

Protease formulation for treatment of microbial infections

Technical field

5 The present invention relates to a stable protease formulation for the treatment of microbial infections.

Background

10 Bacterial, fungus, and viral infections can be non-hostile relatively mild short-term infections to aggressive life-threatening infections. Microbial adhesion on the host cell membrane is the most common mode of establishing an infection in the host. Microbial surface protein recognizes cell surface protein that initiates microbial infection via adhesion, internalization, colonization, and secretion of toxic molecules enabling bacterial cell development and growth within the host tissue (Meena et al. 2020). A good strategy to treat and prevent microbial infections could be to reduce or prevent
15 the first step in infection, microbial adhesion.

Surfaces of mammalian bodies that are exposed to the external environment are colonized by microbes, the microbiota. Some of the endogenous microbes can, under certain circumstances, cause disease as well as some of the exogenous pathogens
20 that live in the external environment, or on other animals, can overcome our antimicrobial defences and do us harm if we come into contact with them. A good strategy to convert a pathogenic dysbiotic microbiota to a healthy symbiotic microbiota could be to target biofilm structures and the adhesion mechanism to reduce overgrowth of pathogens and to increase microbial diversity. Several studies have identified
25 differences in the microbes present in diseased skin versus those present in healthy skin (Rosenthal et al. 2011) and indicates that an imbalance of microorganisms, termed dysbiosis, exists in numerous pathologies.

30 The traditional approach for elimination of skin infection in animals has been to attempt to target the offending agent with non-specific antimicrobials such as chlorhexidine. As more information is obtained about the commensal skin microbiome, questions should be raised about whether such a nonspecific approach is ideal (Rosenthal et al). It is likely that e.g. chlorhexidine that are commonly used in the treatment of pyoderma have efficacy against much of the core microbiome and therefore may suppress both
35 pathogenic and beneficial components.

The unrestrained use of antibiotics has led to increasing antibiotic resistance, one of the most urgent health risks of our time, leading to an emerging rise of non-treatable human infections (WHO, 2019). Antibiotics are also commonly found in veterinary
5 treatment of animals, where many of the drugs used are the same ones used in the treatment of bacterial infections in humans, and antibiotic resistance are associated with treatment failures and subsequent poor animal health. Companion animals are also able to acquire and exchange multidrug resistant pathogens with humans, and may serve as a reservoir of antimicrobial resistance for in-contact people (Buckland *et*
10 *al*, 2016).

It has become increasingly important to develop new antimicrobials capable of withstanding the repertoire of bacterial resistance mechanisms. Targeting bacterial virulence properties such as adhesion and colonisation instead of viability is considered
15 a valuable alternative strategy to antibiotic therapy (Asadi *et al*, 2019). Adhesion of the pathogen to host cells or tissue is the first step during bacterial infection and is also the first step to generate colonisation (biofilms). Biofilm-producing bacteria are believed to account for up to 80 percent of all bacterial infections (Bryers, 2008). Biofilms are communities of microorganisms protected by a self-synthesised layer of complex
20 polysaccharides, proteins, lipids, and extracellular DNA. When bacteria act as biofilm, their resistance to both immune system and antibiotics can be increased by up to 1000 times (Gilbert *et al*, 1997). An anti-adhesion therapy would also be less likely to result in the emergence of mutations leading to resistance, due to reduced impact on the level of selective pressure on the bacterial population.

25 Antibiotics are commonly employed systemically to inhibit the growth of pathogenic bacteria and will during treatment act on a range of bacterial species in our commensal microbiota, which is of importance for our health. This can lead to a reorganisation of resident communities, including decreased bacterial diversity and outgrowth by
30 previously minor contributors in gut and skin microbial populations. The human skin and mucosa, for example, is a complex barrier organ made of a symbiotic relationship between microbial communities in constant dialogue with the host's innate and adaptive immune system, a well-controlled but delicate equilibrium, mandatory for a healthy skin. Antibiotic-induced alterations of healthy microbiota can persist for several
35 months or years post-treatment (Shaw *et al*, 2019).

Enzymes, and particularly proteases, are very suitable to reduce or inhibit microbial adhesion, such as bacterial adhesion, and to dissolve the biofilm without killing microbials. Proteases have been used in medicine for several decades and is an
5 established and well-tolerated class of therapeutic agents (Craik *et al*, 2011).

There is an urgent need to develop new highly stable topical antimicrobial compositions, such as antibacterial compositions, that target microbial/bacterial adhesion, which is based on non-toxic compounds, that do not induce resistance and
10 that has no or low impact on patients' natural microbiota.

Summary

The inventors of the present invention have developed a novel formulation. Said formulation is a stable protease formulation capable of preventing and reducing biofilm formation as well as to increase microbial diversity in/on a subject or biological surface.
15 Furthermore, said formulation does not induce microbial resistance and can therefore be used instead of antibiotics. Thus, the formulation described herein carry high potential for clinical use and provides a solution to the challenge of reducing the use of antibiotics.

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Therefore, in one aspect, the disclosure relates to a composition comprising:

- i. 0.01 to 0.2 % w/w protease;
- ii. 50 to 70 % w/w glycerol;
- iii. 0.0 to 0.1 % w/w hyaluronic acid; and
25 iv. 0.01 to 0.2 % w/w salt comprising a divalent cation.

In another aspect, the disclosure relates to said composition for use in medicine.

In yet another aspect, the disclosure relates to a composition comprising

- 30 i. a protease;
- ii. glycerol;
- iii. hyaluronic acid; and
- iv. a divalent cation,

for use in the treatment of conditions selected from the group consisting of microbial
35 infections, dermatological conditions and oral conditions, in and/or on a mammal.

In one aspect, the disclosure relates to a composition comprising

- i. a protease;
- ii. glycerol;
- 5 iii. hyaluronic acid; and
- iv. calcium dichloride,

for cosmetic use, in and/or on a mammal.

In another aspect, the disclosure relates to a method for prevention and/or reduction of
10 biofilm, wherein the method comprises administration of a composition according to
any one of the preceding claims.

In yet another aspect, the disclosure relates to a method for increasing microbial
diversity, wherein the method comprises administration of the said composition.

15 In one aspect, the disclosure relates to a method of manufacturing the composition
described herein above, comprising mixing the components at 25°C, in the following
order:

- i. glycerol
- 20 ii. buffer;
- iii. hyaluronic acid; and
- iv. protease.

25 **Description of Drawings**

Figure 1. Formulation optimisation increases the stability of trypsin in solution.

Plots displaying the effect of various formulations on the remaining activity of trypsin
after storage at (40 ± 2°C/100% RH). Formulation composition (N16) contains glycerol
60 w/w%, CaCl₂ 0.1 w/w%, Collagen 0.3 w/w% and 0.04 w/w% trypsin at pH 8.5.
30 Formulation composition (N15) contains glycerol 50 w/w%, CaCl₂ 0.1 w/w%, Collagen
0 w/w% and 0.04 w/w% trypsin at pH 8.5. The (Buffer) solution contained 0.04 w/w%
trypsin at pH 7.5. Non-linear regression was used to estimate half-life.

Figure 2. Formulation A significantly decreases biofilm formation *in vitro*. Bar
35 plots displaying the effect on preformed biofilms from **(A)** *Pseudomonas aeruginosa*,

(B) *Staphylococcus pseudintermedius* and (C) *Malassezia pachydermatis* following treatment with a negative control formulation (n=12) and the test Formulation A (n=6). Formulation A significantly decreases the number of colony-forming units (CFU)/peg compared to the control formulation. Data is plotted as means with error bars (95 % confidence intervals).

Figure 3. The number of observed species increased after enzyme treatment. Plot displaying the number of species that were observed in the enzyme treated group (n=7) and in one of the dogs having a *Proteus* infection. The control group (n=6) did not receive any treatment at all. In the end of the study, the group with the largest number of observed species were the group treated with enzyme. No significant change in observed species could be seen in the control group. Data is plotted as means with error bars (standard error).

Figure 4. There was a clear correlation between abundance of clinically relevant pathogens and microbial diversity. The abundance of clinically relevant pathogens were plotted against a microbial diversity score, the Shannon index, a quantitative measure that reflects how many different species there are in a dataset, which showed a clear correlation ($r^2=0.8$) between the two parameters. This indicates that a low abundance of clinically relevant pathogens is correlated with a high microbial diversity score.

Figure 5. The abundance of clinical relevant pathogens was significantly decreased in both treatment groups. Plot displaying the change in relative abundance of clinically relevant pathogens from day 0 to day 28 of the study period for all three patient groups. The control group displayed an increase of clinically relevant pathogens, while both treatment groups displayed a decrease. The group treated with enzyme displayed a slightly larger decrease ($p=0.028$) compared to the other treatment group, which received chlorhexidine treatment ($p=0.048$). Data is plotted as means with error bars (standard error).

Figure 6. The alpha diversity was primarily increased in the enzyme group. Plots displaying the alpha diversity, i.e. the average species diversity in the treated area, on (A) day 0 and (B) day 28. At day 28, the mean alpha diversity had increased from 197 to 404 (median; from 53 to 479) in the enzyme treatment group and for the

chlorhexidine group a small non-significant increase from 359 to 392 (median; from 345 to 388) was observed. There was no apparent difference in the control group (mean; from 300 to 312, median; from 300 to 192). Data is plotted as means with error bars (standard error).

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Figure 7. The gingivitis score and periodontal score/grade was decreased in the treatment group (n=5). Plots displaying the decrease in **(A)** gingivitis score and **(B)** periodontal score after treatment with Formulation A. This suggests a decreased gingivitis and in general, an improved dental health among the dogs treated with Formulation A. Data is plotted as means with error bars (standard error).

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Figure 8. There was a prolonged positive effect displayed in the dogs treated with Formulation A. The gingivitis status testing was continued also after the study ended, and the score continued to stay low even more than 60 days post-initiation. This effect appeared to persist for more than 60 days, suggesting that the non-healthy microbiota had been replaced by a healthier microbiota.

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Detailed description

Definitions

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The term "protease" as used herein means an enzyme that cleaves a polypeptide of any length, short or long, at a peptide bond, for example by hydrolysing the peptide bond. The substrate of a protease is thus a polypeptide. The substrate can be a naturally occurring polypeptide or a synthetic polypeptide.

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The term "biofilm" denotes an extracellular matrix in which microorganisms are dispersed and/or form colonies. The biofilm typically is made of polysaccharides and other macromolecules. In addition, in the present disclosure, the phrases "preventing a biofilm" "reducing a biofilm", or "decreasing a biofilm", and like phrases, means the prevention of biofilm growth, reduction in the rate of biofilm growth, partial eradication of existing biofilm, and/or complete eradication of existing biofilm.

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The term "stability" as used herein refers to *in vivo* stability and storage stability (e.g., storage stability at room temperature).

As generally used herein "pharmaceutically acceptable" refers to those compounds, materials, compositions, and/or dosage forms which are, within the scope of sound medical judgment, suitable for use in contact with the tissues, organs, and/or bodily fluids of human beings and animals without excessive toxicity, irritation, allergic response, or other problems or complications commensurate with a reasonable benefit/risk ratio.

The term "microbial infection" refers to the undesired proliferation or presence of invasion of pathogenic microbes in a host organism. This includes the excessive growth of microbes that are normally present in or on the body of a mammal or other organism. More generally, a microbial infection can be any situation in which the presence of a microbial population(s) is damaging to a host mammal. Thus, a microbial infection exists when excessive numbers of a microbial population are present in or on a mammal's body, or when the effects of the presence of a microbial population(s) is damaging the cells or other tissue of a mammal.

The term "microbial diversity" as used herein refers to a measure of how many different types of microbial species there are in a within a particular environment, such as the mouth or on the skin of a mammal, while at the same time taking into account the overall representation of the species relative to other species.

In the context of the present disclosure, the term "w/w" means a "weight/weight". The expression "% w/w" is synonymous to "wt%" or "weight-percent". By way of example, 10 g of a composition comprising 50 % w/w of A comprises 5 g A.

In the context of the present disclosure, phrases such as "a composition comprising X to Y % of A" are taken to mean a composition comprising A in the range of X to Y %, both thresholds included. That is, the composition does not comprise less than X % of A, and the composition does not comprise more than Y % of A, provided X is smaller than Y.

Stable formulation

The present disclosure relates to a stable protease formulation for treatment of microbial infections. Proteases are very suitable in treatments against microbial infections, biofilm, and are non-toxic and well-tolerated as a therapeutic agent. It is also

an advantage with a formulation that do not induce resistance and that can be used instead of antibiotics.

Therefore, in one embodiment, the disclosure relates to a composition comprising:

- 5
- i. 0.01 to 0.2 % w/w protease;
 - ii. 50 to 70 % w/w glycerol;
 - iii. 0.0 to 0.1 % w/w hyaluronic acid; and
 - iv. 0.01 to 0.2 % w/w salt comprising a divalent cation.

10 In one embodiment of the present disclosure, the divalent cation of said composition is calcium. In a specific embodiment of the present disclosure, the composition comprises 0.01 to 0.2 % of a calcium salt. In a specific embodiment of the present disclosure, the composition comprises 0.01 to 0.2 % calcium chloride.

15 In another embodiment, the composition further comprises 0.1 to 0.4 % w/w buffer, e.g. a pH buffer. In one embodiment of the present disclosure, the buffer may be a mixture of two or more buffer systems.

In yet another embodiment, the buffer of said composition is selected from the group consisting of Tris (2-Amino-2-(hydroxymethyl)propane-1,3-diol) and MOPS (3-Morpholinopropane-1-sulfonic acid). A person skilled in the art will appreciate the use
20 of other pharmaceutically acceptable buffers.

In one embodiment, the composition further comprises a monosaccharide. For example, the monosaccharide may be a pentose or a hexose. The monosaccharide
25 may be present in its cyclic form, in its linear form, or as a mixture of both forms. In one embodiment of the present disclosure, the composition comprises 0.0 to 0.2 % w/w monosaccharide, such as 0.01 to 0.2 % w/w monosaccharide. For dermal application, e.g. for the application of a gel, the composition may advantageously comprise a monosaccharide. Thus, in one embodiment, the composition is a dermal composition
30 and comprises a monosaccharide, such as 0.01 to 0.2 % w/w monosaccharide. In another embodiment, the composition comprises essentially no monosaccharide, such as 0.0 % w/w monosaccharide. In another embodiment, the monosaccharide of said composition is mannose. In a specific embodiment of the disclosure, the composition comprises 0.0 to 0.2 % w/w mannose, such as 0.01 to 0.2 % w/w mannose.

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In one embodiment of the disclosure, the composition further comprises collagen. The collagen may be any collagen extracted and purified from tissues of animals and fishes and shellfishes. The extraction and decomposition method or the modification degree of the resulting collagen is not specifically limited. In a specific embodiment of the present disclosure, the collagen may be modified, such as chemically modified. In a specific embodiment, the collagen is gelatin. In one embodiment of the disclosure, the composition comprises 0.0 to 0.3 % w/w collagen, such as 0.01 to 0.3 % w/w collagen. For cosmetic use, the formulation may advantageously comprise collagen. Collagen may improve stability of dermal compositions, and may have further advantageous properties for dermal use. Thus, in one embodiment of the disclosure, the composition comprises collagen, such as 0.01 to 0.3 % collagen. In another embodiment of the disclosure, the composition comprises essentially no collagen, such as 0.0 % w/w collagen.

In one embodiment, the protease of said composition is trypsin. Trypsin is a serine protease which predominantly cleaves peptide chains at the carboxyl side of the amino acids lysine and arginine. In one embodiment of the disclosure, the composition comprises 0.01 to 0.2 % w/w protease, such as 0.01 to 0.2 % w/w trypsin. In a further embodiment of the disclosure, the composition comprises 0.02 to 0.09 % w/w protease, such as 0.02 to 0.09 w/w trypsin. In a specific embodiment of the disclosure, the composition comprises essentially 0.04 % w/w protease, such as 0.04 % w/w protease.

In one embodiment of the disclosure, the composition comprises one or more of a diol, a triol, or a polyol, such as glycerol, propylene glycol, or a sugar alcohol. In one embodiment, the composition comprises a diol. In a specific embodiment, the composition of the disclosure comprises 50 to 80 % w/w of the diol, such as 50 to 75 % w/w, such as 50 to 70 % w/w, such as 50 to 62 % w/w. In a specific embodiment, the composition comprises 50 to 80 % w/w glycerol, such as 50 to 75 % w/w, such as 50 to 70 % w/w, such as 50 to 62 % w/w. In another specific embodiment of the disclosure, the composition comprises essentially 60 % diol, such as glycerol. The role of the diol, triol, and/or polyol is in part to reduce the water activity in the composition. Reducing the water activity in turn deactivates the protease. Deactivation of the protease improves its stability in the composition.

In one embodiment of the disclosure, the composition comprises a glycosaminoglycan, such as hyaluronic acid. As used herein "hyaluronic acid" encompasses hyaluronic acid and its hyaluronate salts, including, but not limited to, sodium hyaluronate, potassium hyaluronate, magnesium hyaluronate and calcium hyaluronate. In a specific
5 embodiment, the composition comprises 0.0 to 0.1 % w/w glycosaminoglycan, such as hyaluronic acid. In another specific embodiment, the composition comprises 0.002 to 0.1 % w/w glycosaminoglycan, such as hyaluronic acid. In another specific embodiment, the composition comprises 0.0 to 0.04 % w/w glycosaminoglycan, such as hyaluronic acid. In another specific
10 embodiment, the composition comprises 0.002 to 0.04 % w/w glycosaminoglycan, such as hyaluronic acid. In another specific embodiment, the composition comprises essentially 0.02 % w/w glycosaminoglycan, such as hyaluronic acid. In one embodiment of the present disclosure, the composition comprises 0.0 to 0.1 % w/w hyaluronic acid, such as 0.002 to 0.1 % w/w.

15 In another embodiment of the disclosure, the composition comprises or consists essentially of:

- i. 0.02 to 0.09 % w/w protease;
- ii. 50 to 62 % w/w glycerol;
- iii. 0.0 to 0.04 % w/w hyaluronic acid;
- 20 iv. 0.01 to 0.1 % w/w CaCl_2 ;
- v. 0.1 to 0.4 % w/w Tris or MOPS;
- vi. 0.0 to 0.2 % w/w mannose; and
- vii. 0.0 to 0.3 % w/w collagen.

25 In another embodiment of the disclosure, said composition comprises or consists essentially of:

- i. 0.04 % w/w protease;
- ii. 60 % w/w glycerol;
- iii. 0.2 % w/w hyaluronic acid;
- 30 iv. 0.01 %w/w calcium dichloride; and
- v. 0.1 % w/w Tris or MOPS.

In another embodiment of the disclosure, said composition comprises or consists essentially of:

- 35 i. 0.04 % w/w protease;

- ii. 60 % w/w glycerol;
- iii. 0.02 % w/w hyaluronic acid;
- iv. 0.01 %w/w calcium dichloride; and
- v. 0.1 % w/w Tris or MOPS.

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In yet another embodiment of the present disclosure, the combination of components of said composition does not exceed 100%.

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In another embodiment, the pH of said composition is between 6.0 to 7.5, such as 6.2 to 7.4. pH of solutions can be assessed in a number of ways. For example, pH can be assessed using a pH electrode or a pH indicator.

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In one embodiment, the composition further comprises water. In a specific embodiment of the disclosure, at least part of the water of the composition originates from the buffer.

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The presence of free water can activate and/or destabilize the protease. Water in the composition may be deactivated by other components, such as for instance glycerol and other alcohols describes herein. Accordingly, in one embodiment, the composition comprises less than 50 % w/w water, such as 20 to 50 % w/w water. To improve the stability of the protease, it is important that the water activity in the composition is low. Upon applying the composition to e.g. a wound, the skin, or in the oral cavity, water from said wound, skin, or oral cavity mixes with the composition, increasing the water activity. This in turn activates the protease.

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In another embodiment, the composition is the form of a spray, a gel, a cream, lotion, an ointment, a foam, or a dental chew.

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In one embodiment, the disclosure relates to said composition for use in medicine. Specifically, one embodiment provides for the disclosed composition for use in medicine.

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In another embodiment, the disclosure provides a composition comprising

- i. a protease;
- ii. glycerol;
- iii. hyaluronic acid; and
- iv. a divalent cation,

for use in the treatment of conditions selected from the group consisting of microbial infections, dermatological conditions and oral conditions, or for cosmetic use, in and/or on a mammal.

5 In one embodiment, the disclosure provides a composition comprising

- i. a protease;
- ii. glycerol;
- iii. hyaluronic acid; and
- iv. a divalent cation,

10 for use in the treatment of a condition selected from the group consisting of microbial infections, dermatological conditions and oral conditions, in a mammal.

In one embodiment of the present disclosure, the microbial infection is selected from the group consisting of bacterial infections, fungal infections, yeast infections and viral infections.

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In another embodiment, the dermatological condition is a skin infection. In yet another embodiment, the dermatological condition is an inflammatory skin disease. In another embodiment, the inflammatory skin disease is selected from the group consisting of atopic dermatitis, psoriasis, rosacea and acne.

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In one embodiment, the oral condition is an oral infection. In one embodiment, the oral condition is a dental infection. In another embodiment, the mammal is a dog, a cat, a horse or a human.

25 In one embodiment, the protease of said composition is trypsin.

In one embodiment of this invention, the divalent cation of said composition is calcium.

In another embodiment, the composition further comprises 0.1 to 0.4 %w/w buffer.

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In yet another embodiment, the buffer of said composition buffer is selected from the group consisting of Tris and MOPS.

In one embodiment, the composition further comprises a monosaccharide.

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In another embodiment, the monosaccharide of said composition is mannose.

In one embodiment of this invention, the composition further comprises collagen.

5 In another embodiment, the pH of said composition is between 6.0 to 7.5, such as 6.2 to 7.4.

In one embodiment, the composition further comprises water.

10 In another embodiment, the composition is the form of a spray, a gel, a cream, lotion, an ointment, a foam, or a dental chew.

In one embodiment of the present disclosure, the composition is for topical use.

15 In another embodiment, the present disclosure relates to a method for prevention and/or reduction of biofilm, wherein the method comprises administration of a composition according to any one of the preceding claims.

In one embodiment, the biofilm is a biofilm in or on a mammal.

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In another embodiment, the biofilm is a biofilm of a biological surface.

In one embodiment, the present disclosure relates to a method for increasing microbial diversity, wherein the method comprises administration of the said composition.

25 It is known in the art that the skin barrier and skin microbiota is essential for protection of the body against external aggressions, and that factors that alter the composition of skin microbiota and the skin barrier function inducing a state of dysbiosis (unbalanced state) which has been evidenced in some chronic inflammatory skin diseases such as atopic dermatitis, psoriasis, rosacea or acne (Schommer *et al*, 2013). Hence,
30 compositions that stimulate the skin microbiome and microbial diversity could be a great use in the treatment of such conditions, and for cosmetic purposes.

Furthermore, the stable formulation of the present disclosure were found to have similar positive effect on reduction of clinically relevant pathogens as chlorhexidine.

35 However, it is known in the art that chlorhexidine can cause allergic reactions

(especially so in eczema patients), that cross-resistance to antibiotics have been reported and that it has limited effect against fungi and mycobacteria. Furthermore, chlorhexidine for dental use can cause side effects such as staining of teeth, increased calculus formation and negative effects on the oral (healthy) microbiome. Therefore, a person skilled in the art will appreciate that the use of the formulation of the present disclosure would be beneficial over chlorhexidine.

In another embodiment, the disclosure relates to a method of manufacturing the said composition comprising mixing the components at 25°C, in the following order:

- i. glycerol
- ii. buffer;
- iii. hyaluronic acid; and
- iv. protease.

One embodiment of the present disclosure provides for a method of treating a condition selected from the group consisting of microbial infections, dermatological conditions and oral conditions, in and/or on a mammal, said method comprising administering a composition comprising:

- i. a protease;
- ii. glycerol;
- iii. hyaluronic acid; and
- iv. a divalent cation,

to a subject in need thereof.

Examples

Example 1: Stability experiments

Methods

A fractional factorial design of 5 parameters, and their concentrations, was used to study the most important factors influencing trypsin stability (Table 1).

Table 1. The components that were tested when studying the stability of trypsin.

Formulation Variable	Settings	Units
Glycerol	50 – 62	wt %
CaCl ₂	0.01 – 0.1	wt%
pH	6 – 8.5	pH
Enzyme	0.02 – 0.09	wt%
Collagen	0 – 0.3	wt%

Various combinations were mixed and the pH was adjusted using MOPS buffer. Two different formulation were tested: Formulation composition N16 which contained glycerol 60 w/w%, CaCl₂ 0.1 w/w%, Collagen 0.3 w/w% and 0.04 w/w% trypsin at pH 8.5, and Formulation composition N15 containing glycerol 50 w/w%, CaCl₂ 0.1 w/w%, Collagen 0 w/w% and 0.04 w/w% trypsin at pH 8.5. A buffer solution was used as a control, containing 0.04 w/w% trypsin at pH 7.5. The test formulation, Formulation A, was also evaluated under same conditions.

The formulation mixes were put into sealed polypropylene bottles. Products from each formulation composition were placed on stability at a stressed condition (40 ± 2°C/100% RH) for up to 411 days. Samples were pulled at predetermined time points and analysed directly by measuring the enzyme activity. The trypsin activity was measured at 30°C using Z-Gly-Pro-Arg-pNA as substrate in a spectrophotometer at 405 nm. The decrease in absorbance per minute was used as activity parameter. The half-life or remaining activity relative to time point zero was determined and used to describe the stability. Non-linear regression was used to estimate half-life.

Results

The half-life for trypsin increased from 2 days in buffer to more than 400 days in the most stable formulations. The half-life could not be accurately determined in the most stable formulations since more than 88% of the enzyme activity still remained after 411 days. The remaining activity after storage at 40 °C for 411 days varied from 20 to 88% for the tested combinations (Figure 1). Formulation A was also very stable with a remaining activity of >90% after storage at 40 °C for 119 days. Its shelf life at room temperature 25-30 °C was estimated to be more than 3 years.

Conclusion

It is possible to dramatically improve the stability of trypsin in solution at stressed conditions of 40 °C. The expected shelf life at room temperature of optimised formulations is expected to be more than 3 years.

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Example 2: Anti-biofilm activity

The aim of this study was to evaluate the test formulation, Formulation A, whose stability was tested in Example 1, against preformed biofilms as well as against biofilm formation using the MBEC™ Assay by following the ASTM E2799-12 standard. The
10 ASTM E2799-12 is a standard test method for testing disinfectant efficacy against *Pseudomonas aeruginosa*, *Staphylococcus pseudintermedius* and *Malassezia pachydermatis* biofilm using the MBEC assay.

Methods

15 A mature biofilm was established on pegs under batch conditions with very low shear produced by gentle rotation of the device on an orbital shaker. At the end of 24-48 hours of growth, the pegs containing the biofilm were rinsed to remove planktonic cells and the peg lid was placed in a receiver plate. The wells in the receiver plate were filled according to an experimental design that contained the appropriate sterility, growth,
20 and neutralizer controls as well as the disinfectants. In this study, the disinfectants were a formulation described in Table 2 and a negative control, which was a carrier solution with no active agent (Table 3). After a specified contact time, the peg lid was placed in a receiver plate containing neutralizer, and the entire device was placed in a sonicator to remove the biofilm and disaggregate the clumps. Samples from each well
25 were then diluted, plated and the viable cells counted. The log reduction in viable cells was calculated by subtracting the mean log density for the treated biofilm from the mean log density for the untreated controls. The results were measured in colony-forming units (CFU)/peg.

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Table 2. Composition of Formulation A (w/w%)

Component	Conc. (w/w%)
Glycerol	60
Tris	0.1
CaCl ₂ x 2H ₂ O	0.01
Water	39
Hyaluronic acid	0.02
Trypsin	0.04
pH	7.5

Table 3. Summary of experimental settings used in Example 2.

Sample	Sample description	Contact time	Test dilutions %	Sample size
Formulation A	Test material	4 hours for killing the biofilm	33% of supplied stock solution	n=6
carrier solution (no active agent)	CAMHB – Negative Control	4 hours for killing the biofilm	N/A	n=6

5 CAMHB: Cation-adjusted Mueller-Hinton broth

Results

The number of CFU/peg were significantly decreased for all microbes analysed in the study; *Pseudomonas aeruginosa*, *Staphylococcus pseudintermedius* and *Malassezia*
 10 *pachydermatis*, after treatment Formulation A compared to the control formulation (Figure 2).

Conclusion

Formulation A successfully decreased the biofilm formed by tested microbes.

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Example 3: Microbial diversity in the nasal folds of French bulldogs (clinical study)

The aim of this clinical study was to compare two different solutions on the effect of the natural microbiota in skin folds of dogs.

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Methods

The study was a randomized, single-blind and prospective study, including 19 privately owned dogs (French bulldogs). The two solutions tested were Formulation A (enzyme) with 0.002 w/w% denatonium benzoate added and a 2% chlorhexidine solution.

- 5 Denatonium benzoate is a bittering agent, which was added to Formulation A to prevent dogs from licking on the wound or treated area. The included dogs were randomised into three different groups; one control group that did not receive any treatment (n=6) and two treatment groups; enzyme treatment group (n=7) and a chlorhexidine treatment group (n=6).

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The following inclusion criteria were used:

- i. Tentatively diagnosed with nasomaxillary fold dermatitis (intertrigo) based on history and clinical examination
- ii. Lack of concurrent systemic illness or metabolic conditions, unless the condition is stable in the opinion of the primary investigator
- 15 iii. Dog not pregnant or lactating
- iv. Signed informed animal owner consent

The following exclusion criteria were used:

- 20 i. Clinical lesions consistent with deep pyoderma in nasal folds (such as fistula, phlegmon and furuncle)
- ii. Ulcerations in nose folds
- iii. Dogs that did not allow being touched on the nose folds
- iv. Any other form of topical treatment in the nose folds or any form of antibiotic treatment two weeks prior to or during the study period
- 25 v. Changes in ongoing treatment with oral corticosteroids from two weeks prior to the inclusion and throughout the study period
- vi. Injections of corticosteroids from 90 days prior to inclusion and throughout the study period

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At day 0, the dog underwent general clinical examination. Cytology samples were taken from bilateral nasomaxillary folds. Swabs were collected from the dogs' nasomaxillary folds and sent for next generation DNA sequencing (MiDOG, CA, USA) according to manufacturer's instructions. On day 0, the pet owners were also informed of how to apply the test solution (enzyme treatment solution or chlorhexidine) on each

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nose fold twice daily (morning and evening) until the return visit to the clinic, except for the day of the visit. Two bottles of test product were provided to the owners, and they were instructed to store the test product at room temperature.

5 At day 14, the same tests and assessments as on day 0 were performed. Possible side effects were also noted at this visit. The pet owners were informed about continuing the treatment according to previous introductions another week and that treatment should not take place the day of the return visit and to return all spray bottles to the clinic at next visit.

10 At day 28, the same tests and assessments were performed as on previous visits to the clinic. Again, possible side effects were recorded. The weight of the test solution bottles is recorded. The pet owners were instructed to discontinue all topical product application and reassess patient in two weeks.

15 At day 42 (14 days after treatment was discontinued), the last collection of test samples and assessments were performed.

15 Results

20 All dogs showed a low diversity at both bacterial and fungal phylum level, but high bacterial diversity and low fungal diversity at the species level. The main skin bacterial phyla inhabiting the skin fold of the 19 dogs were *Firmicutes*, *Actinobacteria* and *Proteobacteria*. The main skin fungal phyla were *Ascomycota* and *Basidiomycota*. The topical treatment increased the diversity of bacterial and fungal compositions over time, shown as an increase in microbial diversity score (Figure 3). For the enzyme treatment group (Formulation A) the increase was 38%, for the chlorhexidine group 11% and for the control group it was < 5%. A clear correlation ($r^2=0.8$) between abundance of clinically relevant pathogens and microbial diversity was seen, indicating that a low abundance of clinically relevant pathogens is correlated with a high microbial diversity score, the Shannon index, a quantitative measure that reflects how many different species there are in a dataset (Figure 4). In both treatment groups the abundance of clinically relevant pathogens decreased significantly (enzyme treatment group; $p=0.028$, chlorhexidine group; $p=0.048$) compared to the control (Figure 5). The alpha diversity, i.e. the average species diversity in the treated area increased primarily for the enzyme treatment group from day 0 (Figure 6A) to day 28 (Figure 6B).

Conclusion

Based on the results of this study, we conclude that topical therapy using Formulation A both reduces the relative abundance of clinically relevant pathogens and increases microbial diversity in the nasal folds of French bulldogs. This indicates that formulation A could be an alternative to chlorhexidine.

Example 4: Dental health of dogs (clinical study)

The aim of this study was to evaluate the treatment effect of Formulation A in order to improve the dental status of dogs.

Methods

This clinical study was an open label prospective study on 5 privately owned dogs. The following inclusion criteria were used:

- i. An OraStrip® Dental Diagnostic test score (Gingivitis Score) ≥ 3 (meaning a risk of $\geq 97\%$ of a moderate to severe active periodontal disease)
- ii. Good to excellent health
- iii. Free of disease or infection
- iv. Recent dental > 6-8 weeks ago
- v. Signed informed animal owner consent

The OraStrip® Dental Diagnostic test (CET OraStrip, Virbac, Ft. Worth, TX) is a test to detect periodontal disease and to assess the risk of moderate to severe periodontal disease that can be performed at the clinic. The diagnostic test determines level of thiol, an organosulfur compound produced by the bacteria associated with periodontal disease and as a measure of level of gingivitis where 0 is no thiol detected (no gingivitis) and 1–5 indicates active and increasing gingivitis.

The following exclusion criteria were used in the study:

- i. Multiple tooth extractions
- ii. Major dental procedures
- iii. Daily medication
- iv. Any other form of dental health treatment during the study period
- v. Non-compliance with the study protocol
- vi. Side effects of treatment, after recording of data. Possible side effects were handled by the physician

At day 0, the dog underwent a relevant clinical examination at the clinic. The clinical examination included a dental status evaluation using the OraStrip® Dental Diagnostic test (on saliva) according to the manufacturer's instructions. A periodontal scoring was also performed in order to assess the degree of gingivitis and periodontitis (Table 4).

- 5 Photos were taken of the dogs' teeth and gum. The pet owners were informed of how the Formulation A should be applied twice daily (morning and evening) until the return visit, and supplied with an instructions document. Each owner was supplied with one bottle of Formulation A, and informed that it should be stored at room temperature.

10 **Table 4.** Periodontal scoring of the dogs.

PD Score / Stage	Classification	Clinical manifestation
0	Clinically normal	No gingival inflammation or periodontitis clinically evident
1	Slight gingivitis	Gingivitis only without attachment loss. The height and architecture of the alveolar margin are normal
2	Early periodontitis	≤ 25% of attachment loss at most. There is a stage 1 furcation involvement in multi-rooted teeth and early radiologic signs of periodontitis. The loss of periodontal attachment is less than 25%*
3	Moderate periodontitis	25-50% of attachment loss*, or a stage 2 furcation involvement in multi-rooted teeth
4	Advanced periodontitis	≥ 50% of attachment loss*, or a stage 3 furcation involvement in multi-rooted teeth

* As measured either by probing of the clinical attachment level, or radiographic determination of the distance of the alveolar margin from the cemento-enamel junction relative to the length of the root

- 15 At day 14, the periodontal grading and the OraStrip® Dental Diagnostic test were performed and recorded in the same way as at day 0. Again, a photo was taken of the dogs' teeth and gum, and the owner is informed about continuing the treatment according to previous instructions.

- 20 At day 28, the same tests and grading were performed as on previous visits. At this visit, possible side effects were also noted and the weight of the bottle was recorded.

Results

At day 28 of the treatment period, the OraStrip® test shows the level of gingivitis had decreased for the group treated with Formulation A, indicating a decreased risk of a more severe periodontal disease (Figure 7A). The periodontal score had also
5 decreased for the dogs treated with Formulation A, suggesting an improved dental health among the treated dogs (Figure 7B). The level of gingivitis was also tested after the treatment period ended, and the score continued to stay low even more than 60 days after treatment stop (Figure 8).

Conclusion

Formulation A appears to increase the dental health of the dogs, with decreased gingivitis and periodontal score. Furthermore, this effect appears to persist for more than 60 days. This suggests that the non-healthy microbiota has been replaced by a healthier microbiota.

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