aib, Acc, or Lys; A_{31} is Leu, Cha, Acc, or Lys; A_{32} is His; A_{33} is Thr; and A_{34} is Ala; or a pharmaceutically acceptable salt thereof. Two preferred groups of peptides of the immediately foregoing group of peptides is where at least one of A_7 and A_{11} is Cha; or where at least one of A_{16} or A_{19} is Aib, or a pharmaceutically acceptable salt thereof.

The following are examples of peptides of this invention as encompassed by formula (II); [Cha^{7}]hPTH(1–34)NH_2; [Cha^{11}]hPTH(1–34)NH_2; [Cha^{15}]hPTH(1–34)NH_2, [$Cha^{7,11}$]hPTH(1–34)NH_2; [$Cha^{7,11}$, $Nle^{8,18}$, Tyr^{34}]hPTH(1–34)NH_2; [Cha^{23}]hPTH(1–34)NH_2; [Cha^{24}]hPTH(1–34)NH_2; [$Nle^{8,18}$, Cha^{27}]hPTH(1–34)NH_2, [Cha^{28}]hPTH(1–34)NH_2; [Cha^{31}]hPTH(1–34)NH_2; [Cha^{27}]hPTH(1–34)NH_2; [$Cha^{27,29}$]hPTH(1–34)NH_2; [Cha^{28}]bPTH(1–34)NH_2; [Cha^{28}]hPTH(1–34)NH_2; [$Cha^{24,28,31}$]hPTH(1–34)NH_2; [Aib^{16}]hPTH(1–34)NH_2; [Aib^{19}]hPTH(1–34)NH_2; [Aib^{34}]hPTH(1–34)NH_2; [$Aib^{16,19}$]hPTH(1–34)NH_2; [$Aib^{16,19,34}$]bPTH(1–34)NH_2; [$Aib^{16,34}$]hPTH(1–34)NH_2; [$Aib^{19,34}$]hPTH(1–34)NH_2; [$Cha^{7,11}$, $Nle^{8,18}$, $Aib^{16,19}$, Tyr^{34}]hPTH(1–34)NH_2; [$Cha^{7,11}$, $Nle^{8,18,31}$, $Aib^{16,19}$, Tyr^{34}]hPTH(1–34)NH_2; [Cha^{7}, Aib^{16}]hPTH(1–34)NH_2;[Cha^{11}, Aib^{16}]hPTH(1–34)$_2$, [Cha^{7}, Aib^{34}]hPTH(1–34)NH_2; [Cha^{11}, Aib^{34}]hPTH(1–34)NH_2; [Cha^{27}, Aib^{16}]hPTH(1–34)NH_2; [Cha^{27}, Aib^{34}]hPTH(1–34)NH_2; [Cha^{28}, Aib^{16}]hPTH(1–34)NH_2; [Cha^{28}, Aib^{34}]hPTH(1–34)NH_2; [Nle^{31}]hPTH(1–34)NH_2, [$hArg^{27}$]hPTH(1–34)NH_2; [Dap^{1}, $Nle^{8,18}$, Tyr^{34}]hPTH(1–34)NH_2; [Nle^{31}]bPTH(1–34)NH_2; [Nle^{31}]hPTH(1–34)NH_2, [$hArg^{27}$]bPTH(1–34)NH_2; [$hArg^{27}$]hPTH(1–34)NH_2; [$Cha^{7,11}$, Aib^{19}, Lys^{30}]hPTH(1–34)NH_2; [Aib^{12}]hPTH(1–34)NH_2, [$Cha^{24,28,31}$, Lys^{30}]hPTH(1–34)NH_2; [$Cha^{28,31}$]hPTH(1–34)NH_2, [$Cha^{7,11}$, $Nle^{8,18}$, Aib^{34}]hPTH(1–34)NH_2; [Aib^{3}]hPTH(1–34)NH_2, [Cha^{8}]hPTH(1–34)NH_2; [Cha^{15}]hPTH(1–34)NH_2; [$Cha^{7,11}$, Aib^{19}]hPTH(1–34)NH_2;[$Cha^{7,11}$, Aib^{16}]hPTH(1–34)NH_2;[Aib^{17}]hPTH(1–34)NH_2; [Cha^{5}]hPTH(1–34)NH_2; [$Cha^{7,11,15}$]hPTH(1–34)NH_2; [$Cha^{7,11}$, $Nle^{8,18}$, Aib^{19}, Tyr^{34}]hPTH(1–34)NH_2; [$Cha^{7,11}$, $Nle^{8,18}$, Aib^{19}, Lys^{30}, Tyr^{34}]hPTH(1–34)NH_2; [$Cha^{7,11,15}$]hPTH(1–34)NH_2; [Aib^{17}]hPTH(1–34)NH_2; [$Cha^{7,11}$, Leu^{27}]hPTH(1–34)NH_2; [$Cha^{7,11,15}$, Leu^{27}]hPTH(1–34)NH_2, [$Cha^{7,11,27}$]hPTH(1–34)NH_2; [$Cha^{7,11,15,27}$]hPTH(1–34)NH_2; [Trp^{15}]hPTH(1–34)NH_2; [Nal^{15}]hPTH(1–34)NH_2; [Trp^{15}, Cha^{23}hPTH(1–34)NH_2; [$Cha^{15,23}$]hPTH(1–34)NH_2; [$Phe^{7,11}$]hPTH(1–34)NH_2; [$Nal^{7,11}$]hPTH(1–34)NH_2, [$Trp^{7,11}$]hPTH(1–34)NH_2, [$Phe^{7,11,15}$]hPTH(1–34)NH_2; [$Nal^{7,11,15}$]hPTH(1–34)NH_2; [$Trp^{7,11,15}$]hPTH(1–34)NH_2; and [$Tyr^{7,11,15}$]hPTH(1–34)NH_2.

The following are specific examples of peptides encompassed by one or more of formulas (III) to (V), hereinabove. [Cha^{7}]hPTHrP(1–34)NH_2; [Cha^{11}]hPTHrP(1–34)NH_2; [$Cha^{7,11,15}$]hPTHrP(1–34)NH_2;[Aib^{16}, Tyr^{34}hPTHrP(1–34)NH_2; [Aib^{19}]hPTHrP(1–34)NH_2, [$Aib^{16,19}$]hPTHrP(1–34)NH_2; [$Cha^{7,11}$, Aib^{16}hPTHrP(1–34)NH_2; [$Cha^{7,11}$, Aib^{19}]hPTHrP(1–34)NH_2; [Cha^{22}, $Leu^{23,28,31}$, $Glu^{25,29}$, $Lys^{26,30}$]hPTHrP(1–34)NH_2; [$Glu^{22,25,29}$, $Leu^{23,28,31}$, $Lys^{26,27,30}$]hPTHrP(1–34)NH_2; [$Cha^{22,23}$, $Glu^{25,29}$, $Leu^{28,31}$, $Lys^{26,30}$]hPTHrP(1–34)NH_2; [$Glu^{22,25}$, $Leu^{23,28,31}$, Aib^{29}, $Lys^{26,30}$]hPTHrP(1–34)NH_2; [$Glu^{22,25,29}$, Lys23,26,30, $Leu^{28,31}$]hPTHrP(1–34)NH_2; [$Glu^{22,25,29}$, $Leu^{23,28,31}$, Lys^{26}, Cha^{30}]hPTHrP(1–34)NH_2; [$Glu^{22,25,29}$, $Leu^{23,28,31}$, Lys^{26}, Aib^{30}]hPTHrP(1–34)NH_2; [$Glu^{22,25,29}$, $Leu^{23,31}$, $Lys^{26,28,30}$]hPTHrP(1–34)NH_2; [$Cha^{22,23,24,27,28,31}$, $Glu^{25,29}$, $Lys^{26,30}$]hPTHrP(1–34)NH_2; [$Glu^{22,25,29}$, $Cha^{23,24,28,31}$, $Lys^{26,27,30}$]hPTHrP(1–34)NH_2; [$Glu^{22,25,29}$, $Cha^{23,24,27,31}$, $Lys^{26,28,30}$]hPTHrP(1–34)NH_2; [$Glu^{22,25,29}$, $Lys^{23,26,30}$, $Cha^{24,27,28,31}$]hPTHrP(1–34)NH_2; [Cha^{22}, $Leu^{23,28,31}$, $Glu^{25,29}$, $Lys^{26,27,30}$]hPTHrP(1–34)NH_2; [Cha^{22}, $Leu^{23,31}$, $Glu^{25,29}$, $Lys^{26,28,30}$]hPTHrP(1–34)NH_2; [Cha^{22}, $Leu^{23,26,30}$, $Glu^{25,29}$, $Leu^{28,31}$]hPTHrP(1–34)NH_2; [Cha^{22}, $Leu^{23,28,31}$, Glu^{25}, $Lys^{26,30}$, Aib^{29}]hPTHrP(1–34)NH_2; [Cha^{22}, $Leu^{23,28,31}$, $Glu^{25,29}$, Lys^{26}, Aib^{30}]hPTHrP(1–34)NH_2; [$Glu^{22,25}$, $Leu^{23,28,31}$, $Lys^{26,27,30}$, Aib^{29}]hPTHrP(1–34)NH_2; [$Glu^{22,25}$, $Lys^{23,26,30}$, $Leu^{28,31}$, Aib^{29}]hPTHrP(1–34)NH_2; [$Glu^{22,25}$, $Leu^{23,31}$, $Lys^{26,28,30}$, Aib^{29}]hPTHrP(1–34)NH_2; [$Cha^{7,11}$, $Glu^{22,25,29}$, $Leu^{23,28,31}$, $Lys^{26,30}$]hPTHrP(1–34)NH_2; [$Cha^{7,11,22}$, $Leu^{23,28,31}$, $Glu^{25,29}$, $Lys^{26,30}$]hPTHrP(1–34)NH_2; [$Cha^{7,11}$, $Glu^{22,25,29}$, $Leu^{23,28,31}$, $Lys^{26,27,30}$]hPTHrP(1–34)NH_2; [$Cha^{7,11,22,23}$, $Glu^{25,29}$, $Leu^{28,31}$, $Lys^{26,30}$]hPTHrP(1–34)NH_2; [$Cha^{7,11}$, $Glu^{22,25,29}$, $Leu^{23,31}$, $Lys^{26,28,30}$]hPTHrP(1–34)NH_2; [$Cha^{7,11}$, $Glu^{22,25}$, $Leu^{23,28,31}$, Aib^{29}, $Lys^{26,30}$]hPTHrP(1–34)NH_2; [$Cha^{7,11}$, $Glu^{22,25,29}$, $Leu^{23,28,31}$, Lys^{26}, Aib^{30}]hPTHrP(1–34)NH_2; [Cha^{15}, $Glu^{22,25,29}$, $Leu^{23,28,31}$, $Lys^{26,30}$]hPTHrP(1–34)NH_2; [$Cha^{15,22}$, $Leu^{23,28,31}$, $Glu^{25,29}$, $Lys^{26,30}$]hPTHrP(1–34)NH_2; [Cha^{15}, $Glu^{22,25,29}$, $Leu^{23,28,31}$, $Lys^{26,27,30}$]hPTHrP(1–34)NH_2; [$Cha^{15,22,23}$, $Glu^{25,29}$, $Leu^{28,31}$, $Lys^{26,30}$]hPTHrP(1–34)NH_2; [Cha^{15}, $Glu^{22,25}$, $Leu^{23,28,31}$, Aib^{29}, $Lys^{26,30}$]hPTHrP(1–34)NH_2; [Cha^{15}, $Glu^{22,25,29}$, $Lys^{23,26,30}$, $Leu^{28,31}$]hPTHrP(1–34)NH_2; [Cha^{15}, $Glu^{22,25,29}$, $Leu^{23,28,31}$, Lys^{26}, Aib^{30}]hPTHrP(1–34)NH_2; [Cha^{15} $Glu^{22,28,29}$, $Leu^{23,31}$, $Lys^{26,28,30}$]hPTHrP(1–34)NH_2; [$Cha^{15,30}$, $Glu^{22,25,29}$, $Leu^{23,28,31}$, Lys^{26}]hPTHrP(1–34)NH_2; [$Cha^{7,8,22}$, $Leu^{23,28,31}$, $Glu^{25,29}$, $Lys^{26,30}$]hPTHrP(1–34)NH_2; [$Cha^{7,8}$, $Glu^{22,25,29}$, $Leu^{23,28,31}$, $Lys^{26,27,30}$]hPTHrP(1–34)NH_2; [$Cha^{7,8,22,23}$, $Glu^{25,29}$, $Leu^{28,31}$, $Lys^{26,30}$]hPTHrP(1–34)NH_2; [$Cha^{7,8}$, $Glu^{22,25,29}$, $Leu^{23,28,31}$, $Lys^{26,30}$]hPTHrP(1–34)NH_2; [$Cha^{7,8}$, $Glu^{22,25}$, $Leu^{23,28,31}$, Aib^{29}, $Lys^{26,30}$]hPTHrP(1–34)NH_2; [$Cha^{7,8}$, $Glu^{22,25,29}$, $Lys^{23,26,30}$, $Leu^{28,31}$]hPTHrP(1–34)NH_2; [$Cha^{7,8}$, $Glu^{22,25,29}$, $Leu^{23,28,31}$, Lys^{26}, Aib^{30}]hPTHrP(1–34)NH_2. [$Cha^{7,8}$, $Glu^{22,25,29}$, $Leu^{23,31}$, $Lys^{26,28,30}$]hPTHrP(1–34)NH_2; [$Cha^{7,8,30}$, $Glu^{22,25,29}$, $Leu^{23,28,31}$, Lys^{26}]hPTHrP(1–34)NH_2; [Ser^{1}, Ile^{5}, $Cha^{7,11,22}$, Met^{8}, Asn^{10}, His^{14}, $Leu^{23,28,31}$, $Glu^{25,29}$, $Lys^{26,30}$]hPTHrP(1–34)NH_2; [Ser^{1}, Ile^{5}, $Cha^{7,11}$, Met^{8}, Asn^{10}, His^{14}, $Glu^{22,25,29}$, $Leu^{23,28,31}$, Lys^{26}, Aib^{30}]hPTHrP(1–34)NH_2; [Ser^{1}, Ile^{5}, $Cha^{7,11}$, Met^{8}, Asn^{10}, His^{14}, $Glu^{22,25}$, $Leu^{23,28,31}$, Aib^{29}, $Lys^{26,30}$]hPTHrP(1–34)NH_2; [Ser^{1}, Ile^{5}, $Cha^{7,11}$, Met^{8}, Asn^{10}, His^{14}, $Glu^{22,25,29}$, $Leu^{23,31}$, $Lys^{26,28,30}$]hPTHrP(1–34)NH_2; [Ser^{1}, Ile^{5}, $Cha^{7,11}$, Met^{8}, Asn^{10}, His^{14}, $Glu^{22,25}$, $Leu^{23,28,31}$, Aib^{29}, $Lys^{26,30}$]hPTHrP(1–34)NH_2; [Ser^{1}, Ile^{5}, $Cha^{7,11}$, Met^{8}, Asn^{10}, His^{14}, $Glu^{22,25,29}$, $Leu^{23,28,31}$, Lys^{26}, Aib^{30}]PTHrP(1–34)NH_2; [Ser^{1}, Ile^{5}, $Cha^{7,11,22,23}$, Met^{8}, Asn^{10}, His^{14}, $Glu^{25,29}$, $Leu^{28,31}$, $Lys^{26,30}$]hPTHrP(1–34)NH_2; [Ser^{1}, Ile^{5}, $Cha^{7,11}$, Met^{8}, Asn^{10}, His^{14}]hPTHrP(1–34)NH_2; [Ser^{1}, Ile^{5}, Met^{8}, Asn^{10}, Leu^{11}, His^{14}, Aib^{16}]hPTHrP(1–34)NH_2; [Ser^{1}, Ile^{5}, Met^{8}, Asn^{10}, $Leu^{11,28,31}$, His^{14}, $Cha^{22,23}$, $Glu^{25,29}$, $Lys^{26,30}$]hPTHrP(1–34)NH_2; [Ser^{1}, Ile^{5}, $Cha^{7,11}$, Met^{8}, Asn^{10}, His^{14}, $Glu^{22,25,29}$, $Leu^{23,28,31}$, $Lys^{26,30}$]hPTHrP(1–34)NH_2; [Ser^{1}, Ile^{5}, Met^{8}, Asn^{10}, His^{14} Cha^{15}, $Glu^{22,25,29}$, $Leu^{23,28,31}$, $Lys^{26,30}$]hPTHrP(1–34)NH_2; [Ser^{1}, Ile^{5}, $Cha^{7,8}$, Asn^{10}, His^{14}, $Glu^{22,25,29}$, $Leu^{23,28,31}$, $Lys^{26,30}$]hPTHrP(1–34)NH_2; [$Glu^{22,25,29}$, $Leu^{23,28,31}$, $Lys^{24,26,30}$]hPTHrP(1–34)NH_2; [Aib^{23}, $Leu^{23,28,31}$, $Glu^{25,29}$, $Lys^{26,30}$]hPTHrP(1–34)NH_2; [$Glu^{22,29}$, $Leu^{23,28,31}$, Aib^{25}, $Lys^{26,30}$]hPTHrP(1–34)NH_2; [$Glu^{22,24,29}$, $Leu^{23,28,31}$, Aib^{26}, Lys^{30}]hPTHrP(1–34)NH_2; [$Glu^{22,25,29}$, $Leu^{23,28}$, $Lys^{26,30,31}$]hPTHrP(1–34)NH_2; [Ser^{1}, Ile^{5}, Met^{8}, Asn^{10}, $Leu^{11,23,28,31}$, His^{14}, Cha^{22}, $Glu^{25,29}$, $Lys^{26,30}$]hPTHrP(1–34)NH_2, [Ser^{1}, Ile^{5}, Met^{8}, Asn^{10}, $Leu^{11,2831}$, His^{14}, $Glu^{22,25,29}$,

$Lys^{23,26,30}$PTHrP(1–34)NH_2; [Ser^1, Ile^5, Met^8, Asn^{10}, $Leu^{11,23,28,31}$, His^{14}, $Glu^{22,25,29}$, $Lys^{26,27,30}$]hPTHrP(1–34)NH_2; [Ser^1, Ile^5, Met^8, Asn^{10}, $Leu^{11,23,31}$, His^{14}, $Glu^{22,25,29}$, $Lys^{26,28,30}$]hPTHrP(1–34) NH_2; [Ser^1, Ile^5, Met^8 Asn^{10}, $Leu^{11,23,28,31}$, His^{14}, $Glu^{22,25}$, Aib^{29}, $Lys^{26,30}$]hPTHrP (1–34)NH_2; [Ser^1, Ile^5, Met^8, Asn^{10}, $Leu^{11,23,28,31}$, His^{14}, $Glu^{22,25,29}$, Lys^{26}, Aib^{30}]hPTHrP (1–34)NH_2; or [Ser^1, Ile^5, Met^8]hPTHrP(1–34)NH_2 [$Glu^{22,25}$, Ahc^{23}, $Lys^{26,30}$, $Leu^{28,31}$, Aib^{29}]hPTHrP(1–34) NH_2; [$Glu^{22,25}$, $Leu^{23,28,31}$, Lys26,30, Ahc^{27}, Aib^{29}]hPTHrP (1–34)NH_2; [$Glu^{22,25}$, $Leu^{23,28}$, $Lys^{26,30}$, Aib^{29}, Ahc^{31}]hPTHrP(1–34)NH_2, [$Glu^{22,25}$, Cha^{23}, Lys26, 30, $Leu^{28,31}$, Aib^{29}]hPTHrP(1–34) NH_2; [$Glu^{22,25}$, Cha^{23}, $Lys^{26,30}$ Leu^{28}, Aib^{29}]hPTHrP(1–34) NH_2; [$Glu^{22,25}$, Cha^{23}, $Lys^{26,30}$, Aib^{29}]hPTHrP(1–34)NH_2, [Ahc^{22}, $Leu^{23,28,31}$, Glu^{25}, $Lys^{26,30}$, Aib^{29}]hPTHrP(1–34) NH_2; [$Glu^{22,25}$, $Leu^{23,28,31}$, Lys^{26}, Aib^{29}, Ahc^{30}]hPTHrP (1–34)NH_2; [$Glu^{22,25}$, Cha^{23}, $Lys^{26,30}$, Aib^{29}, Leu^{31}] hPTHrP(1–34)NH_2; [$Glu^{22,25}$, $Leu^{23,28,31}$, Ahc^{24}, $Lys^{26,30}$, Aib^{29}]hPTHrP(1–34) NH_2; [Glu22,25, $Leu^{23,31}$, $Lys^{26,30}$, Ahc^{28}, Aib^{29}]hPTHrP (1–34)NH_2; [$Glu^{22,23}$, $Leu^{23,28,3}$, Lys^{26}, $Aib^{29,30}$]hPTHrP (1–34)NH_2; [$Aib^{22,29}$, $Leu^{23,28,31}$, Glu^{25}, $Lys^{26,30}$]hPTHrP (1–34)NH_2; [$Glu^{22,25}$, $Leu^{23,28,31}$, $Aib^{26,29}$, Lys^{30}]hPTHrP (1–34)NH_2; [Cha^{22}, Ahc^{23}, $Glu^{25,29}$, $Lys^{26,30}$, $Leu^{28,31}$]hPTHrP(1–34)NH_2; [Cha^{22}, $Leu^{23,28,31}$, Ahc^{24}, $Glu^{25,29}$, $Lys^{26,30}$]hPTHrP(1–34)NH_2; [Cha^{22}, $Leu^{23,28,31}$, $Glu^{25,29}$, $Lys^{26,30}$, Ahc^{27}]hPTHrP(1–34) NH_2; [Cha^{22}, $Leu^{23,31}$, $Glu^{25,29}$, $Lys^{26,30}$, Ahc^{28}]hPTHrP (1–34)NH_2; [Cha^{22}, $Leu^{23,28,31}$, $Glu^{25,29}$, Lys^{26}, Leu^{28}, Ahc^{30}]hPTHrP(1–34) NH_2, [$Cha^{22,23}$, $Glu^{25,29}$, $Lys^{26,30}$, Leu^{31}]hPTHrP(1–34) NH_2; [Cha^{22}, $Leu^{23,28}$, $Glu^{25,29}$, $Lys^{26,30}$, Ahc^{31}]hPTHrP (1–34)NH_2; [$Cha^{22,23}$, $Glu^{25,29}$, $Lys^{26,30}$, Leu^{31}]hPTHrP (1–34)NH_2; [$Cha^{22,23}$, $Glu^{25,29}$, $Lys^{26,30}$, Leu^{28}]hPTHrP (1–34)NH_2; [$Cha^{22,23}$, $Glu^{25,29}$, $Lys^{26,30}$]hPTHrP(1–34) NH_2; [Glu^{22}, $Leu^{23,28,31}$, $Aib^{25,29}$, $Lys^{26,30}$]hPTHrP(1–34) NH_2; [$Glu^{22,29}$, Ahc^{23}, Aib^{25}, $Lys^{26,30}$, $Leu^{28,31}$]hPTHrP (1–34)NH_2; [Ahc^{22}, $Leu^{23,21,31}$, Aib^{25}, $Lys^{26,30}$, Glu^{29}]hPTHrP(1–34)NH_2; [$Aib^{22,25}$, $Leu^{23,28,31}$, $Lys^{26,30}$, Glu^{29}]hPTHrP (1–34)NH_2; [$Glu^{22,29}$, $Leu^{23,28,31}$, Ahc^{24}, Aib^{25}, $Lys^{26,30}$]hPTHrP(1–34)NH_2; [$Glu^{22,29}$, $Leu^{23,28,31}$, $Aib^{25,26}$, Lys^{30}]hPTHrP(1–34)NH_2; [$Glu^{22,29}$, $Leu^{23,28,31}$, Aib^{25}, $Lys^{26,30}$, Ahc^{27}]hPTHrP(1–34) NH_2; [$Glu^{22,29}$, $Leu^{23,31}$, Aib^{25}, $Lys^{26,30}$, Ahc^{28}]hPTHrP (1–34)NH_2; [$Glu^{22,29}$, $Leu^{23,28}$, Aib^{25}, $Lys^{26,30}$, Ahc^{31}] hPTHrP(1–34)NH_2; [$Glu^{22,29}$, $Leu^{23,28,31}$, $Aib^{25,30}$, Lys^{26}] hPTHrP(1–34)NH_2; [$Glu^{22,29}$, $Leu^{23,28,31}$, Aib^{25}, Lys^{26}, Ahc^{30}]hPTHrP(1–34) NH_2; [$Glu^{22,29}$, Cha^{23}, Aib^{25}, $Lys^{26,30}$, $Leu^{28,31}$]hPTHrP (1–34)NH_2; [$Glu^{22,29}$, Cha^{23}, Aib^{25}, $Lys^{26,30}$, Leu^{31}]hPTHrP(1–34)NH_2; [$Glu^{22,29}$, Cha^{23}, Aib^{25}, $Lys^{26,30}$]hPTHrP(1–34)NH_2; [$Glu^{22,29}$, Cha^{23}, Aib^{25}, $Lys^{26,30}$, Leu^{28}]hPTHrP(1–34)NH_2; [$Glu^{22,25,29}$, Cha^{23}, Lys^{26}, $Leu^{28,31}$, Aib^{30}]hPTHrP(1–34)NH_2; [$Glu^{22,25,29}$, Cha^{23}, Lys^{26}, Aib^{30}, Leu^{31}]hPTHrP(1–34)NH_2;[$Glu^{22,25,29}$, Cha^{23}, Lys^{26}, Aib^{30}]hPTHrP(1–34)NH_2;[$Glu^{22,25,29}$, Cha^{23}, Lys^{26}, Leu^{28}, Aib^{30}]hPTHrP(1–34)NH_2; [$Glu^{22,25,29}$, $Leu^{23,28,31}$, Lys^{26}, Ahc^{27}, Aib^{30}]hPTHrP(1–34)NH_2; [$Glu^{22,25,29}$, $Leu^{23,28,31}$, Ahc^{24}, Lys^{26}, Aib^{30}]hPTHrP(1–34) NH_2; [Ahc^{22}, $Leu^{23,28,31}$, $Glu^{25,29}$, Lys^{26}, Aib^{30}]hPTHrP (1–34)NH_2; [$Aib^{22,30}$, $Leu^{23,28,31}$, $Glu^{25,29}$, Lys^{26}]hPTHrP (1–34)NH_2; [$Glu^{22,25}$, $Leu^{23,28}$, $Lys^{26,30,31}$, Aib^{29}]hPTHrP (1–34)NH_2; [Cha^{22}, $Leu^{23,28}$, $Glu^{25,29}$, $Lys^{26,30,31}$]hPTHrP (1–34)NH_2; [Ahc^{22}, $Leu^{23,28}$, $Glu^{25,29}$, $Lys^{26,30,31}$]hPTHrP (1–34)NH_2; [$Glu^{22,25,29}$, $Leu^{23,28}$, $Lys^{26,30,31}$, Ahc^{30}] hPTHrP(1–34) NH_2; [$Glu^{22,25,29}$, $Leu^{23,28,31}$, Lys^{26}, Ahc^{30}] hPTHrP(1–34) NH_2; [Ahc^{22}, $Leu^{23,28,31}$, $Glu^{25,29}$, $Lys^{26,30}$] hPTHrP(1–34) NH_2; [$Glu^{22,25,29}$, $Leu^{23,28}$, $Lys^{26,30,31}$, Ahc^{27}]hPTHrP (1–34)NH_2.

In another aspect, the present invention is directed to a method of treating osteoporosis in a patient in need thereof, which comprises administering to said patient a compound of formula (I), (II), (III), (IV) or (V) or a pharmaceutically acceptable salt thereof, as defined hereinabove.

In another aspect, the present invention is directed to a method of treating osteoporosis in a patient in need thereof, which comprises administering to said patient a combination of a bisphosphonate or calcitonin and a compound of formula (I), (II), (III), (IV) or (V) or a pharmaceutically acceptable salt thereof, as defined hereinabove.

In another aspect, the present invention is directed to a pharmaceutical composition comprising a compound of formula (I), (II), (III), (IV) or (V) or a pharmaceutically acceptable salt thereof and a pharmaceutically acceptable carrier or diluent.

In another aspect, the present invention is directed to a pharmaceutical composition comprising a compound of formula (I), (II), (III), (IV) or (V) or a pharmaceutically acceptable salt thereof as defined hereinabove, a bisphosphonate or calcitonin and a pharmaceutically acceptable carrier or diluent.

In another aspect, the present invention is directed to a method of treating osteoporosis in a patient in need thereof, which comprises administering to said patient a peptide of the formula [$Glu^{22,25}$, $Leu^{23,28,31}$, Aib^{29}, $Lys^{26,30}$]hPTHrP (1–34)NH_2 or a pharmaceutically acceptable salt thereof.

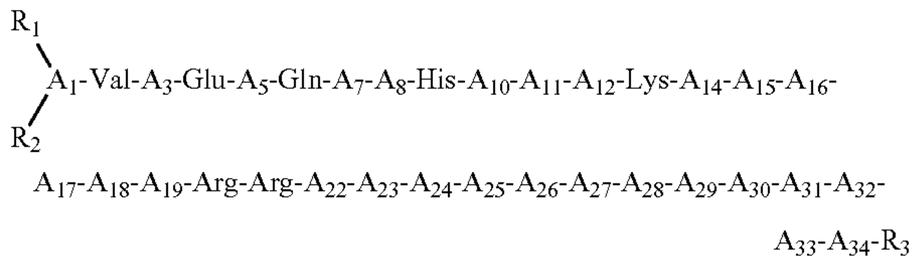
In another aspect, the present invention is directed to a method of treating osteoporosis in a patient in need thereof, which comprises administering to said patient a combination of a bisphosphonate or calcitonin and a peptide of the formula [$Glu^{22,25}$, $Leu^{23,28,31}$, Aib^{29}, $Lys^{26,30}$]hPTHrP (1–34) NH_2 or a pharmaceutically acceptable salt thereof.

In another aspect, the present invention is directed to a pharmaceutical composition comprising a peptide of the formula [$Glu^{22,25}$, $Leu^{23,28,31}$, Aib^{29}, $Lys^{26,30}$]hPTHrP (1–34) NH_2 or a pharmaceutically acceptable salt thereof and a pharmaceutically acceptable carrier or diluent.

In another aspect, the present invention is directed to a pharmaceutical composition comprising a peptide of the formula [$Glu^{22,25}$, $Leu^{23,28,31}$, Aib^{29}, $Lys^{26,30}$]hPTHrP (1–34) NH_2 or a pharmaceutically acceptable salt thereof, a bisphosphonate or calcitonin, and a pharmaceutically acceptable carrier or diluent.

In another aspect, the present invention is directed to a method of treating osteoporosis in a patient in need thereof, which comprises administering to said patient a peptide of the formula (VI):

[SEQ ID NO: 4]



wherein

A_1 is Ala, Ser, or Dap;

A_3 is Ser or Aib;

A_5 is His, Ile, Acc, or Cha;

A_7 is Leu, Cha, Nle, β-Nal, Trp, Pal, Acc, Phe, or p-X-Phe in which X is OH, a halogen, or CH_3;

A_8 is Leu, Met, Acc, or Cha;

A_{10} is Asp or Asn;

A_{11} is Lys, Leu, Cha, Acc, Phe, or β-Nal;

A_{12} is Gly, Acc, or Aib;

A_{14} is Ser or His;
A_{15} is Ile, Acc, or Cha;
A_{16} is Gln or Aib;
A_{17} is Asp or Aib;
A_{18} is Leu, Aib, Acc, or Cha;
A_{19} is Arg or Aib;
A_{22} is Phe, Glu, Aib, Acc, or Cha;
A_{23} is Phe, Leu, Lys, Acc, or Cha;
A_{24} is Leu, Lys, Acc, or Cha;
A_{25} is His, Lys, Aib, Acc, or Glu;
A_{26} is His, Aib, Acc, or Lys;
A_{27} is Leu, Lys, Acc, or Cha;
A_{28} is Ile, Leu, Lys, Acc, or Cha;
A_{29} is Ala, Glu, Acc, or Aib;
A_{30} is Glu, Leu, Nle, Cha, Aib, Acc, or Lys;
A_{31} is Ile, Leu, Cha[.], Lys, Acc, or deleted;
A_{32} is His or deleted;
A_{33} is Thr or deleted;
A_{34} is Ala or deleted;

each of R_1 and R_2 is, independently, H, C_{1-12} alkanyl, C_{7-20} phenylalkyl, C_{11-20} naphthylalkyl, C_{1-12}, hydroxyalkyl, C_{2-12} hydroxyalkenyl, C_{7-20} hydroxyphenylalkyl, or C_{11-20} hydroxynaphthylalkyl; or one and only one of R_1 and R_2 is COE_1 in which E_1 is C_{1-12} alkyl, C_{2-12} alkyl C_{2-12} alkenyl, C_{7-20} phenylalkyl, C_{11-20} naphthylalkyl, C_{1-12} hydroxyalkyl, C_{2-12} hydroxyalkenyl, C_{7-20} hydroxyphenylalkyl, or C_{11-20} hydroxynaphthylalkyl; and

R_3 is OH, NH_2, C_{1-12} alkoxy, or NH—Y—CH_2-Z in which Y is a C_{1-12} hydrocarbon moiety and Z is H, OH, CO_2H or $CONH_2$;

provided that at least one of A_5, A_7, A_8, A_{11}, A_{12}, A_{15}, A_{18}, A_{22}, A_{23}, A_{24}, A_{25}, A_{26}, A_{27}, A_{28}, A_{29}, A_{30}, or A_{31} is Acc; or a pharmaceutically acceptable salt thereof.

In another aspect, the present invention is directed to a method of treating osteoporosis in a patient in need thereof, which comprises administering to said patient a combination of bisphosphonate or calcitonin and a peptide of formula (VI), as defined hereinabove.

In another aspect, the present Invention is directed to a pharmaceutical composition comprising a pharmaceutically acceptable carrier or diluent and a peptide of formula (VI), as defined hereinabove.

In another aspect, the present invention is directed to a pharmaceutical composition comprising a bisphosphonate or calcitonin, a pharmaceutically acceptable carrier or diluent, and a peptide of formula (VI), as defined hereinabove.

With the exception of the N-terminal amino acid, all abbreviations (e.g. Ala or A_1) of amino acids in this disclosure stand for the structure of —NH—CH(R)—CO—, wherein R is a side chain of an amino acid (e.g., CH_3 for Ala). For the N-terminal amino acid, the abbreviation stands for the structure of ═N—CH(R)—CO—, wherein R is a side chain of an amino acid, β-Nal, Nle, Dap, Cha, Nva, Amp, Pal, Ahc, and Aib are the abbreviations of the following α-amino acids[;] : β-(2-naphthyl)alanine, norleucine, α,β-diaminopropionic acid, cyclohexylalanine, norvaline, 4-amino-phenylalanine, β-(3-pyridinyl)alanine, 1-amino-1-cyclo-hexanecarboxylic acid, and α-aminoisobutyric acid, respectively. What is meant by Acc is an amino acid selected from the group of 1-amino-1-cyclopropanecarboxylic acid[.], 1-amino-1-cyclobutanecarboxylic acid[;], 1-amino-1-cyclopentanecarboxylic acid[;], 1-amino-1-cyclohexanecarboxylic acid[;], 1-amino-1-cycloheptanecarboxylic acid, 1-amino-1-cyclooctanecarboxylic acid, and 1-amino-1-cyclononanecarboxylic acid. In the above formula, hydroxyalkyl, hydroxyphenyl-alkyl, and hydroxynaphthylalkyl may contain 1–4 hydroxy substituents. Also, COE_1 stands for [—C═O.E_1] —*C═O.E*$_1$. Examples of [—C═O.E_1] —*C═O.E*$_1$ include, but are not limited to, acetyl and phenylpropionyl.

A peptide of this invention is also denoted herein by another format, e.g., [$Ahc^{7,11}$]hPTH(1–34)NH_2, with the substituted amino acids from the natural sequence placed between the second set of brackets (e.g., Ahc^7 for Leu^7, and Ahc^{11} for Leu^{11} in hPTH). The abbreviation hPTH stands for human PTH; hPTHrP for human PTHrP, rPTH for rat PTH, and bPTH for bovine PTH. The numbers between the parentheses refer to the number of amino acids present in the peptide (e.g., hPTH(1–34) is amino acids 1 through 34 of the peptide sequence for human PTH). The sequences for hPTH (1–34), hPTHrP(1–34), bPTH(1–34), and rPTH(1–34) are listed in Nissenson, et al., Receptor, 3 193 (1993). The designation of "NH_2" in PTH(1–34)NH_2 indicates that the C-terminus of the peptide is amidated. PTH(1–34), on the other hand, has a free acid C-terminus.

Each of the peptides of the invention is capable of stimulating the growth of bone in a subject (i.e., a mammal such as a human patient). Thus, it is useful in the treatment of osteoporosis and bone fractures when administered alone or concurrently with antiresorptive therapy, e.g., bisphosphonate and calcitonin.

The peptides of this invention can be provided in the form of pharmaceutically acceptable salts. Examples of such salts include, but are not limited to, those formed with organic acids (e.g., acetic, lactic, maleic, citric, malic, ascorbic, succinic, benzoic, methanesulfonic, toluenesulfonic, or pamoic acid), inorganic acids (e.g., hydrochloric acid, sulfuric acid, or phosphoric acid), and polymeric acids (e.g., tannic acid, carboxymethyl cellulose, polylactic, polyglycolic, or copolymers of polylactic-glycolic acids).

A therapeutically effective amount of a peptide of this invention and a pharmaceutically acceptable carrier substance (e.g., magnesium carbonate, lactose, or a phospholipid with which the therapeutic compound can form a micelle) together form a therapeutic composition (e.g., a pill, tablet, capsule, or liquid) for administration (e.g., orally, intravenously, transdermally, pulmonarily, vaginally, subcutaneously, nasally, iontophoretically, or by intratracheally) to a subject. The pill, tablet, or capsule that is to be administered orally can be coated with a substance for protecting the active composition from the gastric acid or intestinal enzymes in the stomach for a period of time sufficient to allow it to pass undigested into the small intestine. The therapeutic composition can also be in the form of a biodegradable or nonbiodegradable sustained release formulation for subcutaneous or intramuscular administration. See, e.g., U.S. Pat. Nos. 3,773,919 and 4,767,628 and PCT Application No WO 94/15587. Continuous administration can also be achieved using an implantable or external pump (e.g., INFUSAID™ pump). The administration can also be conducted intermittently, e.g., single daily injection, or continuously at a low dose, e.g., sustained release formulation.

The dose of a peptide of the present invention for treating the above-mentioned diseases or disorders varies depending upon the manner of administration, the age and the body weight of the subject, and the condition of the subject to be treated, and ultimately will be decided by the attending physician or veterinarian.

Also contemplated within the scope of this invention is a peptide covered by the above generic formula for use in treating diseases or disorders associated with deficiency in bone growth or the like, e.g. osteoporosis or fractures.

Other features and advantages of the present invention will be apparent from the detailed description and from the claims.

DETAILED DESCRIPTION OF THE INVENTION

Based on the description herein, the present invention can be utilized to its fullest extent. The following specific examples are to be construed as merely illustrative, and should not be construed as a limitation of the remainder of the disclosure in any way whatsoever. Further, all publications cited herein are incorporated by reference.

Structure

PTH(1–34) and PTHrP(1–34) have been reported to have two amphophilic alpha helical domains. See, e.g., Barden, et al., Biochem., 32:7126 (1992). The first "-helix is formed between amino acid residues 4 through 13, while the second "-helix is formed between amino acid residues 21 through 29. Some peptides of this invention contain the substitution of Acc for one or more residues within or near these two regions of PTH(1–34) and PTHrP(1–34), e.g., Ahc^{7} and Ahc^{11} within the first "-helix or Ahc^{27} and Ahc^{28} within the second "-helix; or Cha^{7} and Cha^{11} within the first α-helix or Cha^{27} and Cha^{28} within the second α-helix.

Synthesis

The peptides of the invention can be prepared by standard solid phase synthesis. See, e.g., Stewart, J. M., et al., Solid Phase Synthesis (Pierce Chemical Co., 2d ed. 1984). The following is a description of how [$Glu^{22,25}$, Leu23,28, Lys26,30, Aib^{29}, or Ahc^{31}]hPTH(1–34)NH_2 was prepared. Other peptides of the invention can be prepared in an analogous manner by a person of ordinary skill in the art.

1-[N-tert-Butoxycarbonyl-amino]-1-cyclohexanecarboxylic acid(Boc-Ahc-OH) was synthesized as follows.

19.1 g (0.133 mol) of 1-amino-1-cyclohexanecarboxylic acid (Acros Organics, Fisher Scientific, Pittsburgh, Pa) was dissolved in 200 ml of dioxane and 100 ml of water. To it was added 67 mg of 2N NaOH. The solution was cooled in an ice-water bath. 32.0 g (0.147 mol) of di-tert-butyl-dicarbonate was added to this solution. The reaction mixture was stirred overnight at room temperature. Dioxane was then removed under reduced pressure. 200 ml of ethyl acetate was added to the remaining aqueous solution The mixture was cooled in an ice-water bath. The pH of the aqueous layer was adjusted to about 3 by adding 4N HCl. The organic layer was separated. The aqueous layer was extracted with ethyl acetate (1×100 ml). Two organic layers were combined and washed with water (2×150 ml), dried over anhydrous $MgSO_4$, filtered and concentrated to dryness under reduced pressure. The residue was recrystallized in ethyl acetate/hexanes, 9.2 g of a pure product was obtained, 29% yield. Other protected Acc amino acids can be prepared in an analogous manner by a person or ordinary skill in the art.

The peptide was synthesized on an Applied Biosystems (Foster City, Calif.) model 430A peptide synthesizer which was modified to do accelerated Boc-chemistry solid phase peptide synthesis. See Schnoize, et al., Int. J. Peptide Protein Res., 90:180 (1992). 4-Methylbenz-hydrylamine (MBHA) resin (Peninsula, Belmont, Calif.) with the substitution of 0.93 mmol/g was used. The Boc amino acids (Bachem, Calif., Torrance, Calif.; Nova Biochem., LaJolla, Calif.) was used with the following side chain protection: Boc-Ala-OH, Boc-Arg(Tos)-OH, Boc-Asp(OcHex)-OH, Boc-Glu (OcHex)-OH, Boc-His(DNP)-OH, Boc-Val-OH, Boc-Leu-OH, Boc-Gly-OH, Boc-Gln-OH, Boc-Ile-OH, Boc-Lys (2ClZ)-OH, Boc-Ahc-OH, Boc-Thr(Bzl)-OH, Boc-Ser (Bzl)-OH; and Boc-Aib-OH. The synthesis was carried out on a 0.14 mmol scale. The Boc groups were removed by treatment with 100% TFA for 2×1 min. Boc amino acids (2.5 mmol) were pre-activated with HBTU (2.0 mmol) and DIEA (1.0 mL) in 4 mL of DMF and were coupled without prior neutralization of the peptide-resin TFA salt. Coupling times were 5 min except for the Boc-Aib-OH, and its following residue Boc-Leu-OH, and Boc-Ahc-OH, and its following residue Boc-Lys(2Clz)-OH, wherein the coupling times for these four residues were 2 hrs.

At the end of the assembly of the peptide chain, the resin was treated with a solution of 20% mercaptoethanol/10% DIEA in DMF for 2×30 min. to remove the DNP group on the His side chain. The N-terminal Boc group was then removed by treatment with 100% TFA for 2×2 min. The partially-deprotected peptide-resin was washed with DMF and DCM and dried under reduced pressure. The final cleavage was done by stirring the peptide-resin in 10 mL of HF containing 1 mL of anisole and dithiothreitol (24 mg) at OEC for 75 min. HF was removed by a flow of nitrogen. The residue was washed with ether (6×10 mL) and extracted with 4N HOAc (6×10 mL).

The peptide mixture in the aqueous extract was purified on a reversed-phase preparative high pressure liquid chromatography (HPLC) using a reversed phase VYDAC™ C_{18} column (Nest Group, Southborough, Mass.) The column was eluted with a linear gradient (10% to 45% of solution B over 130 min.) at a flow rate of 10 mL/min (Solution A=0.1% aqueous TFA; Solution B=acetonitrile containing 0.1% of TFA). Fractions were collected and checked on analytical HPLC. Those containing pure product were combined and lyophilized to dryness, 85 mg of a white solid was obtained. Purity was >99% based on analytical HPLC analysis. Electro-spray mass spectrometer analysis gave the molecular weight at 3972.4 (in agreement with the calculated molecular weight of 3972.7).

The synthesis and purification of [Cha^{22}, $Leu^{23,2831}$, Glu^{25}, $Lys^{26,30}$, Ahc^{27}, Aib^{29}]hPTHrP(1–34)NH_2 was carried out in the same manner as the above synthesis of [$Glu^{22,25}$, $Leu^{23,28}$, $Lys^{26,30}$, Aib^{29}, Ahc^{31}]hPTHrP(1–34) NH_2 The protected amino acid Boc-Cha-OH was purchased from Bachem, Calif. The purity of the final product was >99%, and the electron-spray mass spectrometer gave the molecular weight at 3997.2 (calculated molecular weight is 3996.8).

The following is a description of how [Aib^{34}]hPTH(1–34) NH_2 was prepared. The peptide, [Aib^{34}]hPTH(1–34)NH_2, was synthesized on an Applied Biosystems (Foster City, Calif.) model 430A peptide synthesizer which was modified to do accelerated Boc-chemistry solid phase peptide synthesis. See Schnoize, et al. Int. J. Peptide Protein Res., 90:180 (1992). 4-Methylbenz-hydrylamine (MBHA) resin (Peninsula, Belmont, Calif.) with the substitution of 0.93 mmol/g was used. The Boc amino acids (Bachem, Calif. Torrance, Calif.; Nova Biochem., LaJolla, Calif.) were used with the following side chain protection: Boc-Arg(Tos)-OH, Boc-Asp(OcHxl)-OH, Boc-Asn(Xan)-OH, Boc-Glu (OcHxl)-OH. Boc-His(DNP)-OH, Boc-Asn-GH, Boc-Val-OH, Boc-Leu-OH, Boc-Ser-OH, Boc-Gly-Oh, Boc-Met-OH, Boc-Gln-OH, Boc-Ile-OH, Boc-Lys(2ClZ)-OH, Boc-Ser(Bzl)-OH, and Boc-Trp(Fm)-OH The synthesis was carried out on a 0.14 mmol scale. The Boc groups were removed by treatment with 100% TFA for 2×1 min. Boc amino acids (2.5 mmol) were pre-activated with HBTU (2.0 mmol) and DIEA (1.0 mL) in 4 mL of DMF and were coupled without prior neutralization of the peptide-resin TFA salt. Coupling times were 5 min except for the Boc-

Aib-OH and the following residue. Boc-Asn(Xan)-OH, wherein the coupling times were 20 min.

At the end of the assembly of the peptide chain, the resin was treated with a solution of 20% mercaptoethanol/10% DIEA in DMF for 2×30 min. to remove the DNP group on the His side chain. The N-terminal Boc group was then removed by treatment with 100% TFA for 2×2 min. After neutralization of the peptide-resin with 10% DIEA in DMF (1×1 min.). the formyl group on the side chain of Trp was removed by treatment with a solution of 15% ethanolamine/15% water/70% DMF for 2×30 min. The partially-deprotected peptide-resin was washed with DMF and DCM and dried under reduced pressure. The final cleavage was done by stirring the peptide-resin in 10 mL of HF containing 1 mL of anisole at 0° C. for 75 min. HF was removed by a flow of nitrogen. The residue was washed with ether (6×10 mL) and extracted with 4N HOAc (6×10 mL).

The peptide mixture in the aqueous extract was purified on a reversed-phase preparative high pressure liquid chromatography (HPLC) using a reversed phase VYDAC™ C_{18} column (Nest Group, Southborough, Mass.). The column was eluted with a linear gradient (10% to 45% of solution B over 130 min.) at a flow rate of 10 mL/min (Solution A=0.1% aqueous TFA; Solution B=acetonitrile containing 0.1% of TFA). Fractions were collected and checked on analytical HPLC. Those containing pure product were combined and lyophilized to dryness. 62.3 mg of a white solid was obtained Punty was >99% based on analytical HPLC analysis. Electro-spray mass spectrometer analysis gave the molecular weight at 4054.7 (in agreement with the calculated molecular weight of 4054.7).

The synthesis and purification of [$Cha^{7,11}$]hPTH(1–34)NH_2 was carried out in the same manner as the above synthesis of [Aib^{34}]hPTH(1–34)NH_2. The protected amino acid Boc-Cha-OH was purchased from Bachem, Calif. The purity of the final product was >98%, and the electron-spray mass spectrometer gave the molecular weight at 4197 0 (calculated molecular weight is 4196.9).

The following is a description of how [$Glu^{22,25}$, $Leu^{23,28}$, $Lys^{26,30}$, Aib^{29}, Ahc^{31}]hPTH(1–34)NH_2 was prepared. Other peptides of the invention can be prepared in an analogous manner by a person of ordinary skill in the art.

1-[N-tert-Butoxycarbonyl-amino]-1-cyclohexanecarboxylic acid (Boc-AHC-OH) was synthesized as follows:

19.1 g (0.133 mol) of 1-amino-1-cyclohexanecarboxylic acid (Acros Organics, Fisher Scientific, Pittsburgh, Pa) was dissolved in 200 ml of dioxane and 100 ml of water To it was added 67 mg of 2N NaOH. The solution was cooled in an ice-water bath. 32.0 g (0.147 mol) of di-tert-butyl-dicarbonate was added to this solution. The reaction mixture was stirred overnight at room temperature. Dioxane was then removed under reduced pressure. 200 ml of ethyl acetate was added to the remaining aqueous solution. The mixture was cooled in an ice-water bath. The pH of the aqueous layer was adjusted to about 3 by adding 4N HCl. The organic layer was separated. The aqueous layer was extracted with ethyl acetate (1×100 ml). Two organic layers were combined and washed with water (2×150 ml), dried over anhydrous $MgSO_4$, filtered and concentrated to dryness under reduced pressure. The residue was recrystallized in ethyl acetate/hexanes 9.2 g of a pure product was obtained, 29% yield. Other protected Acc amino acids can be prepared in an analogous manner by a person or ordinary skill in the art.

The peptide was synthesized on an Applied Biosystems (Foster City, Calif.) model 430A peptide synthesizer which was modified to do accelerated Boc-chemistry solid phase peptide synthesis. See Schnoize, et al., Int. J. Peptide Protein Res., 90:180 (1992). 4-Methylbenz-hydrylamine (MBHA) resin (Peninsula, Belmont, Calif.) with the substitution of 0.93 mmol/g was used. The Boc amino acids (Bachem, Calif., Torrance, Calif.; Nova Biochem., LaJolla, Calif.) was used with the following side chain protection: Boc-Ala-OH, Boc-Arg(Tos)-OH, Boc-Asp(OcHex)-OH, Boc-Glu(OcHex)-OH, Boc-His(DNP)-OH, Boc-Val-OH. Boc-Leu-OH, Boc-Gly-OH, Boc-Gln-OH, Boc-Ile-OH, Boc-Lys(2ClZ)-OH, Boc-Ahc-OH, Boc-Thr(Bzl)-OH, Boc-Ser(Bzl)-OH; and Boc-Aib-OH. The synthesis was carried out on a 0.14 mmol scale. The Boc groups were removed by treatment with 100% TFA for 2×1 min. Boc amino acids (2.5 mmol) were pre-activated with HBTU (2.0 mmol) and DIEA (1.0 mL) in 4 mL of DMF and were coupled without prior neutralization of the peptide-resin TFA salt. Coupling times were 5 min except for the Boc-Aib-OH, and its following residue Boc-Leu-OH, and Boc-Ahc-OH, and its following residue Boc-Lys(2Clz)-OH, wherein the coupling times for these four residues were 2 hrs.

At the end of the assembly of the peptide chain, the resin was treated with a solution of 20% mercaptoethanol/10% DIEA in DMF for 2×30 min. to remove the DNP group on the His side chain. The N-terminal Boc group was then removed by treatment with 100% TFA for 2×2 min. The partially-deprotected peptide-resin was washed with DMF and DCM and dried under reduced pressure. The final cleavage was done by stirring the peptide-resin in 10 mL of HF containing 1 mL of anisole and dithiothreitol (24 mg) at 0° C. for 75 min. HF was removed by a flow of nitrogen. The residue was washed with ether (6×10 mL) and extracted with 4N HOAc (6×10 mL).

The peptide mixture in the aqueous extract was purified on a reversed-phase preparative high pressure liquid chromatography (HPLC) using a reversed phase VYDAC™ C_{18} column (Nest Group, Southborough, Mass.) The column was eluted with a linear gradient (10% to 45% of solution B over 130 min.) at a flow rate of 10 mL/min (Solution A=0.1% aqueous TFA; Solution B=acetonitrile containing 0.1% of TFA). Fractions were collected and checked on analytical HPLC. Those containing pure product were combined and lyophilized to dryness. 85 mg of a white solid was obtained. Purity was >99% based on analytical HPLC analysis. Electro-spray mass spectrometer analysis gave the molecular weight at 3972.4 (in agreement with the calculated molecular weight of 3972.7).

The synthesis and purification of [Cha^{22}, $Leu^{23,28,31}$, Glu^{25}, $Lys^{26,30}$, Ahc^{27}, Aib^{29}]hPTHrP(1–34)NH_2 was carried out in the same manner as the above synthesis of [$Glu^{22,25}$, $Leu^{23,28}$, $Lys^{26,30}$, Aib^{29}, Ahc^{31}]hPTHrP(1–34)NH_2. The protected amino acid Boc-Cha-OH was purchased from Bachem, Calif. The purity of the final product was >99%, and the electron-spray mass spectrometer gave the molecular weight at 3997.2 (calculated molecular weight is 3996.8).

The full names for the abbreviations used above are as follows: Boc for t-butyloxycarbonyl, HF for hydrogen fluoride, Fm for formyl, Xan for xanthyl, Bzl for benzyl, Tos for tosyl, DNP for 2,4-dinitrophenyl, DMF for dimethylformamide, DCM for dichloromethane, HBTU for 2-(1H-Benzotnazol-1-yl)-1,1,3,3-tetramethyl uronium hexafluorophosphate, DIEA for diisopropylethylamine. HOAc for acetic acid, TFA for trifluoroacetic acid, 2ClZ for 2-chlorobenzyloxycarbonyl, and OcHex for O-cyclohexyl.

The substituents R_1 and R_2 of the above generic formula may be attached to the free amine of the N-terminal amino acid by standard methods known in the art. For example,

alkyl groups, e.g., C_{1-12} alkyl, may be attached using reductive alkylation. Hydroxyalkyl groups, e.g., C_{1-12} hydroxyalkyl, may also be attached using reductive alkylation wherein the free hydroxy group is protected with a t-butyl ester. Acyl groups, e.g., COE_1, may be attached by coupling the free acid, e.g., E_1COOH, to the free amine of the N-terminal amino acid by mixing the completed resin with 3 molar equivalents of both the free acid and diisopropylcarbodiimide in methylene chloride for one hour and cycling the resulting resin through steps (a) to (f) in the above wash program. If the free acid contains a free hydroxy group, e.g., p-hydroxyphenylpropionic acid, then the coupling should be performed with an additional 3 molar equivalents of HOBT.

Other peptides of this invention can be prepared in an analogous manner by a person of ordinary skill in the art.

Functional Assays

A. Binding to PTH Receptor

The peptides of the invention were tested for their ability to bind to the PTH receptor present on the SaOS-2 (human osteosarcoma cells). SaOS-2 cells (American Type Culture Collection. Rockville, Md.; ATCC #HTB 85) were maintained in RPMI 1640 medium (Sigma, St. Louis, Mo.) supplemented with 10% fetal bovine serum (FBS) and 2 mM glutamine at 37EC in a humidified atmosphere of 5% CO_2 in air. The medium was changed every three or four days and the cells were subcultured every week by trypsinization.

SaOS-2 cells were maintained for four days until they had reached confluence. The medium was replaced with 5% FBS in RPMI 1640 medium and incubated for 2 hrs at room temperature with 10×10^4 cpm mono-^{125}I-[Nle^{8,18}, Tyr^{34}(3-^{125}I)]bPTH(1–34)NH_2 in the presence of a competing peptides of the invention at various concentrations between 10^{-11}M to 10^{-4}M. The cells were washed four times with ice-cold PBS and lysed with 0.1 M NaOH, and the radioactivity associated with the cells was counted in a scintillation counter. Synthesis of mono-^{125}I-[Nle^{8,18}, Tyr^{34}(3-^{125}I)bPTH (1–34)NH_2 was carried out as described in Goldman, M. E., et al., Endocrinol., 123:1468 (1988).

The binding assay was conducted with various peptides of the invention, and the Kd value (half maximal inhibition of binding of mono-^{125}I-[Nle^{8,18}, Tyr^{34}(3-^{125}I)]bPTH(1–34)NH_2) for each peptide was calculated.

As shown in Table I, all of the tested peptides had a high binding affinity for the PTH receptor on the SaOS-2 cell.

B. Stimulation of Adenylate Cyclase Activity

The ability of the peptides of the invention to induce a biological response in SaOS-2 cells were measured. More specifically, any stimulation of the adenylate cyclase was determined by measuring the level of synthesis of cAMP (adenosine 3',5'-monophosphate) as described previously in Rodan, et al., J. Clin. Invest. 72: 1511 (1983) and Goldman, et al., Endocrinol., 123:1468 (1988). Confluent SAOS-2 cells in 24 wells plates were incubated with 0.5:Ci [^{3}H] adenine (26.9 Ci/mmol, New England Nuclear, Boston, Mass.) in fresh medium at 37EC for 2 hrs, and washed twice with Hank's balanced salt solution (Gibco, Gaithersburg, Md.). The cells were treated with 1 mM IBMX [isobutylmethyl-xanthine, Sigma, St. Louis, Mo.] in fresh medium for 15 min, and the peptides of the invention were added to the medium to incubate for 5 min. The reaction was stopped by the addition of 1.2 M trichloroacetic acid (TCA) (Sigma, St. Louis, Mo.) followed by sample neutralization with 4N KOH. cAMP was isolated by the two-column chromatographic method (Salmon, et al. 1974, Anal. Biochem. 58, 541). The radioactivity was counted in a scintillation counter (Liquid Scintillation Counter 2200CA, PACKARD, Downers Grove, Ill.).

The respective EC_{50} values (half maximal stimulation of adenylate cyclase) for the tested peptides were calculated and shown in Table I. All tested peptides were found to be potent stimulators of adenylate cyclase activity, which is a biochemical pathway indicative as a proximal signal for osteoblast proliferation (e.g. bone growth).

TABLE I

PEPTIDE	Kd (μM)	EC_{50} (nM)
[Cha^{7,11}]hPTH(1-34)NH_2	0.01	0.6
[Cha^{23}]hPTH(1-34)NH_2	0.2	20
[Cha^{24}]hPTH(1-34)NH_2	0.1	10
[Nle^{8,18}, Cha^{27}]hPTH(1-34)NH_2;	0.05	2
[Cha^{28}]hPTH(1-34)NH_2	0.05	2.5
[Cha^{31}]hPTH(1-34)NH_2	0.03	4
[Aib^{10}]hPTH(1-34)NH_2,	0.004	0.7
[Aib^{19}]hPTH(1-34)NH_2;	0.005	0.6
[Aib^{34}]hPTH(1-34)NH_2,	0.007	3
[Nle^{31}]hPTH(1-34)NH_2,	0.004	0.7
[hArg^{27}]hPTH(1-34)NH_2	0.007	1
[Dap, Nle^{8,18}, Tyr^{34}]hPTH(1-34)NH_2	0.150	10
[Cha^{24,28,31}, Lys^{30}]hPTH(1-34)NH_2;	0.5	7
[Cha^{7,11}, Nle^{8,18}, Tyr^{34}]hPTH(1-34)NH_2	0.006	0.6
[Cha^{7,11}, Nle^{8,18}, Aib^{16,19}, Tyr^{34}]hPTH(1-34)NH_2	0.005	1.5
[Cha^{7,11}, Nle^{8,18,31}, Aib^{16,19}, Tyr^{34}]hPTH(1-34)NH_2	0.04	4
[Cha^{11}]hPTH(1-34)NH_2	0.005	2
[Cha^{28,31}]hPTH(1-34)NH_2	0.06	7
[Cha^{7,11}, Nle^{8,18,31}, Aib^{34}]hPTH(1-34)NH_2	0.03	1.5
[Cha^{15}]hPTH(1-34)NH_2	0.005	1.3
[Cha$^{7\ 11}$, Aib^{19}]hPTH(1-34)NH_2	0.007	0.5
[Cha$^{7\ 11}$, Aib^{16}]hPTH(1-34)NH_2	0.004	1.1
[Aib$^{16\ 19}$]hPTH(1-34)NH_2	0.004	0.6
[Aib^{17}]hPTH(1-34)NH_2	0.005	2
[Aib^{3}]hPTH(1-34)NH_2	0.004	1.1
[Cha^{7,11}, Aib^{19}, Lys^{30}]hPTH(1-34)NH_2	0.004	2
[Cha^{7}]hPTH(1-34)NH_2	0.02	2.3
[Cha^{24,28,31}]hPTH(1-34)NH_2	1.0	30
[Aib^{17}]hPTH(1-34)NH_2	0.05	3
[Cha^{7,11,15}]hPTH(1-34)NH_2	0.01	1.4

TABLE II

PEPTIDE	Kd (μM)	EC_{50} (nM)
[Glu^{22,25}, Leu^{23,28}, Lys^{26,30}, Aib^{29}, Ahc^{31}]hPTHrP(1-34)NH_2	0.200	3.7
[Glu^{22,25}, Ahc^{23}, Lys^{26,30}, Leu^{28,31}, Aib^{29}]hPTHrP(1-34)NH_2	0.070	3.9
[Glu^{22,25}, Leu^{23,28,31}, Lys^{26,30}, Ahc^{27}, Aib^{29}]hPTHrP(1-34)NH_2	0.230	3.0
[Glu^{22,25,29}, Leu^{23,28,31}, Lys^{26}, Ahc^{30}]hPTHrP(1-34)NH_2	0.230	20
[Cha^{22}, Leu^{23,28,31}, Glu^{25}, Lys^{26,30}, Ahc^{27}, Aib^{29}]hPTHrP(1-34)NH_2	0.060	2.0
[Glu^{22,25}, Leu^{23,28,31}, Ahc^{24}, Lys^{26,30}, Aib^{29}]hPTHrP(1-34)NH_2	0.006	0.5
[Glu^{22,29}, Leu^{23,28,31}, Aib^{25}, Lys^{26,30}, Ahc^{27}]hPTHrP(1-34)NH_2		5
[Glu^{22}, Leu^{23,28,31}, Aib^{25,29}, Lys^{26,30}, Ahc^{27}]hPTHrP(1-34)NH_2		2
Ahc^{22}, Leu^{23,28,31}, Glu^{25}, Lys^{26,30}, Aib^{29}]hPTHrP(1-34)NH_2		0.3
[Glu^{22,25}, Leu^{23,31}, Lys^{26,30}, Ahc^{28}, Aib^{29}]hPTHrP(1-34)NH_2		0.5
[Cha^{22}, Ahc^{23}, Glu^{25}, Lys^{26,30}, Leu^{28,31}, Aib^{29}]hPTHrP(1-34)NH_2		0.4

Other Embodiments

It is to be understood that while the invention has been described in conjunction with the detailed description thereof, that the foregoing description is intended to illustrate and not limit the scope of the invention, which is defined by the scope of the appended claims. Other aspects, advantages, and modifications are within the claims.

SEQUENCE LISTING

<160> NUMBER OF SEQ ID NOS: 5

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<220> FEATURE:
<221> NAME/KEY: VARIANT
<222> LOCATION: 3
<223> OTHER INFORMATION: Xaa = Ser, Thr, or Aib
<220> FEATURE:
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<222> LOCATION: 5
<223> OTHER INFORMATION: Xaa = Leu, Nle, Ile, Cha, beta-Nal, Trp, Pal, Acc, Phe or p-X-Phe in which X is OH, a halogen, or CH3
<220> FEATURE:
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<222> LOCATION: 8
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<220> FEATURE:
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<220> FEATURE:
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<222> LOCATION: 15
<223> OTHER INFORMATION: Xaa = Leu, Nle, Ile, Cha, beta-Nal, Trp, Pal, Acc, Phe, or p-X-Phe, in which X is OH, a halogen, or CH3
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<220> FEATURE:
<221> NAME/KEY: VARIANT
<222> LOCATION: 22
<223> OTHER INFORMATION: Xaa = Acc or Glu
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<223> OTHER INFORMATION: Xaa = Leu, Acc, or Cha
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<222> LOCATION: 31
<223> OTHER INFORMATION: Xaa = Val, Leu, Nle, Acc, Cha, or deleted
<220> FEATURE:
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<221> NAME/KEY: VARIANT
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<223> OTHER INFORMATION: Xaa = Asn or deleted
<220> FEATURE:
<221> NAME/KEY: VARIANT
<222> LOCATION: 34
<223> OTHER INFORMATION: Xaa = Phe, Tyr, Amp, Aib, or deleted
<220> FEATURE:
<223> OTHER INFORMATION: This sequence may also encompass a deletion peptide with a length of 30, 31, 32 or 33 amino acids

<400> SEQUENCE: 1

Xaa Val Xaa Glu Xaa Gln Xaa Xaa His Asn Xaa Xaa Lys His Xaa Xaa
1 5 10 15

Xaa Xaa Xaa Arg Xaa Xaa Xaa Xaa Arg Lys Xaa Xaa Xaa Xaa Xaa Xaa
20 25 30

Xaa Xaa

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<220> FEATURE:
<221> NAME/KEY: VARIANT
<222> LOCATION: 3
<223> OTHER INFORMATION: Xaa = Ser, Thr, or Aib
<220> FEATURE:
<221> NAME/KEY: VARIANT
<222> LOCATION: 5
<223> OTHER INFORMATION: Xaa = Leu, Nle, Ile, Cha, beta-Nal, Trp, Pal, Phe, or p-X-Phe in which X is OH, a halogen, or CH3
<220> FEATURE:
<221> NAME/KEY: VARIANT
<222> LOCATION: 7
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<220> FEATURE:
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<222> LOCATION: 8
<223> OTHER INFORMATION: Xaa = Met, Nva, Leu, Val, Ile, Cha, or Nle
<220> FEATURE:
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<222> LOCATION: 11

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<223> OTHER INFORMATION: Xaa = Leu, Nle, Ile, Cha, beta-Nal, Trp, Pal, Phe, or p-X-Phe in which X is OH, a halogen, or CH3
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<222> LOCATION: 12
<223> OTHER INFORMATION: Xaa = Gly or Aib
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<222> LOCATION: 15
<223> OTHER INFORMATION: Xaa = Leu, Nle, Ile, Cha, beta-Nal, Trp, Pal, Phe, or p-X-Phe in which X is OH, a halogen, or CH3
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<221> NAME/KEY: VARIANT
<222> LOCATION: 16
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<220> FEATURE:
<221> NAME/KEY: VARIANT
<222> LOCATION: 17
<223> OTHER INFORMATION: Xaa = Ser, Thr, or Aib
<220> FEATURE:
<221> NAME/KEY: VARIANT
<222> LOCATION: 18
<223> OTHER INFORMATION: Xaa = Met, Nva, Leu, Val, Ile, Nle, Cha, or Aib
<220> FEATURE:
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<222> LOCATION: 19
<223> OTHER INFORMATION: Xaa = Glu or Aib
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<223> OTHER INFORMATION: Xaa = Val, Cha, or Met
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<222> LOCATION: 23
<223> OTHER INFORMATION: Xaa = Trp or Cha
<220> FEATURE:
<221> NAME/KEY: VARIANT
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<223> OTHER INFORMATION: Xaa = Leu or Cha
<220> FEATURE:
<221> NAME/KEY: VARIANT
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<223> OTHER INFORMATION: Xaa = Lys, Aib, Leu, hArg, Gln, or Cha
<220> FEATURE:
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<222> LOCATION: 28
<223> OTHER INFORMATION: Xaa = Leu or Cha
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<223> OTHER INFORMATION: Xaa = Asp or Lys
<220> FEATURE:
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<222> LOCATION: 31
<223> OTHER INFORMATION: Xaa = Val, Nle, Cha, or deleted
<220> FEATURE:
<221> NAME/KEY: VARIANT
<222> LOCATION: 32
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<220> FEATURE:
<221> NAME/KEY: VARIANT
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<223> OTHER INFORMATION: Asn or deleted
<220> FEATURE:
<221> NAME/KEY: VARIANT
<222> LOCATION: 34
<223> OTHER INFORMATION: Xaa = Phe, Tyr, Amp, Aib, or deleted
<220> FEATURE:
<223> OTHER INFORMATION: This sequence may also encompass a deletion peptide with a length of 30, 31, 32 or 33 amino acids

<400> SEQUENCE: 2

Xaa Val Xaa Glu Xaa Gln Xaa Xaa His Asn Xaa Xaa Lys His Xaa Xaa
1 5 10 15

Xaa Xaa Xaa Arg Xaa Glu Xaa Xaa Arg Lys Xaa Xaa Gln Xaa Xaa Xaa
20 25 30

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Xaa Xaa

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<222> LOCATION: 3
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<222> LOCATION: 5
<223> OTHER INFORMATION: Xaa = His, Ile, or Cha
<220> FEATURE:
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<223> OTHER INFORMATION: Xaa = Leu, Cha, Ile, beta-Nal, Trp, Pal, Phe,
or p-X-Phe in which X is OH, a halogen, or CH3
<220> FEATURE:
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<222> LOCATION: 8
<223> OTHER INFORMATION: Xaa = Leu, Met or Cha
<220> FEATURE:
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<222> LOCATION: 10
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<220> FEATURE:
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<222> LOCATION: 11
<223> OTHER INFORMATION: Xaa = Lys, Leu, Cha, Phe, or beta-Nal
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<222> LOCATION: 12
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<220> FEATURE:

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<223> OTHER INFORMATION: Xaa = His, Aib, or Lys
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<222> LOCATION: 27
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<220> FEATURE:
<221> NAME/KEY: VARIANT
<222> LOCATION: 28
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<220> FEATURE:
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<222> LOCATION: 30
<223> OTHER INFORMATION: Xaa = Glu, Cha, Aib, or Lys
<220> FEATURE:
<221> NAME/KEY: VARIANT
<222> LOCATION: 31
<223> OTHER INFORMATION: Xaa = Ile, Leu, Cha, Lys, or deleted
<220> FEATURE:
<221> NAME/KEY: VARIANT
<222> LOCATION: 32
<223> OTHER INFORMATION: Xaa = His or deleted
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<221> NAME/KEY: VARIANT
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<223> OTHER INFORMATION: Xaa = Ala or deleted
<220> FEATURE:
<223> OTHER INFORMATION: This sequence may also encompass a deletion peptide with a length of 30, 31, 32 or 33 amino acids

<400> SEQUENCE: 3

Xaa Val Xaa Glu Xaa Gln Xaa Xaa His Xaa Xaa Xaa Lys Xaa Xaa Xaa
1 5 10 15

Xaa Xaa Xaa Arg Arg Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
20 25 30

Xaa Xaa

<210> SEQ ID NO 4
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<221> NAME/KEY: VARIANT
<222> LOCATION: 5
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<220> FEATURE:
<221> NAME/KEY: VARIANT
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<223> OTHER INFORMATION: Xaa = Leu, Cha, Ile, beta-Nal, Trp, Pal, Phe, or p-X-Phe in which X is OH, a halogen, or CH3
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<222> LOCATION: 8
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<220> FEATURE:
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<220> FEATURE:
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<222> LOCATION: 31
<223> OTHER INFORMATION: Xaa = Ile, Leu, Cha, Lys, Acc, or deleted
<220> FEATURE:

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<221> NAME/KEY: VARIANT
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<223> OTHER INFORMATION: Xaa = His or deleted
<220> FEATURE:
<221> NAME/KEY: VARIANT
<222> LOCATION: 33
<223> OTHER INFORMATION: Xaa = Thr or deleted
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<222> LOCATION: 34
<223> OTHER INFORMATION: Xaa = Ala or deleted
<220> FEATURE:
<223> OTHER INFORMATION: This sequence may also encompass a deletion peptide with a length of 30, 31, 32 or 33 amino acids

<400> SEQUENCE: 4

Xaa Val Xaa Glu Xaa Gln Xaa Xaa His Xaa Xaa Xaa Lys Xaa Xaa Xaa
1 5 10 15

Xaa Xaa Xaa Arg Arg Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
20 25 30

Xaa Xaa

<210> SEQ ID NO 5
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<220> FEATURE:
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<222> LOCATION: 7
<223> OTHER INFORMATION: Xaa = Leu, Cha, Ile, beta-Nal, Trp, Pal, Acc, Phe, or p-X-Phe in which X is OH, a halogen, or CH3
<220> FEATURE:
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<223> OTHER INFORMATION: Xaa = Leu, Met, Acc, or Cha
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<223> OTHER INFORMATION: Xaa = Asp or Asn
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<223> OTHER INFORMATION: Xaa = Lys, Leu, Cha, Acc, Phe, or beta-Nal
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<223> OTHER INFORMATION: Xaa = Ser or His
<220> FEATURE:
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<223> OTHER INFORMATION: Xaa = Gln or Aib
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<222> LOCATION: 17
<223> OTHER INFORMATION: Xaa = Asp or Aib
<220> FEATURE:
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<222> LOCATION: 18
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<222> LOCATION: 19
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<221> NAME/KEY: VARIANT
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<223> OTHER INFORMATION: Xaa = Phe, Glu, Aib, Acc, or Cha
<220> FEATURE:
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<222> LOCATION: 27
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<223> OTHER INFORMATION: Xaa = Ile, Leu, Lys, Acc, or Cha
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<220> FEATURE:
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<223> OTHER INFORMATION: Xaa = His or deleted
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<221> NAME/KEY: VARIANT
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<220> FEATURE:
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<222> LOCATION: 34
<223> OTHER INFORMATION: Xaa = Ala or deleted
<220> FEATURE:
<223> OTHER INFORMATION: This sequence may also encompass a deletion
peptide with a length of 30, 31, 32 or 33 amino acids

<400> SEQUENCE: 5

Xaa Val Xaa Glu Xaa Gln Xaa Xaa His Xaa Xaa Xaa Lys Xaa Xaa Xaa
1 5 10 15

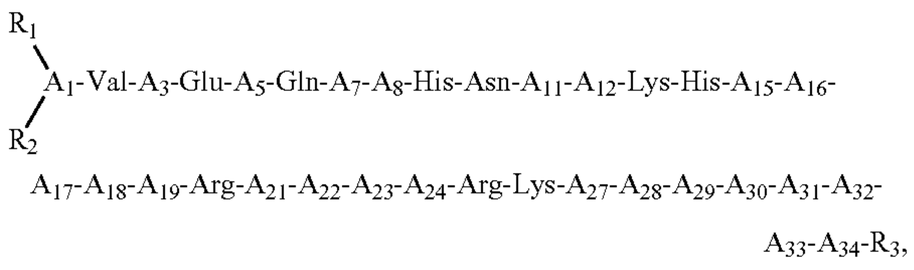
Xaa Xaa Xaa Arg Arg Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
20 25 30

Xaa Xaa

What is claimed is:

1. A peptide of formula (I):

[SEQ ID NO: 1]



R_1
\
A_1-Val-A_3-Glu-A_5-Gln-A_7-A_8-His-Asn-A_{11}-A_{12}-Lys-His-A_{15}-A_{16}-
/
R_2

A_{17}-A_{18}-A_{19}-Arg-A_{21}-A_{22}-A_{23}-A_{24}-Arg-Lys-A_{27}-A_{28}-A_{29}-A_{30}-A_{31}-A_{32}-A_{33}-A_{34}-R_3,

wherein

A_1 is Ser, Ala, or Dap;

A_3 is Ser, Thr, or Aib;

A_5 is Leu, Nle, Ile, Cha, β-Nal, Trp, Pal, Acc, Phe or p-X-Phe, in which X is OH, a halogen, or CH_3;

A_7 is Leu, Nle, Ile, Cha, β-Nal, Trp, Pal, Acc, Phe, or p-X-Phe in which X is OH, a halogen, or CH_3;

A_8 is Met, Nva, Leu, Val, Ile, Cha, Acc or Nle;

A_{11} is Leu, Nle, Ile, Cha, β-Nal, Trp, Pal, Acc, Phe or p-X-Phe in which X is OH, a halogen, or CH_3;

A_{12} is Gly, Acc or Aib;

A_{15} is Leu, Nle, Ile, Cha, β-Nal, Trp, Pal, **[**ACC**]** *Acc*, Phe, or p-X-Phe in which X is OH, a halogen, or CH_3;

A_{16} is Ser, Asn, Ala, or Aib;

A_{17} is Ser, Thr, or Aib;

A_{18} is Met, Nva, Leu, Val, Ile, Nle, Acc, Cha, or Aib;

A_{19} is Glu or Aib;

A_{21} is Val, Acc, Cha, or Met;

A_{22} is Acc or Glu;

A_{23} is Trp, Acc, or Cha;

A_{24} is Leu, Acc, or Cha;

A_{27} is Lys, Aib, Leu, hArg, Gln, Acc, or Cha;

A_{28} is Leu, Acc, or Cha;

A_{29} is Gln;

A_{30} is Asp or Lys;

A_{31} is Val, Leu, Nle, Acc, Cha, or deleted;

A_{32} is His or deleted;

A_{33} is Asn or deleted;

A_{34} is Phe, Tyr, Amp, Aib, or deleted;

each of R_1 and R_2 is, independently, H, $C_{1\text{-}12}$ alkyl, $C_{2\text{-}12}$ alkenyl, $C_{7\text{-}20}$ phenylalkyl, $C_{11\text{-}20}$ naphthylalkyl, $C_{1\text{-}12}$ hydroxyalkyl, $C_{2\text{-}12}$ hydroxyalkenyl, $C_{7\text{-}20}$ hydroxyphenylalkyl, or **[**$C_{7\text{-}20}$**]** $C_{11\text{-}20}$ hydroxynaphthylalkyl; or one and only one of R_1 and R_2 is COE_1 in which E_1 is $C_{1\text{-}12}$ alkyl, $C_{2\text{-}12}$ alkenyl, $C_{7\text{-}20}$ phenylalkyl, $C_{11\text{-}20}$ naphthylalkyl, $C_{1\text{-}12}$ hydroxyalkyl, $C_{2\text{-}12}$ hydroxyalkenyl, $C_{7\text{-}20}$ hydroxy-phenylalkyl, or **[**$C_{11\text{-}120}$**]** $C_{11\text{-}20}$ hydroxynaphthylalkyl; and

R_3 is OH, NH_2, $C_{1\text{-}12}$ alkoxy, or NH—Y—CH_2-Z in which Y is a $C_{1\text{-}12}$ hydrocarbon moiety and Z is H, OH, CO_2H, or $CONH_2$;

provided that at least one of A_5, A_7, A_8, A_{11}, A_{12}, A_{15}, A_{18}, A_{21}, A_{22}, A_{23}, A_{24}, A_{27}, A_{28}, and A_{31} is Acc; or a pharmaceutically acceptable salt thereof.

2. A peptide of claim **1**, wherein

A_3 is Ser;

A_5 is Ile or Acc;

A_7 is Leu, Acc, or Cha;

A_8 is Acc, Met, Nva, Leu, Val, Ile, or Nle;

A_{11} is Leu, Acc, or Cha;

A_{12} is Acc or Gly;

A_{15} is Leu, Acc, or Cha;

A_{16} is Asn or Aib;

A_{17} is Ser or Aib;

A_{18} is Acc, Met, or Nle;

A_{21} is Val or Acc;

A_{27} is Lys, hArg, Acc, or Cha;

A_{31} is Val, Leu, Nle, Acc, or Cha;

A_{32} is His;

A_{33} is Asn;

A_{34} is Phe, Tyr, Amp, or Aib;

or a pharmaceutically acceptable salt thereof.

3. A peptide of claim **2**, wherein

A_5 is Ile or Ahc;

A_7 is Leu, Ahc, or Cha;

A_8 is Ahc, Met, or Nle;

A_{11} is Leu, Ahc, or Cha;

A_{12} is Ahc or Gly;

A_{15} is Leu, Ahc, or Cha;

A_{18} is Met or Ahc;

A_{21} is Val or Ahc;

A_{22} is Glu or Ahc;

A_{23} is Trp, Ahc, or Cha;

A_{24} is Leu, Ahc, or Cha;

A_{27} is Lys, hArg, Ahc, or Cha;

A_{28} is Leu, Ahc, or Cha;

A_{31} is Val, Leu, Nle, Ahc, or Cha;

R_1 is H;

R_2 is H; and

R_3 is NH_2;

or a pharmaceutically acceptable salt thereof.

4. A peptide of claim **3**, wherein at least one of A_7, A_{11}, A_{15}, A_{23}, A_{24}, A_{27}, A_{28}, **[**or**]** *and* A_{31} is Cha.

5. A peptide of claim **3**, wherein at least one of A_{16}, A_{17}, A_{19}, **[**or**]** *and* A_{34} is Aib.

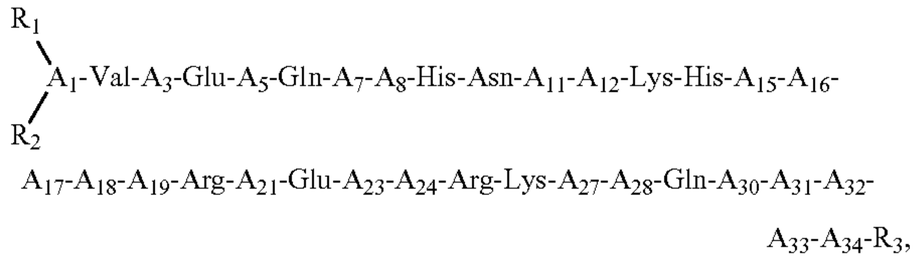
6. A peptide of the formula**[**:**]** : [$Cha^{22,23}$, Glu^{25}, $Lys^{26,30}$, Leu^{28}, Aib^{29}]hPTHrP(1–34)NH_2; [$Cha^{22,23}$, Glu^{25}, $Lys^{26,30}$, Aib^{29}]hPTHrP(1–34)NH_2; [$Glu^{22,25}$, $Leu^{23,28,31}$, Lys^{26}, Nle^{30}]hPTHrP(1–34)NH_2; [$Glu^{22,25}$, $Leu^{23,28,30,31}$, Lys^{26}, Aib^{29}]hPTHrP(1–34)NH_2**[**, **]]** ; [$Glu^{22,25,29}$, $Leu^{23,28,30,31}$, Lys^{26}]hPTHrP(1–34)NH_2; **[**[$Glu^{22,25,28}$, $Leu^{23,28,31}$, Lys^{26}, Nle^{30}]hPTHrP(1–34)NH_2;**]** [*Glu*^{22,25,29}, *Leu*^{23,28,31}, *Lys*^{26}, *Nle*^{30}]*hPTHrP(1–34)*NH_2; [Ser^1, Ile^5, Met^8, Asn^{10}, $Leu^{11,23,28,31}$, His^{14}, Cha^{15}, $Glu^{22,25}$, $Lys^{26,30}$, Aib^{29}]hPTHrP(1–34)NH_2; [$Glu^{22,25}$, Cha^{23}, Lys^{26}, $Leu^{28,31}$, Aib^{29}, Nle^{30}]hPTHrP(1–34)NH_2, [$Cha^{22,23}$, Glu^{25}, $Lys^{26,30}$, $Leu^{28,31}$, Aib^{29}]hPTHrP(1–34)NH_2; [Cha^{22}, $Leu^{23,28,31}$, $Glu^{25,29}$, Lys^{26}, Nle^{30}]hPTHrP(1–34)NH_2; [$Cha^{7,11,15}$] hPTHrP(1–34)NH_2; [$Cha^{7,8,15}$]hPTHrP(1–34)NH_2; [Glu^{22}, Cha^{23}, $Aib^{25,29}$, $Lys^{26,30}$, $Leu^{28,31}$]hPTHrP(1–34)NH_2; [Glu^{22}, Cha^{23}, $Aib^{25,29}$, Lys^{26}, Leu^{28}]hPTHrP(1–34)NH_2; [Glu^{22}, $Leu^{23,28}$, $Aib^{25,29}$, Lys^{26}]hPTHrP(1–34)NH_2; [Aib^{29}]hPTHrP(1–34)NH_2; [$Glu^{22,25}$, Cha^{23}, Lys^{26}, $Leu^{28,31}$, Aib^{29}, Nle^{30}]hPTHrP(1–34)NH_2; [$Glu^{22,25}$, Cha^{23}, $Lys^{26,30}$, Aib^{29}, Leu^{31}]hPTHrP(1–34)NH_2; **[**[Glu22,25, $Leu^{23,28,31}$, Lys^{26}, $Aib^{29,31}$]hPTHrP(1–34)NH_2**]** [*Glu*^{22,25}, *Leu*^{23,28,31}, *Lys*^{26}, *Aib*^{29,30}]*hPTHrP(1–34)*NH_2; [$Glu^{22,25}$, $Leu^{23,28,31}$, Lys^{26}, Aib^{29}]hPTHrP(1–34)NH_2; [$Glu^{22,25}$, $Leu^{23,28,31}$, $Aib^{26,29}$, Lys^{30}]hPTHrP(1–34)NH_2; [$Glu^{22,25}$, Cha^{23}, $Lys^{26,30}$, $Leu^{28,31}$, Aib^{29}]hPTHrP(1–34)NH_2;

[Glu^{22,25}, Cha^{23}, Lys^{26,30}, Aib^{29}]hPTHrP(1–34)NH$_2$; [[Glu^{22,25}, Cha^{23}, Lys26,30, Leu^{28}, Aib^{29}]hPTHrP(1–34)NH$_2$] [*Glu^{22,25}, Cha^{23}, Lys^{26,30}, Leu^{28}, Aib^{29}]hPTHrP(1–34)NH$_2$*; or [Leu^{27}, Aib^{29}]hPTHrP(1–34)NH$_2$; or a pharmaceutically acceptable salt thereof.

7. A peptide of the formula: [Glu^{22,25}, Leu^{23,28,31}, Lys^{26}, Ahc^{27}, Aib^{29}, Nle^{30}]hPTHrP(1–34)NH$_2$; [[Ser^{1}, Ile^{5}, Cha^{7,11}, Met^{8}, Asn^{10,} *His*^{14}, Glu^{22,25}, Leu^{23,28,31}, Lys^{26,30}, Ahc^{27}, Aib^{29}]hPTHrP(1–34)NH$_2$] [*Ser^{1}, Ile^{5}, Cha^{7,11}, Met^{8}, Asn^{10}, His^{14}, Glu^{22,25}, Leu^{23,28,31}, Lys^{26,30}, Ahc^{27}, Aib^{29}]hPTHrP(1–34)NH$_2$*; [(Ser^{1}, Ile^{5}, Met^{8}, Asn^{10}, Leu^{11,23,28,31}, His^{14}, Cha^{15}, Glu^{22,25}, Lys^{26,30}, Ahc^{27}, Aib^{29}]hPTHrP(1–34)NH$_2$] [*Ser^{1}, Ile^{5}, Met^{8}, Asn^{10}, Leu^{11,23,28,31}, His^{14}, Cha^{15}, Glu^{22,25}, Lys^{26,30}, Ahc^{27}, Aib^{29}]hPTHrP(1–34)NH$_2$*; [[Glu^{22}, Ahc^{23}, Aib^{25,29}, Lys^{26,30}, Leu^{28,31}]hPTHrP(1–34)NH^2] [*Glu^{22}, Ahc^{23}, Aib^{25,29}, Lys^{26,30}, Leu^{28,31}]hPTHrP(1–34) NH$_2$*; [Glu^{22,25}, Leu^{23,28,31}, Lys^{26,30}, Ahc^{29}]hPTHrP(1–34)NH$_2$; [Cha^{22}, Leu^{23,28,31}, Ahc^{24}, Glu^{25}, Lys^{26,30}, Aib^{29}]hPTHrP(1–34)NH$_2$; [Glu^{22,25}, Leu^{23,28,31}, Ahc^{24,27}, Lys^{26,30}, Aib^{29}]hPTHrP(1–34)NH$_2$; [Ahc^{22,24,27}, Leu^{23,28,31}, Glu^{25}, Lys^{26,30}, Aib^{29}]hPTHrP (1–34)NH$_2$; [Cha^{22}, Leu^{23,28,31}, Aib^{25,29}, Lys^{26,30}, Ahc^{27}]hPTHrP(1–34)NH$_2$; [Ahc^{22,27}, Leu^{23,28,31}, Aib^{25,29}, Lys^{26,30}]hPTHrP(1–34)NH$_2$; [Glu^{22}, Leu^{23,28,31}, Ahc^{24,27}, Lys^{25,26,30}, Aib^{29}]hPTHrP(1–34)NH$_2$; [Glu^{22}, Leu^{23,28}, Ahc^{24,27}, Lys^{25,26,30}, Aib^{29}]hPTHrP(1–34)NH$_2$; [Glu^{22}, Cha^{23}, Ahc^{24,27}, Lys^{25,26,30}, Leu^{28}, Aib^{29}]hPTHrP(1–34) NH$_2$; [[Glu^{22,25}, Cha^{23}, Ahc^{24,27}, Lys^{25,26,30}, Leu^{28,31}, Aib^{29}]hPTHrP(1–34)NH$_2$] [*Glu^{22,25}, Cha^{23}, Ahc^{24,27}, Lys^{26,30}, Leu^{28,31}, Aib^{29}]hPTHrP(1–34)NH$_2$*; [Glu^{22}, Cha^{23}, Ahc^{24,27}, Lys^{25,26,30}, Leu^{28,31}, Aib^{29}]hPTHrP(1–34)NH$_2$; [Glu^{22}, Leu^{23,28,31}, Ahc^{24,27}, Lys^{25,26,30}, Aib^{29}]hPTHrP(1–34)NH$_2$; [Glu^{22}, Leu^{23,28}, Ahc^{24,27}, Lys^{25,26}, Aib^{29}]hPTHrP(1–34)NH$_2$; [[Glu^{22}, Cha^{23}, Ahc, Lys^{25,26}, Leu^{28,31}, Aib^{29}]hPTHrP(1–34)NH$_2$] [*Glu^{22}, Cha^{23}, Ahc^{24,27}, Lys^{25,26}, Leu^{28,31}, Aib^{29}]hPTHrP(1–34)NH$_2$*; [[Glu^{22}, Cha^{23}, Ahc^{24,27}, Lys^{25,26}, Leu^{28,31}, Aib^{29}]hPTHrP(1–34)NH$_2$] [*Glu^{22}, Cha^{23}, Ahc^{24,27}, Lys^{25,26}, Leu^{28}, Aib^{29}]hPTHrP(1–34)NH$_2$*; [Glu^{22}, Leu^{23,28}, Lys^{25,26}, Ahc^{27}, Aib^{29}]hPTHrP(1–34)NH$_2$; [[Glu^{22}, Leu^{23,28,31}, Lys^{25,26}, Ahc^{27}, Aib^{29}]hPTHrP(1–34)NH$_2$] [*Glu^{22}, Leu^{23,28,31}, Lys^{25,26}, Ahc^{27}, Aib^{29}]hPTHrP(1–34)NH$_2$*; [Glu^{22}, Leu^{23,28,31}, Lys^{25,26,30}, Ahc^{27}, Aib^{29}]hPTHrP(1–34)NH$_2$; [Glu^{22}, Leu^{23,28}, Lys^{25,26,30}, Ahc^{27}, Aib^{29}]hPTHrP(1–34) NH$_2$; [Glu^{22,25}, Leu^{23,28,31}, Lys^{26}, Aib^{29}, Ahc^{30}]hPTHrP (1–34)NH$_2$; [Aib^{22,29}, Leu^{23,28,31}, Glu^{25}, Lys^{26,30}]hPTHrP (1–34)NH$_2$; [[Cha^{22}, Ahc^{23}, Glu^{25,29}, Lys^{26,30}, Leu^{26,31}]hPTHrP(1–34)NH$_2$] [*Cha^{22}, Ahc^{23}, Glu^{25,29}, Lys^{26,30}, Leu^{28,31}]hPTHrP(1–34)NH$_2$*; [Cha^{22}, Leu^{23,28,31}, Ahc^{24}, Glu^{25,29}, Lys^{26,30}]hPTHrP(1–34) NH$_2$; [[Cha^{22}, Leu^{23,28,31}, Glu^{25,29}, Lys^{26,30}]hPTHrP(1–34) NH$_2$;] [Cha^{22}, Leu^{23,28,31}, Glu^{25,29}, Lys^{26,30}, Ahc^{27}]hPTHrP (1–34)NH$_2$; [Cha^{22}, Leu^{23,31}, Glu^{25,29}, Lys^{26,30}, Ahc^{28}] hPTHrP(1–34)NH$_2$; [Cha^{22}, Leu^{23,28,31}, Glu^{25,29}, Lys^{26}, Ahc^{30}]hPTHrP(1–34)NH$_2$; [Cha^{22}, Leu^{23,28}, Glu^{25,29}, Lys^{26,30}, Ahc^{31}]hPTHrP(1–34) NH$_2$; [Glu^{22,29}, Ahc^{23}, Aib^{25}, Lys^{26,30}, Leu^{28,31}]hPTHrP (1–34)NH$_2$; [Ahc^{22}, Leu^{23,28,31}, Aib^{25}, Lys^{26,30}, Glu^{29}]hPTHrP(1–34)NH$_2$; [(Glu^{22,29}, Leu^{23,28,31}, Ahc^{24}, Aib^{25}, Lys^{26,30}]hPTHrP (1–34)NH$_2$] [*Glu^{22,29}, Leu^{23,28,31}, Ahc^{24}, Aib^{25}, Lys^{26,30}] hPTHrP(1–34)NH$_2$*; [Glu^{22,29}, Leu^{23,31}, Aib^{25}, Lys^{26,30}, Ahc^{28}]hPTHrP(1–34)NH$_2$; [Glu^{22,29}, Leu^{23,28}, Aib^{25}, Lys^{26,30}, Ahc^{31}]hPTHrP(1–34)NH$_2$; [[Glu^{22,25,29}, Leu^{23,28,31}, Aib^{25}, Lys^{26}, Ahc^{30}]hPTHrP(1–34)NH$_2$] [*Glu^{22,25,29}, Leu^{23,28,31}, Aib^{25}, Lys^{26}, Ahc^{30}]hPTHrP(1–34) NH$_2$*; [Glu^{22,25,29}, Leu^{23,28,31}, Ahc^{24}, Lys^{26}, Aib^{30}] hPTHrP (1–34)NH$_2$; [Ahc^{22}, Leu^{23,28,31}, Glu^{25,29}, Lys^{26}, Aib^{30}]hPTHrP(1–34)NH$_2$; [Ahc^{22}, Leu^{23,28}, Glu^{25,29}, Lys^{26,30,31}]hPTHrP(1–34)NH$_2$; [[Glu^{22,25,29}, Leu^{23,28}, Lys^{26,31}, Ahc^{30}]hPTHrP(1–^{34})NH$_2$] [*Glu^{22,25,29}, Leu^{23,28}, Lys^{26,31}, Ahc^{30}]hPTHrP(1–34)NH$_2$*; [Glu^{22,25,29}, Leu^{23,28}, Lys^{26,30,31}, Ahc^{27}]hPTHrP(1–34)NH$_2$; [Ahc^{22}, Cha^{23}, Glu^{25}, Lys^{26,30}, Leu^{28,31}, Aib^{29}]hPTHrP(1–34)NH$_2$; [Ahc^{22}, Cha^{23}, Lys^{25,26,30}, Leu^{28,31}, Aib^{29}]hPTHrP(1–34)NH$_2$; [Ahc^{22}, Cha^{23}, Lys^{25,26}, Leu^{28,31}, Aib^{29}]hPTHrP(1–34) NH$_2$; [Ahc^{22}, Leu^{23,28}, Lys^{25,26}, Aib^{29}]hPTHrP(1–34)NH$_2$; [Ahc^{22}, Leu^{23,28}, Arg^{25}, Lys^{26}, Aib^{29}]hPTHrP(1–34)NH$_2$; [Ahc^{22,24}, Leu^{23,28,31}, Glu^{25}, Lys^{26,30}, Aib^{29}]hPTHrP(1–34) NH$_2$; [Ahc^{22,24}, Leu^{23,28,31}, Lys^{25,26,30}, Aib^{29}]hPTHrP (1–34)NH$_2$; [Ahc^{22,24}, Leu^{23,28,31}, Lys^{25,26}, Aib^{29}]hPTHrP (1–34)NH$_2$; [[Ahc^{22,24}, Leu^{23,28}, Lys^{25,26}, Aib29]hPTHrP (1–34)NH$_2$] [*Ahc^{22,24}, Leu^{23,28}, Lys^{25,26}, Aib^{29}]hPTHrP (1–34)NH$_2$*; [Ahc^{22,24}, Leu^{23,28}, Arg^{25}, Lys^{26}, Aib^{29}] hPTHrP(1–34)NH$_2$; [[Glu^{22}, Leu^{23,28}, Lys^{23,28,31}, Ahc^{24}, Lys^{25,26,30}, Aib^{29}]hPTHrP(1–34)NH$_2$] [*Glu^{22}, Leu^{23,28,31}, Ahc^{24}, Lys^{25,26,30}, Aib^{29}]hPTHrP(1–34)NH$_2$*; [[Glu^{22}, Leu^{23,28,1}, Ahc^{24}, Lys^{25,26}, Aib^{29}]hPTHrP(1–34)NH$_2$] [*Glu^{22}, Leu^{23,28,31}, Ahc^{24}, Lys^{25,26}, Aib^{29}]hPTHrP(1–34) NH$_2$*; [Glu^{22}, Leu^{23,28}, Ahc^{24}, Lys^{25,26}, Aib^{29}]hPTHrP (1–34)NH$_2$; [Glu^{22}, Leu^{23,28,31}, Ahc^{24}, Arg^{25}, Lys^{26,30}, Aib^{29}]hPTHrP(1–34) NH$_2$; [Glu^{22}, Leu^{23,28,31}, Ahc^{24}, Arg^{25}, Lys^{26}, Aib^{29}]hPTHrP(1–34)NH$_2$; [Glu^{22}, Leu^{23,28}, Ahc^{24}, Arg^{25}, Lys^{26}, Aib^{29}]hPTHrP(1–34)NH$_2$; [Glu^{22}, Ahc^{23}, Aib^{25,29}, Lys^{26,30}, Leu^{28,31}]hPTHrP(1–34)NH$_2$; [Glu^{22}, Ahc^{23}, Aib^{25,29}, Lys^{26}, Leu^{28}]hPTHrP(1–34)NH$_2$; [Glu^{22}, Ahc^{23,31}, Aib^{25,29}, Lys^{26}, Leu^{28}]hPTHrP(1–34)NH$_2$; [Glu^{22}, Leu^{23,28}, Aib^{25,29}, Lys^{26,30}, Ahc^{31}]hPTHrP(1–34) NH$_2$; [[Glu^{2}, Leu^{23,28}, Aib^{25,29}, Lys^{26}, Ahc^{13}]hPTHrP (1–34)NH$_2$] [*Glu^{22}, Leu^{23,28}, Aib^{25,29}, Lys^{26}, Ahc^{31}]hPTHrP (1–34)NH$_2$*; [Glu^{22,25}, Leu^{23,28}, Ahc^{24,31}, Lys^{26,30}, Aib^{29}] hPTHrP(1–34)NH$_2$; [Glu^{22}, Leu^{23,28}, Ahc^{24,31}, Lys^{25,26}, Aib^{29}]hPTHrP(1–34) NH$_2$; [[Glu^{22}, Leu^{23,28,31}, Ahc^{24}, Aib^{25,29}, Lys^{26,30},]hPTHrP (1–34)NH$_2$] [*Glu^{22}, Leu^{23,28,31}, Ahc^{24}, Aib^{25,29}, Lys^{26,30}]hPTHrP(1–34)NH$_2$*; or a pharmaceutically acceptable salt thereof.

8. A peptide of formula (II):

[SEQ ID NO: 2]



R_1 / R_2

A_1-Val-A_3-Glu-A_5-Gln-A_7-A_8-His-Asn-A_{11}-A_{12}-Lys-His-A_{15}-A_{16}-A_{17}-A_{18}-A_{19}-Arg-A_{21}-Glu-A_{23}-A_{24}-Arg-Lys-A_{27}-A_{28}-Gln-A_{30}-A_{31}-A_{32}-A_{33}-A_{34}-R_3,

wherein

A_1 is Ser, Ala, or Dap;

A_3 is Ser, Thr, or Aib;

A_5 is Leu, Nle, Ile, Cha, β-Nal, Trp, Pal, Phe or p-X-Phe, in which X is OH, a halogen, or CH_3;

A_7 is Leu, Nle, Ile, Cha, β-Nal, Trp, Pal, Phe, or p-X-Phe in which X is OH, a halogen, or CH_3;

A_8 is Met, Nva, Leu, Val, Ile, Cha, or Nle;

A_{11} is Leu, Nle, Ile, Cha, β-Nal, Trp, Pal, Phe or p-X-Phe in which X is OH, a halogen, or CH_3;

A_{12} is Gly or Aib;

A_{15} is Leu, Nle, Ile, Cha, β-Nal, Trp, Pal, Phe, or p-X-Phe in which X is OH, a halogen, or CH_3;

A_{16} is Ser, Asn, Ala, or Aib,

A_{17} is Ser, Thr, or Aib;

A_{18} is Met, Nva, Leu, Val, Ile, Nle, Cha, or Aib;

A_{19} is Glu or Aib;
A_{21} is Val, Cha, or Met;
A_{23} is Trp or Cha;
A_{24} is Leu or Cha;
A_{27} is Lys, Aib, Leu, hArg, Gln, or Cha;
A_{28} is Leu or Cha;
A_{30} is Asp or Lys;
A_{31} is Val, Nle, Cha, or deleted;
A_{32} is His or deleted;
A_{33} is Asn or deleted;
A_{34} is Phe, Tyr, Amp, Aib, or deleted;
each of R_1 and R_2 is independently, H, C_{1-12} alkyl, C_{2-12} alkenyl, C_{7-20} phenylalkyl, C_{11-20} naphthylalkyl, C_{1-12} hydroxyalkyl, C_{2-12} hydroxyalkenyl, C_{7-20} hydroxyphenylalkyl, or C_{11-20} hydroxynaphthylalkyl; or one and only one of R_1 and R_2 is COE_1 in which E_1 is C_{1-12} alkyl, C_{2-12} alkenyl, C_{7-20} phenylalkyl, C_{11-20} naphthylalkyl, C_{1-12} hydroxyalkyl, C_{2-12} hydroxyalkenyl, C_{7-20} hydroxy-phenylalkyl, or C_{11-20} hydroxynaphthylalkyl; and
R_3 is OH, NH_2, C_{1-12} alkoxy, or **[**NH—Y—C_2-Z**]** *NH—Y—CH_2-—Z* in which Y is a C_{1-12} hydrocarbon moiety and Z is H, OH, CO_2H, or $CONH_2$;
provided that at least one of A_5, A_7, A_8, A_{11}, A_{15}, A_{18}, A_{21}, A_{23}, A_{24}, A_{27}, A_{28}, and A_{31} is Cha, or at least one of A_3, A_{12}, A_{16}, A_{17}, A_{18}, A_{19}, and A_{34} is Aib; or a pharmaceutically acceptable salt thereof;
provided that if A_{12} is Gly, then at least one of A_5 is Leu, Nle, β-Nal, Trp, Pal, Phe or p-X-Phe; A_{15} is Nle, Ile, β-Nal, Trp, Pal, Phe, or p-X-Phe; A_7 is Ile; or A_{11} is Ile or Pal;
provided that if A_{12} is Aib and A_1 is Dap, then at least one of A_5 is Leu, Nle, β-Nal, Trp, Pal, Phe or p-X-Phe; A_7 is Ile; A_{11} is Ile or Pal; or A_{15} is Nle, Ile, β-Nal, Trp, Pal, Phe, or p-X-Phe; and further
provided that the peptide is not [Aib^{12}, Tyr^{34}]hPTH(1–34)NH_2 or [Cha^8]hPTH(1–34)NH_2.

9. A peptide of claim **8** wherein at least one of A_7, A_{11}, A_{15}, A_{23}, A_{24}, A_{27}, A_{28}, and A_{31} is Cha; or a pharmaceutically acceptable salt thereof.

10. A peptide of claim **9**, wherein
A_3 is Ser;
A_5 is Ile;
A_7 is Leu or Cha;
A_8 is Met, Nva, Leu, Val, Ile, or Nle;
A_{11} is Leu or Cha;
A_{12} is Gly;
A_{15} is Leu or Cha;
A_{16} is Asn or Aib;
A_{17} is Ser;
A_{18} is Met or Nle;
A_{21} is Val;
A_{27} is Lys, hArg, or Cha;
[A_{32} is His;**]**
A_{31} is Val, Nle, or Cha;
A_{32} is His;
A_{33} is Asn;
A_{34} is Phe, Tyr, Amp, or Aib;
R_1 is H;
R_2 is H; and
R_3 is NH_2;
or a pharmaceutically acceptable salt thereof.

11. A peptide of claim **10**, wherein at least one of A_7 and A_{11} is Cha; or a pharmaceutically acceptable salt thereof.

12. A peptide of the formula: [$Cha^{7,11,15}$]hPTH(1–34)NH_2; or a pharmaceutically acceptable salt thereof.

13. A peptide of claim **10**, wherein at least one of A_{15}, A_{23}, A_{24}, A_{27}, A_{28}, and A_{31} is Cha; or a pharmaceutically acceptable salt thereof.

14. A peptide of claim **8**, wherein at least one of A_3, A_{12}, A_{16}, A_{17}, A_{18}, A_{19} and A_{34} is Aib; or a pharmaceutically acceptable salt thereof.

15. A peptide of claim **14**, wherein
A_3 is Ser or Aib;
A_5 is Ile;
A_7 is Leu or Cha;
A_8 is Met, Nva, Leu, Val, Ile, or Nle;
A_{11} is Leu or Cha;
A_{15} is Leu or Cha;
A_{16} is Asn or Aib;
A_{18} is Met, Aib, or Nle;
A_{21} is Val;
A_{27} is Lys, Aib, Leu, hArg, or Cha;
A_{31} is Val, Nle, or Cha;
A_{32} is His;
A_{33} is Asn;
A_{34} is Phe, Tyr, Amp, or Aib;
R_1 is H;
R_2 is H; and
R_3 is NH_2;
or a pharmaceutically acceptable salt thereof.

16. A peptide of claim **9**, wherein at least one of A_3, A_{12}, A_{16}, A_{17}, A_{19}, and A_{34} is Aib; or a pharmaceutically acceptable salt thereof.

17. A peptide of the formula: [Aib^{17}]hPTH(1–34)NH_2 or [Aib^{12}]hPTH(1–34)NH_2; or a pharmaceutically acceptable salt thereof.

18. A peptide of claim **8** wherein at least one of A_7, A_{11}, A_{15}, A_{23}, A_{24}, A_{27}, A_{28}, and A_{31} is Cha and at least one of A_3, A_{12}, A_{16}, A_{17}, A_{18}, A_{19}, and A_{34} is Aib; or a pharmaceutically acceptable salt thereof.

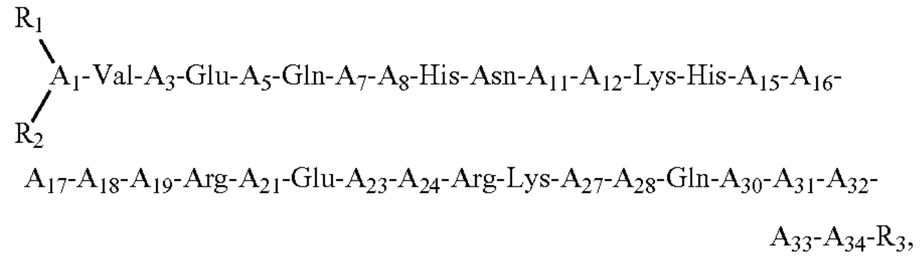
19. A peptide of claim **18**, wherein
A_3 is Ser or Aib;
A_5 is Ile;
A_7 is Leu or Cha;
A_8 is Met, Nva, Leu, Val, Ile, or Nle;
A_{11} is Leu or Cha;
A_{15} is Leu or Cha;
A_{16} is Asn or Aib;
A_{18} is Met, Aib, or Nle;
A_{21} is Val;
A_{27} is Lys, Aib, Leu, hArg, or Cha;
A_{31} is Val, Nle, or Cha;
A_{32} is His;
A_{33} is Asn;
A_{34} is Phe, Tyr, Amp, or Aib;
R_1 is H;
R_2 is H; and
R_3 is NH_2;
or a pharmaceutically acceptable salt thereof.

20. A peptide of claim **19**, wherein at least one of A_7 and A_{11} is Cha and at least one of A_{16}, A_{19}, and A_{34} is Aib; or a pharmaceutically acceptable salt thereof.

21. A peptide of claim **19**, wherein at least one of A_{24}, A_{28}, and A_{31} is Cha and at least one of A_{16} and A_{17} is Aib; or a pharmaceutically acceptable salt thereof.

22. A peptide of formula (III):

[SEQ ID NO: 2]



wherein

A_1 is Ser or Dap;

A_3 is Ser, Thr, or Aib;

A_5 is Leu, Nle, Ile, Cha, β-Nal, Trp, Pal, Phe or p-X-Phe, in which X is OH, a halogen, or CH_3;

A_7 is Leu, Nle, Ile, Cha, β-Nal, Trp, Pal, Phe, or p-X-Phe in which X is H, OH, a halogen, or CH_3;

A_8 is Met, Nva, Leu, Val, Ile, Cha, or Nle;

A_{11} is Leu, Nle, Ile, Cha, β-Nal, Trp, Pal, Phe or p-X-Phe in which X is OH, a halogen, or CH_3;

A_{12} is Gly or Aib;

A_{15} is Leu, Nle, Ile, Cha, β-Nal, Trp, Pal, Phe, or p-X-Phe in which X is OH, a halogen, or CH_3;

A_{16} is Ser, Asn, Ala, or Aib;

A_{17} is Ser, Thr, or Aib;

A_{18} is Met, Nva, Leu, Val, Ile, Nle, Cha, or Aib;

A_{19} is Glu or Aib;

A_{21} is Val, Cha, or Met;

A_{23} is Trp or Cha;

A_{24} is Leu or Cha;

A_{27} is Lys, Aib, Leu, hArg, Gln, or Cha

A_{28} is Leu or Cha;

A_{30} is Asp or Lys;

A_{31} is Val, Nle, Cha, or deleted;

A_{32} is His or deleted;

A_{33} is Asn or deleted;

A_{34} is Phe, Tyr, Amp, Aib, or deleted;

each of R_1 and R_2 is, independently, H, $C_{1\text{-}12}$ alkyl, $C_{2\text{-}12}$ alkenyl, $C_{7\text{-}20}$ phenylalkyl, $C_{11\text{-}20}$ naphthylalkyl, $C_{1\text{-}12}$ hydroxyalkyl, $C_{2\text{-}12}$ hydroxyalkenyl, $C_{7\text{-}20}$ hydroxyphenylalkyl, or $C_{11\text{-}20}$ hydroxynaphthylalkyl, or one and only one of R_1 and R_2 is COE_1 in which E_1 is $C_{1\text{-}12}$ alkyl, $C_{2\text{-}12}$ alkenyl, $C_{7\text{-}20}$ phenylalkyl, $C_{11\text{-}20}$ naphthylalkyl, $C_{1\text{-}12}$ hydroxyalkyl, $C_{2\text{-}12}$ hydroxyalkenyl, $C_{7\text{-}20}$ hydroxy-phenylalkyl, or $C_{1\text{-}120}$ hydroxynaphthylalkyl; and

R_3 is OH, NH_2, $C_{1\text{-}12}$ alkoxy, or NH'Y—CH_2-Z in which Y is a $C_{1\text{-}12}$ hydrocarbon moiety and Z is H, OH, CO_2H, or $CONH_2$; or a pharmaceutically acceptable salt thereof;

provided that at least one of A_1 is Dap; A_7 is β-Nal, Trp, Pal, Phe, or p-X-Phe; A_{15} is β-Nal, Trp, Pal, Phe, or p-X-Phe; A_{27} is hArg; or A_{31} is Nle;

provided that if A_{12} is Gly, then at least one of A_5 is Leu, Nle, β-Nal, Trp, Phe or p-X-Phe; A_{11} is Pal or p-X-Phe; or A_{15} is Nle, Ile, β-Nal, Trp, Pal, Phe or p-X-Phe;

provided that if A_{12} is Aib, then at least one of A_1 is Ser, A_5 is Leu, Nle, β-Nal, Trp, Phe or p-X-Phe; A_{11} is Pal or p-X-Phe; or A_{15} is Nle, Ile, β-Nal, Trp, Pal, Phe or p-X-Phe;

and also provided that said peptide of formula (III) is not [$hArg^{27}$]hPTH(1–34)NH_2.

23. A peptide of claim 22, wherein

A_1 is Ser or Dap;

A_3 is Ser or Aib;

A_8 is Met, Nva, Leu, Val, Ile, or Nle;

A_{16} is Asn or Aib;

A_{18} is Met, Aib, or Nle;

A_{21} is Val;

A_{27} is Lys, Aib, Leu, hArg, or Cha;

A_{31} is Val, Nle, or Cha;

A_{32} is His;

A_{33} is Asn;

A_{34} is Phe, Tyr, Amp, or Aib;

R_1 is H;

R_2 is H; and

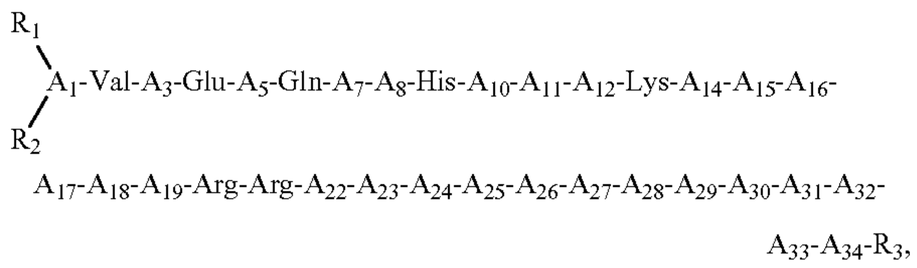
R_3 is NH_2;

or a pharmaceutically acceptable salt thereof.

24. A peptide of the formula **[**[Nle^{31}]±hPTH±(1–34)NH_2**]** [*Nle^{31}]hPTH(1–34)NH_2*; or a pharmaceutically acceptable salt thereof.

25. A peptide of formula (IV):

[SEQ ID NO: 3]



wherein

A_1 is Ala, Ser, or Dap;

A_3 is Ser or Aib;

A_5 is His, Ile, or Cha;

A_7 is Leu, Cha, Nle, β-Nal, Trp, Pal, Phe, or p-X-Phe in which X is OH, a halogen, or CH_3;

A_8 is Leu, Met, or Cha;

A_{10} is Asp or Asn;

A_{11} is Lys, Leu, Cha, Phe or β-Nal;

A_{12} is Gly or Aib;

A_{14} is Ser or His;

A_{15} is Ile, or Cha;

A_{16} is Gln or Aib;

A_{17} is Asp or Aib;

A_{18} is Leu, Aib, or Cha;

A_{19} is Arg or Aib;

A_{22} is Phe, Glu, Aib, or Cha;

A_{23} is Phe, Leu, Lys, or Cha;

A_{24} is Leu, Lys or Cha;

A_{25} is His, Aib, or Glu;

A_{26} is His, Aib, or Lys;

A_{27} is Leu, Lys, or Cha;

A_{28} is Ile, Leu, Lys or Cha;

A_{29} is Ala, Glu or Aib;

A_{30} is Glu, Cha, Aib or Lys;

A_{31} is Ile, Leu, Cha, Lys or deleted;

A_{32} is His or deleted;

A_{33} is Thr or deleted;

A_{34} is Ala, Aib or deleted;

each or R_1 and R_2 is, independently, H, C_{1-12} alkyl, C_{2-12} alkenyl, C_{7-20} phenylalkyl, C_{11-20} naphthylalkyl, **[**C_{1-20} hydroxyalkyl**]** *C_{1-12} hydroxyalkyl*, C_{2-12} hydroxyalkenyl, C_{7-20} hydroxyphenylalkyl, or C_{11-20} hydroxynaphthylalkyl; or one and only one of R_1 and R_2 is COE_1 in which E_1 is C_{1-12} alkyl, C_{2-12} alkenyl, C_{7-20} phenylalkyl, C_{11-20} naphthylalkyl, C_{1-12} hydroxyalkyl, C_{2-12} hydroxyalkenyl, C_{7-20} hydroxyphenylalkyl, or C_{11-20} hydroxynaphthylalkyl; and

R_3 is OH, NH_2, C_{1-12} alkoxy, or NH—Y—CH_2-Z in which Y is a C_{1-12} hydrocarbon moiety and Z is H, OH, CO_2H, or $CONH_2$;

provided that at least one of A_5,A_7, A_8, A_{11}, A_{15}, A_{18}, A_{22}, A_{23}, A_{24}, A_{27}, A_{28}, A_{30}, or A_{31} is Cha, or at least one of A_3, A_{12}, A_{16}, A_{17}, A_{18}, A_{19}, A_{22}, A_{25}, A_{26}, A_{29}, A_{30}, or A_{34} is Aib; or a pharmaceutically acceptable salt thereof;

provided that if A_{12} is Gly, then at least one of A_1 is Ser; A_5 is Ile; A_8 is Met; A_{10} is Asn; A_{11} is Leu; A_{14} is His; A_{22} is Aib; A_{23} is Leu or Lys; A_{24} is Lys; A_{25} is Aib or Glu; A_{27} is Lys; A_{28} is Leu or Lys; A_{29} is Glu or Aib; A_{30} is Cha or Aib; A_{31} is Leu or Lys; or A_{34} is Aib and

provided that said peptide is not **[**[Glu22,25, Leu23,28, 31, Aib^{29}, $Lys^{26,30}$]hPTHrP(1–34)NH_2**]** [*$Glu^{22,25}$, $Leu^{23,28,31}$, Aib^{29}, $Lys^{26,30}$]hPTHrP(1–34)NH_2*.

26. A peptide of claim **25**, wherein A_{22} is Phe or Cha; A_{23} is Phe or Cha; A_{25} is His; A_{26} is His; A_{27} is Leu or Cha; A_{28} is Ile or Cha; A_{29} is Ala; A_{30} is Glu or Lys; A_{31} is Ile or Cha; A_{32} is His; and A_{33} is Thr; or a pharmaceutically acceptable salt thereof.

27. A peptide of claim **26**, wherein at least one of A_7 and A_{11} is Cha; or a pharmaceutically acceptable salt thereof.

28. A peptide of claim **26**, wherein at least one of A_{16} **[**or**]** *and* A_{19} is Aib; or a pharmaceutically acceptable salt thereof.

29. A peptide of claim **25** wherein A_{22} is Glu, Aib, or Cha; A_{23} is Leu, Lys, or Cha; A_{25} is Aib or Glu; A_{26} is Aib or Lys; A_{28} is Leu, Lys, or Cha; A_{29} is Glu or Aib; A_{30} is Cha, Aib, or Lys; A_{31} is Leu, Cha, or Lys; A_{32} is His; A_{33} is Thr; and A_{34} is Ala; or a pharmaceutically acceptable salt thereof.

30. A peptide of claim **28**, wherein at least one of A_7 and A_{11} is Cha; or a pharmaceutically acceptable salt thereof.

31. A peptide of claim **29**, wherein at least one of A_{16} or A_{19} is Aib; or a pharmaceutically acceptable salt thereof.

32. A method of treating osteoporosis in a patient in need thereof, which comprises administering to said patient a peptide of the formula:

[SEQ ID NO: 1]

R_1
\
A_1-Val-A_3-Glu-A_5-Gln-A_7-A_8-His-Asn-A_{11}-A_{12}-Lys-His-A_{15}-A_{16}-
/
R_2

A_{17}-A_{18}-A_{19}-Arg-A_{21}-A_{22}-A_{23}-A_{24}-Arg-Lys-A_{27}-A_{28}-A_{29}-A_{30}-A_{31}-A_{32}-A_{33}-A_{34}-R_3,

wherein

A_1 is Ser, Ala or Dap;

A_3 is Ser, Thr or Aib;

A_5 is Leu, Nle, Ile, Cha, β-Nal, Trp, Pal, Acc, Phe or p-X-Phe in which X is OH, a halogen or CH_3;

A_7 is Leu, Nle, **[**Tie**]** *Ile*, Cha, β-Nal, Trp, Pal, *Acc,* Phe or p-X-Phe in which X is OH, a halogen or CH_3;

A_8 is Met, Nva, Leu, Val, **[**tie**]** *Ile*, Cha, Acc or Nle;

A_{11} is Leu, Nle, Ile, Cha, β-Nal, Trp, Pal, Acc, Phe or p-X-Phe in which X is OH, a halogen, or CH_3;

A_{12} is Gly, Acc or Aib;

A_{15} is Leu, Nle, Ile, Cha, β-Nal, Trp, Pal, Acc, Phe, or p-X-Phe in which X is OH, a halogen or CH_3;

A_{16} is Ser, Asn, Ala or Aib;

A_{17} is Ser, Thr or Aib;

A_{18} is Met, Nva, Leu, Val, Ile, Nle, Acc, Cha or Aib;

A_{19} is Glu or Aib;

A_{21} is Val, Acc, Cha or Met;

A_{22} is Acc or Glu;

A_{23} is Trp, Acc or Cha;

A_{24} is Leu, Acc or Cha;

A_{27} is Lys, Aib, Leu, hArg, Gln, Acc or Cha;

A_{28} is Leu, Acc or Cha;

A_{29} is Gln, Acc or Aib;

A_{30} is Asp or Lys;

A_{31} is Val, Leu, Nle, Acc, Cha, or deleted;

A_{32} is His or deleted;

A_{33} is Asn or deleted;

A_{34} is Phe, Tyr, Amp, Aib, or deleted;

each of R_1 and R_2 is, independently, H, C_{1-12} alkyl, C_{2-12} alkenyl, C_{7-20} phenylalkyl, C_{11-20} naphthylalkyl, C_{1-12} hydroxyalkyl, C_{2-12} hydroxyalkenyl, C_{7-20} hydroxyphenylalkyl or C_{11-20} hydroxynaphthylalkyl; or one and only one of R_1 and R_2 is COE_1 in which E_1 is C_{1-12} alkyl, C_{2-12} alkenyl, C_{7-20} phenylalkyl, C_{11-20} naphthylalkyl, C_{1-12} hydroxyalkyl, C_{2-12} hydroxyalkenyl, C_{7-20} hydroxy-phenylalkyl, or C_{11-20} hydroxynaphthylalkyl; and

R_3 is OH, NH_2, C_{1-12} alkoxy or NH—Y—CH_2-Z in which Y is a C_{1-12} hydrocarbon moiety and Z is H, OH, CO_2H or $CONH_2$;

provided that at least one of A_5, A_7, A_8, A_{11}, A_{12}, A_{15}, A_{18}, A_{21}, *A_{22}*, A_{23}, A_{24}, A_{27}, A_{28}, A_{29} and A_{31} is Acc; or a pharmaceutically acceptable salt thereof.

33. The method according to claim **32**, further comprising administering to said patient a bisphosphonate or calcitonin.

34. A method of treating osteoporosis in a patient in need thereof, which comprises administering to said patient a peptide of the formula:

[SEQ ID NO: 2]

R_1
\
A_1-Val-A_3-Glu-A_5-Gln-A_7-A_8-His-Asn-A_{11}-A_{12}-Lys-His-A_{15}-A_{16}-
/
R_2

A_{17}-A_{18}-A_{19}-Arg-A_{21}-Glu-A_{23}-A_{24}-Arg-Lys-A_{27}-A_{28}-Gln-A_{30}-A_{31}-A_{32}-A_{33}-A_{34}-R_3,

wherein

A_1 is Ser, Ala, or Dap;

A_3 is Ser, Thr, or Aib;

A_5 is Leu, Nle, Ile, Cha, β-Nal, Trp, Pal, Phe or p-X-Phe, in which X is OH, a halogen or CH_3;

A_7 is Leu, Nle, Ile, Cha, β-Nal, Trp, Pal, Phe or p-X-Phe in which X is OH, a halogen or CH_3;

A_8 is Met, Nva, Leu, Val, Ile, Cha, or Nle;

A_{11} is Leu, Nle, Ile, Cha, β-Nal, Trp, Pal, Phe or p-X-Phe in which X is OH, a halogen or CH_3;

A_{12} is Gly or Aib;

A_{15} is Leu, Nle, Ile, Cha, β-Nal, Trp, Pal, Phe or p-X-Phe in which X is OH, a halogen or CH_3;

A_{16} is Ser, Asn, Ala or Aib;

A_{17} is Ser, Thr or Aib;

A_{18} is Met, Nva, Leu, Val, Ile, Nle, Cha or Aib;

A_{19} is Glu or Aib;

A_{21} is Val, Cha or Met;

A_{23} is Trp or Cha;

A_{24} is Leu or Cha;

A_{27} is Lys, Aib, Leu, hArg, Gln or Cha;

A_{28} is Leu or Cha;

A_{30} is Asp or Lys;

A_{31} is Val, Nle, Cha or deleted;

A_{32} is His or deleted;

A_{33} is Asn or deleted;

A_{34} is Phe, Tyr, Amp, Aib or deleted;

each of R_1 and R_2 is, independently, H, C_{1-12} alkyl, C_{2-12} alkenyl, C_{7-20} phenylalkyl, C_{11-20} naphthylalkyl, C_{1-12} hydroxyalkyl, C_{2-12} hydroxyalkenyl, C_{7-20} hydroxyphenylalkyl or C_{11-20} hydroxynaphthylalkyl; or one and only one of R_1 and R_2 is COE_1 in which E_1 is C_{1-12} alkyl, C_{2-12} alkenyl, C_{7-20} phenylalkyl, C_{11-20} naphthylalkyl, C_{1-12} hydroxyalkyl, C_{2-12} hydroxyalkenyl, C_{7-20} hydroxy-phenylalkyl or C_{11-20} hydroxynaphthylalkyl; and

R_3 is OH, NH_2, C_{1-12} alkoxy or NH—Y—CH_2-Z in which Y is a C_{1-12} hydrocarbon moiety and Z is H, OH, CO_2H or $CONH_2$;

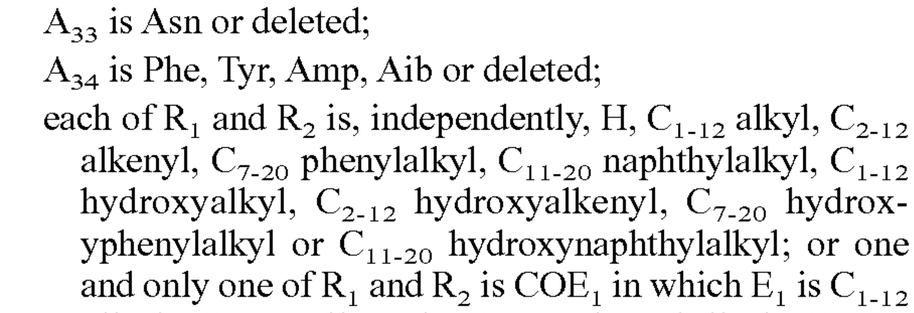
provided that at least one of A_5, A_7, A_8, A_{11}, A_{15}, A_{18}, A_{21}, A_{23}, A_{24}, A_{27}, A_{28}, and A_{31} is Cha or at least one of A_3, A_{12}, A_{16}, A_{17}, A_{18}, A_{19}, and A_{34} is Aib; or a pharmaceutically acceptable salt thereof; and

provided that the peptide is not [Aib^{12}, Tyr^{34}]hPTH(1–34)NH_2 or [Cha^8]hPTH(1–34)NH_2.

35. The method according to claim **34**, further comprising administering to said patient a bisphosphonate or calcitonin.

36. A method of treating osteoporosis in a patient in need thereof, which comprises administering to said patient a peptide of the formula:

[SEQ ID NO: 2]



R_1
\
A_1-Val-A_3-Glu-A_5-Gln-A_7-A_8-His-Asn-A_{11}-A_{12}-Lys-His-A_{15}-A_{16}-
/
R_2

A_{17}-A_{18}-A_{19}-Arg-A_{21}-Glu-A_{23}-A_{24}-Arg-Lys-A_{27}-A_{28}-Gln-A_{30}-A_{31}-A_{32}-

A_{33}-A_{34}-R_3,

wherein

A_1 is Ser or Dap;

A_3 is Ser, Thr, or Aib;

A_5 is Leu, Nle, Ile, Cha, β-Nal, Trp, Pal, Phe or p-X-Phe, in which X is OH, a halogen, or CH_3;

A_7 is Leu, Ile, Nle, Cha, β-Nal, Trp, Pal, Phe or p-X-Phe in which X is H, OH, a halogen or CH_3;

A_8 is Met, Nva, Leu, Val, Ile, Cha, or Nle;

A_{11} is Leu, Nle, Ile, Cha, β-Nal, Trp, Pal, Phe or p-X-Phe in which X is OH, a halogen or CH_3;

A_{12} is Gly or Aib;

A_{15} is Leu, Nle, Ile, Cha, β-Nal, Trp, Pal, Phe or p-X-Phe in which X is OH, a halogen or CH_3;

A_{16} is Ser, Asn, Ala or Aib;

A_{17} is Ser, Thr or Aib;

A_{18} is Met, Nva, Leu, Val, Ile, Nle, Cha or Aib;

A_{19} is Glu or Aib;

A_{21} is Val, Cha or Met;

A_{23} is Trp or Cha;

A_{24} is Leu or Cha;

A_{27} is Lys, Aib, Leu, hArg, Gln or Cha;

A_{28} is Leu or Cha;

A_{30} is Asp or Lys;

A_{31} is Val, Nle, Cha or deleted;

A_{32} is His or deleted;

A_{33} is Asn or deleted;

A_{34} is Phe, Tyr, Amp, Aib or deleted;

each of R_1 and R_2 is, independently, H, C_{1-12} alkyl, C_{2-12} alkenyl, C_{7-20} phenylalkyl, C_{11-20} naphthylalkyl, C_{1-12} hydroxyalkyl, C_{2-12} hydroxyalkenyl, C_{7-20} hydroxyphenylalkyl or C_{11-20} hydroxynaphthylalkyl; or one and only one of R_1 and R_2 is COE_1 in which E_1 is C_{1-12} alkyl, C_{2-12} alkenyl, C_{7-20} phenylalkyl, C_{11-20} naphthylalkyl, C_{1-12} hydroxyalkyl, C_{2-12} hydroxyalkenyl, C_{7-20} hydroxy-phenylalkyl, or C_{11-20} hydroxynaphthylalkyl; and

R_3 is OH, NH_2, C_{1-12} alkoxy or NH—Y—CH_2-Z in which Y is a C_{1-12} hydrocarbon moiety and Z is H, OH, CO_2H or $CONH_2$;

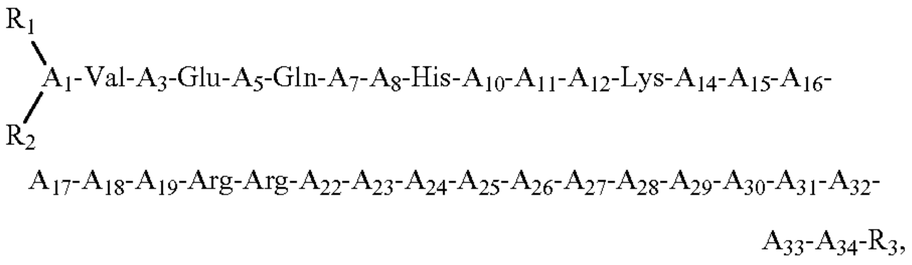
provided that at least one of A_1 is Dap; A_7 is β-Nal, Trp, Pal, Phe, or p-X-Phe; A_{15} is β-Nal, Trp, Pal, Phe, or p-X-Phe; A_{27} is hArg; or A_{31} is Nle; or a pharmaceutically acceptable salt thereof; and

provided that said peptide is not [$hArg^{27}$]hPTH(1–34)NH_2.

37. The method according to claim **[34]** *36*, further comprising administering to said patient a bisphosphonate or calcitonin.

38. A method of treating osteoporosis in a patient in need thereof, which comprises administering to said patient a peptide of the formula:

[SEQ ID NO: 3]



R_1
\
A_1-Val-A_3-Glu-A_5-Gln-A_7-A_8-His-A_{10}-A_{11}-A_{12}-Lys-A_{14}-A_{15}-A_{16}-
/
R_2

A_{17}-A_{18}-A_{19}-Arg-Arg-A_{22}-A_{23}-A_{24}-A_{25}-A_{26}-A_{27}-A_{28}-A_{29}-A_{30}-A_{31}-A_{32}-

A_{33}-A_{34}-R_3,

wherein

A_1 is Ala, Ser or Dap;

A_3 is Ser or Aib;

A_5 is His, Ile, or Cha;

A_7 is Leu, Cha, Nle, β-Nal, Trp, Pal, Phe or p-X-Phe in which X is OH, a halogen or CH_3;

A_8 is Leu, Met or Cha;

A_{10} is Asp or Asn;

A_{11} is Lys, Leu, Cha, Phe or β-Nal;

A_{12} is Gly or Aib;

A_{14} is Ser or His;

A_{15} is Ile or Cha;
A_{16} is Gln or Aib;
A_{17} is Asp or Aib;
A_{18} is Leu, Aib or Cha;
A_{19} is Arg or Aib;
A_{22} is Phe, Glu, Aib or Cha;
A_{23} is Phe, Leu, Lys or Cha;
A_{24} is Leu, Lys or Cha;
A_{25} is His, Aib or Glu;
A_{26} is His, Aib or Lys;
A_{27} is Leu, Lys or Cha;
A_{28} is Ile, Leu, Lys or Cha;
A_{29} is Ala, Glu or Aib;
A_{30} is Glu, Cha, Aib or Lys;
A_{31} is Ile, Leu, Cha, Lys or deleted;
A_{32} is His or deleted;
A_{33} is Thr or deleted;
A_{34} is Ala, Aib or deleted;
each of R_1 and R_2 is, independently, H, C_{1-12} alkyl, C_{7-20} phenylalkyl, C_{11-20} naphthylalkyl, C_{1-12} hydroxyalkyl, C_{2-12} hydroxyalkenyl, C_{7-20} hydroxyphenylalkyl or C_{11-20} hydroxynaphthylalkyl; or one and only one of R_1 and R_2 is COE_1 in which E_1 is C_{1-12} alkyl, C_{2-12} alkenyl, C_{7-20} phenylalkyl, C_{11-20} naphthylalkyl, C_{1-12} hydroxyalkyl, C_{2-12} hydroxyalkenyl, C_{7-20} hydroxyphenylalkyl, or C_{11-20} hydroxynaphthylalkyl; and
R_3 is OH, NH_2, C_{1-12} alkoxy or NH—Y—CH_2-Z in which Y is a C_{1-12} hydrocarbon moiety and Z is H, OH, CO_2H or $CONH_2$;
provided that at least one of A_5, A_7, A_8, A_{11}, A_{15}, A_{18}, A_{22}, A_{23}, A_{24}, A_{27}, A_{28}, A_{30}, or A_{31} is Cha; or at least one of A_3, A_{12}, A_{16}, A_{17}, A_{18}, A_{19}, A_{22}, A_{25}, A_{26}, A_{29}, A_{30} or A_{34} is Aib; or a pharmaceutically acceptable salt thereof; and
provided that said peptide is not **[**[Glu22,25, Leu23,28, 31, Aib^{29}, $Lys^{26,30}$]hPTHrP(1–34)NH_2**]** [*$Glu^{22,25}$, $Leu^{23,28,31}$, Aib^{29}, $Lys^{26,30}$]hPTHrP(1–34)NH_2*.

39. The method according to claim **38**, further comprising administering to said patient a bisphosphonate or calcitonin.

40. A method of treating osteoporosis in a patient in need thereof, which comprises administering to said patient a peptide of the formula:

[SEQ ID NO: 4]

R_1
\
A_1-Val-A_3-Glu-A_5-Gln-A_7-A_8-His-A_{10}-A_{11}-A_{12}-Lys-A_{14}-A_{15}-A_{16}-
/
R_2

A_{17}-A_{18}-A_{19}-Arg-Arg-A_{22}-A_{23}-A_{24}-A_{25}-A_{26}-A_{27}-A_{28}-A_{29}-A_{30}-A_{31}-A_{32}-

A_{33}-A_{34}-R_3,

wherein
A_1 is Ala, Ser or Dap;
A_3 is Ser or Aib;
A_5 is His, Ile, or Cha;
A_7 is Leu, Cha, Nle, β-Nal, Trp, Pal, Phe or p-X-Phe in which X is OH, a halogen or CH_3;
A_8 is Leu, Met or Cha;
A_{10} is Asp or Asn;
A_{11} is Lys, Leu, Cha, Phe or βNal;
A_{12} is Gly or Aib;
A_{14} is Ser or His;
A_{15} is Ile or Cha;
A_{16} is Gln or Aib;
A_{17} is Asp or Aib;
A_{18} is Leu, Aib or Cha;
A_{19} is Arg or Aib;
A_{22} is Phe, Glu, Aib, Acc or Cha;
A_{23} is Phe, Leu, Lys, Acc or Cha;
A_{24} is Leu, Lys, Acc or Cha;
A_{25} is His, Aib or Glu;
A_{26} is His, Aib or Lys;
A_{27} is Leu, Lys, Acc or Cha;
A_{28} is Ile, Leu, Lys, Acc or Cha;
A_{29} is Ala, Glu or Aib;
A_{30} is Glu, Cha, Aib, Acc or Lys;
A_{31} is Ile, Leu, Cha, Lys, Acc or deleted;
A_{32} is His or deleted;
A_{33} is Thr or deleted;
A_{34} is Ala or deleted;
each of R_1 and R_2 is, independently, H, C_{1-12} alkyl, C_{7-20} phenylalkyl, C_{11-20} naphthylalkyl, C_{1-12} hydroxyalkyl, C_{2-12} hydroxyalkenyl, C_{7-20} hydroxyphenylalkyl or C_{11-20} hydroxynaphthylalkyl; or one and only one of R_1 and R_2 is COE_1 in which E_1 is C_{1-12} alkyl, C_{2-12} alkenyl, C_{7-20} phenylalkyl, C_{11-20} naphthylalkyl, C_{1-12} hydroxyalkyl, **[**C_{2-2}**]** C_{2-20} hydroxyalkenyl, C_{7-20} hydroxyphenylalkyl or C_{11-20} hydroxynaphthylalkyl; and
R_3 is OH, NH_2, **[**C1-12**]** C_{1-12} alkoxy or NH—Y—CH_2-Z in which Y is a C_{1-12} hydrocarbon moiety and Z is H, OH, CO_2H or $CONH_2$;
provided that at least one of A_{23}, A_{24}, A_{28} or A_{31} is Lys; or a pharmaceutically acceptable salt thereof.

41. The method according to claim **40**, further comprising administering to said patient a bisphosphonate or calcitonin.

42. A pharmaceutical composition comprising the peptide of claim **1** or a pharmaceutically acceptable salt thereof and a pharmaceutically acceptable carrier or diluent.

43. A pharmaceutical composition comprising the peptide of claim **42** or a pharmaceutically acceptable salt thereof, a bisphosphonate or calcitonin, and a pharmaceutically acceptable carrier or diluent.

44. A pharmaceutical composition comprising the peptide of claim **8** or a pharmaceutically acceptable salt thereof and a pharmaceutically acceptable carrier or diluent.

45. A pharmaceutical composition comprising the peptide of claim **44** or a pharmaceutically acceptable salt thereof, a bisphosphonate or calcitonin, and a pharmaceutically acceptable carrier or diluent.

46. A pharmaceutical composition comprising the peptide of claim **22** or a pharmaceutically acceptable salt thereof and a pharmaceutically acceptable carrier or diluent.

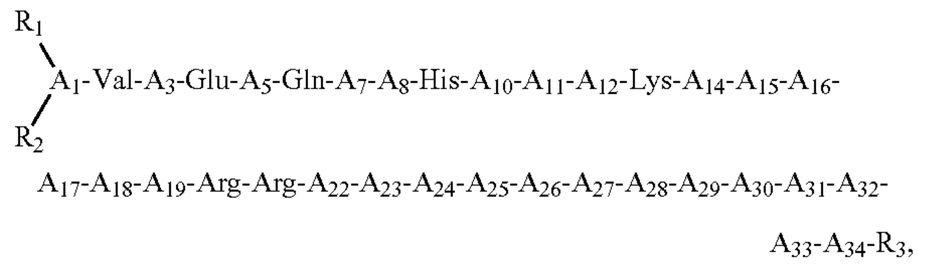
47. A pharmaceutical composition comprising the peptide of claim **46** or a pharmaceutically acceptable salt thereof, a bisphosphonate or calcitonin, and a pharmaceutically acceptable carrier or diluent.

48. A pharmaceutical composition comprising the peptide of claim **25** or a pharmaceutically acceptable salt thereof and a pharmaceutically acceptable carrier or diluent.

49. A pharmaceutical composition comprising the peptide of claim **48** or a pharmaceutically acceptable salt thereof, a bisphosphonate or calcitonin, and a pharmaceutically acceptable carrier or diluent.

50. A method of treating osteoporosis in a patient in need thereof, which comprises administering to said patient a peptide of the formula (VI):

[SEQ ID NO: 5]



wherein

A_1 is Ala, Ser, or Dap;

A_3 is Ser or Aib;

A_5 is His, Ile, Acc, or Cha;

A_7 is Leu, Cha, Nle, β-Nal, Trp, Pal, Acc, Phe or p-X-Phe in which X is OH, a halogen, or CH_3;

A_8 is Leu, Met, Acc, or Cha;

A_{10} is Asp or Asn;

A_{11} is Lys, Leu, Cha, Acc, Phe, or β-Nal;

A_{12} is Gly, Acc, or Aib;

A_{14} is Ser or His;

A_{15} is Ile, Acc, or Cha;

A_{16} is Gln or Aib;

A_{17} is Asp or Aib;

A_{18} is Leu, Aib, Acc, or Cha;

A_{19} is Arg or Aib;

A_{22} is Phe, Glu, Aib, Acc, or Cha;

A_{23} is Phe, Leu, Lys, Acc, or Cha;

A_{24} is Leu, Lys, Acc, or Cha;

A_{25} is His, Lys, Aib, Acc, or Glu;

A_{26} is His, Aib, Acc, or Lys;

A_{27} is Leu, Lys, Acc, or Cha;

A_{28} is Ile, Leu, Lys, Acc, or Cha;

A_{29} is Ala, Glu, Acc, or Aib;

A_{30} is Glu, Leu, Nle, Cha, Aib, Acc, or Lys;

A_{31} is Ile, Leu, Cha, Lys, Acc, or deleted;

A_{32} is His or deleted;

A_{33} is Thr or deleted;

A_{34} is Ala or deleted;

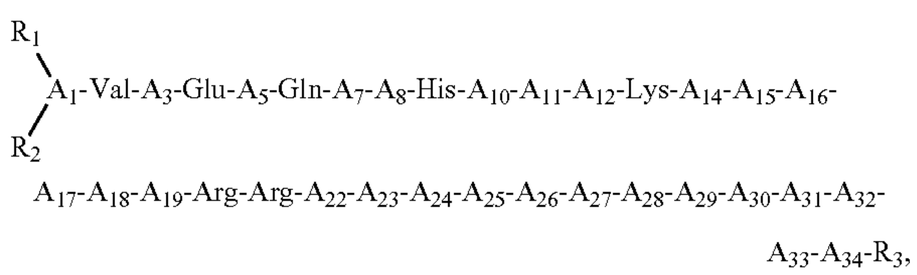
each of R_1 and R_2 is, independently, H, C_{1-12} alkyl, C_{7-20} phenylalkyl, C_{11-20} naphthylalkyl, C_{1-12} hydroxyalkyl, C_{2-12} hydroxyalkenyl, C_{7-20} hydroxyphenylalkyl, or C_{11-20} hydroxynaphthylalkyl; or one and only one of R_1 and R_2 is COE_1 in which E_1 is C_{1-12} alkyl, [C_{2-12} alkyl] C_{2-12} alkenyl, C_{7-20} phenylalkyl, C_{11-20} naphthylalkyl, C_{1-12} hydroxyalkyl, C_{2-12} hydroxyalkenyl, C_{7-20} hydroxyphenylalkyl, or C_{11-20} hydroxynaphthylalkyl; and

R_3 is OH, NH_2, C_{1-12} alkoxy, or NH—Y—CH_2-Z in which Y is a C_{1-12} hydrocarbon moiety and Z is H, OH, CO_2H or $CONH_2$;

provided that at least one of A_5, A_7, A_8, A_{11}, A_{12}, A_{15}, A_{18}, A_{22}, A_{23}, A_{24}, A_{25}, A_{26}, A_{27}, A_{28}, A_{29}, A_{30}, or A_{31} is Acc; or a pharmaceutically acceptable salt thereof.

51. A method of treating osteoporosis in a patient in need thereof, which comprises administering to said patient a combination of a bisphosphonate or calcitonin and a peptide of the formula (VI):

[SEQ ID NO: 5]



wherein

A_1 is Ala, Ser, or Dap;

A_3 is Ser or Aib;

[As] A_5 is His, Ile, Acc, or Cha;

A_7 is Leu, Cha, Nle, β-Nal, Trp, Pal, Acc, Phe, or p-X-Phe in which X is OH, a halogen, or CH_3;

A_8 is Leu, Met, Acc, or Cha;

A_{10} is Asp or Asn;

A_{11} is Lys, Leu, Cha, Acc, Phe, or β-Nal;

A_{12} is Gly, Acc, or Aib;

A_{14} is Ser or His;

A_{15} is Ile, Acc, or Cha;

A_{16} is Gln or Aib;

A_{17} is Asp or Aib;

A_{18} is Leu, Aib, Acc, or Cha;

A_{19} is Arg or Aib;

A_{22} is Phe, Glu, Aib, Acc, or Cha;

A_{23} is Phe, Leu, Lys, Acc, or Cha;

A_{24} is Leu, Lys, Acc, or Cha;

A_{25} is His, Lys, Aib, Acc, or Glu;

A_{26} is His, Aib, Acc, or Lys;

A_{27} is Leu, Lys, Acc, or Cha;

A_{28} is Ile, Leu, Lys, Acc, or Cha;

A_{29} is Ala, Glu, Acc, or Aib;

A_{30} is Glu, Leu, Nle, Cha, Aib, Acc, or Lys;

A_{31} is Ile, Leu, Cha, Lys, Acc, or deleted;

A_{32} is His or deleted;

A_{33} is Thr or deleted;

A_{34} is Ala or deleted;

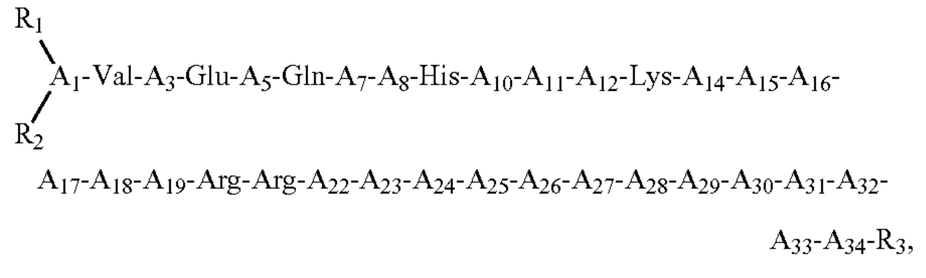
each of R_1 and R_2 is, independently, H, C_{1-12} alkyl, C_{7-20} phenylalkyl, C_{11-20} naphthylalkyl, C_{1-12} hydroxyalkyl, C_{2-12} hydroxyalkenyl, C_{7-20} hydroxyphenylalkyl, or C_{11-20} hydroxynaphthylalkyl; or one and only one of R_1 and R_2 is COE_1 in which E_1 is C_{1-12} alkyl, [C_{2-12} alkyl] C_{2-12} alkenyl, C_{7-20} phenylalkyl, C_{11-20} naphthylalkyl, C_{1-12} hydroxyalkyl, C_{2-12} hydroxyalkenyl, C_{7-20} hydroxyphenylalkyl, or C_{11-20} hydroxynaphthylalkyl; and

R_3 is OH, NH_2, C_{1-12} alkoxy, or NH—Y—CH_2-Z in which Y is a C_{1-12} hydrocarbon moiety and Z is H, OH, CO_2H or $CONH_2$;

provided that at least one of A_5, A_7, A_8, A_{11}, A_{12}, A_{15}, A_{18}, A_{22}, A_{23}, A_{24}, A_{25}, A_{26}, A_{27}, A_{28}, A_{29}, A_{30}, or A_{31} is Acc; or a pharmaceutically acceptable salt thereof.

52. A pharmaceutical composition comprising a bisphosphonate or calcitonin, a pharmaceutically acceptable carrier or diluent, and a peptide of the following formula (VI):

[SEQ ID NO: 5]



$$\begin{array}{l} R_1 \\ \quad \backslash \\ \quad A_1\text{-Val-}A_3\text{-Glu-}A_5\text{-Gln-}A_7\text{-}A_8\text{-His-}A_{10}\text{-}A_{11}\text{-}A_{12}\text{-Lys-}A_{14}\text{-}A_{15}\text{-}A_{16}\text{-} \\ \quad / \\ R_2 \end{array}$$

$$A_{17}\text{-}A_{18}\text{-}A_{19}\text{-Arg-Arg-}A_{22}\text{-}A_{23}\text{-}A_{24}\text{-}A_{25}\text{-}A_{26}\text{-}A_{27}\text{-}A_{28}\text{-}A_{29}\text{-}A_{30}\text{-}A_{31}\text{-}A_{32}\text{-}A_{33}\text{-}A_{34}\text{-}R_3,$$

wherein

A_1 is Ala, Ser, or Dap;

A_3 is Ser or Aib;

A_5 is His, Ile, Acc, or Cha;

A_7 is Leu, Cha, Nle, β-Nal, Trp, Pal, Acc, Phe, or p-X-Phe in which X is OH, a halogen, or CH_3;

A_8 is Leu, Met, Acc, or Cha;

A_{10} is Asp or Asn;

A_{11} is Lys, Leu, Cha, Acc, Phe, or β-Nal;

A_{12} is Gly, Acc, or Aib;

A_{14} is Ser or His;

A_{15} is Ile, Acc, or Cha;

A_{16} is Gln or Aib;

A_{17} is Asp or Aib;

A_{18} is Leu, Aib, Acc, or Cha;

A_{19} is Arg or Aib;

A_{22} is Phe, Glu, Aib, Acc, or Cha;

A_{23} is Phe, Leu, Lys, Acc, or Cha;

A_{24} is Leu, Lys, Acc, or Cha;

A_{25} is His, Lys, Aib, Acc, or Glu;

A_{26} is His, Aib, Acc, or Lys;

A_{27} is Leu, Lys, Acc, or Cha;

A_{28} is Ile, Leu, Lys, Acc, or Cha;

A_{29} is Ala, Glu, Acc, or Aib;

A_{30} is Glu, Leu, Nle, Cha, Aib, Acc, or Lys;

A_{31} is Ile, Leu, Cha, Lys, Acc, or deleted;

A_{32} is His or deleted;

A_{33} is Thr or deleted;

A_{34} is Ala or deleted;

each of R_1 and R_2 is, independently, H, $C_{1\text{-}12}$ alkyl, $C_{7\text{-}20}$ phenylalkyl, $C_{11\text{-}20}$ naphthylalkyl, $C_{1\text{-}12}$ hydroxyalkyl, $C_{2\text{-}12}$ hydroxyalkenyl, $C_{7\text{-}20}$ hydroxyphenylalkyl, or $C_{11\text{-}20}$ hydroxynaphthylalkyl; or one and only one of R_1 and R_2 is COE_1 in which E_1 is $C_{1\text{-}12}$ alkyl, **[**$C_{2\text{-}12}$ alkyl,**]** $C_{2\text{-}12}$ alkenyl, $C_{7\text{-}20}$ phenylalkyl, $C_{11\text{-}20}$ naphthylalkyl, $C_{1\text{-}12}$ hydroxyalkyl, $C_{2\text{-}12}$ hydroxyalkenyl, $C_{7\text{-}20}$ hydroxyphenylalkyl, or $C_{11\text{-}20}$ hydroxynaphthylalkyl; and

R_3 is OH, NH_2, $C_{1\text{-}12}$ alkoxy, or NH—Y—CH_2-Z in which Y is a $C_{1\text{-}12}$ hydrocarbon moiety and Z is H, OH, CO_2H or $CONH_2$;

provided that at least one of A_5, A_7, A_8, A_{11}, A_{12}, A_{15}, A_{18}, A_{22}, A_{23}, A_{24}, A_{25}, A_{26}, A_{27}, A_{28}, A_{29}, A_{30}, or A_{31} is Acc; or a pharmaceutically acceptable salt thereof.

* * * * *

UNITED STATES PATENT AND TRADEMARK OFFICE

CERTIFICATE OF CORRECTION

PATENT NO. : RE 40,850 E
APPLICATION NO. : 11/523812
DATED : July 14, 2009
INVENTOR(S) : Zheng Xin Dong

It is certified that error appears in the above-identified patent and that said Letters Patent is hereby corrected as shown below:

Claim 22, column 45, lines 51-53, that portion of the claim which reads

hydroxyalkenyl, C_{7-20} hydroxy-phenylalkyl, or C_{1-120} hydroxynaphthylalkyl; and
R_3 is OH, NH_2, C_{1-12} alkoxy, or NH'Y-CH_2-Z in which Y is a C_{1-12} hydrocarbon

should read

hydroxyalkenyl, C_{7-20} hydroxy-phenylalkyl, or [C_{1-120} hydroxynaphthylalkyl] C_{11-20} hydroxynaphthylalkyl; and
R_3 is OH, NH_2, C_{1-12} alkoxy, or [NH'Y-CH_2-Z] NH-Y-CH_2-Z in which Y is a C_{1-12} hydrocarbon

Signed and Sealed this

Twentieth Day of April, 2010

David J. Kappos
Director of the United States Patent and Trademark Office