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CA 2235443 C 2011/11/01

(11)(21) 2 235 443

(12) BREVET CANADIEN CANADIAN PATENT

(13) **C** 

(22) Date de dépôt/Filing Date: 1998/06/18

(41) Mise à la disp. pub./Open to Public Insp.: 1998/12/20

(45) Date de délivrance/Issue Date: 2011/11/01 (30) Priorité/Priority: 1997/06/20 (DE19726167.1)

(51) Cl.Int./Int.Cl. *C12N 15/17* (2006.01), *A61K 38/28* (2006.01), *C07K 14/62* (2006.01), *C12P 21/02* (2006.01), *A61K 38/00* (2006.01)

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(54) Titre: NOUVEAUX DERIVES D'INSULINE A ACTION RAPIDE

(54) Title: NOVEL INSULIN DERIVATIVES HAVING A RAPID ONSET OF ACTION

### (57) Abrégé/Abstract:

The present invention relates to insulin derivatives which in comparison to human insulin, have an accelerated onset of action, to a process for their preparation and to their use, in particular in pharmaceutical preparations for the treatment of diabetes mellitus. In particular, the present invention relates to insulin derivatives or physiologically tolerable salts thereof in which asparagine (Asn) in position B3 of the B chain is replaced by a naturally occurring basic amino acid residue and at least one amino acid residue in the positions B27, B28 or B29 of the B chain is replaced by another naturally occurring amino acid residue, it optionally being possible for asparagine (Asn) in position 21 of the A chain to be replaced by Asp, Gly, Ser, Thr or Ala and for phenylalanine (Phe) in position B1 of the B chain and the amino acid residue in position B30 of the B chain to be absent.





# Abstract

Novel insulin derivatives having a rapid onset of action

The present invention relates to insulin derivatives which in comparison to human insulin, have an accelerated onset of action, to a process for their preparation and to their use, in particular in pharmaceutical preparations for the treatment of diabetes mellitus.

In particular, the present invention relates to insulin derivatives or physiologically tolerable salts thereof in which asparagine (Asn) in position B3 of the B chain is replaced by a naturally occurring basic amino acid residue and at least one amino acid residue in the positions B27, B28 or B29 of the B chain is replaced by another naturally occurring amino acid residue, it optionally being possible for asparagine (Asn) in position 21 of the A chain to be replaced by Asp, Gly, Ser, Thr or Ala and for phenylalanine (Phe) in position B1 of the B chain and the amino acid residue in position B30 of the B chain to be absent.

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Description

Novel insulin derivatives having a rapid onset of action

The present invention relates to insulin derivatives which, in comparison to human insulin, have an accelerated onset of action, to a process for their preparation and to their use, in particular in pharmaceutical preparations for the treatment of diabetes mellitus.

Approximately 120 million people worldwide suffer from diabetes mellitus. Among these are approximately 12 million type I diabetics, for whom the administration of insulin is the only therapy possible at present. The affected people are assigned insulin injections, as a rule several times daily, for life. Although type II diabetes, from which approximately 100 million people suffer, is not fundamentally accompanied by an insulin deficiency, in a large number of cases, however, treatment with insulin is regarded as the most favorable or only possible form of therapy.

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With increasing length of the disease, a large number of the patients suffer from so-called diabetic late complications. These are essentially microand macrovascular damage, which depending on the type and extent, result in kidney failure, blindness, loss of extremities or an increased risk of heart/circulation disorders.

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As a cause, chronically increased blood glucose levels are primarily held responsible, since even with careful adjustment of the insulin therapy a normal blood glucose profile, such as would correspond to physiological regulation, is not achieved (Ward, J. D. (1989) British Medical Bulletin 45, 111-126; Drury, P.L. et al. (1989) British Medical Bulletin 45, 127-147; Kohner, E.M. (1989) British Medical Bulletin 45, 148-173)

In healthy people, insulin secretion is closely dependent on the glucose concentration of the blood. Increased glucose levels, such as occur after meals, are rapidly compensated by an increased release of insulin. In the fasting state, the plasma insulin level falls to a basal value, which is sufficient to guarantee a continuous supply of insulin-sensitive organs and tissue with glucose. An optimization of the therapy, the so-called intensified insulin therapy, is today primarily aimed at keeping variations in the blood glucose concentration, especially deviations upward, as low as possible (Bolli, G. B. (1989) Diabetes Res. Clin. Pract. 6, p. 3 - p. 16; Berger, M. (1989) Diabetes Res. Clin. Pract. 6, p. 25 - p. 32). This leads to a significant decrease in the occurrence and the progression of diabetic late damage (The Diabetes Control and Complications Trial Research Group (1993) N. Engl. J. Med. 329, 977-986).

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From the physiology of insulin secretion, it can be deduced that for an improved, intensified insulin therapy using subcutaneously administered preparations, two insulin preparations having different pharmacodynamics are needed. To compensate the blood glucose rise after meals, the insulin must flow in rapidly and must only act for a few hours. For the basal supply, in particular in the night, a preparation should be available which acts for a long time, has no pronounced maximum and only infuses very slowly.

The preparations based on human and animal insulins only fulfill the demands of an intensified insulin therapy, however, in a restricted manner. After subcutaneous administration, rapidly acting insulins (unmodified insulins) pass too slowly into the blood and to the site of action and have too long an overall duration of action. The result is that the postprandial glucose levels are too high and the blood glucose begins to fall severely several hours after the meal (Kang, S. et al. (1991) Diabetes Care 14, 142-148; Home, P.J. et al. (1989) British Medical Bulletin 45, 92-110; Bolli, G. B. (1989) Diabetes Res. Clin. Pract. 6, p. 3 - p. 16). The available basal

insulins in turn, especially NPH insulins, have too short a duration of action and have a too severely pronounced maximum.

Beside the possibility of affecting the profile of action by means of pharmaceutical principles, the alternative presents itself today of designing insulin derivatives, with the aid of genetic engineering, which achieve specific properties such as onset and duration of action solely by means of their structural properties. By the use of suitable insulin derivatives, a significantly better adjustment of the blood glucose more closely adapted to the natural conditions could therefore be achieved.

Insulin derivatives having an accelerated onset of action are described in EP 0 214 826, EP 0 375 437 and EP 0 678 522. EP 0 214 826 relates, inter alia, to substitutions of B27 and B28, but not in combination with the substitution of B3. EP 0 678 522 describes insulin derivatives which have various amino acids, preferably proline, but not glutamic acid, in the position B29. EP 0 375 437 encompasses insulin derivatives having lysine or arginine in B28, which can optionally be additionally modified in B3 and/or A21.

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EP 0 419 504 discloses insulin derivatives which are protected against chemical modifications by changing asparagine in B3 and at least one further amino acid in the positions A5, A15, A18 or A21. Combinations with modifications in positions B27, B28 or B29 are, however, not described. An indication that these compounds have modified pharmacodynamics resulting in a more rapid onset of action is not given.

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WO 92/00321 describes insulin derivatives in which at least one amino acid of the positions B1-B6 is replaced by lysine or arginine. According to WO 92/00321, insulins of this type have a prolonged action. Combinations with modifications of the positions B27, 28, 29, however, are not disclosed.

The object of the present invention is to prepare insulin derivatives which after administration, in particular after subcutaneous administration, have an onset of action which is accelerated in comparison with human insulin.

Insulin derivatives are derivatives of naturally occurring insulins, namely human insulin (see SEQ ID NO 1 = A chain of human insulin; see SEQ ID NO 2 = B chain of human insulin, sequence listing) or animal insulins which differ from the corresponding, otherwise identical naturally occurring insulin by substitution of at least one naturally occurring amino acid residue and/or addition of at least one amino acid residue and/or organic residue.

It is further an object of the present invention to provide a process for the preparation of the insulin derivatives having the property mentioned, the corresponding intermediates and their precursors.

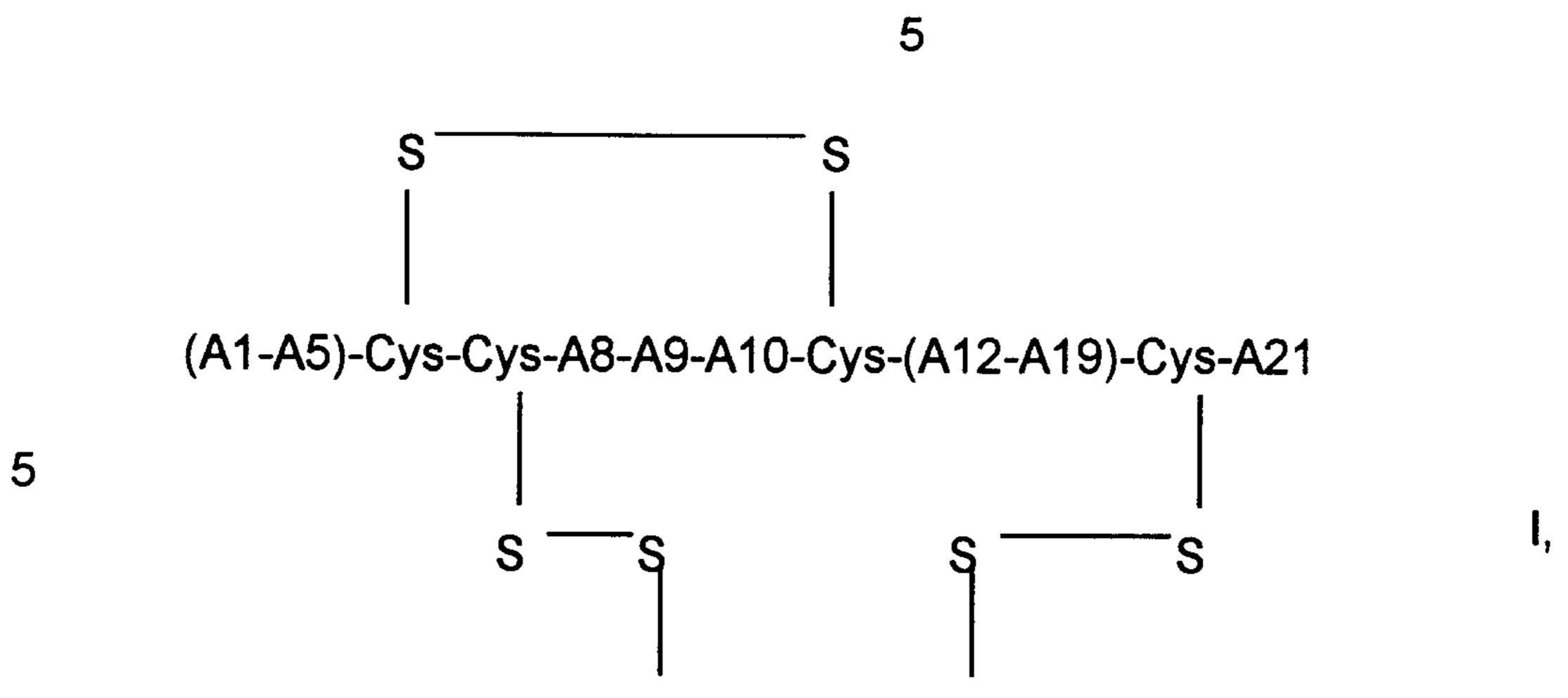
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The object is achieved by an insulin derivative or a physiologically tolerable salt thereof in which asparagine (Asn) in position B3 of the B chain is replaced by a naturally occurring basic amino acid residue and at least one amino acid residue in the positions B27, B28 or B29 of the B chain is replaced by another naturally occurring amino acid residue, it optionally being possible for asparagine (Asn) in position 21 of the A chain to be replaced by Asp, Gly, Ser, Thr or Ala and for phenylalanine (Phe) in position B1 of the B chain and the amino acid residue in position B30 of the B chain to be absent.

Preferably, the insulin derivative or its physiologically tolerable salt is of formula I



B1-Val-B3-Glu-His-Leu-Cys-(B8-B18)-Cys-(B20-B26)-B27-B28-B29-B30

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- (A1-A5) are the amino acid residues in the positions A1 to A5 of the A chain of human insulin (cf. SEQ ID NO 1) or animal insulin,
- (A12-A19) are the amino acid residues in the positions A12 to A19 of the A chain of human insulin (cf. SEQ ID NO 1) or animal insulin,
  - (B8-B18) are the amino acid residues in the positions B8 to B18 of the B chain of human insulin (cf. SEQ ID NO 2) or animal insulin,
  - (B20-B26) are the amino acid residues in the positions B20 to B26 of the B chain of human insulin (cf. SEQ ID NO 2) or animal insulin,
- A8, A9, A10 are the amino acid residues in the positions A8, A9 and A10 of the A chain of human insulin (cf. SEQ ID NO 1) or animal insulin,
  - A21 is Asn, Asp, Gly, Ser, Thr or Ala,
  - is -OH or the amino acid residue in position B30 of the B chain of human insulin (cf. SEQ ID NO 2) or animal insulin,

B1 is a phenylalanine residue (Phe) or a hydrogen atom, B3 is a naturally occurring basic amino acid residue,

B27, B28

5 and B29

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are the amino acid residues in the positions B27, B28 and B29 of the B chain of human insulin (cf. SEQ ID NO 2) or animal insulin or in each case are another naturally occurring amino acid residue, where at least one of the amino acid residues in the positions B27, B28 and B29 of the B chain is replaced by another naturally occurring amino acid residue.

Of the twenty naturally occurring amino acids which are genetically encodable, the amino acids glycine (Gly), alanine (Ala), valine (Val), leucine (Leu), isoleucine (Ile), serine (Ser), threonine (Thr), cysteine (Cys), methionine (Met), asparagine (Asn), glutamine (Gln), phenylalanine (Phe), tyrosine (Tyr), tryptophan (Trp) and proline (Pro) are designated here as neutral amino acids, the amino acids arginine (Arg), lysine (Lys) and histidine (His) are designated as basic amino acids and the amino acids aspartic acid (Asp) and glutamic acid (Glu) are designated as acidic amino acids.

Preferably, the insulin derivative or its physiologically tolerable salt according to the present invention is a derivative of bovine insulin, porcine insulin or human insulin, namely an insulin derivative or a physiologically tolerable salt thereof of the formula I, which is distinguished in that

A8 is alanine (Ala),

A9 is serine (Ser),

A10 is valine (Val) and

30 B30 is alanine (Ala) (amino acid residues A8 to A10 and B30 of bovine insulin),

A8 is threonine (Thr),

A9 is serine (Ser) and

- A10 is isoleucine (Ile) (amino acid residues A8 to A10 of the insulins of man or pigs), where
- B30 is alanine (Ala) (amino acid residue B30 of porcine insulin) or
- 5 B30 is threonine (Thr) (amino acid residue B30 of human insulin, cf. SEQ ID NO 2).

Particularly preferably, an insulin derivative or a physiologically tolerable salt thereof of the formula I with the amino acid residues A8 to A10 and B30 of human insulin is one which is furthermore distinguished in that

- (A1-A5) are the amino acid residues in the positions A1 to A5 of the A chain of human insulin (cf. SEQ ID NO 1),
- (A12-A19) are the amino acid residues in the positions A12 to A19 of the A chain of human insulin (cf. SEQ ID NO 1),
- (B8-B18) are the amino acid residues in the positions B8 to B18 of the B chain of human insulin (cf. SEQ ID NO 2) and
- (B20-B26) are the amino acid residues in the positions B20 to B26 of the B chain of human insulin (cf. SEQ ID NO 2).

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Further preferred embodiments of the present invention are an insulin derivative or a physiologically tolerable salt thereof of the formula I, wherein the amino acid residue in position B1 of the B chain is a phenylalanine residue (Phe) or

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an insulin derivative or a physiologically tolerable salt thereof of the formula I, wherein the amino acid residue in position B3 of the B chain is a histidine (His), lysine (Lys) or arginine residue (Arg).

Further preferred embodiments of the present invention are an insulin derivative or a physiologically tolerable salt thereof of the formula I, wherein at least one of the amino acid residues in the positions B27, B28 and B29 of the B chain is replaced by a naturally occurring amino acid

residue which is selected from the group consisting of the neutral or of the acidic amino acids,

an insulin derivative or a physiologically tolerable salt thereof of the formula I, wherein at least one of the amino acid residues in the positions B27, B28 and B29 of the B chain is a naturally occurring amino acid residue which is selected from the group consisting of isoleucine (IIe), aspartic acid (Asp) and glutamic acid (Glu), preferably wherein at least one of the amino acid residues in the positions B27, B28 of the B chain is replaced by a naturally occuring amino acid residue which is selected from the group consisting of the neutral amino acids, or particularly preferably wherein at least one of the amino acid residues in the positions B27, B28 and B29 of the B chain is an isoleucine residue (IIe), or

an insulin derivative or a physiologically tolerable salt thereof of the formula I, wherein at least one of the amino acid residues in the positions B27, B28 and B29 of the B chain is a naturally occurring amino acid residue which is selected from the group consisting of the acidic amino acids, preferably wherein at least one of the amino acid residues in the positions B27, B28 and B29 of the B chain is an aspartic acid residue (Asp), preferably wherein the amino acid residue in position B27 or B28 of the B chain is an aspartic acid residue (Asp), or wherein at least one of the amino acid residues in the positions B27, B28 and B29 of the B chain is a glutamic acid residue (Glu).

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A preferred embodiment of the present invention is also an insulin derivative or a physiologically tolerable salt thereof of the formula I, wherein the amino acid residue in position B29 of the B chain is an aspartic acid residue (Asp).

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Further preferred embodiments are an insulin derivative or a physiologically tolerable salt thereof of the formula I, wherein the amino acid residue in position B27 of the B chain is a glutamic acid residue (Glu),

an insulin derivative or a physiologically tolerable salt thereof of the formula I, wherein the amino acid residue in position B28 of the B chain is a glutamic acid residue (Glu), or

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an insulin derivative or a physiologically tolerable salt thereof of the formula I, wherein the amino acid residue in position B29 of the B chain is a glutamic acid residue (Glu).

Very particularly preferably, an insulin derivative or a physiologically tolerable salt thereof is one which is distinguished in that the B chain has the sequence

Phe Val Lys Gln His Leu Cys Gly Ser His Leu Val Glu Ala Leu Tyr Leu Val Cys Gly Glu Arg Gly Phe Phe Tyr Thr Pro Glu Thr

(SEQ ID NO 3), for example Lys (B3), Glu (B29)-human insulin, or

an insulin derivative or a physiologically tolerable salt thereof which is distinguished in that the amino acid residue in position B27 of the B chain is an isoleucine residue (IIe), preferably an insulin derivative or a physiologically tolerable salt thereof which is distinguished in that the B chain has the sequence

25 Phe Val Lys Gln His Leu Cys Gly Ser His Leu Val Glu Ala Leu
Tyr Leu Val Cys Gly Glu Arg Gly Phe Phe Tyr Ile Pro Lys Thr

(SEQ ID NO 5), for example Lys (B3), Ile (B27)-human insulin, or an insulin derivative or a physiologically tolerable salt thereof of the formula I, wherein the amino acid residue in position B28 of the B chain is an isoleucine residue (Ile), preferably an insulin derivative or a physiologically tolerable salt thereof which is distinguished in that the B chain has the sequence

Phe Val Lys Gln His Leu Cys Gly Ser His Leu Val Glu Ala Leu

5 Tyr Leu Val Cys Gly Glu Arg Gly Phe Phe Tyr Thr lle Lys Thr

(SEQ ID NO 4), for example Lys (B3), Ile (B28)-human insulin.

Particularly preferably, an insulin derivative or a physiologically tolerable salt thereof of the formula I, which is distinguished in that the amino acid residue in position B28 of the B chain is an isoleucine residue (IIe) and the amino acid residue in position A21 is an asparagine residue (Asp), is preferably one wherein the A chain has the sequence

Gly Ile Val Glu Gln Cys Cys Thr Ser Ile Cys Ser Leu Tyr Gln Leu Tyr Gln Leu Glu Asn Tyr Cys Asp (SEQ ID NO.: 9)

and the B chain has the sequence

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20 Phe Val Lys Gln His Leu Cys Gly Ser His Leu Val Glu Ala Leu Tyr Leu Val Cys Gly Glu Arg Gly Phe Phe Pyr Thr Ile Lys Thr (SEQ ID NO.: 10)

(Lys (B3), Ile (B28), Asp (A21)-human insulin).

The insulin derivatives of the formula I can preferably be prepared by genetic engineering.

The object set at the outset is accordingly further achieved by a process for the preparation of an insulin derivative or of a physiologically tolerable salt thereof of the formula I, comprising the construction of a replicable expression vehicle which contains a DNA sequence which codes for a precursor of the insulin derivative in which the amino acid residue in

position A1 of the A chain is linked to the amino acid residue B30 of the B chain via a peptide chain of the formula II

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in which  $R^1_n$  is a peptide chain having n amino acid residues and n is an integer from 0 to 34, and the B chain is prolonged in position B1 by a peptide chain of the formula III

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in which  $R^2_m$  is a peptide chain having m amino acid residues, m is an integer from 0 to 40, preferably from 0 to 9, and p is 0, 1 or 2, where for p = 0 the peptide chain  $R^2_m$  preferably ends with Lys, expression in a host cell and release of the insulin derivative from its precursor using chemical and/or enzymatic methods.

Preferably, the process is one wherein the host cell is a bacterium, particularly preferably one wherein the bacterium is E. coli.

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Preferably, the process is one wherein the host cell is a yeast, particularly preferably one wherein the yeast is Saccharomyces cerevisiae.

For the preparation of an insulin derivative having the amino acid sequences SEQ ID NO.: 9 (A chain) and SEQ ID NO.: 10 (B chain), the precursor of this insulin derivative preferably has the sequence

Met Ala Thr Thr Ser Thr Gly Asn Ser Ala Arg Phe Val Lys Gln His Leu Cys Gly Ser His Leu Val Glu Ala Leu Tyr Leu Val Cys Gly Glu Arg Gly Phe Phe Tyr Thr Ile Lys Thr Arg Arg Glu Ala Glu Asp Pro Gln Val Gly Gln Val Glu Leu Gly Gly Gly Pro Gly Ala Gly Ser Leu Gln Pro Leu Ala Leu Glu Gly Ser

Leu Gln Lys Arg Gly Ile Val Glu Gln Cys Cys Thr Ser Ile Cys Ser Leu Tyr Gln Ley Glu Asn Tyr Cys Asp (SEQ ID NO.: 11),

a Lys (B3), Ile (B28), Asp (A21)-preproinsulin.

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For the preparation of an insulin derivative having the amino acid sequence SEQ ID NO 3, the precursor of this insulin derivative preferably has the sequence

10 Met Ala Thr Thr Ser Thr Gly Asn Ser Ala Arg

Phe Val Lys Gln His Leu Cys Gly Ser His Leu Val Glu Ala Leu
Tyr Leu Val Cys Gly Glu Arg Gly Phe Phe Tyr Thr Pro Glu Thr

Arg Arg Glu Ala Glu Asp Pro Gln Val Gly Gln Val Glu Leu Gly

Gly Gly Pro Gly Ala Gly Ser Leu Gln Pro Leu Ala Leu Glu Gly

15 Ser Leu Gln Lys Arg

Gly lle Val Glu Gln Cys Cys Thr Ser lle Cys Ser Leu Tyr Gln Leu Glu Asn Tyr Cys Asn (Lys (B3), Glu (B29)-preproinsulin)

20 (SEQ ID NO 6).

For the preparation of an insulin derivative having the amino acid sequence SEQ ID NO 5, the precursor of this insulin derivative preferably has the sequence

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Met Ala Thr Thr Ser Thr Gly Asn Ser Ala Arg

Phe Val Lys Gln His Leu Cys Gly Ser His Leu Val Glu Ala Leu

Tyr Leu Val Cys Gly Glu Arg Gly Phe Phe Tyr Ile Pro Lys Thr

Arg Arg Glu Ala Glu Asp Pro Gln Val Gly Gln Val Glu Leu Gly

Gly Gly Pro Gly Ala Gly Ser Leu Gln Pro Leu Ala Leu Glu Gly Ser Leu Gln Lys Arg

Gly lie Val Glu Gln Cys Cys Thr Ser lie Cys Ser Leu Tyr Gln

Leu Glu Asn Tyr Cys Asn

(Lys (B3), Ile (B27)-preproinsulin)

(SEQ ID NO 8).

For the preparation of an insulin derivative having the amino acid sequence SEQ ID NO 4, the precursor of this insulin derivative preferably 10 has the sequence

Met Ala Thr Thr Ser Thr Gly Asn Ser Ala Arg

Phe Val Lys Gln His Leu Cys Gly Ser His Leu Val Glu Ala Leu

Tyr Leu Val Cys Gly Glu Arg Gly Phe Phe Tyr Thr lle Lys Thr 15

Arg Arg Glu Ala Glu Asp Pro Gln Val Gly Gln Val Glu Leu Gly

Gly Gly Pro Gly Ala Gly Ser Leu Gln Pro Leu Ala Leu Glu Gly

Ser Leu Gln Lys Arg

Gly lle Val Glu Gln Cys Cys Thr Ser lle Cys Ser Leu Tyr Gln

Leu Glu Asn Tyr Cys Asn 20

(Lys (B3), Ile (B28)-preproinsulin)

(SEQ ID NO 7).

25 The present invention accordingly also relates to said precursors of the preferred insulin derivatives, namely the peptides having the sequence numbers SEQ ID NO.: 11, SEQ ID NO 6, SEQ ID NO 7 and SEQ ID NO 8, the DNA sequences which code for said precursors, the expression vehicles which comprise these DNA sequences and the host cells which 30

are transformed using these expression vehicles.

The insulin derivatives of the formula I are mainly prepared by genetic engineering by means of site-directed mutagenesis according to standard methods.

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To do this, a gene structure coding for the desired insulin derivative of the formula I is constructed and expressed in a host cell - preferably in a bacterium such as E coli or a yeast, in particular Saccharomyces cerevisiae - and - if the gene structure codes for a fusion protein - the insulin derivative of the formula I is released from the fusion protein; analogous methods are described, for example, in EP-A-0 211 299, EP-A-0 227 938, EP-A-0 229 998, EP-A-0 286 956 and the DE patent application P 38 21 159.

The removal of the fusion protein component can be carried out chemically by cell disruption by means of cyanogen halide (see EP-A-0 180 920).

In the preparation by means of a preproinsulin precursor which has a fusion protein component (presequence) according to US 5,358,857, the removal of the fusion protein component takes place in a later stage together with the removal of the C peptide.

The insulin precursor is then subjected to oxidative sulfitolysis according to the method described, for example, by R.C. Marshall and A.S. Inglis in "Practical Protein Chemistry - A Handbook" (Publisher A. Darbre) 1986, pages 49 - 53 and then renatured in the presence of a thiol with formation of the correct disulfide bridges, e.g. according to the method described by G.H. Dixon and A.C. Wardlow in Nature (1960), pages 721 - 724.

The insulin precursors, however, can also be folded directly (EP-A-0 600 372; EP-A-0 668 292).

The C peptide is removed by means of tryptic cleavage - e.g. according to the method of Kemmler et al., J.B.C. (1971), pages 6786 - 6791 and the

insulin derivative of the formula I is purified by means of known techniques such as chromatography - e.g. EP-A-0 305 760 - and crystallization.

If n in formula II is 0, tryptic cleavage serves to sever the peptide bond between A and B chains.

In this process, the B chain C terminal ends with arginine or two arginine residues. These can be removed enzymatically by means of carboxy-peptidase B.

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The insulin derivatives according to the invention have full biological activity. This was shown by intravenous administration to rabbits and the blood glucose fall resulting therefrom (Examples 5 and 6).

The more rapid onset of action after subcutaneous administration was 15 shown in fasting dogs using the euglycemic clamp technique (Example 7). 0.3 IU/kg was administered. The reference preparation was human insulin. In the clamp technique, the blood glucose value is measured at short time intervals after insulin injection and exactly the amount of glucose to compensate the fall is infused. This has the advantage that no counter-20 regulation occurs with the animals, as would be the case with a severe fall in the blood glucose after the administration of insulin. The amount and the time course of the infused glucose characterize the action of the insulin. Lys(B3), Glu(B29)- (SEQ ID NO 3) and Lys(B3), Ile(B28)- (SEQ ID NO 4) insulin have a clearly more rapid onset of action than human insulin. The 25 maximum action (glucose infusion rate) is achieved after 100 minutes with human insulin, after 80 minutes, however, with Lys(B3), Glu(B29)-insulin (SEQ ID NO 3) and already after 60 minutes with Lys(B3)-, Ile(B28)-insulin (SEQ ID NO 4). Therefore these analogs, when they are injected shortly before a meal, should compensate the postprandial rise in the blood 30 glucose better than human insulin.

The insulin derivatives described are therefore suitable both for the therapy of type I and of type II diabetes mellitus, preferably in combination with a basal insulin.

- The present invention therefore also relates to the use of the insulin derivative and/or its physiologically tolerable salt of the formula I for the production of a pharmaceutical preparation which has an insulin activity with a rapid onset of action.
- A suitable carrier medium which is physiologically acceptable and compatible with the insulin derivative is a sterile aqueous solution which is made isotonic with blood in the customary manner, e.g. by means of glycerol, sodium chloride, glucose, and in addition contains one of the customary preservatives, e.g. phenol, m-cresol or p-hydroxybenzoic acid ester. The carrier medium can additionally contain a buffer substance, e.g. sodium acetate, sodium citrate, sodium phosphate. For adjustment of the pH, dilute acids (typically HCI) or alkalis (typically NaOH) are used. The preparation can furthermore contain zinc ions.
- The insulin derivatives can be employed in the pharmaceutical preparations even in the form of their physiologically tolerable salts, as alkali metal or as ammonium salts. Any desired proportion of one or more insulin derivatives of the formula I or an insulin derivative of the formula I can be present in a mixture of other of these insulin derivatives independently of one another in each case in dissolved, amorphous and/or crystalline form.

It is sometimes advantageous to add to the preparation according to the invention a suitable amount of a suitable stabilizer which prevents the precipitation of protein under thermomechanical stress on contact with various materials. Such stabilizers are disclosed, for example, in EP-A-18609, DE-A 32 40 177 or in WO-83/00288.

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The present invention further relates to a pharmaceutical preparation which comprises at least one insulin derivative and/or a physiologically tolerable salt thereof of the formula I, preferably in dissolved, amorphous and/or crystalline form.

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The insulin derivatives according to the invention have a rapid onset of action. In practical insulin therapy, it is customary under certain circumstances to mix rapid-acting insulins with preparations which contain a depot auxiliary (e.g. NPH insulin). Depending on the composition, preparations result from this whose profiles of action correspond to the superimposed individual profiles provided the individual components in the mixture are stable and are not mutually affected. When mixing an insulin derivative with human NPH insulin, however, it is to be expected that, particularly on long-term storage, an exchange takes place between the dissolved derivative and the crystalline NPH insulin. As a result of this both the pharmacodynamics of the depot insulin and those of the dissolved rapidly acting insulin are changed in an unforeseeable manner. In order to avoid this, it is sensible to prepare the rapidly acting derivative using a depot auxiliary - for example as NPH insulin. This depot form of the insulin derivative can then be mixed as desired with the dissolved rapidly acting form without the composition of one or the other form changing in the course of storage due to exchange.

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Although the invention in essence relates to rapidly acting insulin derivatives, it accordingly, however, also comprises the possibility of preparing derivatives of this type as a depot form for the purpose of miscibility, the depot auxiliary preferably being protamine sulfate and the insulin derivative and/or its physiologically tolerable salt being present with the protamine sulfate in a cocrystallizate.

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The present invention further relates to an injectable solution which comprises the pharmaceutical preparations described above in dissolved form.

Examples

Example 1:

Construction of Lys (B3)-proinsulin as a starting point for the plasmids relevant to the invention corresponding to Examples 2-4

US 5,358,857 describes the vector pINT 90d and the PCR primers Tir and Insu 11. These components serve as starting materials for the construction of a plasmid pINT 125d, which codes for the desired Lys (B3)-proinsulin.

Additionally, the primers Insu 35 having the sequence

5' TTT GTG AAG CAG CAC CTG 3'

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and Insu 36 having the sequence

5' CAG GTG CTG CTT CAC AAA 3'

are synthesized.

A PCR reaction is carried out using the primers Tir and Insu 36 and a second reaction is carried out using the primers Insu 11 and Insu 35. The template used for this is pINT 90d DNA.

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The products of the two PCR reactions are partially complementary, such that when they are combined in a third PCR reaction with the primers Tir and Insu 11 they afford a fragment which codes for a proinsulin variant which contains the B chain lysine in position 3. This PCR fragment is precipitated in ethanol for purification, dried and then digested with the restriction enzymes Nco 1 and Sal 1 according to the instructions of the manufacturer. The reaction mixture is separated by gel electrophoresis and the desired Nco 1/Sal 1 fragment is isolated.

The application cited describes a plasmid pINT 91d which codes for a mini-proinsulin. If the sequence coding for mini-proinsulin is excised by means of Nco 1 and Sal 1 and the residual plasmid DNA is isolated, this residual plasmid DNA can be reacted with the shown Nco 1/Sal 1 PCR fragment in a T 4 ligase reaction to give the plasmid pINT 125d. This is transformed by E. coli K12, replicated therein and reisolated. After verification of the plasmid structure by means of DNA sequence and restriction analysis, pINT 125d DNA is used as template DNA for the introduction of further mutations into this proinsulin variant.

Example 2: Construction of Lys (B3) Glu (B29)-proinsulin

For the preparation of the mutein, the primers 329a having the sequence

5' TTC TAC ACA CCC GAG ACC CGC GGC ATC G - 3'

and 329b having the sequence

5' GCC GCG GGT CTC GGG TGT GTA GAA GAA GC 3'

are synthesized.

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The template used is DNA of the plasmids pINT125d and pINT91d.

Primer 329a is reacted with the primer Insu 11 on the template pINT91d and primer 329b is reacted with Tir (see above example) on the template pINT125d in a PCR reaction. Since both PCR products are partially complementary, they can be combined in a direct PCR reaction and reacted again with the primers Tir and Insu 11. A DNA fragment results which codes for the desired mutein. This fragment is double-digested using the restriction enzymes Nco 1 and Sal 1 and the resulting Nco 1/Sal 1 fragment is inserted into the pINT 91d residual plasmid DNA in a T4 ligase reaction.

The plasmid pINT 329 results, which after amplification in E. coli K12 by means of restriction and DNA sequence analysis is verified with respect to the desired structure.

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The proinsulin derivative encoded by the plasmid is characterized by the two amino acid replacements and a C-bonding member which consists of the amino acid arginine.

Example 3: Construction of Lys (B3) Ile (B27)-proinsulin

The construction is carried out according to the preceding example using the primer pairs

15 KB3 JB 27A

5' TTC TAC ATC CCC AAG ACC CGC CG 3'

and Insu 11

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and also

K B3 J 27B

5' CTT GGG GAT GTA GAA GAA GCC TCG 3'

25

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and Tir.

The template used in both PCR reactions is DNA of the plasmid pINT125d. The PCR products of both reactions are combined in a third reaction, as described in Example 1, and the product is cloned corresponding to the example.

The plasmid pINT332 results.

Example 4: Construction of Lys (B3) Ile (B28)-proinsulin

The construction is carried out according to Example 3 using the primer pairs:

**KB3 JB 28A** 

5' TAC ACA ATC AAG ACC CGC CGG GAG - 3'

and Insu 11

10

and also

KB J B28B

5' GGT CTT GAT TGT GTA GAA GAA GCC TCG - 3'

15 and Tir.

The plasmid pINT 333 results.

Expression of the constructed insulin variants

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The plasmids pINT 329, 332 and 333 are each transformed by way of example by E. coli K12 W3110. Recombinant bacteria which contain plasmids which encode the respective variants are then fermented according to Example 4 of the US patent having the patent number 5227293 and the desired raw material for the production of the respective insulin variants is thus produced.

Example 5: Construction of Lys (B3), Ile (B28), Asp (A21)-proinsulin

Construction is carried out as in Example 3. Instead of pINT125d, however, the template serving for the PCR reaction is DNA of the plasmid pINT333, which was constructed in Example 4. The following primer pair is used here:

P-pint 365

# 5'-TTTTTTGTCGACTATTAGTCGCAGTAGTTCTACCAGCTG-3' and Tir.

# 5 The plasmid pINT365 results.

Example 6: Biological activity of Lys(B3), Glu(B29)-insulin after intravenous administration to rabbits

Time [h]	Human insulin	Lys(B3), Glu(B29)- insulin
0	100	100
0.25	89.17	89.47
0.5	67.56	58.32
0.75	73.24	66.59
1	73.13	68.21
1.5	78.12	71.95
2	89.47	80.88
3	107.01	94.2
4	104.55	99.78

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8 rabbits received the indicated insulins intravenously (0.2 IU/kg). In the course of the following four hours, the blood glucose concentration was determined at the times indicated and calculated in % of the starting value at time 0. The mean values show no significant differences in the biological activity between human insulin and Lys(B3),Glu(B29)-insulin.

Example 7: Biological activity of Lys(B3), Ile(B27)- and Lys(B3), Ile(B28)- insulin after intravenous administration to rabbits

6 rabbits received the indicated insulins intravenously (0.2 IU/kg). In the course of the following four hours, the blood glucose concentration was

determined at the times indicated and calculated in % of the starting value at time 0. The mean values show no significant differences in the biological activity between human insulin, Lys(B3),Ile(B27)- and Lys(B3),Ile(B28)-insulin.

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Time [h]	H insulin	Lys(B3), Ile(B27)- insulin	Lys(B3), lle(B28)- insulin
0	100	100	100
0.33	67.8	62.6	63.3
0.66	54.9	60.6	55.8
1	55.2	66.8	59.3
1.5	63	79.2	66.7
2	77.8	90.9	81.6
3	91.5	96.3	97.2
4	99.5	96	101.6

Example 8: Pharmacodynamics of Lys(B3),Glu(B29)-insulin and Lys(B3),Ile(B28)-insulin after subcutaneous administration to dogs

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Four dogs in each case received subcutaneous injections of the indicated insulins (0.3 IU/kg). The blood glucose was kept at 3.7 to 4 mmol/l by continuous infusion of glucose. The mean glucose infusion rate  $\pm$  SEM from the time of injection (t = 0) through 240 minutes is shown.

# WITH RAPIDLY ACTING INSULIN DER GLUCOSE CLAMP IN FASTING DOGS

Characteristics of the glucose infusion profiles

Dose: 1 x 0.3 IU/kg s.c. at to

	Incremer	crement phase	tmax	Decreme	Decrement phase
Preparation	Inflection point	Slope of the	(min)	Inflection point	Slope of the
	(min)	inflection point		(min)	inflection point
H. insulin, Hoechst	43	0.144	100	156	- 0.065
Lys(B3),Glu(B29) insulin	33	0.227	80	127	- 0.091
Lys(B3),lle(B28) insulin	16	0.267	60	104	- 0.102

# WITH RAPIDLY ACTING INSULIN DER SLUCOSE CLAMP IN FASTING DOGS

Dose: 1 x 0.3 iU/kg s.c. at to

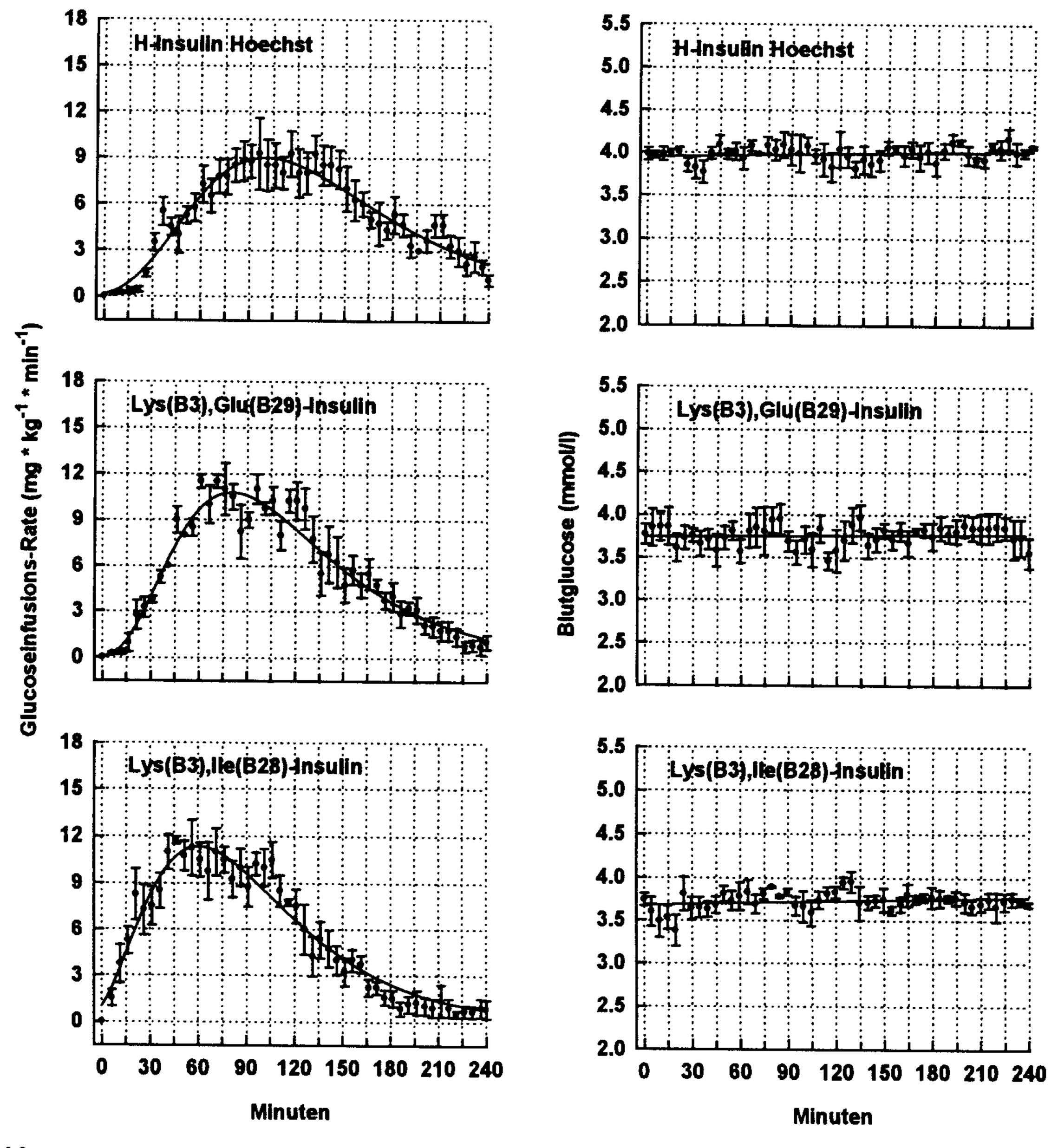
(mean ± sem, n = 4)

					· · · · · · · · · · · · · · · · · · ·						_																								-									
	Se	/ or	sem	0.07	0.17	0.14	0.18	0.13	0.12	0.09	9.0	2 C	0.16	0.14	0.11	90'0	0.0	0.02	0. c	. c	0.15	0.10	0.08	0.0	0.07	0.20	0.08	0.07	0.14 0.05	800	0.13	0.05	0.05 6.05	7.0	900	0.08	0.11	0.08	0.08	60.0	0.17	0.10	9 6	0.03
ا ا	d gluco	mmo	mean	3.74	3.60 3.49	3.53		3.64	3.65	3.63	3.68	3 73	3.78	3.83	3.69	3.81	3.89	3.78	5.83 6.7	80.0	.59	1.74	26.8	50.0	2 5	.70	.71	.73	. 6	69	80									·				
B28) insul	<b>60</b>	Time	min	0		5	-			_	<del>4</del> 5				20				3 8			•		125	3 8	135	9 9	145	155	_	165 3	_		2 2 2 2 2					210 3	215 3	220 3	225	3 %	240
(B3).lle(B	•	- <b>k</b> g-	sem	8 8	53	68	- 3 €	25	15	12	2 8	28	. g	85	<b>S</b>	<del></del>	<u> </u>	7 5	\$ Z	2 2	5	<u> </u>	2 2	2 C	1 0	<u></u>	<u>ඉ</u>	400	<u> </u>	<u>.</u>	4	<u> </u>	<b>2</b> 5	2 49	4	<u>ე</u>	<b>~</b>	<u> </u>	4 (		- ¢	0 00		
Lys	`	mg.min	nean	83	¥ 15	25 0.	25	. <del></del>	<b>8</b>	8 :	25	5.	. <del></del>	75	•	<u>0</u>	2 22	<b>.</b>	2.7 7.0 7.0 7.0	28			75 0.2%					<u> </u>	<u> </u>							<b></b>		0.6	7.0	0.0	20.0	2 C	9.0	3 0.6
	Glucose			0.	— w			_		4- 1	- 6	_	<del>-</del>		<b>*</b>	9.0	<u>ن</u> ج	2 ∘	<u> </u>	10	10	<b>∞</b>	<u> </u>		<del></del>		. <u>.</u>			<u> </u>	<del></del>	·	3 2	- 0	7.	1.2	•	ф О	4.0	5 C	4 6 -	90	9	0.8
	_	<u>=</u>	min	ő		<del>"</del>	7 8	<u>س</u>	· አ	4 ,	<del>4</del> <del>2</del> <del>2</del>	- S	6	99	7	9 7		- 8 8	- S		5	= ;	116	126	13.	136	142	- t	156	162	166	171	187	186	191		202	506	212	277	226	231	(1)	240
	ose	mol/l	sem	0.12	0.17	$\sim$	0.0	0.08	0.15	0.12	0. J	0.11	0.14	0.20	0.23	0.28	رن د م	5 5	0.14	0.16	0.20	0.16	0.10	0.23	0.21	0.15	0.15		0.11	0.11	0.15	9.6	0.00	0.14	0.10	0.14	0.13	0.14	0.17	0.18		0.23	' <b>_</b> *	0.18
lii	ood glucc	Ē	mean	3.77	3.86 3.86	3.86	3.73	3.76	3.66	3.71	3.73 3.73	3.81	3.57	3.81	3.85	3.87	2) (2) 4) (3)	2.62	3.56	3.71	3.59	3.84	3.46 2.46	3.70	3.88	3.97		2.5	3.71	3.81	3.67	3.81 8.93	3.5	3.85	3.78	3.81	88.	3.85	S 4		85	22.	.74	.56
329) -insu	面	Time	ے		ი 6		22 52	8	ဗ္တ (	<del>4</del>			8					3 8		_			1 1 C		_	_		_	155	160	ss (	5 t	2 8	185	961	ψ,			•			230		
Ĭ					<u> </u>	-	· · · · · · · · · · · · · · · · · · ·		-	- <u>-</u>			<del></del>	—	i	· <u>-</u> ·	"4	•				··· ·			·		<del></del>				• •			<del>, , , , , , , , , , , , , , , , , , , </del>	<i></i> :-	9244	·		·			<del></del>	· ·	
3,6	e T	9	E	O 9	<b>့</b>	0 4	o uo	7	<del>-</del> (	<b>5</b> N	<b>-</b> 0	ဖ	က	<b>~</b>	m c	<b>.</b>	ם נס		) <del></del>	_	•	•			•																_			
Lys(B3),G	usion rate	.min .kg	in sem	0.00	0.0	0.60	0.65	0.22	0.41	0.0	00.0	0.56	0.43	1.22	0.43	- C	2.5 2.5	2 2	9.0	0.41	0.89	96.0	1 2 4 4 2 4	1.34	1.52	44.	1.85 7.65	1.07	96.0	0.82	1.03	- 4.0 - 4.0	9.0	0.87	0.41	 20. (	0.51	9 6	0.62	2. C	98.0	0.37	0.51	0.47
.ys(B3),	cose infusion rate	mg.minkg	mean	0.00	ŧ (Q	36.	c.	ຸດ	<b></b>	0 0	<b>,</b>		0	0	0 6	- c	8.25			10	2		o ro														•			1.70	0.78	.85	.76	1.07 0.47
.ys(B3),	3	<b>a</b> )	nean	0.00	ŧ (Q	0.95	3.25	3.75	5.25	3 8	8 8	8.50	11.50	10.00	11.50	3 5	2.00 2.00 2.00 2.00	2 0	3 2 8	9.75	10.25	8.00	10.25	9.75	7.75	5.50	6.75	4.75	5.75	4.75	5.50	4.75	. 8	2.88	3.25	3.13	2.13			221 1.41 0.58	78	.85	36 0.76	240 1.07 0.47
.ys(B3),	Glucose infu	Time	sem min mean	0.00	.08 11 0.36	.04 0.95	08 3.25	.13 31 3.75	13 36 5.25	3 8	.06 51 8.00	.10 56 8.50	.13 61 11.50	.06 67 10.00	71 11.50	40 40 40 40 40 40 40 40 40 40 40 40 40 4	15 86 875	18 91	2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2	0 101 9.75	106 10.25	111 8.00	10.25	2 126 9.75	131 7.75	5.50	3 141 6.75	151 4.75	156 5.75	2 161 4.75	166 5.50	10 171 4.70	14 181 4.00	186 2.88	11 3.25	03 13	201 2.13	20.7	.06 04 04 04 04 04 04 04	11 221 1.70	12 226 0.78	.85	5 236 0.76	40 1.07
.ys(B3),	ucose infu	<b>a</b> )	sem min mean	.98 0.07 0 97 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0	98 0.08 11 0.36	.99 16 0.95 16 0.95 17 16 0.95 17 16 17 17 17 17 17 17 17 17 17 17 17 17 17	5 0.08 26 3.25	.82 0.13 31 3.75	7 0.13 36 5.25	9 41 5.00	98 51 8.00	01 0.10 56 8.50	93 61 11.50	08 0.06 67 10.00	98 11.50	0.10	2.00 2.00 2.00 2.00	3 0 18	96 11.00	3 0.10 101 9.75	3 0.08 10.25	2 0.19 111 8.00	10.77 10.25 10.71 10.25	0.12 126 9.75	131 7.75	0.16 136 5.50	0.13 141 6.75	151 4 75	0.04 156 5.75	0.12 161 4.75	166 5.50	17.1 4.75 0.10	0.14 181 4.00	0.09 186 2.88	0.11 191 3.25	3 0.03 196 3.13	8 0.11 201 2.13	0.00	2 1.79 0.06 0.07 0.04	5 0.11 221 1.41	12 226 0.78	.13 231 0.85	.05 236 0.76	2 240 1.07
Lys(B3),	Blood glucose infu	mol/l Time	n mean sem min mean	3.98 0.07 0.00	98 0.08 11 0.36	3.99 0.04 16 0.95	3.85 0.08 26 3.25	3.82 0.13 3.75	7 0.13 36 5.25	9 41 5.00	3.98 0.06 51 8.00	4.01 0.10 56 8.50	93 61 11.50	4.08 0.06 67 10.00	3.98 0.01 71 11.50	4.00	4.03	4.03	3.99 0.22 96 11.00	4.08 0.10 101 9.75	3.96 0.08 10.25	3.92 0.19 111 8.00	3.83 U.17 170 10.25 4.04 10.21 10.25	0.12 126 9.75	3.81 0.09 131 7.75	0.16 136 5.50	3 141 6.75	151 4 75	0.04 156 5.75	0.12 161 4.75	166 5.50	17.1 4.75 0.10	14 181 4.00	0.09 186 2.88	89 4.12 0.11 191 3.25	4 4.13 0.03 196 3.13	99 3.98 0.11 201 2.13	04 3.91 0.00 2.00	23   3.92   0.06   2.17   1.79	19 4.05 0.11 221 1.41	24 4.17 0.12 226 0.78	9 4.00 0.13 231 0.85	4 4.01 0.05 236 0.76	39 4.08 0.02 240 1.07
hst Lys(B3),	Blood glucose infu	e Time	mean sem min mean	3.98 0.07 0.00	3.98 0.08 11 0.36	3.99 0.04 16 0.95	3.85 0.08 26 3.25	3.82 0.13 3.75	7 0.13 36 5.25	9 41 5.00	3.98 0.06 51 8.00	4.01 0.10 56 8.50	3.93 6.13	4.08 0.06 67 10.00	3.98 0.01 71 11.50	4.00	4.03	4.03	3.99 0.22 96 11.00	4.08 0.10 101 9.75	3.96 0.08 10.25	3.92 0.19 111 8.00	3.83 U.17 170 10.25 4.04 10.21 10.25	4 3.95 0.12 126 9.75	3.81 0.09 131 7.75	0.16 136 5.50	0.13 141 6.75	151 4.75	0.04 156 5.75	0.12 161 4.75	166 5.50	17.1 4.75 0.10	0.14 181 4.00	4 4.03 0.09 186 2.88	9 4.12 0.11 191 3.25	4 4.13 0.03 196 3.13	99 3.98 0.11 201 2.13	04 3.91 0.00 2.00	23   3.92   0.06   2.17   1.79	5 0.11 221 1.41	24 4.17 0.12 226 0.78	29 4.00 0.13 231 0.85	4 4.01 0.05 236 0.76	39 4.08 0.02 240 1.07
hst Lys(B3),	on rate Blood glucose Glucose infu	nin kg Time mmol/l Time	sem min mean sem min mean	3.98 0.07 0.00	08 3.98 0.08 11 0.36	19 14 3.99 0.04 16 0.95	25 24 3.85 0.08 26 3.25 (	3.82 0.13 3.75	7 0.13 36 5.25	9 41 5.00	3.98 0.06 51 8.00	89 54 4.01 0.10 56 8.50	3.93 6.13	4.08 0.06 67 10.00	3.98 0.01 71 11.50	4.00	4.03	4.03	3 3.99 0.22 96 11.00	.68   4.08   0.10   101   9.75	4 104 3.96 0.08 106 10.25	6 109 3.92 0.19 111 8.00 13.92 1.0.0	3.83 U.17 170 10.25 4.04 10.21 10.25	4 3.95 0.12 126 9.75	9 129 3.81 0.09 131 7.75	0 134 3.92 0.16 136 5.50	0.13 141 6.75	151 4.75	29 154 4.05 0.04 156 5.75	79 159 3.97 0.12 161 4.75	50 164 4.04 0.09 166 5.50	39 109 3.90 0.10 171 4.70 54 174 4.01 0.10 176 2.75	0.14 181 4.00	72 184 4.03 0.09 186 2.88	72 189 4.12 0.11 191 3.25	0.00 194 4.13 0.03 196 3.13	.72 199 3.98 10.11 201 2.13	. 72   2.04   3.91   0.05   2.06   2.	23   3.92   0.06   2.17   1.79	04 219 4.05 0.11 221 1.41	68 224 4.17 0.12 226 0.78	9 4.00 0.13 231 0.85	.47 234 4.01 0.05 236 0.76	36 239 4.08 0.02 240 1.07
hst Lys(B3),	Blood glucose Glucose infu	e Time	mean sem min mean sem min mean	00 3.98 0.07 0.00 0.00 0.00 0.00 0.00 0.00 0.0	23 0.08 11 0.36 5.36 0.08 11 0.36 6.36 11 0.36	35 0.19 14 3.99 0.04 16 0.95 14 0.05 15 16 0.95 17 18 0.95 18 18 18 18 18 18 18 18 18 18 18 18 18	50 0.25 24 3.85 0.08 26 3.25	50 0.56 29 3.82 0.13 3.75	50 0.90 34 3.77 0.13 36 5.25 5.25 5.25 5.25 5.25 5.25 5.25 5.2	30 0.50 39 0.07 41 6.00 00 00 00 00 00 00 00 00 00 00 00 00	25 0.54 49 3.98 0.06 51 8.00 8.00	75 0.89 54 4.01 0.10 56 8.50	25 1.19 61 11.50 61 11.50 Fig. 11.50	50   1.09   64   4.08   0.06   67   10.00	75   1.08   69   3.98   0.01   71   11.50   75   74   74   75   74   75   75   75	10.10	75 1.09 0.12 75 1.29 84 4.09 0.15 86 8.25	75 1.08 4.03 0.18 9.00	5   2.33   94   3.99   0.22   96   11.00	50 1.68 4.08 0.10 101 9.75	50 1.44 104 3.96 0.08 106 10.25	.00 1.06 1.09 3.92 0.19 111 8.00	7. 114   3.83   0.17   110.25   10.25   10.25   119   4.04   10.21   10.25   1	00 1.37 124 3.95 0.12 126 9.75	.25   1.19   1.29   3.81   0.09   1.15   7.75	.50 1.30 134 3.92 0.16 136 5.50	.50   1.25   139   3.86   0.13   141   6.75	151 4.75	5 1.29 1.54 4.05 0.04 1.56 5.75	0.79 159 3.97 0.12 161 4.75	0.50 164 4.04 0.09 166 5.50	39 109 3.90 0.10 171 4.70 54 174 4.01 0.10 176 2.75	1.19   3.88   0.14   4.00	7 0.72 184 4.03 0.09 186 2.88	3 0.72 189 4.12 0.11 191 3.25	0 0.00 194 4.13 0.03 196 3.13	7 0.72   199   3.98   0.11   201   2.13   7.05   7.	7 00.72   2.91   0.05   2.06	57   0.72   5.92   0.05   1.79   3.97   1.79   3.97	33 10.72 1.03 1.00 00 00 1.00 1.00 1.70 1.70 1.70 1.70 1.70	17 0.68 4.17 0.12 226 0.78	98 229 4.00 0.13 231 0.85	00 0.47 234 4.01 0.05 236 0.76	17 0.36 239 4.08 0.02 240 1.07
hst Lys(B3),	infusion rate Blood glucose Glucose infu	min kg Time mmol/l Time	mean sem min mean sem min mean	0.00 0.00 3.98 0.07 0.00 0.00 0.00 0.00 0.00 0.00 0.0	23 0.08 11 0.36 5.36 0.08 11 0.36 6.36 11 0.36	0.35 0.19 14 3.99 0.04 16 0.95 10 0.44 0.44 0.95 10 0.05 10 0.	50 0.25 24 3.85 0.08 26 3.25	50 0.56 29 3.82 0.13 3.75	50 0.90 34 3.77 0.13 36 5.25 5.25 5.25 5.25 5.25 5.25 5.25 5.2	30 0.50 39 0.07 41 6.00 00 00 00 00 00 00 00 00 00 00 00 00	25 0.54 49 3.98 0.06 51 8.00 8.00	75 0.89 54 4.01 0.10 56 8.50	25 1.19 61 11.50 61 11.50 Fig. 11.50	50   1.09   64   4.08   0.06   67   10.00	75   1.08   69   3.98   0.01   71   11.50   75   74   74   75   74   75   75   75	10.10	8 75 84 4.09 0.12 0.1 8 75 84 4.09 0.15 86 8.25	75 1.08 4.03 0.18 9.00	9.25   2.33   94   3.99   0.22   96   11.00	8.50   1.68   99   4.08   0.10   101   9.75	50 1.44 104 3.96 0.08 106 10.25	.00 1.06 1.09 3.92 0.19 111 8.00	.25   1.47   3.83   0.17   10.25   10.	00 1.37 124 3.95 0.12 126 9.75	.25   1.19   1.29   3.81   0.09   1.15   7.75	.50 1.30 134 3.92 0.16 136 5.50	.50   1.25   139   3.86   0.13   141   6.75	00 1.46 1.49 1.06 0.08 151 4.75	25 1.29 154 4.05 0.04 156 5.75	00 0.79 159 3.97 0.12 161 4.75	0.50 164 4.04 0.09 166 5.50	1.39   3.90   0.10   1/1   4./0	1.19   3.88   0.14   4.00	7 0.72 184 4.03 0.09 186 2.88	33 0.72 189 4.12 0.11 191 3.25	96   3.00   0.00   194   4.13   0.03   3.13	01   3.67   0.72   199   3.98   0.11   201   2.13	06   4.67   0.72   2.04   3.91   0.05   2.06	11   4.67   0.72   5.92   1.79	21 3.53 0.74 2.19 4.05 0.04 1.41	26   2.00   2.24   4.17   0.12   0.78   0.78	67         0.98         4.00         0.13         231         0.85	36 2.00 0.47 234 4.01 0.0 <del>5</del> 236 0.76	40 1.17 0.36 2.39 4.08 0.02 2.40 1.07

# GLUCOSE-CLAMP BEIM NÜCHTERNEN HUND MIT SCHNELL WIRKSAMEN INSULINDERIVATEN

Dosis: 1 x 0.3 IU/kg s.c. bei t<sub>0</sub>

 $(mean \pm sem, n = 4)$ 



# Key:

- 1 GLUCOSE CLAMP IN FASTING DOGS WITH RAPIDLY ACTING INSULIN DERIVATIVES
- 5 Dose: 1 × 0.3 IU/kgs.c.at t<sub>0</sub>
  - 3 H-insulin, Hoechst
  - Glucose infusion rate (mg.kg<sup>-1</sup>.min<sup>-1</sup>)
  - 5 Minutes

### SEQUENCE LISTING

### (1) GENERAL INFORMATION:

- (i) APPLICANT:
  - (A) NAME: Sanofi-Aventis Deutschland GmbH
  - (B) STREET: -
  - (C) CITY: Frankfurt
  - (D) STATE: -
  - (E) COUNTRY: Germany
  - (F) POSTAL CODE (ZIP): 65926
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  - (H) TELEFAX: 069-35-7175
  - (I) TELEX: -
- (ii) TITLE OF INVENTION: NOVEL INSULIN DERIVATIVES HAVING A RAPID ONSET OF ACTION
- (iii) NUMBER OF SEQUENCES: 11
- (iv) COMPUTER READABLE FORM:
  - (A) MEDIUM TYPE: Floppy disk
  - (B) COMPUTER: IBM PC compatible
  - (C) OPERATING SYSTEM: PC-DOS/MS-DOS
  - (D) SOFTWARE: PatentIn Release #1.0, Version #1.30 (EPO)
- (vi) CURRENT APPLICATION DATA:
  - (A) APPLICATION NUMBER: CA 2,235,443
  - (B) FILING DATE: 18-JUN-1998
  - (C) CLASSIFICATION:
- (vii) PRIOR APPLICATION DATA:
  - (A) APPLICATION NUMBER: DE 19726167.1
  - (B) FILING DATE: 20-JUN-1997
  - (C) CLASSIFICATION:
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- (2) INFORMATION FOR SEQ ID NO: 1:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 21 amino acids
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS: single
    - (D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: peptide
  - (ix) FEATURE:
    - (A) NAME/KEY: Peptide
    - (B) LOCATION: 1..21

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:

Gly Ile Val Glu Gln Cys Cys Thr Ser Ile Cys Ser Leu Tyr Gln Leu 1 15

Glu Asn Tyr Cys Asn 20

- (2) INFORMATION FOR SEQ ID NO: 2:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 30 amino acids
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS: single
    - (D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: peptide
  - (ix) FEATURE:
    - (A) NAME/KEY: Peptide
    - (B) LOCATION: 1..30
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:

Phe Val Asn Gln His Leu Cys Gly Ser His Leu Val Glu Ala Leu Tyr 1 15

Leu Val Cys Gly Glu Arg Gly Phe Phe Tyr Thr Pro Lys Thr 20 30

- (2) INFORMATION FOR SEQ ID NO: 3:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 30 amino acids
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS: single
    - (D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: peptide
  - (ix) FEATURE:
    - (A) NAME/KEY: Peptide
    - (B) LOCATION:1..30
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:

Phe Val Lys Gln His Leu Cys Gly Ser His Leu Val Glu Ala Leu Tyr 1 15

Leu Val Cys Gly Glu Arg Gly Phe Phe Tyr Thr Pro Glu Thr 20 30

- (2) INFORMATION FOR SEQ ID NO: 4:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 30 amino acids
    - (B) TYPE: amino acid

- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: peptide
- (ix) FEATURE:
  - (A) NAME/KEY: Peptide
  - (B) LOCATION: 1..30
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4:

Phe Val Lys Gln His Leu Cys Gly Ser His Leu Val Glu Ala Leu Tyr 1 15

Leu Val Cys Gly Glu Arg Gly Phe Phe Tyr Thr Ile Lys Thr 20 30

- (2) INFORMATION FOR SEQ ID NO: 5:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 30 amino acids
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS: single
    - (D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: peptide
  - (ix) FEATURE:
    - (A) NAME/KEY: Peptide
    - (B) LOCATION: 1..30
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 5:

Phe Val Lys Gln His Leu Cys Gly Ser His Leu Val Glu Ala Leu Tyr 5

Leu Val Cys Gly Glu Arg Gly Phe Phe Tyr Ile Pro Lys Thr 20 30

- (2) INFORMATION FOR SEQ ID NO: 6:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 97 amino acids
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS: single
    - (D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: peptide
  - (ix) FEATURE:
    - (A) NAME/KEY: Peptide
    - (B) LOCATION:1..97
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 6:

Met Ala Thr Thr Ser Thr Gly Asn Ser Ala Arg Phe Val Lys Gln His 1 10 15

Leu Cys Gly Ser His Leu Val Glu Ala Leu Tyr Leu Val Cys Gly Glu 20 25 30

Arg Gly Phe Phe Tyr Thr Pro Glu Thr Arg Arg Glu Ala Glu Asp Pro 35 40

Gln Val Gly Glu Leu Gly Gly Gly Pro Gly Ala Gly Ser Leu 50

Gln Pro Leu Ala Leu Glu Gly Ser Leu Gln Lys Arg Gly Ile Val Glu 65 75 80

Gln Cys Cys Thr Ser Ile Cys Ser Leu Tyr Gln Leu Glu Asn Tyr Cys 85 90

Asn

## (2) INFORMATION FOR SEQ ID NO: 7:

- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 97 amino acids
  - (B) TYPE: amino acid
  - (C) STRANDEDNESS: single
  - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: peptide
- (ix) FEATURE:
  - (A) NAME/KEY: Peptide
  - (B) LOCATION: 1..97
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 7:

Met Ala Thr Thr Ser Thr Gly Asn Ser Ala Arg Phe Val Lys Gln His 1 10

Leu Cys Gly Ser His Leu Val Glu Ala Leu Tyr Leu Val Cys Gly Glu 20 30

Arg Gly Phe Phe Tyr Thr Ile Lys Thr Arg Arg Glu Ala Glu Asp Pro 35

Gln Val Gly Glu Leu Gly Gly Gly Pro Gly Ala Gly Ser Leu 50

Gln Pro Leu Ala Leu Glu Gly Ser Leu Gln Lys Arg Gly Ile Val Glu 65 70 75 80

Gln Cys Cys Thr Ser Ile Cys Ser Leu Tyr Gln Leu Glu Asn Tyr Cys 85 90

Asn

## (2) INFORMATION FOR SEQ ID NO: 8:

- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 97 amino acids
  - (B) TYPE: amino acid
  - (C) STRANDEDNESS: single
  - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: peptide
- (ix) FEATURE:
  - (A) NAME/KEY: Peptide
  - (B) LOCATION: 1..97
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 8:

Met Ala Thr Thr Ser Thr Gly Asn Ser Ala Arg Phe Val Lys Gln His 1 10

Leu Cys Gly Ser His Leu Val Glu Ala Leu Tyr Leu Val Cys Gly Glu
20 30

Arg Gly Phe Phe Tyr Ile Pro Lys Thr Arg Arg Glu Ala Glu Asp Pro 35

Gln Val Gly Glu Leu Gly Gly Gly Pro Gly Ala Gly Ser Leu 50 60

Gln Pro Leu Ala Leu Glu Gly Ser Leu Gln Lys Arg Gly Ile Val Glu 65 75 80

Gln Cys Cys Thr Ser Ile Cys Ser Leu Tyr Gln Leu Glu Asn Tyr Cys 85 90

Asn

- (2) INFORMATION FOR SEQ ID NO: 9:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 21 amino acids
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS: single
    - (D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: peptide
  - (ix) FEATURE:
    - (A) NAME/KEY: Peptide
    - (B) LOCATION:1..21
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 9:
  - Gly Ile Val Glu Gln Cys Cys Thr Ser Ile Cys Ser Leu Tyr Gln Leu 1 10 15

Glu Asn Tyr Cys Asp

(2) INFORMATION FOR SEQ ID NO: 10:

- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 30 amino acids
  - (B) TYPE: amino acid
  - (C) STRANDEDNESS: single
  - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: peptide
- (ix) FEATURE:
  - (A) NAME/KEY: Peptide
  - (B) LOCATION: 1..30
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 10:

Phe Val Lys Gln His Leu Cys Gly Ser His Leu Val Glu Ala Leu Tyr 1 15

Leu Val Cys Gly Glu Arg Gly Phe Phe Tyr Thr Ile Lys Thr 20 30

- (2) INFORMATION FOR SEQ ID NO: 11:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 97 amino acids
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS: single
    - (D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: peptide
  - (ix) FEATURE:
    - (A) NAME/KEY: Peptide
    - (B) LOCATION: 1..97
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 11:

Met Ala Thr Thr Ser Thr Gly Asn Ser Ala Arg Phe Val Lys Gln His 1 10

Leu Cys Gly Ser His Leu Val Glu Ala Leu Tyr Leu Val Cys Gly Glu 25

Arg Gly Phe Phe Tyr Thr Ile Lys Thr Arg Arg Glu Ala Glu Asp Pro 35

Gln Val Gly Glu Leu Gly Gly Gly Pro Gly Ala Gly Ser Leu 50

Gln Pro Leu Ala Leu Glu Gly Ser Leu Gln Lys Arg Gly Ile Val Glu 65 75 80

Gln Cys Cys Thr Ser Ile Cys Ser Leu Tyr Gln Leu Glu Asn Tyr Cys 85 90

Asp

# We claim:

1. An insulin derivative or a physiologically tolerable salt thereof comprising an A chain and a B chain;

wherein the A chain has the sequence:

Gly Ile Val Glu Gln Cys Cys Thr Ser Ile Cys Ser Leu Tyr Gln Leu Glu Asn Tyr Cys Asn (SEQ ID NO:1); and wherein the B chain has the sequence:

Phe Val Lys Gln His Leu Cys Gly Ser His Leu Val Glu Ala Leu Tyr Leu Val Cys Gly Glu Arg Gly Phe Phe Tyr Thr Pro Glu Thr (SEQ ID NO 3).

2. An insulin derivative or a physiologically tolerable salt thereof comprising an A chain and a B chain;

wherein the A chain has the sequence:

Gly Ile Val Glu Gln Cys Cys Thr Ser Ile Cys Ser Leu Tyr Gln Leu Glu Asn Tyr Cys Asn (SEQ ID NO:1); and wherein the B chain has the sequence:

Phe Val Lys Gln His Leu Cys Gly Ser His Leu Val Glu Ala Leu Tyr Leu Val Cys Gly Glu Arg Gly Phe Phe Tyr Thr Ile Lys Thr (SEQ ID NO 4).

3. An insulin derivative or a physiologically tolerable salt thereof comprising an A chain and a B chain;

wherein the A chain has the sequence:

Gly lle Val Glu Gln Cys Cys Thr Ser lle Cys Ser Leu Tyr Gln Leu Glu Asn Tyr Cys Asp (SEQ ID NO:9); and wherein the B chain has the sequence:

Phe Val Lys Gln His Leu Cys Gly Ser His Leu Val Glu Ala Leu Tyr Leu Val Cys Gly Glu Arg Gly Phe Phe Tyr Thr Ile Lys Thr (SEQ ID NO 4).

- 4. A human insulin derivative or a physiologically tolerable salt thereof comprising a lysine amino acid residue at B3 and a glutamic acid amino acid residue at B29.
- 5. A human insulin derivative or a physiologically tolerable salt thereof comprising a lysine amino acid residue at B3 and an isoleucine amino acid residue at B28.
- 6. A human insulin derivative or a physiologically tolerable salt thereof comprising a lysine amino acid residue at B3, an isoleucine amino acid residue at B28 and an aspartic acid amino acid residue at A21.
- 7. A process for the preparation of an insulin derivative or of a physiologically tolerable salt thereof as claimed in any one of claims 1 to 6, comprising
- (a) constructing a replicable expression vehicle which contains a DNA sequence which codes for a precursor of the insulin derivative, in which the amino acid residue in position A1 of the A chain is linked to the amino acid residue B30 of the B chain via a peptide chain of the formula II

in which  $R_n^1$  is a peptide chain having n amino acid residues and n is an integer from 0 to 34, and the B chain is modified by covalent linkage of the amino acid at position B1 to a peptide chain of the formula III

$$Met-R^2_{m}-(Arg)_{p}-$$

in which  $R_m^2$  is a peptide chain having m amino acid residues, m is an integer from 0 to 40 and p is 0, 1 or 2;

- (b) expressing the DNA sequence which codes for the precursor of the insulin derivative in a host cell; and
- (c) releasing the insulin derivative from its precursor using chemical and/or enzymatic methods.

- 8. The process as claimed in claim 7, wherein the host cell is a bacterium.
- 9. The process as claimed in claim 8, wherein the bacterium is *Escherichia coli*.
- 10. The process as claimed in claim 7, wherein the host cell is a yeast.
- 11. The process as claimed in claim 10, wherein the yeast is *Saccharomyces* cerevisiae.
- 12. The process as claimed in any one of claims 7 to 11 for the preparation of an insulin derivative as claimed in claim 1, wherein the precursor of the insulin derivative has the sequence:

Met Ala Thr Thr Ser Thr Gly Asn Ser Ala Arg
Phe Val Lys Gln His Leu Cys Gly Ser His Leu Val Glu Ala Leu
Tyr Leu Val Cys Gly Glu Arg Gly Phe Phe Tyr Thr Pro Glu Thr
Arg Arg Glu Ala Glu Asp Pro Gln Val Gly Gln Val Glu Leu Gly
Gly Gly Pro Gly Ala Gly Ser Leu Gln Pro Leu Ala Leu Glu Gly
Ser Leu Gln Lys Arg
Gly Ile Val Glu Gln Cys Cys Thr Ser Ile Cys Ser Leu Tyr Gln
Leu Glu Asn Tyr Cys Asn
(SEQ ID NO 6).

13. The process as claimed in any one of claims 7 to 11 for the preparation of an insulin derivative as claimed in claim 2, wherein the precursor of the insulin derivative has the sequence:

Met Ala Thr Thr Ser Thr Gly Asn Ser Ala Arg
Phe Val Lys Gln His Leu Cys Gly Ser His Leu Val Glu Ala Leu
Tyr Leu Val Cys Gly Glu Arg Gly Phe Phe Tyr Thr Ile Lys Thr
Arg Arg Glu Ala Glu Asp Pro Gln Val Gly Gln Val Glu Leu Gly
Gly Gly Pro Gly Ala Gly Ser Leu Gln Pro Leu Ala Leu Glu Gly

Ser Leu Gln Lys Arg
Gly Ile Val Glu Gln Cys Cys Thr Ser Ile Cys Ser Leu Tyr Gln
Leu Glu Asn Tyr Cys Asn
(SEQ ID NO 7).

14. The process as claimed in any one of claims 7 to 11 for the preparation of an insulin derivative as claimed in claim 3, wherein the precursor of the insulin derivative has the sequence:

Met Ala Thr Thr Ser Thr Gly Asn Ser Ala Arg Phe Val Lys Gln His Leu Cys Gly Ser His Leu Val Glu Ala Leu Tyr Leu Val Cys Gly Glu Arg Gly Phe Phe Tyr Thr Ile Lys Thr Arg Arg Glu Ala Glu Asp Pro Gln Val Gly Gln Val Glu Leu Gly Gly Gly Pro Gly Ala Gly Ser Leu Gln Pro Leu Ala Leu Glu Gly Ser Leu Gln Lys Arg Gly Ile Val Glu Gln Cys Cys Thr Ser Ile Cys Ser Leu Tyr Gln Leu Glu Asn Tyr Cys Asp (SEQ ID NO:11).

- 15. A precursor of the insulin derivative as claimed in claim 12.
- 16. A precursor of the insulin derivative as claimed in claim 13.
- 17. A precursor of the insulin derivative as claimed in claim 14.
- 18. A nucleic acid molecule comprising a DNA sequence which codes for a precursor of the insulin derivative as claimed in claim 15.
- 19. A nucleic acid molecule comprising a DNA sequence which codes for a precursor of the insulin derivative as claimed in claim 16.
- 20. A nucleic acid molecule comprising a DNA sequence which codes for a precursor of the insulin derivative as claimed in claim 17.
- 21. An expression vehicle comprising the nucleic acid molecule as claimed in claim 18.

- 22. An expression vehicle comprising the nucleic acid molecule as claimed in claim 19.
- 23. An expression vehicle comprising the nucleic acid molecule as claimed in claim 20.
- 24. A host cell which is transformed using the expression vehicle as claimed in any one of claims 21 to 23.
- 25. A pharmaceutical preparation, which comprises at least one insulin derivative and/or a physiologically tolerable salt thereof as claimed in any one of claims 1 to 6 and a physiologically acceptable carrier.
- 26. A pharmaceutical preparation as claimed in claim 25, which comprises the insulin derivative and/or the physiologically tolerable salt thereof in dissolved, amorphous and/or crystalline form.
- 27. A pharmaceutical preparation as claimed in claim 25 or 26, which further comprises a depot auxiliary.
- 28. A pharmaceutical preparation as claimed in claim 27, wherein the depot auxiliary is protamine sulfate, where the insulin derivative and/or the physiologically tolerable salt thereof is present with the protamine sulfate in a cocrystallizate.
- 29. An injectable solution having insulin activity, comprising the pharmaceutical preparation as claimed in any one of claims 25 to 28 in dissolved form.
- 30. The use of the insulin derivative and/or its physiologically tolerable salt as claimed in any one of claims 1 to 6 for the production of a pharmaceutical preparation which has an insulin activity with a rapid onset of action.

- 31. The use of the pharmaceutical preparation as claimed in any one of claims 25 to 28 for treating diabetes mellitus.
- 32. The use of the insulin derivative and/or its physiologically tolerable salt as claimed in any one of claims 1 to 6 for treating diabetes mellitus.
- 33. The use of the insulin derivative and/or its physiologically tolerable salt as claimed in claim 4 for treating diabetes mellitus.