Process for the production of biodegradable, functionalised polymer particles and use thereof as pharmaceutical supports

The present invention relates to a process for the production of biodegradable, functionalised polymer particles and to the use thereof as pharmaceutical supports.

Background of the invention

The effectiveness of many active ingredients depends upon whether they are transported exactly to the desired target site in the body as the active ingredients used are frequently able to exert their full action only at a specific location and are ineffective at other locations or even have negative effects there. The latter particularly applies when combatting cancerous conditions and in particular cancerous conditions occurring in the central nervous system (CNS).

One good example of the difficulties which may occur on transport of active ingredients in the body is the blood-brain barrier (BBB). Active ingredients destined for the CNS must first cross this barrier. This network of blood vessels and cells protects the CNS and prevents non-water-soluble substances from entering the CNS. Fat-soluble substances can straightforwardly pass through this barrier by simple diffusion. Active and passive transport systems are, however, necessary for the transport of polar substances and ions. However, since, for example, more than 98% of all newly discovered medicines destined for the CNS are not water-soluble, they cannot cross the BBB and thus cannot reach their target site.

This example is intended to illustrate the difficulties which may occur during transport of active ingredients in the animal organism. In the case of the BBB, for example, it has been attempted to make it more permeable to active ingredients by modifying the membrane's permeability. Permeability of the BBB to active ingredients may, for example, be achieved by artificially increasing osmotic pressure or by administering bradykinin analogues. Sanovich et al. (Sanovich, E. et al., Brain Res. 1995, Vol. 705(1-2), pp. 125-135) and others have, for example, demonstrated increased permeability of the BBB to lanthanum by simultaneous administration with the bradykinin analogue RMP-7. One fundamental disadvantage of opening the BBB by one of the above-stated

mechanisms is that permeability to all substances, which thus also includes toxic substances which can damage the target cells in the CNS, is consequently increased.

Another possibility is to modify the active ingredient chemically in such a manner that it becomes more lipophilic and may thus also more readily pass through the BBB, as has for example been demonstrated for chlorambucil derivatives (Greig, N.H. et al., Cancer Chemother. Pharmacol. 1990, Vol. 25(5), pp. 320-325).

An alternative to both the above-stated systems is to use nano- or microparticles, the use of which does not entail modifying either the permeability of the membrane or the active ingredient. Kreuter (Kreuter, J.J. Anat. 1996, Vol. 189(3), pp. 503-505) demonstrates that the polymer particles used for this purpose are substantially produced by chemical processes, such as emulsion polymerisation, interfacial polymerisation, desolvation, evaporation and solvent precipitation.

According to DE 197 45 950 A1, polymer particles prepared from the most varied substances (polymers (in this case: polybutyl cyanoacrylate), solid or liquid lipids, o/w emulsions, w/o/w emulsions or phospholipid vesicles), to which the pharmaceutical substances are attached, are used to transport these substances into the CNS.

Another critical factor in the transport of nano- and microparticles through a membrane, and in particular the membranes of the BBB, is the size of the polymer particles. Previous research results show that polymer particles up to a size of 270 nm are capable of getting through the BBB (Lockman, P.R. et al., Drug Develop. Indust. Pharmacy 2002, Vol. 28(1), pp. 1-12).

It is thus important for the polymer particles used to be of a specific size. ?According to Lockman (Lockman, P.R. et al., Drug Develop. Indust. Pharmacy 2002, Vol. 28(1), pp. 1-12), the size of the polymer particles produced by the above processes is generally determined by photon correlation spectroscopy. This method, which is based on Brownian motion, measures the polymer particles with a laser beam, the time dependency of light variation being used to determine particle size. However, this

analytical method for the subsequent determination of the size of the polymer particles produced is very time-consuming and costly.

The object of the present invention is to provide a rapidly usable, low cost production or transport system for biologically active substances which permits effective and reliable production or transport of active ingredients, or to at least provide the public with a useful choice.

Summary of the invention

Accordingly, a first aspect of the invention relates to a process for producing polymer particles having surface-bound proteins, the process comprising A) providing a cell comprising one or more genes that code for one or more fusion proteins, the fusion proteins comprising a polymer synthase, and (a) one or more binding domains capable of binding one or more biologically active substances or one or more coupling reagents or a combination thereof, or (b) one or more biologically active proteins, or (c) a combination thereof, the polymer synthase comprising a polymer particle binding domain; and B) cultivating the cell in a culture medium so that the cell produces the fusion protein from the one or more genes and the fusion protein produces polymer particles, wherein the polymer particle binding domain of the fusion protein is covalently bound to a polymer particle; and C) separating the polymer particles from the cultivated cells to produce a composition comprising polymer particles having surface-bound proteins.

Another aspect of the invention relates to an in vitro process for producing polymer particles having surface-bound proteins, the process comprising A) providing a solution suited to polymer particle formation comprising at least one substrate. B) introducing into the solution one or more fusion proteins, the fusion proteins comprising a polymer synthase and (a) one or more binding domains capable of binding one or more biologically active substances or one or more coupling reagents or a combination thereof, or (b) one or more biologically active proteins, or (c) a combination thereof, the polymer synthase of the fusion protein comprising a polymer particle binding domain, and C) maintaining the solution suited to polymer particle formation so that the fusion protein produces polymer particles, wherein the polymer particle binding domain of the

fusion protein is covalently bound to a polymer particle.

Another aspect of the invention relates to a polymer particle comprising a polymer core and one or more surface-bound proteins, the one or more surface-bound proteins comprising A) one or more fusion proteins comprising (i) a polymer synthase comprising a particle binding domain and (ii) one or more binding domains capable of binding one or more biologically active substances or one or more coupling reagents or a combination thereof, or B) one or more fusion proteins comprising (i) a polymer synthase comprising a particle binding domain and (ii) one or more biologically active proteins, or C) a combination thereof, wherein the polymer particle binding domain of the fusion protein is covalently bound to the polymer particle.

Another aspect of the invention relates to a method of binding a biologically active substance comprising A) providing a composition of polymer particles produced by a method of the invention or providing a polymer particle of the invention, wherein optionally a coupling reagent is bound to the binding domain when a binding domain is present, and B) contacting the composition or polymer particle with a sample comprising a biologically active substance so that the binding domain or the biologically active protein or the coupling reagent binds the biologically active substance.

Described is a nucleic acid molecule encoding a fusion protein comprising a polymer synthase and one or more binding domains or one or more biologically active proteins or a combination thereof, wherein (1) the binding domain is selected from one or more enzymes, one or more abzymes, one or more non-catalytic proteins or one or more antibodies, one or more FLAG epitopes, one or more cysteine residues, or a combination of any two or more thereof, (2) the biologically active protein is selected from one or more oligopeptides, one or more enzymes, one or more abzymes, one or more non-catalytic proteins or one or more antibodies, one or more FLAG epitopes, one or more cysteine residues, or a combination of any two or more thereof.

Described is an expression vector comprising a nucleic acid of the invention.

Described is a host cell comprising a nucleic acid, a vector or a polymer particle of the invention.

Another aspect of the invention relates to a polymer particle produced according to a method of the invention.

In the description in this specification reference may be made to subject matter which isnot within the scope of the claims of the current application. That subject matter should be readily identifiable by a person skilled in the art and may assist in putting into practice the invention as defined in the claims of this application.

For the purposes of the present invention, "biologically active substance" is any substance capable of initiating a biological response on the part of the organism. These substances comprise not only enzymes and abzymes, which catalyse a specific reaction in the organism, and proteins, such as for example antibodies, which bring about an indirect response of the organism to the presence of this substance in the organism, but also inorganic and organic molecules which are non-biological in origin, i.e. are not formed by a naturally occurring organism, but are instead produced artificially. Most pharmaceutical active ingredients also belong to the latter group. Depending on their nature, the biologically active substances may also be suitable for binding further biologically active substances.

Described is a process involving introducing at least one inducible gene into a microorganism, wherein the gene codes for a protein which controls the size of the polymer particles, and culturing the microorganism with induction of the above-stated, at least one inducible gene in a culture medium under conditions which are suitable for the production of the biodegradable polymer particles by the microorganism.

By means of these processes, it is possible to produce biocompatible, biodegradable polymer particles which are suitable for transporting biologically active substances and wherein, by optimally controlling the size of the polymer particles formed, said particles may be produced as required in the desired size. Due to the controlled production of polymer particles of a specific size, the yield of polymer particles of the desired size is

increased, so increasing the efficiency of the process and simultaneously helping to reduce costs. Moreover, polymer particles may be produced by the process according to the invention which meet the above-stated particle size requirement for transport through the BBB. Thanks to this process it is above all possible to produce polymer particles which are smaller than the polymer particles of this kind naturally produced by microorganisms. The average size of the polymer particles naturally produced by microorganisms is 300 to 500 nm (Wieczorek, R. et al, J. Bacteriol. 1997, Vol. 177(9), pp. 2425-2435), while the process according to the invention makes it possible to control production of the polymer particles in such a manner that they are considerably smaller than this average value.

The particle size-determining gene is here induced by an upstream inducible promoter, such as for example a BAD promoter, which is induced by arabinose. The microorganisms used for this purpose have no gene for controlling the size of the polymer particles or this gene is inactivated and replaced by the at least one inducible gene described in the process according to the invention, which gene codes for a protein which controls the size of the polymer particles. This gene is here introduced into the cell by means of a vector which is described in greater detail in the experimental section of this description. As a consequence, it is for the first time possible to produce biocompatible, biodegradable polymer particles of a defined size in a microbial process.

These polymer particles are deposited as cytoplasmic inclusions in the cell. The core of these polymer particles consists of polyhydroxyalkyl carboxylates, in particular polyhydroxy alkanoates, and is enclosed by a shell membrane consisting of proteins and phospholipids. The shell membrane consists of lipids and proteins embedded therein. The polyhydroxyalkyl carboxylates, which form the core of these polymer particles, have melting points of 50°C to 176°C, a crystallinity of 30% to 70% and elongation at break values of 5% to 300%.

In order also to be able to use this advantageous process in microorganisms which are more suited to biotechnological cultivation (for example certain forms of *E. coli*, which are classified as GRAS organisms) but, due to their genetic makeup, are unable to form the above-stated polymer particles, at least one further gene which codes for a protein

involved in the formation of the polymer particles is introduced as well as the at least one inducible gene which codes for a protein which controls the size of the polymer particles. Any protein capable of influencing the metabolism leading to formation of the polymer particles and thus the composition of the polymer particles formed may be considered. The at least one further gene which codes for a protein involved in the formation of the polymer particles is here selected such that it codes for a thiolase, a reductase or a polymer synthase. A polymer synthase is taken to be any protein which is capable of catalysing the final step for formation of a polymer. Apart from the polymer synthases described in the present invention, formation of a polymer may, for example, also be undertaken by a lipase. The at least one further gene which codes for a protein involved in the formation of the polymer particles is preferably selected such that it codes for phaA thiolase, phaB ketoacyl reductase or phaC synthase from *Ralstonia eutropha*. Due to the introduction of these additional genes, the cell is enabled to produce proteins which allow it to form the polymer particles. Purposeful selection of the at least one further gene which codes for a protein involved in the formation of the polymer particles also makes it possible to influence the subsequent composition of the polymer particles. Genes which code for proteins involved in the metabolic pathway towards formation of the polymer particles may have different substrate specificities, form different reaction products or block branches in the metabolic pathway in order to exert a purposeful influence on the substrates and molecules involved in the formation of the polymer particles.

In order to allow production of the polymer particles in microorganisms, such as the mutants of the genus *E. coli* described in Example 4 , which have a modified fatty acid metabolism, all that is required, apart from the at least one inducible gene which codes for a protein which controls the size of the polymer particles, is the polymer synthase. Depending on which organism is used, further genes may be introduced into the cell in order to enable production of the polymer particles under the stated conditions. If a cell does not contain all the genes required for formation of the polymer particles, production of the polymer particles may nevertheless proceed if the intermediates produced by the missing proteins are supplied to the cell via the nutrient medium. At least one polymer synthase is, however, always required for formation of the polymer particles.

The properties of the polymer particles may be influenced by controlling the composition thereof. By influencing their properties, it is, for example, possible to influence the rate of biodegradability of the polymer particles. Apart from the above-stated possibility of purposefully selecting the further genes introduced into the cell which code for a protein involved in the formation of the polymer particles, it is particularly preferred for the purposes of influencing the composition of the polymer particles formed *in vivo* to introduce into the cell at least one additional gene which codes for a thiolase and/or a polymer synthase. The differing substrate specificity of the thiolases and polymer synthases results in different intermediate and final products and thus in a different composition of the formed polymer core of the particle.

The principle underlying the production of these polymer particles is illustrated by way of example in Figure 2. Activated precursors for biosynthesis of the polymers may in principle be derived from the central metabolites acetyl CoA or from intermediates of the primary metabolic pathways the citrate cycle, fatty acid β oxidation and *de novo* fatty acid biosynthesis, and from amino acid metabolism. If fatty acids are used as the carbon source, intermediates (acyl CoA, in particular 3-hydroxyacyl CoA), which serve as activated precursors for PHA biosynthesis, are produced by fatty acid β oxidation.

Particle size is controlled in that the at least one inducible gene which codes for a protein which controls the size of the polymer particles is derived from the family of phasin-like proteins and is preferably selected from the group comprising the phasin gene phaP from *Ralstonia eutropha* and the phasin gene phaF from *Pseudomonas oleovorans*. Phasins are amphiphilic proteins with a molecular weight of 14 to 28 kDa which bind tightly to the hydrophobic surface of the polymer particles.

Polymer particles with a different composition of the polymers forming them exhibit different mechanical properties and release biologically active substances, in particular pharmaceutical active ingredients, at different rates. For example, polymer particles composed of C6-C14 3-hydroxy fatty acids exhibit a higher rate of polymer degradation due to the low crystallinity of the polymer. An increase in the molar ratio of polymer constituents with relatively large side chains on the polymer backbone usually reduces

crystallinity and results in more pronounced elastomeric properties. By controlling polymer composition in accordance with the process described in the invention, it is accordingly possible to influence the biodegradability of the polymer particles and thus also the release rate for biologically active substances, in particular pharmaceutical active ingredients.

At least one fatty acid with functional side groups is preferably introduced into the culture medium as a substrate for the formation of the polymer particles, with at least one hydroxy fatty acid and/or at least one mercapto fatty acid and/or at least one β-amino fatty acid particularly preferably being introduced. "Fatty acids with functional side groups" should be taken to mean saturated or unsaturated fatty acids. These also include fatty acids containing functional side groups which are selected from the group comprising methyl groups, alkyl groups, hydroxyl groups, phenyl groups, sulfhydryl groups, primary, secondary and tertiary amino groups, aldehyde groups, keto groups, ether groups, carboxyl groups, O-ester groups, thioester groups, carboxylic acid amide groups, hemiacetal groups, acetal groups, phosphate monoester groups and phosphate diester groups.

Use of the substrates is determined by the desired composition and the desired properties of the polymer particles, which are influenced both genetically by the use of different genes which code for proteins with different substrate specificity and by the additives, substrates and reaction conditions present in the culture medium which are used.

In order to achieve still more accurate control of the size of the polymer particles formed, the substrate is added to the culture medium in a quantity such that it is sufficient to ensure control of the size of the polymer particles. This yields an additional possibility for exerting still more effective control over particle size.

The microorganism used to form the polymer particles in the process according to the invention is selected from the genera comprising *Ralstonia*, *Acaligenes*, *Pseudomonas* and *Halobiforma*. The microorganism used is preferably selected from the group comprising *Ralstonia eutropha*, *Alcaligenes latus*, *Escherichia coli*, *Pseudomonas fragi*,

Pseudomonas putida, *Pseudomonas oleovorans*, *Pseudomonas aeruginosa*, *Pseudomonas fluorescens*, and *Halobiforma haloterrestris*. This group comprises both microorganisms which are naturally capable of producing biocompatible, biodegradable polymer particles and microorganisms, such as for example *E. coli*, which, due to their genetic makeup, are not capable of so doing. The genes required to enable the latter-stated microorganisms to produce the polymer particles are introduced using the process according to the invention. In principle, any culturable microorganism may be used for the production of polymer particles by means of the above-described process, even if the microorganism cannot produce the substrates required to form the polymer particles due to a different metabolism. In such cases, the necessary substrates are added to the culture medium and are then converted into polymer particles by the proteins which have been expressed by the genes which have been introduced into the cell.

In order to obtain the polymer particles from the cells, the cultured microorganisms are disrupted in per se known manner and the polymer particles are then separated from the cell debris. The size range of the polymer particles obtained in this manner may be narrowed still further using standard methods, such as for example exclusion chromatography or density gradient centrifugation, to select the polymer particles of the desired size.

The shell membrane consisting of proteins and lipids of the polymer particles produced according to the invention may also be modified in order to impart to the particles properties which are more favourable to the transport of the active ingredients in the body. To this end, a lipid layer located on the surface of the polymer particles is separated from the polymer particles obtained in the process according to the invention and is replaced by a lipid layer of a different composition.

When replacing the lipid layer by a lipid layer of a different composition, the properties of the new lipid layer which are of significance are those having an influence on the transport of the polymer particles through a biological membrane. If the lipid layer of the shell membrane matches the lipid layer of the target membrane, better particle uptake may be observed (Fernart, L. et al., J. Pharmacol. Exp. There. 1999, Vol. 291(3), pp. 1017-1022). The lipid layer of the polymer particles may accordingly be removed

and be replaced by a lipid layer of a different composition. This is preferably achieved by acetone extraction or phospholipases or non-denaturing detergents are used. A mixture of the appropriate amphiphilic molecules is then added to the now virtually lipid-free polymer particles in order to obtain a lipid layer of the desired composition. The new lipid layer here preferably consists of a mixture of amphiphilic molecules which are selected from the group comprising phospholipids and ether lipids.

In addition to the lipids, it is also possible to remove the proteins located on the surface of the polymer particles, with the exception of polymer synthase, by using the above-stated detergents and to replace them with functionalised proteins. Naked polymer particles may in this manner be prepared for numerous modification possibilities which are described in greater detail below. Polymer particles obtained from the microorganism *H. haloterrestris* are particularly suitable for this type of modification.

Depending on the subsequent application of the polymer particles, particle size is controlled by the at least one inducible gene in such a manner that the polymer particles formed have a diameter of 10 nm to 3 µm, preferably of 1 nm to 900 nm and particularly preferably of 10 nm to 100 nm. In one particular embodiment of the present invention, this size control may also be achieved by controlling the availability of a substrate in the culture medium or by combining the two mechanisms.

For the purposes of functionalising the polymer particles produced using the process according to the invention, it is necessary for the at least one inducible gene introduced in the process which codes for a protein which controls the size of the polymer particles to comprise a polymer particle binding domain and at least one binding domain, wherein the at least one binding domain is capable of binding a biologically active substance and/or a coupling reagent. It is likewise particularly preferred for the at least one further gene which codes for a protein involved in the formation of the polymer particles to comprise a polymer particle binding domain and at least one binding domain, wherein the at least one binding domain is capable of binding a biologically active substance and/or a coupling reagent.

A "binding domain" designates that part of the protein which is obtained by genetic modification of the at least one inducible gene previously introduced into the cell and/or of the at least one further gene which codes for a protein involved in the formation of the polymer particles. The binding domain capable of binding the biologically active substances and/or a coupling reagent is selected from the group comprising oligopeptides, enzymes, abzymes and non-catalytic proteins. This group particularly preferably comprises FLAG epitopes or at least one cysteine. These groups are also designated functional proteins in the experimental section. The coding sequence for the members stated in this group is inserted in the coding sequence of the at least one inducible gene previously introduced into the cell and/or of the at least one further gene which codes for a protein involved in the formation of the polymer particles. In this manner, proteins are produced which are not only involved in the formation of or the control of the formation of the polymer particles, but, thanks to their binding domain obtained by genetic modification, are also capable of binding biologically active substances and/or coupling reagents. The binding domain may also be obtained after production of the polymer particles by chemically modifying the proteins located on the surface with coupling reagents (*c.f.* Example 8).

A "coupling reagent" for this purpose is an inorganic or organic compound which is suitable for binding to itself a biologically active substance or further coupling reagents on one side and the binding domain on the other side.

This structure makes it possible to produce multifunctional polymer particles which are suitable for transporting biologically active substances. The "polymer particle binding domain" preferably consists of part of a protein which enables it to bind to the hydrophobic surface of the polymer particles. The polymer particle domain which comprises part of a protein bound on the surface of the polymer particle is here selected from the group of proteins which comprises a polymer depolymerase, a polymer regulator, a polymer synthase and a particle size-determining protein. These proteins preferably originate from microorganisms which are capable of forming polymer particles, in particular those from the genera *Ralstonia*, *Alcaligenes* and *Pseudomonas*. The particle size-controlling protein is here preferably derived from the family of

phasin-like proteins and the phasin from *R. eutropha* and *P. oleovorans* is still more preferably used.

A "polymer regulator" for the purposes of the invention is a protein which regulates the transcription of the genes phaA, phaB and phaC involved in the formation of the polymer particles. It is withdrawn from transcription regulation by binding to the particle surface. One example of such a regulator is the phasin repressor (phaR) from *R. eutropha*, which binds to the promoter of a phasin-like gene, the expression product of which regulates the size of polymer particles formed, and prevents the gene from being read. Because the phasin repressor is bound on the surface of the polymer particles formed, this site on the promoter is released and transcription of the underlying gene can begin.

The idea of using the binding domain of a polymer synthase for binding coupling reagents and/or biologically active substances arises from the elevated stability of the bond between the polymer particle binding domain of this protein and the core of the polymer particle. The inventor has surprisingly discovered that this bond cannot be detached from the core of the biodegradable polymer particle either by denaturing reagents, such as for example. sodium dodecyl sulfate (SDS), urea, guanidium hydrochloride or dithiothreitol, nor by using acidic conditions. The polymer synthase derived from *R. eutropha*, *P. oleovorans*, *P. putida* or *P. aeruginosa* is preferably used for this purpose.

It is here a particular advantage of the process according to the invention that the genetic engineering modification of the proteins binding to the surface of the polymer particles does not affect the functionality of the proteins involved in the formation of the polymer particles. For example, the functionality of the polymer synthase is retained if a protein is fused with the N-terminal end thereof, resulting in the production of a binding domain for binding biologically active substances and/or coupling reagents. Should the functionality of a protein nevertheless be impaired by the fusion, this shortcoming may be offset by the presence of a further gene which codes for a protein which performs the same function and is present in an active state.

In this manner, it is possible to ensure a stable bond of the biologically active substances and/or coupling reagents bound to the polymer particles via the binding domain of the proteins, in particular polymer synthase.

During genetic engineering modification of the genes which code for proteins which, once expressed, bind to the particle surface, it is also possible to introduce genes with different modifications into the cell. Once these proteins with their different binding domains have been expressed and the polymer particles have been formed, it is possible in this manner to use the different binding domains to multifunctionalise the particle surface. This process enables straightforward and efficient mass production of functionalised polymer particles.

Coupling reagents are used for the subsequent functionalisation of the proteins bound on the surface of the polymer particles, these coupling reagents preferably being selected from the group comprising bis(2-oxo-3-oxazolydinyl)phosphonic chloride (BOP-Cl), bromotrispyrrolidinophosphonium hexafluorophosphate (PyBroP), benzotriazol-1-yl-oxy-trispyrrolidinophosphonium hexafluorophosphate (PyBOP), n-hydroxysuccinimide biotin, 2-(1H-benzotriazol-1-yl)-1,1,3,3-tetramethyluronium hexafluorophosphate (HBTU), dicyclohexylcarbodiimide, disuccinimidyl carbonate, 1-(3-dimethylaminopropyl)-3-ethylcarbodiimide (EDC), bis(2-oxo-3-oxazolydinyl)phosphine, diisopropylcarbodiimide (DIPC), 2-(1H-benzotrioxazolyl)-1,1,3,3-tetramethyluronium tetrafluoroborate (TBTU), 2-(5-norbornene-2,3-dicarboxyimido)-1,1,3,3-tetramethyluronium tetrafluoroborate (TNTU), para-nitrophenylchloroformate, and O-(n-succinimidyl)-1,1,3,3-tetramethyluronium tetrafluoroborate (TSTU).

The biologically active substances used are preferably pesticides, herbicides, pharmaceutically active substances and proteins.

The pharmaceutically active substances are selected from the group comprising dideoxyinosine, floxuridine, 6-mercaptopurine, doxorubicin, daunorubicin, 1-darubicin, cisplatin, methotrexate, taxol, antibiotics, anticoagulants, germicides, antiarrhythmic

agents and active ingredient precursors and derivatives of the listed groups of active ingredients.

The proteins are preferably selected from the group comprising insulin, calcitonin, ACTH, glucagon, somatostatin, somatotropin, somatomedin, parathyroid hormone, erythropoietin, hypothalamic release factors, prolactin, thyroid-stimulating hormone, endorphins, enkephalins, vasopressins, non-naturally occurring opiates, superoxide dismutase, antibodies, interferons, asparaginase, arginase, arginine deaminase, adenosine deaminase, ribonuclease, trypsin, chymotrypsin and pepsin.

Described is a process for the *in vitro* production of biodegradable polymer particles, wherein the process comprises providing a solution suited to polymer particles formation with at least one substrate, introducing into the solution a protein which is suited to controlling the size of the polymer particles and introducing at least one further protein which is involved in the formation of the polymer particles. The *in vitro* process also offers the advantage that the size of the polymer particles may be controlled as early as during production of the polymer particles and subsequent costly and time-consuming size determination and separation of the polymer particles formed into individual size classes is avoided.

The at least one further protein used here, which is involved in the formation of the polymer particles, is a polymer synthase, wherein this polymer synthase is preferably selected from the group comprising the polymer synthase from *R. eutropha*, *P. oleovorans*, *P. putida* and *P. aeruginosa*.

In contrast with *in vivo* synthesis, an *in vitro* synthesis, in which proteins and enzymes isolated from microorganisms are used in the laboratory, is normally very costly as both the enzymes and, in some cases, also the enzyme substrates must be isolated and purified beforehand. In one particular embodiment of the present invention for the *in vitro* synthesis of the biocompatible, biodegradable polymer particles, there is added to the solution suited to polymer particle formation at least one fatty acid, particularly preferably a β-mercapto fatty acid and/or a β-amino fatty acid and an acyl CoA oxidase or other oxidising and activating enzymes for the formation of the polymer particles.

Using these substrates instead of R/S-3-hydroxy fatty acids and acyl CoA synthetase results in a CoA recycling system, in which the acyl CoA oxidase will activate and oxidise the fatty acid while consuming CoA and hydrolysing ATP. During polymerisation, the polymer synthase eliminates CoA, which may then in turn be used by the acyl CoA oxidase. An appreciable reduction in the costs of this *in vitro* process may be achieved as a consequence.

Another advantage of the *in vitro* process for the production of polymer particles is that the at least one substrate is added to the solution suited to polymer particle formation in such a quantity that it is sufficient to ensure control of the size of the polymer particles.

Moreover, the size of the polymer particles formed may also additionally be controlled by adding the polymer synthase to the solution in a quantity which is sufficient to ensure control of the size of the polymer particles formed. As in the *in vivo* process, the *in vitro* process also comprises further possibilities for controlling polymer particle size still more accurately, so increasing the yield of polymer particles of the desired size and making the process more efficient and cost-effective.

In the *in vitro* process, however, particle size is actually controlled by introducing a protein which controls the size of the polymer particles into the solution, which protein is derived from the family of phasin-like proteins and is preferably selected from the group comprising the phasin from *Ralstonia eutropha* and the phasin from *Pseudomonas oleovorans*.

By selecting the at least one substrate as well the enzymes used, it is also possible, as in the *in vivo* process, to regulate the composition of the polymer and so obtain polymer cores with different properties. When more than one substrate is used, it is for example possible, depending on the type of polymer formed from the enzymes, to obtain polymer particles with a different composition of the polymer core. As already described further above, the polymers produced in this manner impart the most varied properties to the polymer particles.

In the *in vitro* process for the production of the polymer particles, the composition of the lipid layer on the surface of the polymer particle is controlled by adding at least one amphiphilic molecule from the group of phospholipids and ether lipids to the solution (In the absence of additional amphiphilic molecules, the particles obtained in vitro are surrounded only by proteins, which constitutes another type of particles.). In this manner, it is possible, for example, to produce polymer particles with a specific surface charge which is specifically adjusted to the biological membrane which is subsequently to be crossed in the body. The advantage of modifying the lipid layer of the shell membrane in the *in vitro* process is that it is possible to dispense with subsequent modification of the lipid layer, as in the *in vivo* process. Since the amphiphilic molecules, such as for example phospholipids or ether lipids, are already added to the starting solution, a lipid layer of the desired composition is obtained from the outset.

Another aspect of the *in vitro* process, which is also used in the *in vivo* process, is that at least one pharmaceutically active substance is added to the solution suited to polymer particle formation. Said substance is incorporated into the polymer particle during polymer particle synthesis and may subsequently be released into the body by diffusion through the particle matrix or by degradation of the polymer particle. The latter-stated variant has the further advantage in the polymer particles produced by the process according to the invention that the rate of biodegradation of the polymer particles may be regulated by the above-described control of the composition of the polymer core. In this manner, continuous release of the active ingredient over a specific period is possible.

Apart from the incorporation of active ingredients into the growing polymer particles, functionalisation proceeds, as in the *in vivo* process, by selecting at least one of the proteins introduced into the solution suited to polymer particle formation in such a manner that the at least one introduced protein comprises a polymer particle binding domain and at least one binding domain, wherein the at least one binding domain is capable of binding a biologically active substance and/or a coupling reagent. The proteins used in this case for the formation of the polymer particles in the *in vitro* process may be obtained from the *in vivo* process described further above and they then exhibit, depending on the type of production described above, the corresponding

properties. They are then added to the solution suited to polymer particle formation, where they are involved in the formation of the polymer particles. As in the *in vivo* process, the polymer particles produced in this manner may also additionally be modified after the production thereof by subsequent modification with the coupling reagents already described further above or by addition of biologically active substances which bind to the binding domain of the proteins which have been bound to the surface of the polymer particles.

Also described is a polymer particle of defined size, with a surface layer of amphiphilic molecules, into which is introduced at least one protein which is selected from the group comprising a polymer depolymerase, a polymer regulator, a polymer synthase and a particle size-influencing protein, wherein the at least one protein comprises a polymer particle binding domain and a binding domain which is capable of binding a biologically active substance and/or a coupling reagent and which protein, in a preferred embodiment, is produced according to the above-described process.

Thanks to their advantageous properties, the polymer particles of the present invention are particularly suitable for the production of a pharmaceutical preparation, a pesticide or a herbicide, wherein the pharmaceutical preparation is preferably suitable for the treatment of diseases of the central nervous system. The possibilities for modification of the polymer particles described by the present process allow the conditions to be met for passage through the BBB.

The control of particle size, control of the composition of the shell membrane and in particular also the functionalisation of particle surface mean that these biodegradable polymer particles are a suitable transport vehicle for biologically active substances of all kinds and moreover enable targeted transport of the polymer particles to their site of action. Multifunctionalisation makes it possible, for example, simultaneously to bind not only at least one pharmaceutical active substance to the particle surface but also an antibody, the binding specificity of which enables precise guidance to the target site. These and further advantages are explained in greater detail in the following exemplary embodiments.

The term "comprising" as used in this specification and claims means "consisting at least in part of". When interpreting statements in this specification and the claims which include that term, the features, prefaced by that term in each statement or claim, all need to be present but other features can also be present. Related terms such as "comprise" and "comprised" are to be interpreted in the same manner.

In this specification where reference has been made to patent specifications, other external documents, or other sources of information, this is generally for the purpose of providing a context for discussing the features of the invention. Unless specifically stated otherwise, reference to such external documents is not to be construed as an admission that such documents, or such sources of information, in any jurisdiction, are prior art, or form part of the common general knowledge in the art.

Description of the Figures

Figure 1 shows a schematic overview of an *in vivo* produced biodegradable polymer particle and the proteins and lipids which are bound to the surface.
The abbreviations relate to the following proteins:
A: polymer depolymerase
B: phasin (name of the coding gene in *R. eutropha:* phaP, in *P. oleovorans:* phaF)
C: polymer synthase
D: phospholipid
E: polymer regulators (for example phasin repressor (PhaR from *R.* eutropha)

Figure 2 shows an example of synthesis of one of the possible biodegradable polymers in *R. eutropha*. The simple polyhydroxy alkanoate polyhydroxybutyric acid (PHB) is produced in a three-stage process starting from the substrate acetyl CoA. The C4 repeat unit in PHB is β-hydroxybutyric acid. The final step in the synthesis results in the formation of the polymer particle with the polymer synthase bound to the surface thereof.

Figures 3 and 4 show the results from the *in vivo* experiments. Figure 3 here describes the behaviour of polymer particle diameter with enhanced phasin expression (increased quantity of inducer in the solution), while Figure 4 describes the behaviour of the

polymer particle count in the cell with enhanced phasin expression (increased quantity of inducer in the solution).

Figure 5 and 6 show the results of the *in vitro* experiments. Figure 5 describes the influence of the quantity ratio of phasin to polymer synthase on polymer particle diameter, while Figure 6 describes the influence of the ratio of substrate to polymer synthase on polymer particle diameter.

Figure 7 shows the vector pBBad22K (Sukchawalit, R. et al, FEMS Microbiol Lett. 1999, Vol. 181(2), pp. 217-223) which is used as the starting plasmid in the construction of plasmids pBBad-P (which bears the gene phaP from *R. eutropha*) and pBBad-F (which bears the gene phaF from *P. oleovorans*).

Figure 8 shows the vector pBBad-P which bears the gene phaP for the expression of the particle size-determining protein phasin from *R. eutropha* .

Figure 9 shows the plasmid pBHR68 (Spiekermann, P. et al., Arch. Microbiol. 1999, Vol. 171, pp. 73-80) which imparts storage of the polymer (polyhydroxybutyric acid), said plasmid bearing the genes phB_{Re}, $phbA_{Re}$ and $phbC_{Re}$ from *Ralstonia eutropha* which form the biosynthesis operon for the expression of phaA thiolase, phaB ketoacyl reductase and phaC synthase.

Figure 10 shows the plasmid pBHR71 (Langenbach, S. et al., FEMS Microbiol. Lett. 1997, Vol. 150, pp. 303-309) which imparts storage of the polymer (polyhydroxy alkanoates), said plasmid bearing the gene phaC1 for expression of polymer synthase.

Figure 11 shows the vector pBBad-F which bears the gene phaF for expression of the particle size-determining phasin-like protein from *P. oleovorans*.

Figure 12 shows the vector pBHR71-Cys which has been constructed starting from plasmid pBHR71 from Figure 10. In this structure, the gene phaC1, which codes for polymer synthase, bears two cysteine residues on the N-terminus in order to enable

direct binding of biologically active substances or the binding of biologically active substances via coupling reagents.

Figure 13 also shows the vector pBHR71-FLAG, which has likewise been constructed starting from plasmid pBHR71 from Figure 10. In this structure, the gene phaC1, which codes for polymer synthase, bears two FLAG epitopes on the N-terminus. This enables not only *in vivo* incorporation of gene sequences for functional proteins via the SpeI restriction site of the FLAG gene sequence but also the attachment of coupling reagents and/or biologically active substances *in vitro* onto the already expressed polymer synthase. The biologically active substances used, which are bound directly or via coupling reagents, impart binding to the target site (for example cell surface) or biological activity, in particular enzymatic activity.

Example 1: Control of the size of the *in vivo* produced biodegradable polymer particles

Example 1.1: Production of polymer particles in *R. eutropha*

The biodegradable polymer particles are produced using a *"knock-out"* mutant of *Ralstonia eutropha* (formerly *Alcaligenes eutrophus*) York et al. (York, G.M. et al. J. Bacteriol. 2001, Vol. 183, pp. 4217-4226) which exhibits a defect with regard to the gene which codes for expression of the surface protein phasin on the polymer particles (phaP(-)), as a consequence of which the organism is no longer capable of expressing the phasin which is coded by the phaP gene. The biodegradable polymer particles may moreover also be produced using microorganisms which do not contain this gene and the further genes required for the biosynthesis of polymer particles. Possible examples of such microorganisms would be *Escherichia coli*, and *Halobiforma haloterrestris*. One exemplary embodiment uses *Escherichia coli* (see Example 2), which is naturally not capable of producing the biodegradable polymer particles already described above.

The latter-stated microorganisms are then transformed with a vector which, in the case of *R. eutropha* , contains the phaP gene which codes for phasin and contains the promoter sequence. This gene is controlled by an inducible promoter, preferably a BAD promoter which is induced by arabinose.

Cloning steps

Cloning is performed using the DNA sequence from *R. eutropha* which codes for phaP, this sequence being listed in the "GenBank" under number AF079155 (Hanley, S.Z. et al., FEBS Letters 1999, Vol. 447, pp. 99-105). This sequence is inserted in the NcoI/BamHI restriction site of the vector pBBad22K (Sukchawalit, R. et al., FEMS Microbiol Lett. 1999, Vol. 181(2), pp. 217-223, *c.f.* Figure 7) which contains an inducible promoter (P_{BAD}). For the purposes of cloning, the corresponding NcoI and BamHI restriction sites are inserted into the sequence which codes for phasin by PCR mutagenesis by means of the primers 5'-aaaggcc**ccatgg**tcctcaccccggaaca-3' (SEQ ID No.1, NcoI restriction site in bold) and 5'-aaaggcc**ggatcc**tcagggcactaccttcatcg-3' (SEQ ID No.2, BamHI restriction site in bold). The resultant DNA fragment of SEQ ID No. 3 is hydrolysed with the restriction enzymes NcoI and BamHI and ligated into the likewise hydrolysed vector pBBad22K (Figure 7). The plasmid, which now contains the nucleotide sequences for the protein phasin, is hereinafter designated pBBad-P (Figure 8). This plasmid is transferred into the microorganism by means of the transformation techniques described in the prior art for the corresponding organism. The inducer of the BAD promoter of pBBad-P is arabinose, which makes it possible to achieve quantitative control of the expression of this gene in the microorganisms used. This control mechanism is used to control the expression of the phasin gene and, by means of the quantity of phasin present in the cell, ultimately to control the size of the polymer particles.

Example 1.2: Production of microorganisms for the production of polymer particles which were not originally capable of forming polymer particles

In microorganisms which were not originally capable of producing polymer particles of this type, an additional or the same vector contains further genes which are involved in the formation of the polymer particles. The number and type of the necessary genes which must be introduced into a microbial organism for production of the polymer particles are determined by the core proteome of the organism used. In the simplest case, for example in *E. coli* , at least one thiolase, a reductase and a polymer synthase are necessary in order to produce polymer particles in the manner shown in Figure 2. If organisms are used which comprise specific mutations, such as for example the *E. coli* strain from Example 4, fewer than the above-stated genes are sufficient to enable

production of the polymer particles. The same applies if the organism is already supplied with certain precursor substrates from the metabolic pathway for polymer synthesis.

If a further plasmid is used to introduce into the cell the genes necessary for formation of the polymer particles, the plasmid pBHR68 (Figure 9), which contains the 5.2 kb SmaI/EcoRI fragment from the chromosomal DNA of *R. eutropha*, is particularly suitable. This plasmid contains the biosynthesis operon for the production of biodegradable polymer particles from *R. eutropha* (Spiekermann, P. et al., Arch. Microbiol. 1999, Vol. 171, pp. 73-80).

Example 1.3: Control of particle size by means of the pBBad-P vector

The microorganisms modified in this manner are incubated at 30°C in Luria Broth medium. Induction of the promoter with arabinose proceeds in the late logarithmic growth phase. 24 h after incubation, the size of the particles in the cells is determined. To this end, the cells are separated from the nutrient medium by centrifugation and disrupted. Microscopic determination of polymer particle size is then performed by tunnelling electron microscopy in combination with analytical gel filtration chromatography. The polymer particle count is determined by investigating the intact cells under a light microscope and counting the number of the polymer particles per cell.

As is demonstrated by the results, by controlling expression it is possible to control both the average size of the polymer particles formed (*c.f.* Figure 3) and the average number thereof (*c.f.* Figure 4) in the individual cells. Increasing the copy number of phasin in a controlled manner brings about a decrease in the average polymer particle diameter with a simultaneous increase in the average number of polymer particles. In this manner, it is possible bring about a distinct increase in the yield of polymer particles with the desired diameter. This makes production of the polymer particles quicker and more cost-effective.

Example 1.3: Control of particle size by substrate availability

Another mechanism for controlling the size of the polymer particles is to regulate the availability of substrates or of the polymer synthase which are required for synthesis of the polymer particles. Availability of the polymer synthase may be controlled by means of antisense methods, by genetic regulation or by the availability of the substrates which are required for formation of the polymer synthase in the nutrient medium.

One example of particle size regulation by the availability of a substrate in the nutrient medium is an experiment in which the concentration of the carbon source present is reduced in order consequently to control the diameter of the polymer particles formed. To this end, an *E. coli* strain which contains the above-stated plasmid pBHR68 (Figure 9) with the biosynthesis operon for the production of biodegradable polymer particles from *R. eutropha* (Figure 8), is grown in M9 medium with 1.5% (w/v) glucose at 30°C. At the beginning of the stationary growth phase, the concentration of the carbon source is reduced to 1/50th of its original value by adding M9 medium without glucose and the microorganisms are incubated under otherwise constant growth conditions for a further 20 h. On completion of the test, the microorganisms contain polymer particles with a diameter of on average 130 nm.

Example 2: Influencing the composition of the *in vivo* produced polymer particles

The composition of the biodegradable polymer particle may be modified by introducing into the cell, in addition to the gene coding for phaP, further genes which code for enzymes which provide substrates for the synthase, in particular thiolases or further polymer synthases. As a result, during formation of the polymer particles, the cell is capable of incorporating a series of different monomers into the growing polymer chain and so produce polymer particles having a core which consists of differently composed polymers.

Examples which may be mentioned in this connection are various polymer synthases which, due to their substrate specificity, incorporate 3-hydroxy fatty acids (C4-C16) differently into the growing polymer chain in both the *in vivo* and the *in vitro* process. It is, for example, possible to use the polymer synthase from *R. eutropha*, which produces biodegradable polymer chains from C4 fatty acids (C4), polymer synthases from

Aeromonas punctata (C4 and C6), *Thiocapsa pfennigii* (C4 and C8) and *P. aeruginosa* (C6 to C14). Simultaneously introducing two or more polymer synthases into the cell also makes it possible to produce polymer particles with the most varied properties. In a continuous batch, new polymer synthases with modified substrate specificity may also be obtained by random mutagenesis (*in vitro* evolution).

The composition of the polymer particles formed may also be influenced in other manners. Providing different carbon sources, precursors of different carbon sources and intermediates of the metabolic pathway leading to formation of the polymer particles likewise have an impact on the nature of the polymer formed. Moreover, the metabolic pathways involved in the formation of the polymer particles may be controlled by inhibitors, the use of *knockout* mutants of the metabolic pathway in question and the expression of enzymes which result in the metabolic pathway in the formation of other intermediate or final products.

The following enzymes are used to influence fatty acid metabolism, these enzymes having the property of modifying the intermediates of fatty acid metabolism and providing different products, for example. fatty acids with different side chains, for the formation of the polymer particles: (R)-specific enoyl-CoA hydratases, transacylases and ketoacyl-CoA/ACP reductases. These enzymes have a different specificity with regard to the chain length of the (R)-hydroxyacyl-CoA substrate provided for the polymer synthase. As a consequence, building blocks with differing side chain lengths are obtained in the polymer, resulting in polymers of different compositions.

However, introducing additional enzymes into the cell is not the only way of influencing the metabolic pathway - inhibitors of the metabolic pathway may also be added to the medium. Examples of such inhibitors are acrylic acid and triclosan (synonyms: TCC or 5-chloro-2-(2,4-dichlorophenoxy)phenol), to name but a few.

Example 3: *In vivo* production of biodegradable polymer particles from (R)-3-hydroxybutyric acid with a highly crystalline core

This experiment was performed using an *E. coli* strain which contains the plasmid pBBad-P and the plasmid pBHR68 already stated in Example 1. Culturing was

performed under the conditions stated in Example 1 with glucose as the carbon source. The polymer particles formed consist of (R)-3-hydroxybutyric acid and have a diameter of 50 to 500 nm, depending on how much inducer (arabinose) is added to the medium. The average size of the polymer particles formed in this manner here varies by only approx. 20 to 50 nm. In order to clarify the control characteristics achieved by the phasin introduced by means of the pBBad-P plasmid, the experiment is repeated with the above-stated *E. coli* strain without the pBBad-P plasmid. In this case, polymer particles with a diameter of 150-250 nm are obtained. In comparison with the control experiment, controlling particle size by means of the phasin enables the production of much larger but above all also much smaller polymer particles.

Further fractionation of the polymer particles by size may then proceed by processes known in the prior art, such as for example, exclusion chromatography, density gradient centrifugation, or ultrafiltration in 5 mM phosphate buffer (pH 7.5).

Example 4: *In vivo* production of biodegradable polymer particles from (R)-3-hydroxy fatty acids with a core of low crystallinity

In this experiment, biodegradable, elastomeric polymer particles having a core of (R)-3-hydroxy fatty acids are produced in *E. coli*. The polymer chains are on average made up of 6 to 14 carbon atoms. This is achieved by regulated expression of the phaF gene from *P. oleovorans* in *E. coli* , which is very similar to the phasin-coding gene phaP from *R. eutropha* and to the gene for polymer synthase from *P. aeruginosa* (phaC). In this Example, the expression product of the gene phaF is used to control particle size. The polymer particles may be produced in *E. coli* cells with modified fatty acid metabolism solely by the polymer synthase (phaC) of the pseudomonads (Langenbach, S. et al., FEMS Microbiol. Lett. 1997, Vol. 150, pp. 303-309). Fatty acid metabolism is modified in such a manner that, when fatty acids are used as the carbon source, CoA-activated intermediates (for example enoyl-CoA) of fatty acid β oxidation are accumulated which in turn act as precursors for polymer synthesis. To this end, fadB mutants of *E. coli* are used (Langenbach, S. et al., FEMS Microbiol. Lett. 1997, Vol. 150, pp. 303-309) or inhibitors are used which correspondingly inhibit fatty acid metabolism (for example acrylic acid; Qi, Q. et al., FEMS Microbiol. Lett. 1998, Vol. 167, pp. 89-94). The otherwise required ketoacyl reductase and the thiolase are no longer necessary when

using the mutants or inhibited microorganisms. Enoyl-CoA hydratases intrinsic to *E. coli* then catalyse the formation of R-3-hydroxyacyl-CoA from enoyl-CoA. This is then converted by the polymer synthase into poly-(R)-3-hydroxy fatty acid, which forms the polymer core of the polymer particle formed.

Cloning

The phaF gene of *P. oleovorans*, which has already been described by Prieto et al. (Prieto, M.A. et al., J. Bacteriol. 1999, Vol. 181(3), pp. 858-868) and is listed in the "GenBank" database under number AJ010393, is transformed into *E. coli* using vector pBBad-F (Figure 11). This vector is obtained on the basis of vector pBBad22K (Figure 7). The individual cloning steps here correspond to those described in Example 1. The phaF gene is cloned into the NcoI/BamHI restriction site of the vector pBBad22K. The primers used for PCR mutagenesis of the above-stated phaF gene have the following sequences: 5'-aaagggc**catgg**ctggcaagaagaattccgagaa-3' (SEQ ID No. 4, NcoI restriction site in bold) and 5'-aaaggg**ggatcc**tcagatcagggtaccggtgcctgtctg-3' (SEQ ID No. 5, BamHI restriction site in bold). The resultant DNA fragment of SEQ ID No. 6 is then cloned into the above-described plasmid pBBad22K. The plasmid which now contains the sequence for phaF is hereinafter designated pBBad-F (Figure 11). In addition to this plasmid, the plasmid pBHR71 containing the nucleotide sequence for polymer synthase (Figure 10; Langenbach, S. et al., FEMS Microbiol. Lett. 1997, Vol. 150, pp. 303-309) is also transformed into *E. coli*.

The carbon source used for the subsequent expression is the fatty acid decanoic acid. Expression of the phasin phaF gene is again controlled by means of the inducer arabinose. The test is here performed as already described in Example 1. Depending on the quantity of inducer previously used, the polymer particles formed in this way have a diameter of 100-500 nm.

Example 4: Control of the size of the *in vitro* produced biodegradable polymer particles

The size of the biodegradable polymer particles formed is also controlled in *in vitro* production by the availability of the polymer synthase, of phasin or phasin-like proteins and the availability of the substrates and metabolic intermediates.

The necessary enzymes and substrates must first be made available for the *in vitro* batch for the production of biodegradable polymer particles. The recombinant polymer synthase from *R. eutropha* or *P. aeruginosa* is used for this test (Gerngross, T.U. and Martin, D.P., Proc. Natl. Acad. Sci. USA 1995, Vol. 92, pp. 6279-6283; Qi, Q. et al., Appl. Microbiol. Biotechnol. 2000, Vol. 54, pp. 37-43), which are purified by affinity chromatography (with His-Tag fusion or Ni^{2+}-NTA agarose). The polymer particle size-determining proteins from *R. eutropha* and *P. aeruginosa* , for controlling particle size, are purified in a similar manner. In order to express this protein in *R. eutropha*, the same vector is used which has already been used in Example 1 to express the phasin gene. The reaction batch for *in vitro* production of the biodegradable polymer particles additionally contains, apart from the polymer synthase and the polymer particle size-determining protein phasin: R-3-hydroxybutyryl-CoA or R-3-hydroxydecanoyl-CoA as substrate for synthesis of the polymer particles, 50 mM phosphate buffer (pH 7.5), 1 mM $MgCl_2$ and 5% glycerol (v/v) for stabilisation. Due to the use of R-3-hydroxybutyryl-CoA or R-3-hydroxydecanoyl-CoA as precursors for the polymer synthase, it is not necessary to use the thiolase and the reductase (*c.f.* Fig. 2).

The results of Fig. 5 show the influence of the quantity ratio of phasin to polymer synthase on polymer particle diameter, while Fig. 6 shows the influence of the ratio of substrate to polymer synthase on polymer particle diameter. These results also show that polymer particle size regulation may be regulated by the quantity of polymer synthase present. Effective regulation of particle size may accordingly be achieved by means of the process according to the invention.

Example 5: Influencing the composition of the *in vitro* produced polymer particles

Example 5.1: Influencing polymer composition

The composition of the biodegradable polymer particle may be modified by adding different substrates to the reaction batch and/or by using in each case different polymer synthases with a different substrate spectrum (*c.f.* Example 2). In this manner, a series of different monomers is incorporated into the growing polymer chain during formation of the polymer particles and polymer particles with a different polymer composition are produced. Due to the use of the polymer synthase from *P. aeruginosa* , R-3-hydroxy

fatty acid building blocks with 6-14 C atoms are incorporated into the growing polymer chain. If, for example, only one substrate is supplied, homopolymeric polymer particles are obtained when using the polymer synthase from *P. aeruginosa*.

Further examples of how the composition of the polymer particles may be modified are stated in Example 2.

Example 5.2: Influencing the membrane composition of the polymer particles

For the purposes of subsequent fusion with the membrane of a cell, into which an active ingredient is to be introduced, for example, it must be possible to control the composition of the phospholipid layer on the surface of biodegradable polymer particles. This regulation is achieved in *in vitro* production (Example 4) of the biodegradable polymer particles right from the provision of the solution suited to polymer particle formation. The membrane can be individually tailored to the corresponding requirements by adding a mixture of different amphiphilic molecules.

Phospholipids are added to the *in vitro* reaction batch. A defined number of molecules of the polymer synthase are now involved in the formation of the polymer particles. As polymerisation proceeds, the polymer particles according to the invention become larger and the polymer synthases on the surface are no longer able completely to shield the surface of the polymer particles. As a result, hydrophobic zones (of polymer) are exposed, to which amphiphilic molecules spontaneously attach themselves.

In the case of *in vivo* produced particles, the phospholipid layer which is already present must first be removed. To this end, acetone extraction is performed or phospholipases or non-denaturing detergents are used to destroy the phospholipid layer. Then, as already described above, a mixture of the appropriate amphiphilic molecules is added to the now virtually lipid-free polymer particles. Negatively charged phospholipids or phosphatidyl choline are preferably used here. In one exemplary embodiment of purposeful control of membrane composition, the polymer particles are suspended in PBS containing 1% octyl glycoside (pH 7.5) and dialysed against an excess of phosphatidyl choline while being stirred. The resultant particle surface is particularly well suited to fusion with brain capillary endothelial cells (BCEC).

Example 5.3: Incorporation of various substances into the growing polymer particles

Uptake of substances into the core of the biodegradable polymer particle has already been qualitatively investigated with the assistance of the lipophilic fluorescent dye Nile Red (Sigma, St. Louis, Mo. USA) or Rhodamine 123 (Spiekermann, P. et al., Arch. Microbiol. 1999, Vol. 171, pp. 73-80). If, during *in vivo* or *in vitro* production, as described above, the fluorescent dye Nile Red is added to the medium or the reaction batch, dyeing of the polymer particles produced can be observed, i.e. dyeing of the polymer particles begins as early as during the synthesis thereof and even before isolation from the cell.

If active ingredients with a pharmaceutical action are added, they may be incorporated into the polymer core of the particle. For example, the nonpolar antitumour agent paclitaxel is added to the reaction batch in an *in vitro* experiment for the production of the polymer particles. The water-insoluble paclitaxel is dissolved and concentrated in the hydrophobic polymer core of the polymer particles. Paclitaxel is here merely added to the solution for the formation of the polymer particles, whereupon the active ingredient is concentrated in the particles. This is verified once the polymer particles have formed by removing them from the reaction batch and investigating the solution for the presence of paclitaxel by means of HPLC. The decrease in paclitaxel concentration in the solution here shows that it has been incorporated into the polymer particles. As the polymer particles biodegrade in the organism, the active ingredient is subsequently released. A reaction batch without polymer particles is used as a control, wherein in this case there is no decrease in the paclitaxel concentration in the solution suited to polymer particle formation.

Example 6: Functionalisation of the particle surface

Biologically active substances can be bound to proteins which are already located on the surface of the polymer particles. All the proteins stated in Figure 1 may be considered. This results in a plurality of *"cross-linking"*-strategies, which enable covalent linkage of the biologically active substance via proteins which are bound to the particle surface.

Example 6.1 Purposeful modification of a surface protein for attachment of a pharmaceutical active ingredient

One example of functionalisation of the polymer particles is the attachment of hydrazone-bound doxorubicin (hydrazone-bound doxorubicin has been described by King, H.D. et al., Biconjugate Chem. 1999, Vol. 10, pp. 279-288) to the polymer synthase PhaC1 from *P. aeruginosa*, which is bound on the surface of the polymer particles and contains two N-terminally inserted cysteine residues. These cysteine residues form the binding domain of the polymer synthase, by means of which the biologically active substances may be bound.

Cloning

The triplets coding for the cysteine residues are cloned by PCR mutagenesis into the gene coding for PhaC1, which is then cloned into the XbaI and BamHI restriction site of the plasmid pBHR71 shown in Figure 10. The restriction sites for XbaI and BamHI are also inserted into the gene in this PCR mutagenesis. The primers used for the PCR mutagenesis of the PhaC1-coding gene have the following sequences: primer for the N-terminus 5'-gggc**tctaga**aataaggagatatacatatgtgttgtaagaacaataacgagctt-3' (SEQ ID No. 7, XbaI restriction site in bold, Cys triplet underlined) and primer for the C-terminus 5'-aaacgc**ggatcc**tttcatcgttcatgca-3' (SEQ ID No. 8, BamHI restriction site in bold). The DNA fragment so obtained of SEQ ID No. 9 is then hydrolysed with XbaI and BamHI and ligated into the similarly hydrolysed plasmid pBHR71. The resultant plasmid, designated pBHR71-Cys (Figure 12), is transformed into *E. coli* by means of known techniques, where it then enables the formation of polymer particles bearing a polymer synthase on the surface thereof, which polymer synthase bears two cysteine residues for the attachment of various substances, in particular pharmaceutical substances.

Example 6.1.1 Attachment of doxorubicin (syn.: hydroxyl daunorubicin)

Hydrazone-mediated *"cross-linking"* may now proceed via these surface-exposed cysteine residues. To this end, the following method is used: 100 mg of the isolated polymer particles (total volume 1 ml) to which the polymer synthase is bound are incubated for 3 h at 37°C in helium-perfused PBS (pH 7.5) and 5 mM dithiothreitol (DTT). This treatment reduces the disulfide bridges in the polymer synthase. The low molecular weight compounds are then removed by 30 minutes' centrifugation at 4°C

and 40,000 x g. The reduced polymer particles are then suspended with 1 ml of PBS buffer (pH 7.5), which contains 10 µmol of the hydrazone-bound doxorubicin, and incubated for 30 min at 4°C. After this period, centrifugation is again performed under the above-stated conditions in order to wash the treated polymer particles. Unbound doxorubicin is detected by subsequent HPLC and the successful attachment to the polymer synthase is verified by the reduced concentration.

Example 6.2 Attachment of biologically active substances, in particular pharmaceutical active ingredients to the binding domain of the polymer particles

Active ingredients may also be bound to the polymer particles by being bound via the binding domain onto the proteins bound to the surface of the polymer particles. To this end, a binding domain must first of all be created. Binding of biologically active substances is achieved by genetic modification of surface-bound proteins of the polymer particles (such as for example polymer depolymerase, phasin or phasin-like proteins, polymer synthase, polymer regulator), such that these proteins form an outwardly directed binding domain by means of which a coupling reagent or a biologically active substance may be bound. On fusion of the above-stated surface proteins of the polymer particles with a protein which enables direct attachment of a coupling reagent or a biologically active substance, it is essential to ensure in this process that, after fusion with the surface protein of the polymer particle, the functionality of both the surface protein and the fused protein is fully retained.

In one exemplary embodiment, a polymer particle with two FLAG epitopes is fused directly onto the N-terminus of the polymer synthase PhaC1 from *P. aeruginosa* . The FLAG epitopes enable the binding of commercial anti-FLAG mAbs (monoclonal antibodies) (Anti-FLAG M2, Sigma-Aldrich) and then of enzyme markers which are intended to prove the successful performance of the process. In this Example, secondary antibody/alkaline phosphatase conjugates (antimouse alkaline phosphatase, Sigma-Aldrich) are used as the enzyme marker. The activity of the alkaline phosphatase on the surface of the polymer particles is then determined photometrically.

Example 6.2.1 Production of a polymer particle with a FLAG-PhaC1 fusion protein

The following oligonucleotides are used for the production of the FLAG-polymer synthase fusion protein: 5'-tatg**actagt**gattataaagatgatgatgataaaca-3' and 5'-tatgtttatcatcatctttataatc**actagt**ca-3' (SEQ ID No. 10 and SEQ ID No. 11, SpeI restriction site in bold, FLAG epitope underlined). In order to obtain double-stranded DNA by hybridisation, the two oligonucleotides are mixed together in equimolar quantity (each 10 µM) and incubated for 30 min at room temperature (RT). The double-stranded DNA formed in this manner codes for the FLAG epitope (DYKDDDDK) and has overhanging ends (TA) which are complementary to the overhanging ends of the NdeI restriction site ($\frac{CA^{\nabla}TATG}{GTAT_{\Delta}AC}$). This DNA fragment is hydrolysed with the restriction enzyme NdeI and cloned into the vector pHBR71 (Figure 10) which has to this end been similarly hydrolysed. The plasmid pBHR71-FLAG (Fig. 13) obtained in this manner contains the gene with SEQ ID No. 12 and imparts expression of a polymer synthase with N-terminal FLAG fusion (this part of the protein now forms the binding domain). Biologically active substances and/or coupling reagents may now be bound by means of this binding domain. The singular SpeI restriction site which is likewise introduced during cloning is additionally available for the insertion of any desired further DNA fragments which code for functional proteins.

The following method is used as an example of the functionality of the above-stated construct. Once the pBHR71-FLAG plasmid has been transformed into *E. coli* strains which already contain the plasmid pBBad-F and exhibit a modified fatty acid metabolism (*c.f.* Example 3), the polymer particles are expressed. The polymer particles are isolated from the cells by disrupting the cells and are washed three times with PBS buffer (pH 7.5). The polymer particles are then incubated for 30 min at RT with monoclonal anti-FLAG antibodies which bind to the FLAG epitopes. The polymer particles are then rewashed, as already described above, and then incubated with secondary alkaline phosphatase conjugate for 30 min at RT in PBS buffer. After the 30 minutes' incubation, these particles are washed in 0.1 M tris-HCl (pH 8.5) and then 2 mg/ml of p-nitrophenyl phosphate are added to the particle suspension as a substrate for the alkaline phosphatase. The activity of the alkaline phosphatase is measured spectrometrically at 410 nm. Polymer particles which contain a polymer synthase

without a binding domain, i.e. without the FLAG epitope, are used as a negative control. Since the added p-nitrophenyl phosphate is not converted in the control, the results of the spectrometric measurements at 410 nm are negative. The mere fact that the polymer particles have actually formed is proof that the incorporation of the FLAG epitope has had no effect on the functionality of the polymer synthase.

This process may also be performed with one of the other above-stated surface proteins of the polymer particles. If two or more surface proteins are simultaneously modified, a plurality of the most varied substances may be bound to the polymer particles, so permitting multifunctionality and making them suitable for many different applications.

While in the previous example a biologically active substance was subsequently bound to the already expressed and formed surface protein of the polymer particle, it is also of course possible to fuse the protein directly with the surface protein and then express it. To this end, the coding sequences of the proteins (for example enzymes) are fused with the C-terminal end of the phaC1 gene of the pBHR71-FLAG plasmid.

DNA fragments are obtained by PCR which in each case code for the protein to be inserted and for the C-terminal fragment of the polymer synthase in order to enable fusion with the phaC1 gene. The two fragments are ligated together by means of a restriction site inserted with overhanging primers. At the 5' end of this hybrid gene, at a distance of 7 nt from the start codon, a ribosomal binding site (GAGGAG) and a restriction site are inserted by means of overhanging primers. If the vector used, such as for example the pBHR71-FLAG vector used in this case, already has a ribosomal binding site, it is no longer necessary additionally to insert a ribosomal binding site. Together with an inserted restriction site at the 3' end of this hybrid gene, purposeful cloning into the expression vector pBHR71-FLAG is now possible. The restriction sites at the 5' and 3' ends of the hybrid gene must be selected such that they do not occur a second time within the hybrid gene, so that cloning into an expression vector can proceed colinearly to the present promoter. In this Example, the lacZ gene from *E. coli* is amplified by means of PCR with primers which contain an SpeI restriction site: 5'-**ggactagt**atgaccatgattacggattcactggc-3' (SEQ ID No.13, SpeI restriction site in bold) and 5'-**ccactagt**tttttgacaccagaccaactggtaatggtagcg-3' (SEQ ID No. 14, SpeI restriction site in

bold). In addition, the stop codon is removed from the sequence of the lacZ gene by using these primers in order to obtain a continuous reading frame. The resultant DNA fragment with SEQ ID No. 15 is cloned directly into the SpeI restriction site of the pHBR71-FLAG plasmid. The fusion protein obtained gave rise to the formation of polymer particles with β-galactosidase activity. The corresponding polymer particles are isolated and β-galactosidase activity is demonstrated under reducing conditions with the substrate o-nitrophenyl-beta-D-galactopyranoside (Calbiochem).

Example 7: Stability of the bond between the surface proteins and the polymer core of the polymer particles

Investigations carried out for the purposes of the invention have revealed that the polymer synthase cannot be detached from the core of the biodegradable polymer particle either by treatment with denaturing reagents, such as for example sodium dodecyl sulfate (SDS), urea, guanidium hydrochloride or dithiothreitol, nor by the use of acidic conditions. This is indicative of the presence of a covalent linkage between the polymer particles and the polymer particle binding domain of the polymer synthase. The elevated stability of the bond enables stable transportation of substances bound to or incorporated into the polymer particles to their target site. The N-terminus fragment of the surface-bound polymer synthase (N-terminus to the beginning of the conserved α/β-hydrolase domain) is extremely variable and may be replaced by functional proteins using genetic engineering methods. In this manner, polymer synthase activity and synthesis of polymer particles are retained (Rehm, B.H.A. et al, Biochem. Biophys. Acta 2002, Vol. 1594, pp. 178-190). As a consequence, surface functionalisation is obtained which exhibits elevated stability. A mixture of different proteins with different binding domains to which the biologically active substances and/or coupling reagents are bound are, if required, applied simultaneously, so giving rise to multifunctionalisation of the polymer particle surface. These proteins are applied *in vitro* by adding the purified proteins with different binding domains to the synthesis batch or *in vivo* by expression of the genes in the corresponding microorganism which in each case code for a protein with a binding domain.

Example 7.1 Further possibilities for modifying the surface proteins of the polymer particles

The C-terminal fragment of the surface protein phasin (PhaP) from *R. eutropha* (amino acid residues from >Ala141) is hydrophilic and may be replaced by functional proteins without preventing anchoring of the phasin via the polymer particle binding domain to the surface of the polymer particles.

This anchoring is based on hydrophobic interactions and is reversible (Hanley, S.Z. et al, FEBS Letters 1999, Vol. 447, pp. 99-105). This C-terminal fragment of the intracellular polymer depolymerases is fused by genetic engineering processes with functional proteins and so enables functionalisation of the surface of the polymer particle by means of subsequent attachment of biologically active substances and/or coupling reagents.

The C-terminus (amino acid residue from >180) of the intracellular polymer depolymerase of *R. eutropha* binds the enzyme to the core of the polymer particles (Saegusa, H. et al., J. Bacteriol. 2001, Vol. 183(1), pp. 94-100). This C-terminal fragment of the intracellular polymer depolymerases is fused by genetic engineering processes with functional proteins and so enables functionalisation of the surface of the polymer particle.

The N-terminus (amino acid residue from <140) of the expression products, bound to the surface of the polymer particles, of the genes *phaI* and *phaF* from *Pseudomonas oleovorans* bind the proteins to the polyester core of the polymer particles (Prieto, M.A. et al., J. Bacteriol. 1999, Vol. 181(3), pp. 858-868). This N-terminal fragment of the expression products of the genes phaF and phaI is fused by genetic engineering processes with a functional protein and the resultant binding domain then enables functionalisation of the surface of the polymer particle by attachment of biologically active substances and/or coupling reagents.

Example 8: Covalent modification of the surface proteins of the polymer particles with a coupling reagent

The proteins shown in Figure 1 on the surface of the biodegradable polymer particles may be treated with specific labelling substances which bind specifically to certain amino acids (for example N-hydroxysuccinimide biotin to lysine). This enables the attachment of biologically active substances, such as for example of biotin by iodoacetamide-mediated linkage to cysteine. Molecules, such as biotin, then effect a further linkage of biologically active substances onto the surface proteins of these polymer particles. These include, for example, avidin or streptavidin which may themselves be bound to enzymes and so permit progressive functionalisation of the surface proteins of the biodegradable polymer particles (Rehm, B.H.A. et al., J. Bacteriol. 1994, Vol. 176, pp. 5639-5647). This functionalisation may proceed by attachment of antibodies or pharmaceutical active ingredients. Molecules with different surface charges may also be attached in order to impart to the polymer particles a specific surface charge which is advantageous for transport through/fusion with certain membranes.

Labelling of lysine residues on the surface proteins of the polymer particles with biotin is achieved by means of n-hydroxysuccinimide biotin. In this experiment, polymer particles which bear the polymer synthase PhaC1 from *P. aeruginosa* are isolated from recombinant *E. coli* which bears the plasmid pHBR71. After isolation, the polymer particles are washed three times in PBS (pH 8.0) washed and n-hydroxysuccinimide biotin (Sigma-Aldrich) is then added to the solution up to a final concentration of 5 mM. The reaction is terminated by washing again after 5 minutes' incubation at 4°C. Detection of the biotin bound to the particle surface is performed with the assistance of the streptavidin-alkaline phosphatase conjugate (Sigma-Aldrich) and o-nitrophenyl phosphate (Calbiochem) as substrate. In the control batch with particles which have not been treated with n-hydroxysuccinimide biotin, the particles do not exhibit alkaline phosphatase activity.

In addition to the examples listed here, there are many other coupling reagents for linking biologically active substances with the assistance of which the surface proteins of the polymer particles produced here may be activated.

WHAT WE CLAIM IS:

1. A process for producing polymer particles having surface-bound proteins, the process comprising:

A) providing a cell comprising one or more genes that code for one or more fusion proteins, the fusion proteins comprising a polymer synthase, and

(a) one or more binding domains capable of binding one or more biologically active substances or one or more coupling reagents or a combination thereof, or

(b) one or more biologically active proteins, or

(c) a combination thereof,

the polymer synthase comprising a polymer particle binding domain; and

B) cultivating the cell in a culture medium so that the cell produces the fusion protein from the one or more genes and the fusion protein produces polymer particles, wherein the polymer particle binding domain of the fusion protein is covalently bound to a polymer particle; and

C) separating the polymer particles from the cultivated cells to produce a composition comprising polymer particles having surface-bound proteins.

2. A process of claim 1, wherein the cell further comprises

(a) one or more genes that code for one or more proteins involved in the formation of polymer particles, the proteins comprising a polymer particle binding domain, or

(b) one or more genes that code for one or more additional fusion proteins, the additional fusion proteins comprising a polymer particle binding domain or a protein involved in the formation of the polymer particles, the protein comprising a polymer particle binding domain, or a combination thereof, and

(i) one or more biologically active proteins or one or more

binding domains capable of binding one or more biologically active substances or one or more coupling reagents or a combination thereof, or

(ii) one or more biologically active proteins and one or more binding domains capable of binding one or more biologically active substances or one or more coupling reagents or a combination thereof, or

(c) a combination thereof.

3. A process according to claim 1 or 2, wherein the cell comprises two or more genes that code for two or more fusion proteins or three or more genes that code for three or more fusion proteins.

4. An *in vitro* process for producing polymer particles having surface-bound proteins, the process comprising

A) providing a solution suited to polymer particle formation comprising at least one substrate;

B) introducing into the solution one or more fusion proteins, the fusion proteins comprising a polymer synthase and

(a) one or more binding domains capable of binding one or more biologically active substances or one or more coupling reagents or a combination thereof, or

(b) one or more biologically active proteins, or

(c) a combination thereof,

the polymer synthase of the fusion protein comprising a polymer particle binding domain, and

C) maintaining the solution suited to polymer particle formation so that the fusion protein produces polymer particles, wherein the polymer particle binding domain of the fusion protein is covalently bound to a polymer particle.

5. A process according to claim 4, further comprising introducing into the solution

(a) one or more proteins involved in the formation of polymer particles, the proteins comprising a polymer particle binding domain, or

(b) one or more additional fusion proteins, the additional fusion proteins comprising a polymer particle binding domain or a protein involved in the formation of the polymer particles, the protein comprising a polymer particle binding domain, or a combination thereof, and

(i) one or more biologically active proteins or one or more binding domains capable of binding one or more biologically active substances or one or more coupling reagents or a combination thereof, or

(ii) one or more biologically active proteins and one or more binding domains capable of binding one or more biologically active substances or one or more coupling reagents or a combination thereof, or

(c) a combination thereof.

6. A process according to claim 4 or 5, wherein the polymer synthase or the fusion protein comprising a polymer synthase is added to the solution in a quantity sufficient to ensure control of the size of the polymer particles.

7. A process according to any one of claims 4 to 6, wherein the solution comprises one or more fatty acids and an acyl CoA oxidase.

8. A process according to any one of claims 4 to 7, wherein one or more amphiphilic molecules from the group of phospholipids and ether lipids is added to the solution to control the composition of a lipid layer on the surface of the polymer particle.

9. A process according to any one of claims 4 to 8, wherein two or more, or three or more different fusion proteins are introduced into the solution.

10. A process according to any one of claims 2 to 9, wherein the protein involved in the formation of polymer particles is selected from the group consisting of a phaA thiolase, a phaB ketoacyl reductase, a polymer depolymerase, a polymer

regulator, a polymer synthase and a particle size-determining protein or a combination thereof.

11. A process according to any one of claims 2 to 10, wherein the polymer particle binding domain comprises the polymer particle binding domain of a protein selected from the group consisting of a polymer depolymerase, a polymer regulator, a polymer synthase and a particle size-determining protein.

12. A process according to any one of claims 1 to 11, wherein the polymer synthase is from a microorganism of the genera *Ralstonia*, *Alcaligenes*, *Pseudomonas*, *Aeromonas* or *Thiocapsa*, or from *Ralstonia eutropha*, *Pseudomonas oleovorans*, *Pseudomonas putida*, *Pseudomonas aeruginosa*, *Aeromonas punctata* or *Thiocapsa pfennigii*.

13. A process according to any one of claims 1 to 12, wherein the culture medium or the solution suited to polymer particle formation comprises at least one fatty acid with a functional side group selected from methyl groups, alkyl groups, hydroxyl groups, phenyl groups, sulfhydryl groups, primary, secondary and tertiary amino groups, aldehyde groups, keto groups, ether groups, carboxyl groups, O-ester groups, thioester groups, carboxylic acid amide groups, hemiacetal groups, acetal groups, phosphate monoester groups and phosphate diester groups, or a mixture of any two or more thereof.

14. A process according to any one of claims 1 to 13, wherein a substrate is added to the culture medium or the solution suited to polymer particle formation in such a quantity that it is sufficient to ensure control of the size of the polymer particles.

15. A process according to any one of claims 1 to 14, wherein a lipid layer located on the surface of the polymer particles is separated from the polymer particles and replaced by a lipid layer of another composition.

16. A process according to any one of claims 1 to 15, wherein the polymer particles have a diameter of 10 nm to 3 µm, 10 nm to 900 nm, 10 nm to 100 nm, 50 nm to 500 nm, or 150 nm to 250 nm.

17. A process according to any one of claims 1 to 16, wherein at least one biologically active substance or at least one dye or a mixture thereof is added to the culture medium or the solution suited to polymer particle formation and incorporated into the particles.

18. A process according to any one of claims 1 to 17, wherein the binding domain or the biologically active protein is selected from the group consisting of one or more oligopeptides, one or more antibodies, one or more enzymes, one or more abzymes, one or more non-catalytic proteins, one or more FLAG epitopes and one or more one cysteine residues, or a combination of any two or more thereof.

19. A process according to any one of claims 1 to 18, further comprising

D) binding a coupling reagent to the binding domain, or

E) binding a coupling reagent to the binding domain, and binding a biologically active substance to the coupling reagent, or

F) binding a biologically active substance to the binding domain, or

G) chemically modifying the polymer synthase or the protein involved in the formation of polymer particles to form at least one binding domain by contacting the polymer synthase or the protein with a coupling reagent, or

H) a combination of any two or more thereof.

20. A process according to any one of claims 1 to 19, wherein the biologically active substance is

(a) able to bind another biologically active substance, or

(b) comprises a pharmaceutically active substance, or

(c) selected from

(i) dideoxyinosine, floxuridine, 6-mercaptopurine, doxorubicin, daunorubicin, 1-darubicin, cisplatin, methotrexate, taxol, antibiotics, anticoagulants, germicides, antiarrhythmic agents and active ingredient precursors or

derivatives thereof, or

(ii) insulin, calcitonin, ACTH, glucagons, somatostatin, somatotropin, somatomedin, parathyroid hormone, erythropoietin, hypothalamic release factors, prolactin, thyroid-stimulating hormone, endophins, enkephalins, vasopressins, non-naturally occurring opiates, superoxide dismutase, antibodies, interferons, asparaginase, arginase, arginine deaminase, adenosine deaminase, ribonuclease, trypsin, chymotrypsin or pepsin.

21. A process according to any one of claims 1 to 20, wherein the biologically active protein is selected from insulin, calcitonin, ACTH, glucagons, somatostatin, somatotropin, somatomedin, parathyroid hormone, erythropoietin, hypothalamic release factors, prolactin, thyroid-stimulating hormone, endophins, enkephalins, vasopressins, non-naturally occurring opiates, superoxide dismutase, antibodies, interferons, asparaginase, arginase, arginine deaminase, adenosine deaminase, ribonuclease, trypsin, chymotrypsin or pepsin.

22. A process according to any one of claims 1 to 21, wherein the coupling reagent is selected from the group consisting of bis(2-oxo-3-oxazolydinyl)phosphonic chloride (BOP-Cl), bromotrispyrrolidinophosphonium hexafluorophosphate (PyBroP), benzotriazol-1-yl-oxy-trispyrrolidinophosphonium hexafluorophosphate (PyBOP), n-hydroxysuccinimide biotin, 2-(1H-benzotriazol-1-yl)-1,1,3,3-tetramethyluronium hexafluorophosphate (HBTU), dicyclohexylcarbodiimide, disuccinimidyl carbonate, 1-(3-dimethylaminopropyl)-3-ethylcarbodiimide (EDC), bis(2-oxo-3-oxazolydinyl)phosphine, diisopropylcarbodiimide (DIPC), 2-(1H-benzotrioxazolyl)-1,1,3,3-tetramethyluronium tetrafluoroborate (TBTU), 2-(5-norbornene-2,3-dicarboxyimido)-1,1,3,3-tetramethyluronium tetrafluoroborate (TNTU), para-nitrophenylchloroformate, and O-(n-succinimidyl)-1,1,3,3-tetramethyluronium tetrafluoroborate (TSTU).

23. A polymer particle comprising a polymer core and one or more surface-bound proteins, the one or more surface-bound proteins comprising

A) one or more fusion proteins comprising (i) a polymer synthase

comprising a particle binding domain and (ii) one or more binding domains capable of binding one or more biologically active substances or one or more coupling reagents or a combination thereof, or

B) one or more fusion proteins comprising (i) a polymer synthase comprising a particle binding domain and (ii) one or more biologically active proteins, or

C) a combination thereof,

wherein the polymer particle binding domain of the fusion protein is covalently bound to the polymer particle

24. A polymer particle of claim 23, wherein the particle further comprises

A) one or more additional fusion proteins comprising (i) a polymer particle binding domain and (ii) one or more binding domains capable of binding one or more biologically active substances or one or more coupling reagents or a combination thereof, or

B) one or more additional fusion proteins comprising (i) a polymer particle binding domain and (ii) one or more biologically active proteins, or

C) one or more additional fusion proteins comprising (i) a protein involved in the formation of polymer particles, the protein comprising a polymer particle binding domain and (ii) one or more binding domains capable of binding one or more biologically active substances or one or more coupling reagents or a combination thereof, or

D) one or more additional fusion proteins comprising (i) a protein involved in the formation of polymer particles, the protein comprising a polymer particle binding domain and (ii) one or more biologically active proteins, or

E) a combination of any two or more of A) to D).

25. A polymer particle according to claim 23 or 24, produced by a process according to any one of claims 1 to 22.

26. A polymer particle according to claims 23, 24 or 25, wherein the protein involved in the formation of polymer particles is selected from the group consisting of a polymer depolymerase, a polymer regulator, a polymer synthase and a particle size-determining protein.

27. A polymer particle according to any one of claims 23 to 26, wherein the polymer particle binding domain comprises the particle binding domain of a protein selected from the group consisting of a polymer depolymerase, a polymer regulator, a polymer synthase and a particle size-determining protein.

28. A polymer particle according to claim 26 or 27, wherein the polymer depolymerase, polymer regulator, polymer synthase or particle size-determining protein is from a microorganism of the genera *Ralstonia, Alcaligenes, Pseudomonas, Aeromonas* or *Thiocapsa,* or from *Ralstonia eutropha, Pseudomonas oleovorans, Pseudomonas putida, Pseudomonas aeruginosa, Aeromonas punctata* or *Thiocapsa pfennigii.*

29. A polymer particle according to any one of claims 23 to 28, wherein the polymer particle has an average diameter of 10 nm to 3 μm, 10 nm to 900 nm, 10 nm to 100 nm, 50 to 500 nm, or 150 to 250 nm.

30. A polymer particle according to any one of claims 23 to 29, wherein the core further comprises (a) at least one biologically active substance, (b) at least one dye or (c) at least one biologically active substance and at least one dye.

31. A polymer particle according to any one of claims 23 to 30, wherein the binding domain or biologically active protein is selected from the group consisting of one or more oligopeptides, one or more antibodies, one or more enzymes, one or more abzymes, one or more non-catalytic proteins, one or more FLAG epitopes and one or more one cysteine residues, or a combination of any two or more thereof.

32. A polymer particle according to any one of claims 23 to 31, wherein a coupling reagent is bound to the binding domain, or a coupling reagent is bound to the binding domain and a biologically active substance is bound to the coupling

reagent, or a biologically active substance is bound to the binding domain, or the protein involved in the formation of polymer particles comprises at least one binding domain comprising a coupling reagent bound to the protein, or the biologically active substance is able to bind another biologically active substance, or the biologically active substance comprises a pharmaceutically active substance .

33. A composition, pharmaceutical preparation, pesticide or herbicide comprising polymer particles produced by a process according to any one of claims 1 to 22 or comprising a polymer particle according to any one of claims 23 to 32.

34. A method of binding a biologically active substance comprising

A) providing a composition of polymer particles produced by a method according to any of one of claims 1 to 22 or providing a polymer particle according to any one of claims 23 to 32, wherein optionally a coupling reagent is bound to the binding domain when a binding domain is present, and

B) contacting the composition or polymer particle with a sample comprising a biologically active substance so that the binding domain or the biologically active protein or the coupling reagent binds the biologically active substance.

35. A method according to claim 34, wherein the biologically active substance is an oligopeptide, enzyme, abzyme, non-catalytic protein or antibody.

Sequence listing

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gcacgcttca tgaccaatcc ggaactgccc gccgagccca aggcctggct ggaacaggcc 1560
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cccggaacct acgtgcatga acgatgaaaa ggatccgcgt tt 1722

<210> 10
<211> 36
<212> DNA
<213> Artificial
<220>
<223> Primer

<400> 10
tatgactagt gattataaag atgatgatga taaaca 36

<210> 11
<211> 36
<212> DNA
<213> Artificial
<220>
<223> Primer

<400> 11
tatgtttatc atcatcatct ttataatcac tagtca 36

<210> 12

<211> 1716

<212> DNA

<213> Artificial

<220>

<223> Sequence contains the DNA coding for PhaC1 from *P. aeruginosa* and the DNA coding for a FLAG epitope

<400> 12
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cttcccaagc aagccgcgga aaacacgctg aacctgaatc cggtgatcgg catccggggc 120

aaggacctgc tcacctccgc gcgcatggtc ctgctccagg cggtgcgcca gccgctgcac 180

agcgccaggc acgtggcgca tttcagcctg gagctgaaga acgtcctgct cggccagtcg 240

gagctacgcc caggcgatga cgaccgacgc ttttccgatc cggcctggag ccagaatcca 300

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agccacagcg acctgtcgcc gcaggacatc agtcgtggcc agttcgtcat caacctgctg 420

accgaggcga tgtcgccgac caacagcctg agcaacccgg cggcggtcaa gcgcttcttc 480

gagaccggcg gcaagagcct gctggacggc ctcggccacc tggccaagga cctggtgaac 540

aacggcggga tgccgagcca ggtggacatg gacgccttcg aggtgggcaa gaacctggcc 600

accaccgagg gcgccgtggt gttccgcaac gacgtgctgg aactgatcca gtaccggccg 660

atcaccgagt cggtgcacga acgcccgctg ctggtggtgc cgccgcagat caacaagttc 720

tacgtcttcg acctgtcgcc ggacaagagc ctggcgcgct tctgcctgcg caacggcgtg 780

cagaccttca tcgtcagttg gcgcaacccg accaagtcgc agcgcgaatg gggcctgacc 840

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gacctcaacc tcctcggcgc ctgctccggc gggatcacca ccgcgaccct ggtcggccac 960

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ttcgaactga atacccaggt cgcgctgttc gccgacgaga agactctgga ggccgccaag 1080

cgtcgttcct accagtccgg cgtgctggag ggcaaggaca tggccaaggt gttcgcctgg 1140

atgcgcccca acgacctgat ctggaactac tgggtcaaca actacctgct cggcaaccag 1200

ccgccggcgt tcgacatcct ctactggaac aacgacacca cgcgcctgcc cgccgcgctg 1260

cacggcgagt tcgtcgaact gttcaagagc aacccgctga accgccccgg cgccctggag 1320

gtctccggca cgcccatcga cctgaagcag gtgacttgcg acttctactg tgtcgccggt 1380

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aagtgcgagt tcatcctctc caacagcggt cacatccaga gcatcctcaa cccaccgggc 1500

aaccccaagg cacgcttcat gaccaatccg gaactgcccg ccgagcccaa ggcctggctg 1560

gaacaggccg gcaagcacgc cgactcgtgg tggttgcact ggcagcaatg gctggccgaa 1620

cgctccggca agacccgcaa ggcgcccgcc agcctgggca acaagaccta tccggccggc 1680

gaagccgcgc ccggaaccta cgtgcatgaa cgatga 1716

<210> 13
<211> 34

<212> DNA

<213> Artificial

<220>

<223> Primer

<400> 13
ggactagtat gaccatgatt acggattcac tggc 34

<210> 14

<211> 41

<212> DNA

<213> Artificial

<220>

<223> Primer

<400> 14
ccactagttt tttgacacca gaccaactgg taatggtagc g 41

<210> 15

<211> 3088

<212> DNA

<213> Artificial

<220>

<223> Sequence contains the lacZ gene from *E. coli*

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gtaatagcga agaggcccgc accgatcgcc cttcccaaca gttgcgcagc ctgaatggcg 180

aatggcgctt tgcctggttt ccggcaccag aagcggtgcc ggaaagctgg ctggagtgcg 240

atcttcctga ggccgatact gtcgtcgtcc cctcaaactg gcagatgcac ggttacgatg 300

cgcccatcta caccaacgtg acctatccca ttacggtcaa tccgccgttt gttcccacgg 360

agaatccgac gggttgttac tcgctcacat ttaatgttga tgaaagctgg ctacaggaag 420

gccagacgcg aattattttt gatggcgtta actcggcgtt tcatctgtgg tgcaacgggc 480

gctgggtcgg ttacggccag gacagtcgtt tgccgtctga atttgacctg agcgcatttt 540

tacgcgccgg agaaaaccgc ctcgcggtga tggtgctgcg ctggagtgac ggcagttatc 600

tggaagatca ggatatgtgg cggatgagcg gcattttccg tgacgtctcg ttgctgcata 660

aaccgactac acaaatcagc gatttccatg ttgccactcg ctttaatgat gatttcagcc 720

gcgctgtact ggaggctgaa gttcagatgt gcggcgagtt gcgtgactac ctacgggtaa 780

cagtttcttt atggcagggt gaaacgcagg tcgccagcgg caccgcgcct ttcggcggtg 840

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cgctggggaa tgaatcaggc cacggcgcta atcacgacgc gctgtatcgc tggatcaaat 1440

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agttggtctg gtgtcaaaaa actagtgg 3088