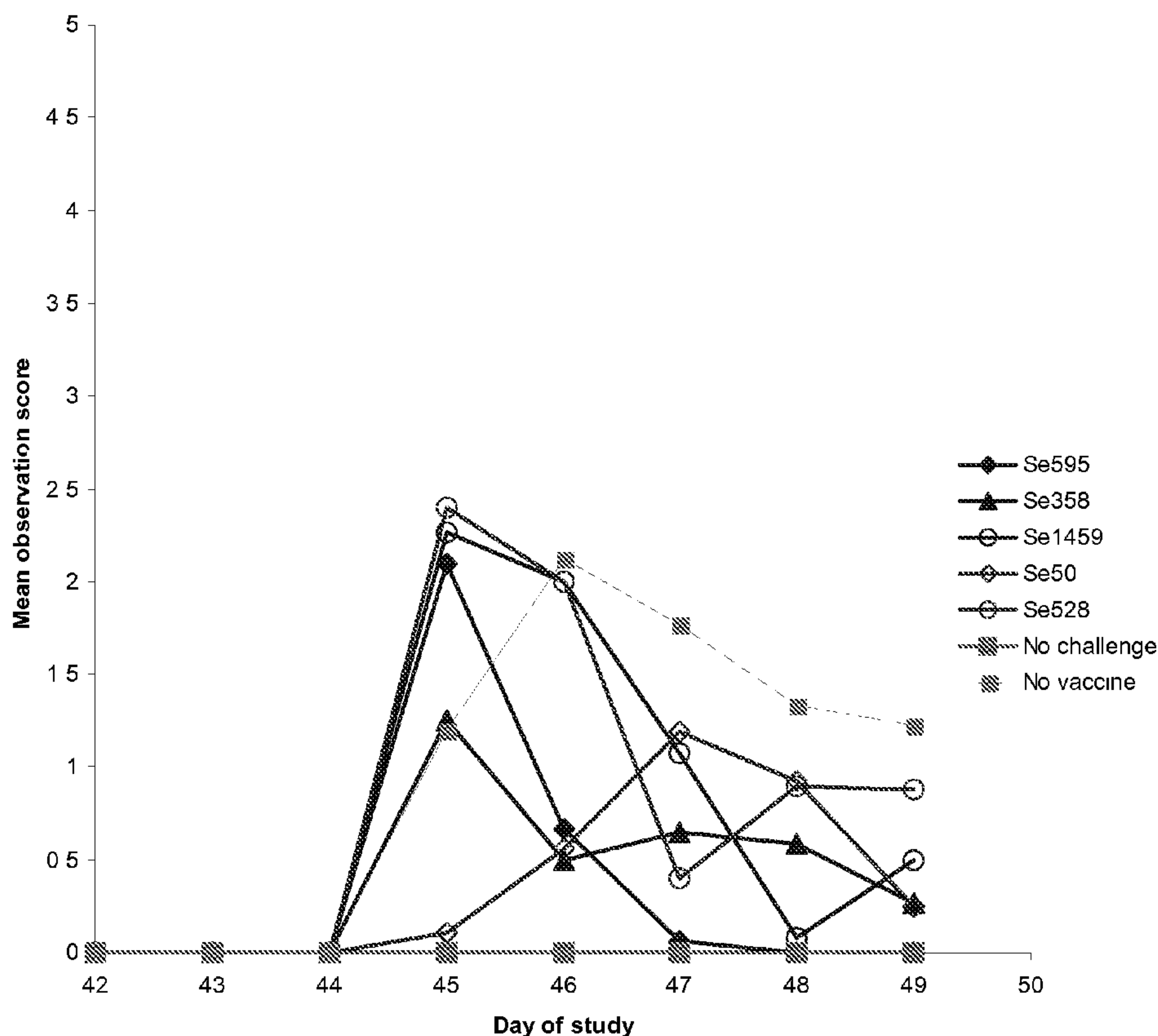




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(57) **Abrégé/Abstract:**

The invention relates to subunit immunogenic or vaccine compositions which may comprise at least one polypeptide of *Streptococcus equi* and methods for preparing and/or formulating such compositions. The invention also relates to the use of such subunit compositions, such as a method for eliciting an immunogenic response or a protective immune response, which may comprise administering the composition to a mammal susceptible to streptococcal infection.

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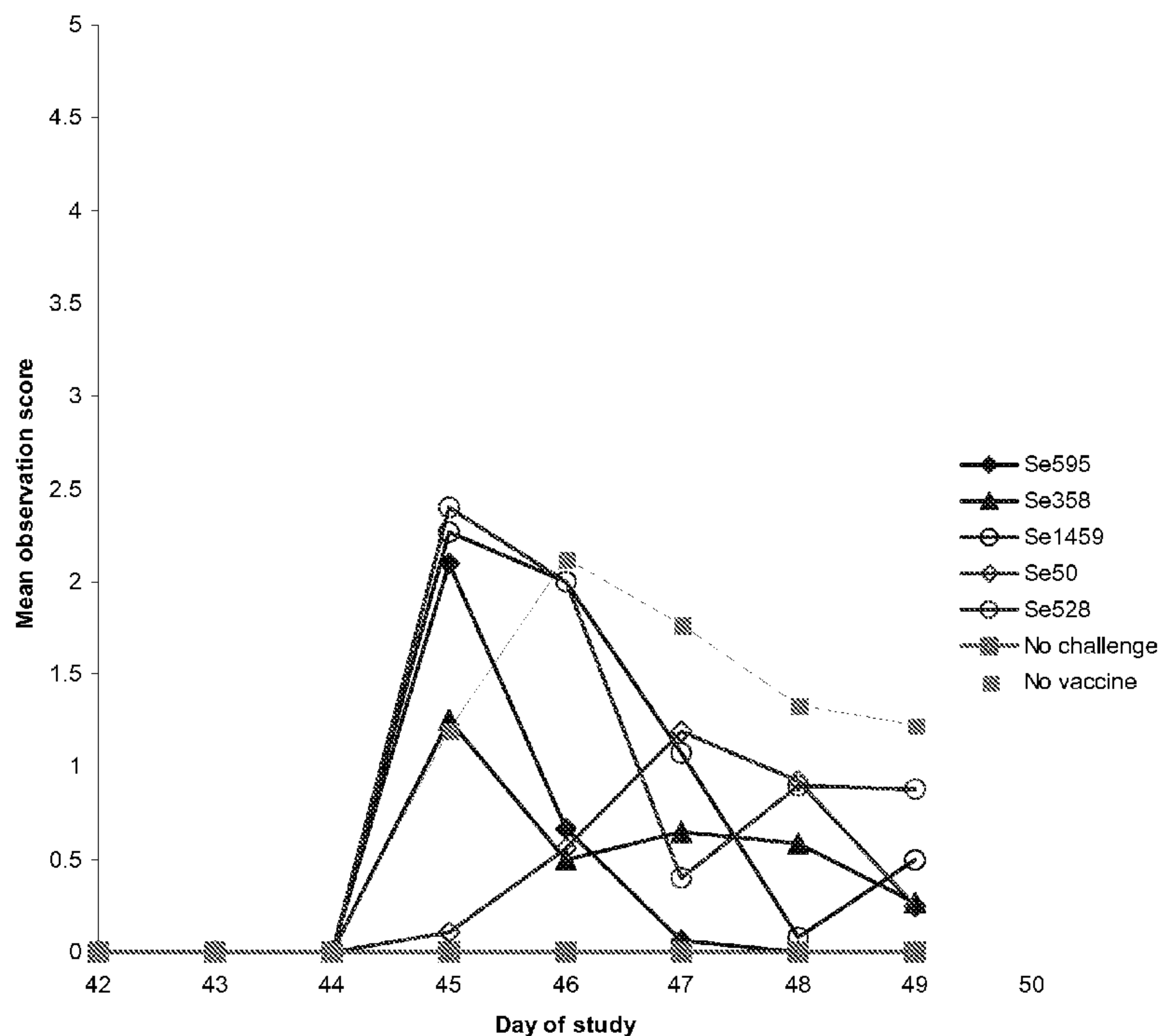
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(57) Abstract: The invention relates to subunit immunogenic or vaccine compositions which may comprise at least one polypeptide of *Streptococcus equi* and methods for preparing and/or formulating such compositions. The invention also relates to the use of such subunit compositions, such as a method for eliciting an immunogenic response or a protective immune response, which may comprise administering the composition to a mammal susceptible to streptococcal infection.

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TITLE OF THE INVENTION

VACCINE AGAINST STREPTOCOCCI

FIELD OF THE INVENTION

5 The present invention relates to polypeptides of streptococcal bacteria, more particularly of *Streptococcus equi*, which may be used to prevent, diagnose and/or treat streptococcal infection.

BACKGROUND OF THE INVENTION

10 *Streptococcus equi* (*S. equi* subsp. *equi*) is a gram-positive coccus that has a large capsule. *S. equi* is the causative agent of strangles in equine, canine and Camelids, notably in horses, donkeys, mules, dogs, camels and dromedaries. Strangles is a highly contagious infection of the upper respiratory tract and associated lymph nodes. In equine, the disease is characterized by fever, thick mucopurulent nasal discharge, swelling and abscessation of the submandibular, submaxillary, and retropharyngeal lymph nodes. In severe cases, the infection may become disseminated with internal and external abscess formation and may ultimately
15 result in death. It can occur in susceptible horses of any age. Transmission occurs through direct contact or indirectly through fomites that have been contaminated by respiratory tract secretions or saliva. Although rare, *S. equi* is a very serious zoonotic pathogen of Camelids (Sechi L A *et al.*, New Microbiol, 1999, 22(4): 383-387), of humans infected by contact with horses (Elsayed S *et al.*, Clin Microbiol Infect, 2003, 9(8): 869-872). In human, mortality was
20 high (25%), especially among older patients and patients with endocarditis, meningitis and disseminated infection (Bradley S F *et al.*, Rev Infect Dis, 1991, 13(2): 270-280; Popescu G A *et al.*, South Med J, 2006, 99(2): 190-191).

After infection, *S. equi* generally has an incubation period 3 to 14 days before the onset of clinical signs. The organism rapidly translocates the mucosa into the lymphatic

channels and from there moves to one or more of the regional lymph nodes leading to inflammation and swelling.

The enlarging lymph nodes soon develop purulent centers with fluctuant liquefaction. Submandibular lymph nodes typically rupture externally through the skin, whilst abscesses
5 formed in the retropharyngeal lymph nodes typically rupture into the guttural pouch. This may take up to 2 weeks to occur after the onset of signs.

The majority (70%) of animals are immune to further attacks of strangles after recovery from a first infection. However, a substantial proportion of animals (30%) can contract the disease a second time, and of these a small proportion can contract the disease
10 more than twice (Todd A.G., J. Comp. Pathol. Ther., 1910, 23, 212-229). Up to 10% of affected animals continue to shed *S. equi* intermittently for prolonged periods after clinical signs have resolved. This carrier state is probably caused by incomplete drainage of exudate from the guttural pouches (empyema) and/or sinuses following rupture of abscesses formed in the retropharyngeal lymph nodes (Newton J R *et al.*, Vet Record, 1997, 140: 84-90).

Drying and hardening of exudate leads to the formation of discrete bodies called chondroids
15 that can remain in the guttural pouch for several years. It is becoming increasingly recognised around the world that sub-clinical persistent carriage of *S. equi* is fundamental to the persistence of this infection between outbreaks (Fintl C *et al.*, Vet Record, 2000, 147: 480-484; Newton J R *et al.*, Vet Record, 1997, 140: 84-90; Newton J R *et al.*, Equine Vet J, 2000,
20 32: 515-526; Timoney J F *et al.*, Vet Record, 1998, 142: 648) and this remains true despite the advent of modern vaccines (Newton R *et al.*, Vet Record, 2005, 156: 291-292).

Protection against the disease is primarily mediated by antibodies of the IgA and IgG subclasses locally produced in the nasopharynx.

Serum antibodies to *Streptococcus equi* (*S. equi*) can be measured by a variety of
25 assays including ELISA (Reif *et al.*, Proc. Am. Assoc. Equine Practitioners, 1982, 27, 33-40), the mouse protection test (Bazely P.L., Aust. Vet. J., 1943, 19, 62-85), and the gel diffusion precipitin test.

Commercially available vaccines consisting of heat-inactivated bacterin or M-protein-rich extracts have been widely used in the field. Although effective in stimulating serum
30 bactericidal antibodies, these vaccines apparently do not stimulate a useful level of nasopharyngeal antibody and so the level of protection stimulated in a horse population is disappointing. An avirulent, genetically modified strain of *S. equi* administered intranasally has recently been shown to stimulate local nasopharyngeal antibodies similar to those found in convalescent immune horses. Vaccinated horses were immune to subsequent experimental

challenge with virulent *S. equi* (Timoney and Galan, 1985, The protective response of the horse to an avirulent strain of Streptococcal equi. In Y. Kimura, S. Kotami and Y. Shokawa, eds., Recent Advances in Streptococci and Streptococcal disease. Reedbooks, Surrey, England, p. 294-295). It therefore appears that an effective strangles vaccine must be able to stimulate a nasopharyngeal immune response.

Bacterin-type vaccines often cause undesirable local and systemic reactions consisting of edema, induration, stiffness, transient fever, and neutrophilia, effects that are probably due to cell wall peptidoglycan. Furthermore, purpura hemorrhagica can develop after vaccination of horses previously sensitised to streptococcal antigens.

Streptococcus zooepidemicus (*S. equi* subsp. *zooepidemicus*) is an important cause of respiratory disease and metritis in equine. *S. zooepidemicus* is an opportunist pathogen that causes purulent respiratory infections of weanling and yearling horses and uterine infections in elderly mares. In the situation of concurrent influenza infection high temperature or transport stress, it can be a devastating and rapid fatal pathogen in the respiratory tract.

Although rare, *S. zooepidemicus* is a very serious zoonotic pathogen of Camelids (Younan M *et al.*, J Vet Med B Infect Dis Vet Public Health, 2005, 52(3): 142-146), of humans infected by contact with horses (Downar J *et al.*, J Clin Microbiol, 2001, 39(6): 2358-2359; Ural O *et al.*, Scand J Infect Dis, 2003, 35(3): 206-207) and of canine animals having respiratory disease, notably already infected by *Bordetella bronchiseptica* (Chalker V J *et al.*, Vet Microbiol, 2003, 95(1-2): 149-156).

S. equi affected mammals may harbor the bacteria for several months in, among other places, the guttural pouches and, thus, shed and act as reservoirs of *S. equi*. Although *S. equi* is quite sensitive to penicillin and other antibiotics, antibiotic treatment is for various reasons mostly ineffective. In order to combat the disease and mitigate serious clinical complications, research has mainly been aimed at developing efficient vaccines.

The development of vaccines has been slow, possibly in part, because vaccines must be able to control infection without inducing a marked inflammatory response.

It is therefore desirable to develop vaccines having a high degree of immunogenicity and which exhibit a good safety profile with limited or no side effects.

Citation or identification of any document in this application is not an admission that such document is available as prior art to the present invention.

SUMMARY OF THE INVENTION

The invention relates to subunit immunogenic or vaccine compositions which may comprise at least one polypeptide of the invention; e.g., polypeptides of *Streptococcus equi*.

The invention therefore further relates to methods for preparing and/or formulating such compositions; e.g., admixing the polypeptides with a suitable veterinary or pharmaceutically acceptable carrier, excipient, diluent or vehicle and/or an adjuvant and/or stabilizer. The invention also relates to the use of such subunit compositions; e.g., a method for eliciting an immunogenic response or a protective immune response, which may comprise administering the composition to a mammal susceptible to streptococcal infection.

The invention relates to recombinant immunogenic or vaccine compositions which may comprise at least one recombinant expression vector encoding at least one polypeptide of the invention, able to express *in vivo* this polypeptide in a mammal susceptible to streptococcal infection. Herein such a immunogenic or vaccine composition which may comprise at least one recombinant expression vector is named a recombinant immunogenic or vaccine composition. The invention therefore further relates to methods for preparing such vectors, e.g., inserting at least one polynucleotide encoding a polypeptide according to the invention into a plasmid vector or viral vector so that the vector contains and expresses the polynucleotide in the host. The invention therefore further relates to methods for formulating such recombinant immunogenic or vaccine compositions; e.g., admixing the vectors with a suitable veterinary or pharmaceutically acceptable carrier, excipient, diluent or vehicle and/or an adjuvant and/or stabilizer. The invention also relates to the use of such recombinant immunogenic or vaccine compositions; e.g., a method for eliciting an immunogenic response or a protective immune response, which may comprise administering the composition to a mammal susceptible to streptococcal infection.

By definition, mammals susceptible to streptococcal infection encompass equine (i.e. horses, mares, foals, donkeys and mules), camelids (i.e. camels, dromedaries), canine (i.e. dogs, bitches, puppies), bovine (i.e. cattle, cows, calves), ovine (i.e. sheeps, ewes, lambs), caprine (i.e. goats), porcine (i.e. pigs, sows, piglets) and human. Mammals susceptible to *Streptococcus equi* infection encompass equine (i.e. horses, mares, foals, donkeys and mules), camelids (i.e. camels, dromedaries), canine (i.e. dogs, bitches, puppies) and human. Mammals susceptible to *Streptococcus zooepidemicus* infection encompass equine (i.e. horses, mares, foals, donkeys and mules), canine (i.e. dogs, bitches, puppies), Camelids (i.e. camels, dromedaries) and human.

The invention also relates to uses recombinant expression vectors; e.g., a method for producing a polypeptide of the invention, which may comprise culturing, growing or propagating prokaryotic or eucaryotic cells transformed to contain the recombinant expression vectors and to express a polynucleotide encoding the polypeptide under conditions

suitable for expression, and harvesting or isolating or separating the polypeptide from cells transformed to express it.

The invention further relates to methods for inducing in an equine, i.e. a horse, a donkey and a mule, an immunogenic or protective immune response against streptococcal bacteria, such as *Streptococcus equi* or *Streptococcus zooepidemicus*, which may comprise administering to the equine a subunit immunogenic or vaccine composition of the invention or a recombinant immunogenic or vaccine composition of the invention.

The invention further relates to methods for inducing in a canine, i.e. a dog, a bitch, a puppy, an immunogenic or protective immune response against streptococcal bacteria, such as *Streptococcus equi* or *Streptococcus zooepidemicus*, which may comprise administering to the canine a subunit immunogenic or vaccine composition of the invention or a recombinant immunogenic or vaccine composition of the invention.

The invention further relates to methods for inducing in a Camelid, i.e. a camel, a dromedary, an immunogenic or protective immune response against streptococcal bacteria, such as *Streptococcus equi* or *Streptococcus zooepidemicus*, which may comprise administering to the Camelid a subunit immunogenic or vaccine composition of the invention or a recombinant immunogenic or vaccine composition of the invention.

The invention further relates to methods for inducing in a human an immunogenic or protective immune response against streptococcal bacteria, such as *Streptococcus equi* or *Streptococcus zooepidemicus*, which may comprise administering to the human a subunit immunogenic or vaccine composition of the invention or a recombinant immunogenic or vaccine composition of the invention.

The invention relates to a polypeptide having an amino acid sequence coded by a polynucleotide selected from the group consisting of polynucleotides identified SEQ ID NOS: 17, 19, 21, 23 and 25. The invention further relates to a polypeptide having an identity which is equal or more than about 70%, 75%, 80%, 85%, 90%, 91%, 92%, 93%, 94%, 95%, 96%, 97%, 98%, or 99% with an amino acid sequence coded by a polynucleotide selected from the group consisting of polynucleotides identified SEQ ID NOS: 17, 19, 21, 23 and 25. The invention relates to a polypeptide selected from the group consisting of polypeptides identified SEQ ID NOS: 18, 20, 22, 24, 26, 48, 50 and 52. The invention further relates to a polypeptide analog having an identity which is equal or more than about 70%, 75%, 80%, 85%, 90%, 91%, 92%, 93%, 94%, 95%, 96%, 97%, 98%, or 99% with a polypeptide selected from the group consisting of polypeptides identified SEQ ID NOS: 18, 20, 22, 24, 26, 48, 50 and 52. The invention further relates to a polypeptide

fragment, containing at least one epitope, which is a part of a polypeptide selected from the group consisting of polypeptides identified SEQ ID NOS: 18, 20, 22, 24, 26, 48, 50 and 52 or analogs thereof, and notably a fragment containing at least one epitope selected from the group consisting of polypeptides identified SEQ ID NOS: 32, 34, 36, 38, 40, 54, 56 and 58.

5 And, the invention relates to uses, compositions and methods involving such polypeptides, analogs thereof or fragments thereof as herein described.

Thus, the invention relates to a recombinant expression vector, wherein said vector contains at least one polynucleotide which codes for a polypeptide selected from the group consisting of polypeptides identified SEQ ID NOS: 18, 20, 22, 24, 26, 48, 50 and 52, and
10 combinations thereof. The invention further relates to a recombinant expression vector, wherein said vector has at least one polynucleotide which codes for a polypeptide analog having an identity which is equal or more than about 70%, 75%, 80%, 85%, 90%, 91%, 92%, 93%, 94%, 95%, 96%, 97%, 98%, or 99% with a polypeptide selected from the group consisting of polypeptides identified SEQ ID NOS: 18, 20, 22, 24, 26, 48, 50 and 52, and
15 combinations thereof. The invention further relates to a recombinant expression vector, wherein said vector has at least one polynucleotide which codes for a polypeptide fragment, containing at least one epitope which is a part of a polypeptide selected from the group consisting of polypeptides identified SEQ ID NOS: 18, 20, 22, 24, 26, 48, 50 and 52 or analogs thereof, and notably a fragment selected from the group consisting of
20 polypeptides identified SEQ ID NOS: 32, 34, 36, 38, 40, 54, 56 and 58. The invention relates to a recombinant expression vector, wherein said vector has at least one polynucleotide selected from the group consisting of polynucleotides identified SEQ ID NOS: 17, 19, 21, 23, 25, 31, 33, 35, 37, 39, 47, 49, 51, 53, 55 and 57, and combinations thereof. The invention further relates to a recombinant expression vector, wherein said
25 vector has at least one polynucleotide having an identity which is equal or more than about 70%, 75%, 80%, 85%, 90%, 91%, 92%, 93%, 94%, 95%, 96%, 97%, 98%, or 99% with a polynucleotide selected from the group consisting of polynucleotides identified SEQ ID NOS: 17, 19, 21, 23, 25, 31, 33, 35, 37, 39, 47, 49, 51, 53, 55 and 57, and combinations thereof. And, the invention relates to uses, compositions and methods involving such
30 vectors as herein described.

Polynucleotides of the invention may be used in hybridization reactions (e.g. Northern or Southern blots, or in nucleic acid microarrays or “gene chips”) or in amplification reactions (e.g. polymerase chain reaction or PCR).

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It should be appreciated that the invention provides polynucleotides which may comprise sequences complementary to those described above.

Polynucleotides according to the invention may be prepared in many ways (e.g. by chemical synthesis, from genomic or cDNA libraries, from the streptococcus itself etc.) and
5 may take various forms (e.g. single stranded, double stranded, primers, probes etc.).

Polynucleotides according to the invention may be labeled e.g. with a radioactive or fluorescent label. This is particularly useful as a primer or as a probe.

In addition, the term "polynucleotide" includes DNA and RNA, and also their analogues, such as those containing modified backbones.

10 Further still, the invention envisions an antibody preparation which may comprise at least one antibody specific to a polypeptide of the invention. This antibody may be polyclonal or monoclonal. The antibody preparation may also comprise a fragment of said antibody.

The invention also involves a diagnostic kit for detecting infection by streptococcal bacteria, including at least one polypeptide according to the invention or an antibody
15 preparation which may comprise at least one antibody specific to a polypeptide of the invention. The invention also involves a diagnostic method for detecting infection by streptococcal bacteria in mammals, which may comprise using a diagnostic kit of the invention and detecting in a sample the polypeptide or an antibody specific to that polypeptide.

20 The invention further concerns a passive immunization method which may comprise administering the antibody preparation to a mammal susceptible to infection by streptococcal bacteria or infected by streptococcal bacteria. In particular, the passive immunization method comprises administering the antibody preparation to an equine susceptible to infection by the streptococcal bacterium or infected by the streptococcal bacterium, notably susceptible to
25 infection by *Streptococcus equi* or infected by *Streptococcus equi*.

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In one aspect, the invention provides an immunogenic composition comprising at least one polypeptide, wherein said polypeptide is a fragment of Se50 protein, said fragment consisting of the amino acid sequence as shown in SEQ ID NO: 32, and a pharmaceutically acceptable excipient, diluent or vehicle.

5 In another aspect, the invention provides an immunogenic composition comprising at least one recombinant vector and at least one polynucleotide inserted therein coding for a polypeptide, wherein said polypeptide is a fragment of Se50 protein, said fragment consisting of the amino acid sequence as shown in SEQ ID NO: 32, and a pharmaceutically acceptable excipient, diluent or vehicle, and said vector is able to express
10 *in vivo* the polypeptide in a mammal susceptible to streptococcal infection caused by *Streptococcus equi*.

In another aspect, the invention provides an immunogenic composition comprising at least one recombinant vector and at least one polynucleotide inserted therein, wherein said polynucleotide consists of the nucleotide sequence as shown in SEQ ID NO: 31
15 or said polynucleotide consists of a nucleotide sequences having a sequence identity which is equal or more than about 95% with SEQ ID NO: 31, and a pharmaceutically acceptable excipient, diluent or vehicle, and said vector is able to express *in vivo* the polypeptide encoded by said polynucleotide in a mammal susceptible to streptococcal infection caused by *Streptococcus equi*, and said encoded peptide elicits an immunogenic response against
20 *Streptococcus equi*.

In another aspect, the invention provides use of at least one polypeptide which is a fragment of Se50 protein, said fragment consisting of the amino acid sequence as shown in SEQ ID NO: 32, in the preparation of a medicament for eliciting an immunogenic response in equine against *Streptococcus equi* infection.

25 In another aspect, the invention provides an antibody preparation comprising at least one antibody specific for the polypeptide as described above, which antibody is polyclonal or monoclonal, or a fragment of said antibody that is specific for said polypeptide, and a pharmaceutically acceptable excipient, diluent or vehicle.

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In another aspect, the invention provides a diagnostic kit for detecting infection by a streptococcal bacterium comprising at least one polypeptide as described above, or the antibody preparation as described above, and a pharmaceutically acceptable excipient, diluent or vehicle.

5 In another aspect, the invention provides a diagnostic method for detecting infection by a streptococcal bacterium, comprising using the diagnostic kit as described above and detecting in a sample the polypeptide or an antibody specific to that polypeptide.

In another aspect, the invention provides use of the antibody preparation as described above in the preparation of a medicament for passive immunization wherein said
10 antibody preparation is formulated for administration to an equine susceptible to infection by *Streptococcus equi* or infected by *Streptococcus equi*.

In another aspect, the invention provides the antibody preparation as described above for use in passive immunization wherein said antibody preparation is formulated for administration to an equine susceptible to infection by *Streptococcus equi* or infected by
15 *Streptococcus equi*.

In another aspect, the invention provides a polypeptide that is a fragment of Se50 protein, said fragment consisting of the amino acid sequence as shown in SEQ ID NO: 32 for use in eliciting an immunogenic response in equine against *Streptococcus equi* infection.

20 In another aspect, the invention provides an immunogenic composition as described above for use in eliciting an immunogenic response in equine against *Streptococcus equi* infection.

It is noted that in this disclosure and particularly in the claims, terms such as “comprises”, “comprised”, “comprising” and the like can have the meaning attributed to it in
25 U.S. Patent law; e.g., they can mean “includes”, “included”, “including”, and the like; and that terms such as “consisting essentially of” and “consists essentially of” have the meaning ascribed to them in U.S. Patent law, e.g., they allow for elements not explicitly recited, but

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exclude elements that are found in the prior art or that affect a basic or novel characteristic of the invention.

These and other embodiments are disclosed or are obvious from and encompassed by, the following Detailed Description.

BRIEF DESCRIPTION OF THE DRAWINGS

The following detailed description, given by way of example, but not intended to limit the invention solely to the specific embodiments described, may best be understood in conjunction with the accompanying drawings, in which:

5 FIG. 1 shows the mean clinical scores of vaccinated and control mice following challenge with *Streptococcus equi*;

 FIG. 2 shows the mean percentage of change in mass of vaccinated and control mice after *Streptococcus equi* challenge;

10 FIG. 3 shows the mean nose dabs score of vaccinated and control mice after *Streptococcus equi* challenge;

 FIG. 4 shows the mean clinical scores of Se1631, Se1681, Sla vaccinated and control mice following challenge with *Streptococcus equi*;

 FIG. 5 shows the mean percentage of change in mass of Se1631, Se1681, Sla vaccinated and control mice after *Streptococcus equi* challenge;

15 FIG. 6 shows the mean nose dabs score of Se1631, Se1681, Sla vaccinated and control mice after *Streptococcus equi* challenge (expressed in colony forming unit (cfu)) and

 FIG. 7 shows the percentage of mice surviving and not losing more than 7% weight of Se1631, Se1681, Sla vaccinated and control mice after *Streptococcus equi* challenge.

DETAILED DESCRIPTION

20 The present invention relates to the identification of polypeptides of *Streptococcus equi* able, when administered to a mammal, to elicit an immunogenic or immune response, and to the identification of polynucleotides encoding these polypeptides. These polypeptides are useful for the production of subunit immunogenic compositions or subunit vaccines. These polynucleotides are useful for the production of DNA immunogenic compositions,
25 DNA vaccines, recombinant viral vector immunogenic compositions or recombinant viral vector vaccines.

 The invention concerns compositions, uses and methods against infections caused by bacteria of the *Streptococcal* family, notably caused by *Streptococcus equi*, e.g. strangles disease in equine, camelid, canine and human, and against infections caused by *Streptococcus*
30 *zooepidemicus* in equine, camelid, canine and human.

 The invention concerns polypeptides, polynucleotides and genes obtained or derived from *Streptococcus equi*. The present invention may relate also to polypeptides, polynucleotides and genes obtained or derived from other streptococci, notably from

Streptococcus zooepidemicus, *Streptococcus suis*, *Streptococcus uberis*, *Streptococcus dysgalactiae*, *Streptococcus agalactiae*, *Streptococcus pneumoniae*, *Streptococcus pyogenes*.

A particular aspect of the invention is a polypeptide selected from the amino acid sequences SEQ ID NOS: 18, 20, 22, 24, 26, 48, 50 and 52.

5 The whole genome of *Streptococcus equi* is available in the Sanger database (http://www.sanger.ac.uk/Projects/S_equi/).

The polypeptides are identified as SEQ ID NOS: 18, 20, 22, 24, 26, 48, 50 and 52 for Se50, Se1459, Se595, Se528, Se358, Se1631, Se1681 and Sla, respectively.

10 A particular aspect of the invention is a polynucleotide selected from the nucleotide sequences SEQ ID NOS: 17, 19, 21, 23 and 25.

The polynucleotides are identified as SEQ ID NOS: 17, 19, 21, 23 and 25 for Se50, Se1459, Se595, Se528 and Se358, respectively.

The practice of the present invention will employ, unless otherwise indicated, conventional techniques of molecular biology, microbiology, recombinant DNA technology, and immunology, which are within the skill of the art. Such techniques are explained fully in the literature. See, e.g., Sambrook et al. (2001) *Molecular Cloning: A Laboratory Manual*, 3rd ed., Cold Spring Harbor Press; *DNA Cloning*, Vols. I and II (D. N. Glover ed. 1985); *Oligonucleotide Synthesis* (M. J. Gait ed. 1984); *Nucleic Acid Hybridization* (B. D. Hames & S. J. Higgins eds. 1984); *Animal Cell Culture* (R. K. Freshney ed. 1986); *Immobilized Cells and Enzymes* (IRL press, 1986); Perbal, B., *A Practical Guide to Molecular Cloning* (1984); the series, *Methods In Enzymology* (S. Colowick and N. Kaplan eds., Academic Press, Inc.); and *Handbook of Experimental Immunology*, Vols. I-IV (D. M. Weir and C. C. Blackwell eds., 1986, Blackwell Scientific Publications).

25 Before describing the present invention in detail, it is to be understood that this invention is not limited to particular DNA, polypeptide sequences or process parameters as such may, of course, vary. It is also to be understood that the terminology used herein is for the purpose of describing particular embodiments of the invention only, and is not intended to be limiting.

30 Unless defined otherwise, all technical and scientific terms used herein have the same meaning as commonly understood by one of ordinary skill in the art to which the invention pertains. Although a number of methods and materials similar or equivalent to those described herein can be used in the practice of the present invention, the preferred materials and methods are described herein.

By the term “complementarity” or “complementary” is meant, for the purposes of the specification or claims, a sufficient number in the oligonucleotide of complementary base pairs in its sequence to interact specifically (hybridize) with a target nucleic acid sequence of the gene polymorphism to be amplified or detected. As known to those skilled in the art, a very high degree of complementarity is needed for specificity and sensitivity involving hybridization, although it need not be 100%. Thus, for example, an oligonucleotide that is identical in nucleotide sequence to an oligonucleotide disclosed herein, except for one base change or substitution, may function equivalently to the disclosed oligonucleotides. A “complementary DNA” or “cDNA” gene includes recombinant genes synthesized by reverse transcription of messenger RNA (“mRNA”).

A “cyclic polymerase-mediated reaction” refers to a biochemical reaction in which a template molecule or a population of template molecules is periodically and repeatedly copied to create a complementary template molecule or complementary template molecules, thereby increasing the number of the template molecules over time.

By the term “detectable moiety” is meant, for the purposes of the specification or claims, a label molecule (isotopic or non-isotopic) which is incorporated indirectly or directly into an oligonucleotide, wherein the label molecule facilitates the detection of the oligonucleotide in which it is incorporated, for example when the oligonucleotide is hybridized to amplified gene polymorphic sequences. Thus, “detectable moiety” is used synonymously with “label molecule”. Synthesis of oligonucleotides can be accomplished by any one of several methods known to those skilled in the art. Label molecules, known to those skilled in the art as being useful for detection, include chemiluminescent, fluorescent or luminescent molecules. Various fluorescent molecules are known in the art which are suitable for use to label a nucleic acid for the method of the present invention. The protocol for such incorporation may vary depending upon the fluorescent molecule used. Such protocols are known in the art for the respective fluorescent molecule.

“DNA amplification” as used herein refers to any process that increases the number of copies of a specific DNA sequence by enzymatically amplifying the nucleic acid sequence. A variety of processes are known. One of the most commonly used is the polymerase chain reaction (PCR) process of Mullis as described in U.S. Pat. Nos. 4,683,195 and 4,683,202. Methods, devices and reagents as described in U.S. Patent Nos. 6,951,726; 6,927,024; 6,924,127; 6,893,863; 6,887,664; 6,881,559; 6,855,522; 6,855,521; 6,849,430; 6,849,404; 6,846,631; 6,844,158; 6,844,155; 6,818,437; 6,818,402; 6,794,177; 6,794,133; 6,790,952; 6,783,940; 6,773,901; 6,770,440; 6,767,724; 6,750,022; 6,744,789; 6,733,999; 6,733,972;

6,703,236; 6,699,713; 6,696,277; 6,664,080; 6,664,064; 6,664,044; RE38,352; 6,650,719;
 6,645,758; 6,645,720; 6,642,000; 6,638,716; 6,632,653; 6,617,107; 6,613,560; 6,610,487;
 6,596,492; 6,586,250; 6,586,233; 6,569,678; 6,569,627; 6,566,103; 6,566,067; 6,566,052;
 6,558,929; 6,558,909; 6,551,783; 6,544,782; 6,537,752; 6,524,830; 6,518,020; 6,514,750;
 5 6,514,706; 6,503,750; 6,503,705; 6,493,640; 6,492,114; 6,485,907; 6,485,903; 6,482,588;
 6,475,729; 6,468,743; 6,465,638; 6,465,637; 6,465,171; 6,448,014; 6,432,646; 6,428,987;
 6,426,215; 6,423,499; 6,410,223; 6,403,341; 6,399,320; 6,395,518; 6,391,559; 6,383,755;
 6,379,932; 6,372,484; 6,368,834; 6,365,375; 6,358,680; 6,355,422; 6,348,336; 6,346,384;
 6,319,673; 6,316,195; 6,316,192; 6,312,930; 6,309,840; 6,309,837; 6,303,343; 6,300,073;
 10 6,300,072; 6,287,781; 6,284,455; 6,277,605; 6,270,977; 6,270,966; 6,268,153; 6,268,143;
 D445,907; 6,261,431; 6,258,570; 6,258,567; 6,258,537; 6,258,529; 6,251,607; 6,248,567;
 6,235,468; 6,232,079; 6,225,093; 6,221,595; D441,091; 6,218,153; 6,207,425; 6,183,999;
 6,183,963; 6,180,372; 6,180,349 ; 6,174,670; 6,153,412; 6,146,834; 6,143,496; 6,140,613;
 6,140,110; 6,103,468; 6,087,097; 6,072,369; 6,068,974; 6,063,563; 6,048,688; 6,046,039;
 15 6,037,129; 6,033,854; 6,031,960; 6,017,699; 6,015,664; 6,015,534; 6,004,747; 6,001,612;
 6,001,572; 5,985,619; 5,976,842; 5,972,602; 5,968,730; 5,958,686; 5,955,274; 5,952,200;
 5,936,968; 5,909,468; 5,905,732; 5,888,740; 5,883,924; 5,876,978; 5,876,977; 5,874,221;
 5,869,318; 5,863,772; 5,863,731; 5,861,251; 5,861,245; 5,858,725; 5,858,718; 5,856,086;
 5,853,991; 5,849,497; 5,837,468; 5,830,663; 5,827,695; 5,827,661; 5,827,657; 5,824,516;
 20 5,824,479; 5,817,797; 5,814,489; 5,814,453; 5,811,296; 5,804,383; 5,800,997; 5,780,271 ;
 5,780,222; 5,776,686; 5,774,497; 5,766,889; 5,759,822; 5,750,347; 5,747,251; 5,741,656;
 5,716,784; 5,712,125; 5,712,090; 5,710,381; 5,705,627; 5,702,884; 5,693,467; 5,691,146;
 5,681,741; 5,674,717; 5,665,572; 5,665,539; 5,656,493; 5,656,461; 5,654,144; 5,652,102;
 5,650,268; 5,643,765; 5,639,871; 5,639,611; 5,639,606; 5,631,128; 5,629,178; 5,627,054;
 25 5,618,703; 5,618,702; 5,614,388; 5,610,017; 5,602,756; 5,599,674; 5,589,333; 5,585,238;
 5,576,197; 5,565,340; 5,565,339; 5,556,774; 5,556,773; 5,538,871; 5,527,898; 5,527,510;
 5,514,568; 5,512,463; 5,512,462; 5,501,947; 5,494,795; 5,491,225; 5,487,993; 5,487,985;
 5,484,699; 5,476,774; 5,475,610; 5,447,839; 5,437,975; 5,436,144; 5,426,026; 5,420,009;
 5,411,876; 5,393,657; 5,389,512; 5,364,790; 5,364,758; 5,340,728; 5,283,171; 5,279,952;
 30 5,254,469; 5,241,363; 5,232,829; 5,231,015; 5,229,297; 5,224,778; 5,219,727; 5,213,961;
 5,198,337; 5,187,060; 5,142,033; 5,091,310; 5,082,780; 5,066,584; 5,023,171 and 5,008,182

may also be employed in the practice of the present invention. PCR involves the use of a
 thermostable DNA polymerase, known sequences as primers, and heating cycles, which
 separate the replicating deoxyribonucleic acid (DNA), strands and exponentially amplify a

gene of interest. Any type of PCR, such as quantitative PCR, RT-PCR, hot start PCR, LAPCR, multiplex PCR, touchdown PCR, etc., may be used. Advantageously, real-time PCR is used. In general, the PCR amplification process involves a cyclic enzymatic chain reaction for preparing exponential quantities of a specific nucleic acid sequence. It requires a small amount of a sequence to initiate the chain reaction and oligonucleotide primers that will hybridize to the sequence. In PCR the primers are annealed to denatured nucleic acid followed by extension with an inducing agent (enzyme) and nucleotides. This results in newly synthesized extension products. Since these newly synthesized sequences become templates for the primers, repeated cycles of denaturing, primer annealing, and extension results in exponential accumulation of the specific sequence being amplified. The extension product of the chain reaction will be a discrete nucleic acid duplex with a termini corresponding to the ends of the specific primers employed.

By the terms “enzymatically amplify” or “amplify” is meant, for the purposes of the specification or claims, DNA amplification, i.e., a process by which nucleic acid sequences are amplified in number. There are several means for enzymatically amplifying nucleic acid sequences. Currently the most commonly used method is the polymerase chain reaction (PCR). Other amplification methods include LCR (ligase chain reaction) which utilizes DNA ligase, and a probe consisting of two halves of a DNA segment that is complementary to the sequence of the DNA to be amplified, enzyme QB replicase and a ribonucleic acid (RNA) sequence template attached to a probe complementary to the DNA to be copied which is used to make a DNA template for exponential production of complementary RNA; strand displacement amplification (SDA); QB replicase amplification (QBR); self-sustained replication (3SR); and NASBA (nucleic acid sequence-based amplification), which can be performed on RNA or DNA as the nucleic acid sequence to be amplified.

A “fragment” of a molecule such as a protein or nucleic acid is meant to refer to any portion of the amino acid or nucleotide genetic sequence.

As used herein, the term “genome” refers to all the genetic material in the chromosomes of a particular organism. Its size is generally given as its total number of base pairs. Within the genome, the term “gene” refers to an ordered sequence of nucleotides located in a particular position on a particular chromosome that encodes a specific functional product (e.g., a protein or RNA molecule). In general, an animal’s genetic characteristics, as defined by the nucleotide sequence of its genome, are known as its “genotype,” while the animal’s physical traits are described as its “phenotype.”

By “hybridization” or “hybridizing,” as used herein, is meant the formation of A-T and C-G base pairs between the nucleotide sequence of a fragment of a segment of a polynucleotide and a complementary nucleotide sequence of an oligonucleotide. By complementary is meant that at the locus of each A, C, G or T (or U in a ribonucleotide) in the fragment sequence, the oligonucleotide sequenced has a T, G, C or A, respectively. The hybridized fragment/ oligonucleotide is called a “duplex.”

A “hybridization complex”, such as in a sandwich assay, means a complex of nucleic acid molecules including at least the target nucleic acid and a sensor probe. It may also include an anchor probe.

A “melting temperature” is meant the temperature at which hybridized duplexes dehybridize and return to their single-stranded state. Likewise, hybridization will not occur in the first place between two oligonucleotides, or, herein, an oligonucleotide and a fragment, at temperatures above the melting temperature of the resulting duplex. It is presently advantageous that the difference in melting point temperatures of oligonucleotide-fragment duplexes of this invention be from about 1°C to about 10°C so as to be readily detectable.

As used herein, the term “nucleic acid molecule” is intended to include DNA molecules (e.g., cDNA or genomic DNA), RNA molecules (e.g., mRNA), analogs of the DNA or RNA generated using nucleotide analogs, and derivatives, fragments and homologs thereof. The nucleic acid molecule can be single-stranded or double-stranded, but advantageously is double-stranded DNA. “DNA” refers to the polymeric form of deoxyribonucleotides (adenine, guanine, thymine, or cytosine) in its either single stranded form, or a double-stranded helix. This term refers only to the primary and secondary structure of the molecule, and does not limit it to any particular tertiary forms. Thus, this term includes double-stranded DNA found, inter alia, in linear DNA molecules (e.g., restriction fragments), viruses, plasmids, and chromosomes. In discussing the structure of particular double-stranded DNA molecules, sequences may be described herein according to the normal convention of giving only the sequence in the 5’ to 3’ direction along the nontranscribed strand of DNA (i.e., the strand having a sequence homologous to the mRNA). An “isolated” nucleic acid molecule is one that is separated from other nucleic acid molecules that are present in the natural source of the nucleic acid.

A “nucleoside” refers to a base linked to a sugar. The base may be adenine (A), guanine (G) (or its substitute, inosine (I)), cytosine (C), or thymine (T) (or its substitute, uracil (U)). The sugar may be ribose (the sugar of a natural nucleotide in RNA) or 2-

deoxyribose (the sugar of a natural nucleotide in DNA). A “nucleotide” refers to a nucleoside linked to a single phosphate group.

As used herein, the term “oligonucleotide” refers to a series of linked nucleotide residues, which oligonucleotide has a sufficient number of nucleotide bases to be used in a PCR reaction. A short oligonucleotide sequence may be based on, or designed from, a genomic or cDNA sequence and is used to amplify, confirm, or reveal the presence of an identical, similar or complementary DNA or RNA in a particular cell or tissue.

Oligonucleotides may be chemically synthesized and may be used as primers or probes.

Oligonucleotide means any nucleotide of more than 3 bases in length used to facilitate detection or identification of a target nucleic acid, including probes and primers.

A “polymerase” is an enzyme that catalyzes the sequential addition of monomeric units to a polymeric chain, or links two or more monomeric units to initiate a polymeric chain. The “polymerase” will work by adding monomeric units whose identity is determined by and which is complementary to a template molecule of a specific sequence. For example, DNA polymerases such as DNA pol 1 and Taq polymerase add deoxyribonucleotides to the 3’ end of a polynucleotide chain in a template-dependent manner, thereby synthesizing a nucleic acid that is complementary to the template molecule. Polymerases may be used either to extend a primer once or repetitively or to amplify a polynucleotide by repetitive priming of two complementary strands using two primers. A “thermostable polymerase” refers to a DNA or RNA polymerase enzyme that can withstand extremely high temperatures, such as those approaching 100°C. Often, thermostable polymerases are derived from organisms that live in extreme temperatures, such as *Thermus aquaticus*. Examples of thermostable polymerases include Taq, Tth, Pfu, Vent, deep vent, ULTma, and variations and derivatives thereof.

A “polynucleotide” refers to a linear chain of nucleotides connected by a phosphodiester linkage between the 3’-hydroxyl group of one nucleoside and the 5’-hydroxyl group of a second nucleoside which in turn is linked through its 3’-hydroxyl group to the 5’-hydroxyl group of a third nucleoside and so on to form a polymer comprised of nucleosides linked by a phosphodiester backbone. A “modified polynucleotide” refers to a polynucleotide in which one or more natural nucleotides have been partially, substantially, or completely replaced with modified nucleotides.

A “primer” is an oligonucleotide, the sequence of at least of portion of which is complementary to a segment of a template DNA which is to be amplified or replicated. Typically primers are used in performing the polymerase chain reaction (PCR). A primer

hybridizes with (or “anneals” to) the template DNA and is used by the polymerase enzyme
uses as the starting point for the replication/amplification process. The primers herein are
selected to be “substantially” complementary to different strands of a particular target DNA
sequence. This means that the primers must be sufficiently complementary to hybridize with
5 their respective strands. Therefore, the primer sequence need not reflect the exact sequence
of the template. For example, a non-complementary nucleotide fragment may be attached to
the 5’ end of the primer, with the remainder of the primer sequence being complementary to
the strand. Alternatively, non-complementary bases or longer sequences can be interspersed
into the primer, provided that the primer sequence has sufficient complementarity with the
10 sequence of the strand to hybridize therewith and thereby form the template for the synthesis
of the extension product.

“Probes” refer to oligonucleotides nucleic acid sequences of variable length, used in
the detection of identical, similar, or complementary nucleic acid sequences by hybridization.
An oligonucleotide sequence used as a detection probe may be labeled with a detectable
15 moiety.

The following are non-limiting examples of polynucleotides: a gene or gene fragment,
exons, introns, mRNA, tRNA, rRNA, ribozymes, cDNA, recombinant polynucleotides,
branched polynucleotides, plasmids, vectors, isolated DNA of any sequence, isolated RNA of
any sequence, nucleic acid probes and primers. A polynucleotide may comprise modified
20 nucleotides, such as methylated nucleotides and nucleotide analogs, uracil, other sugars and
linking groups such as fluororibose and thiolate, and nucleotide branches. The sequence of
nucleotides may be further modified after polymerization, such as by conjugation, with a
labeling component. Other types of modifications included in this definition are caps,
substitution of one or more of the naturally occurring nucleotides with an analog, and
25 introduction of means for attaching the polynucleotide to proteins, metal ions, labeling
components, other polynucleotides or solid support.

An “isolated” polynucleotide or polypeptide is one that is substantially pure of the
materials with which it is associated in its native environment. By substantially free, is meant
at least 50%, at least 55%, at least 60%, at least 65%, at advantageously at least 70%, at least
30 75%, more advantageously at least 80%, at least 85%, even more advantageously at least
90%, at least 91%, at least 92%, at least 93%, at least 94%, at least 95%, at least 96%, at least
97%, most advantageously at least 98%, at least 99%, at least 99.5%, at least 99.9% free of
these materials.

An "isolated" nucleic acid molecule is a nucleic acid molecule separate and discrete from the whole organism with which the molecule is found in nature; or a nucleic acid molecule devoid, in whole or part, of sequences normally associated with it in nature; or a sequence, as it exists in nature, but having heterologous sequences (as defined below) in association therewith.

The term "polynucleotide encoding a protein" as used herein refers to a DNA fragment or isolated DNA molecule encoding a protein, or the complementary strand thereto; but, RNA is not excluded, as it is understood in the art that thymidine (T) in a DNA sequence is considered equal to uracil (U) in an RNA sequence. Thus, RNA sequences for use in the invention, e.g., for use in RNA vectors, can be derived from DNA sequences, by thymidine (T) in the DNA sequence being considered equal to uracil (U) in RNA sequences.

A DNA "coding sequence" or a "nucleotide sequence encoding" a particular protein, is a DNA sequence which is transcribed and translated into a polypeptide in vitro or in vivo when placed under the control of appropriate regulatory elements. The boundaries of the coding sequence are determined by a start codon at the 5' (amino) terminus and a translation stop codon at the 3' (carboxy) terminus. A coding sequence can include, but is not limited to, prokaryotic sequences, cDNA from eukaryotic mRNA, genomic DNA sequences from eukaryotic (e.g., mammalian) DNA, and even synthetic DNA sequences. A transcription termination sequence will usually be located 3' to the coding sequence.

"Homology" refers to the percent identity between two polynucleotide or two polypeptide moieties. Two DNA, or two polypeptide sequences are "substantially homologous" to each other when the sequences exhibit at least about 80%, 81%, 82%, 83%, 84%, 85%, 86%, 87%, 88%, 89%, preferably at least about 90%, 91%, 92%, 93%, 94% and most preferably at least about 95%, 96%, 97%, 98%, 99%, 99.5%, 99.9% sequence identity over a defined length of the molecules. As used herein, substantially homologous also refers to sequences showing complete identity (100% sequence identity) to the specified DNA or polypeptide sequence.

Homology can be determined by hybridization of polynucleotides under conditions that form stable duplexes between homologous regions, followed by digestion with single-stranded-specific nuclease(s), and size determination of the digested fragments. DNA sequences that are substantially homologous can be identified in a Southern hybridization experiment under, for example, stringent conditions, as defined for that particular system. Defining appropriate hybridization conditions is within the skill of the art. See, e.g., Sambrook et al. *supra*; DNA Cloning, *supra*; Nucleic Acid Hybridization, *supra*.

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Two nucleic acid fragments are considered to be "selectively hybridizable" to a polynucleotide if they are capable of specifically hybridizing to a nucleic acid or a variant thereof or specifically priming a polymerase chain reaction: (i) under typical hybridization and wash conditions, as described, for example, in Sambrook et al. *supra* and Nucleic Acid Hybridization, *supra*, (ii) using reduced stringency wash conditions that allow at most about 25-30% basepair mismatches, for example: 2x SSC, 0.1% SDS, room temperature twice, 30 minutes each; then 2x SSC, 0.1% SDS, 37°C once, 30 minutes; then 2 x SSC room temperature twice, 10 minutes each, or (iii) selecting primers for use in typical polymerase chain reactions (PCR) under standard conditions (described for example, in Saiki, et al. (1988) *Science* 239:487-491).

The term "capable of hybridizing under stringent conditions" as used herein refers to annealing a first nucleic acid to a second nucleic acid under stringent conditions as defined below. Stringent hybridization conditions typically permit the hybridization of nucleic acid molecules having at least 70% nucleic acid sequence identity with the nucleic acid molecule being used as a probe in the hybridization reaction. For example, the first nucleic acid may be a test sample or probe, and the second nucleic acid may be the sense or antisense strand of a nucleic acid or a fragment thereof. Hybridization of the first and second nucleic acids may be conducted under stringent conditions, e.g., high temperature and/or low salt content that tend to disfavor hybridization of dissimilar nucleotide sequences. Alternatively, hybridization of the first and second nucleic acid may be conducted under reduced stringency conditions, e.g. low temperature and/or high salt content that tend to favor hybridization of dissimilar nucleotide sequences. Low stringency hybridization conditions may be followed by high stringency conditions or intermediate medium stringency conditions to increase the selectivity of the binding of the first and second nucleic acids. The hybridization conditions may further include reagents such as, but not limited to, dimethyl sulfoxide (DMSO) or formamide to disfavor still further the hybridization of dissimilar nucleotide sequences. A suitable hybridization protocol may, for example, involve hybridization in 6 x SSC (wherein 1 x SSC comprises 0.015 M sodium citrate and 0.15 M sodium chloride), at 65° Celsius in an aqueous solution, followed by washing with 1 x SSC at 65° C. Formulae to calculate appropriate hybridization and wash conditions to achieve hybridization permitting 30% or less mismatch between two nucleic acid molecules are disclosed, for example, in Meinkoth et al. (1984) *Anal. Biochem.* 138: 267-284.

Protocols for hybridization techniques are well known to those of skill in the art and standard molecular biology manuals may be consulted to select a suitable

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hybridization protocol without undue experimentation. See, for example, Sambrook et al. (2001) *Molecular Cloning: A Laboratory Manual*, 3rd ed., Cold Spring Harbor Press.

Typically, stringent conditions will be those in which the salt concentration is less than about 1.5 M sodium ion, typically about 0.01 to 1.0 M Na ion concentration (or other salts) from about pH 7.0 to about pH 8.3 and the temperature is at least about 30° Celsius for short probes (e.g., 10 to 50 nucleotides) and at least about 60° C for long probes (e.g., greater than 50 nucleotides). Stringent conditions may also be achieved with the addition of destabilizing agents such as formamide. Exemplary low stringency conditions include hybridization with a buffer solution of 30 to 35% formamide, 1 M NaCl, 1% SDS (sodium dodecyl sulphate) at 37° Celsius, and a wash in 1-2 x SSC at 50 to 55° Celsius. Exemplary moderate stringency conditions include hybridization in 40 to 45% formamide, 1 M NaCl, 1% SDS at 37° Celsius, and a wash in 0.5-1 x SSC at 55 to 60° Celsius. Exemplary high stringency conditions include hybridization in 50% formamide, 1 M NaCl, 1% SDS at 37° Celsius, and a wash in 0.1 x SSC at 60 to 65° Celsius.

Also encompassed by the present invention is fragments of polypeptide selected from the sequences SEQ ID NOS: 18, 20, 22, 24, 26, 48, 50 and 52. Fragments of polypeptide and epitope determination procedures, such as, generating overlapping peptide libraries (Hemmer B. et al., *Immunology Today*, 1998, 19 (4), 163-168), Pepscan (Geysen H. M. et al., *Proc. Nat. Acad. Sci. USA*, 1984, 81 (13), 3998-4002; Geysen H. M. et al., *Proc. Nat. Acad. Sci. USA*, 1985, 82 (1), 178-182; Van der Zee R. et al., *Eur. J. Immunol.*, 1989, 19 (1), 43-47; Geysen H. M., *Southeast Asian J. Trop. Med. Public Health*, 1990, 21 (4), 523-533; Multipin® Peptide Synthesis Kits de Chiron) and algorithms (De Groot A. et al., *Nature Biotechnology*, 1999, 17, 533-561), can be used in the practice of the invention, without undue experimentation.

The polypeptides or fragments thereof are produced advantageously by *in vitro* expression. Another aspect of the invention is the use of the polynucleotides according to the invention, for the expression and the production of peptides, polypeptides or proteins, or more generally, expression products, e.g., immunogens, antigens or epitopes. In an embodiment, the polypeptides or peptides or proteins encoded by these polynucleotides may be used as subunit immunogens or antigens or epitopes in immunogenic compositions or vaccines. The invention also embraces the immunogenic fragments of these polypeptides, having at least a chain of 8 amino acids of the polypeptide, at least 10, at least 20, such as at least 30, advantageously at least 50 and more advantageously at least 70, e.g., fragments of

the polypeptides containing at least 8 contiguous amino acids of the polypeptide, at least 10 contiguous amino acids of the polypeptide, advantageously at least 20, such as at least 30 and more advantageously at least 50 contiguous amino acids of the polypeptide, and even more advantageously at least 70 contiguous amino acids of the polypeptide. Of course, a fragment
5 is less than the entire polypeptide. A fragment can be combined with other polypeptides, e.g., in fusion polypeptides; for instance, a polypeptide of the invention or fragment thereof can be a portion of a fusion polypeptide which includes another portion (another polypeptide) (i.e. fused to glutathione S-transferase gene (GST) (Chanter N *et al.*, Microb. Pathog., 1999, 27(3): 133-143), or to cholera toxin (Sheoran A S *et al.*, Vaccine, 2002, 20(11-12): 1653-
10 1659)), e.g., an immunogenicity-enhancing portion and/or a secretion-enhancing portion such as a lipoprotein portion that enhances immunogenicity, such as a T-cell epitope peptide or a signal or leader sequence portion. Accordingly, the invention envisions the expression of polypeptides, proteins, antigens, immunogens or epitopes – whether herein identified sequences or fragments thereof or those that are heterologous to the vectors of the invention –
15 as fusions, e.g., as a portion of a fusion polypeptide, e.g., a fusion polypeptide that advantageously includes an immunogenicity enhancing portion such as a lipoprotein portion and/or a secretion-enhancing portion such as a signal or leader sequence portion.

As used herein, the term “antigen” or “immunogen” means a substance that induces a specific immune response in a host animal. The antigen may comprise a whole organism,
20 killed, attenuated or live; a subunit or portion of an organism; a recombinant vector containing an insert with immunogenic properties; a piece or fragment of DNA capable of inducing an immune response upon presentation to a host animal; a protein, a polypeptide, a peptide, an epitope, a hapten, or any combination thereof. Alternately, the immunogen or antigen may comprise a toxin or antitoxin.

25 The term “immunogenic protein or peptide” as used herein also refers includes peptides and polypeptides that are immunologically active in the sense that once administered to the host, it is able to evoke an immune response of the humoral and/or cellular type directed against the protein. Preferably the protein fragment is such that it has substantially the same immunological activity as the total protein. Thus, a protein fragment according to
30 the invention comprises or consists essentially of or consists of at least one epitope or antigenic determinant. The term epitope relates to a protein site able to induce an immune reaction of the humoral type (B cells) and/or cellular type (T cells).

The term “immunogenic protein or peptide” further contemplates deletions, additions and substitutions to the sequence, so long as the polypeptide functions to produce an

immunological response as defined herein. In this regard, particularly preferred substitutions will generally be conservative in nature, *i.e.*, those substitutions that take place within a family of amino acids. For example, amino acids are generally divided into four families: (1) acidic--aspartate and glutamate; (2) basic--lysine, arginine, histidine; (3) non-polar--alanine, valine, leucine, isoleucine, proline, phenylalanine, methionine, tryptophan; and (4) uncharged polar--glycine, asparagine, glutamine, cystine, serine threonine, tyrosine. Phenylalanine, tryptophan, and tyrosine are sometimes classified as aromatic amino acids. It is reasonably predictable that an isolated replacement of leucine with isoleucine or valine, or vice versa; an aspartate with a glutamate or vice versa; a threonine with a serine or vice versa; or a similar conservative replacement of an amino acid with a structurally related amino acid, will not have a major effect on the biological activity. Proteins having substantially the same amino acid sequence as the reference molecule but possessing minor amino acid substitutions that do not substantially affect the immunogenicity of the protein are, therefore, within the definition of the reference polypeptide.

The term "epitope" refers to the site on an antigen or hapten to which specific B cells and/or T cells respond. The term is also used interchangeably with "antigenic determinant" or "antigenic determinant site". Antibodies that recognize the same epitope can be identified in a simple immunoassay showing the ability of one antibody to block the binding of another antibody to a target antigen.

An "immunological response" to a composition or vaccine is the development in the host of a cellular and/or antibody-mediated immune response to a composition or vaccine of interest. Usually, an "immunological response" includes but is not limited to one or more of the following effects: the production of antibodies, B cells, helper T cells, and/or cytotoxic T cells, directed specifically to an antigen or antigens included in the composition or vaccine of interest. Preferably, the host will display either a therapeutic or protective immunological response such that resistance to new infection will be enhanced and/or the clinical severity of the disease reduced. Such protection will be demonstrated by either a reduction or lack of symptoms normally displayed by an infected host, a quicker recovery time and/or a lowered viral titer in the infected host.

The terms "immunogenic" protein or polypeptide as used herein also refers to an amino acid sequence which elicits an immunological response as described above. An "immunogenic" protein or polypeptide, as used herein, includes the full-length sequence of the protein, analogs thereof, or immunogenic fragments thereof. By "immunogenic fragment" is meant a fragment of a protein which includes one or more epitopes and thus

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elicits the immunological response described above. Such fragments can be identified using any number of epitope mapping techniques, well known in the art. See, *e.g.*, *Epitope Mapping Protocols* in *Methods in Molecular Biology*, Vol. 66 (Glenn E. Morris, Ed., 1996) Humana Press, Totowa, N.J. For example, linear epitopes may be determined by *e.g.*,

5 concurrently synthesizing large numbers of peptides on solid supports, the peptides corresponding to portions of the protein molecule, and reacting the peptides with antibodies while the peptides are still attached to the supports. Such techniques are known in the art and described in, *e.g.*, U.S. Pat. No. 4,708,871; Geysen *et al.* (1984) *Proc. Natl. Acad. Sci. USA* 81:3998-4002; Geysen *et al.* (1986) *Molec. Immunol.* 23:709-715.

10 Similarly, conformational epitopes are readily identified by determining spatial conformation of amino acids such as by, *e.g.*, x-ray crystallography and 2-dimensional nuclear magnetic resonance. See, *e.g.*, *Epitope Mapping Protocols*, *supra*.

Synthetic antigens are also included within the definition, for example, polypeptides, flanking epitopes, and other recombinant antigens. See, *e.g.*, Bergmann *et al.* (1993) *Eur. J. Immunol.* 23:2777-2781; Bergmann *et al.* (1996) *J. Immunol.* 157:3242-3249; Suhrbier, A. (1997) *Immunol. and Cell Biol.* 75:402-408; Gardner *et al.* (1998) 12th World AIDS Conference, Geneva, Switzerland, Jun. 28-Jul. 3, 1998. Immunogenic fragments, for purposes of the present invention, will usually include at least about 3 amino acids, preferably at least about 5 amino acids, more preferably at least about 10-15 amino acids, and most
20 preferably 25 or more amino acids, of the molecule. There is no critical upper limit to the length of the fragment, which could comprise nearly the full-length of the protein sequence, or even a fusion protein comprising at least one epitope of the protein.

Accordingly, a minimum structure of a polynucleotide expressing an epitope is that it comprises or consists essentially of or consists of nucleotides to encode an epitope or
25 antigenic determinant of the protein or polyprotein of interest. A polynucleotide encoding a fragment of the total protein or polyprotein, more advantageously, comprises or consists essentially of or consists of a minimum of 21 nucleotides, advantageously at least 42 nucleotides, and preferably at least 57, 87 or 150 consecutive or contiguous nucleotides of the sequence encoding the total protein or polyprotein. Epitope determination procedures, such
30 as, generating overlapping peptide libraries (Hemmer B. *et al.*, *Immunology Today*, 1998, 19 (4), 163-168), Pepscan (Geysen *et al.*, (1984) *Proc. Nat. Acad. Sci. USA*, 81, 3998-4002; Geysen *et al.*, (1985) *Proc. Nat. Acad. Sci. USA*, 82, 178-182; Van der Zee R. *et al.*, (1989) *Eur. J. Immunol.*, 19, 43-47; Geysen H.M., (1990) *Southeast Asian J. Trop. Med. Public Health*, 21, 523-533; Multipin.RTM. Peptide Synthesis Kits de Chiron) and algorithms (De

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Groot A. et al., (1999) Nature Biotechnology, 17, 533-561), and in PCT Application Serial No. PCT/US2004/022605, can be used in the practice of the invention, without undue experimentation. Other documents cited and incorporated herein may also be consulted for methods for determining epitopes of an immunogen or antigen and thus nucleic acid molecules that encode such epitopes.

In one embodiment, the fragments of polypeptide are selected from the sequences SEQ ID NOS: 32, 34, 36, 38, 40, 54, 56 and 58, which are respectively encoded by the nucleotide sequences SEQ ID NOS: 31, 33, 35, 37, 39, 53, 55 and 57. The fragments of polypeptides SEQ ID NOS: 32, 34, 36, 38, 40, 54, 56 and 58 are part of Se50, Se1459, Se595, Se528, Se358, Se1631, Se1681 and Sla, respectively.

The polynucleotides encoding polypeptides according to the present invention, analogs thereof or fragments thereof are inserted into a vector, linked to regulatory elements such as promoter, ribosome binding region and terminator, and start codon and stop codon.

The polypeptides, analogs thereof and fragments thereof can be produced by *in vitro* expression in host cells. The *in vitro* expression vectors are expression vectors used for the *in vitro* expression of proteins in an appropriate cell system. This can be produced in prokaryotic host cells, i.e. in *Escherichia coli* (Mahona F *et al.*, Biochimie 1994, 46(1): 9-14; Watt M A *et al.*, Cell Stress Chaperones 1997, 2(3): 180-90; Frey J Res. Microbiol. 1992, 143(3): 263-9) or in *Lactobacillus* (Seegers J F, Trends Biotechnol. 2002, 20(12): 508-515; Pouwels P H *et al.*, Methods Enzymol. 2001, 336: 369-389), or in eukaryotic host cells, i.e. in yeast (Gerngross T U, Nat Biotechnol. 2004, 22(11): 1409-1414; Malissard M *et al.*, Glycoconj J. 1999, 16(2): 125-139), in insect cells (Oker-Blom C *et al.*, Brief Funct Genomic Proteomic. 2003, 2(3), 244-253), mammal cells or avian cells.

For *Escherichia coli*, different strains can be used, notably BL21(DE3) strain (Novagen or Invitrogen; see Hedayati M A *et al.*, Protein Expr Purif, 2005, 43(2): 133-139) Origami2(DE3) strain (Novagen or Invitrogen), Y1089 (Galan J E *et al.*, Infect Immun, 1987, 55(12): 3181-3187). Vectors are advantageously plasmids, i.e. pGEX plasmids (GE Healthcare), pET plasmids (Novagen or Invitrogen) (Jiang X Y *et al.*, J Biochem Mol Biol. 2006, 39(1): 22-25; Hedayati M A *et al.*, Protein Expr Purif, 2005, 43(2): 133-139; Jedrzejewski M J *et al.*, Protein Expr Purif, 1998, 13(1): 83-89). Vectors can also be a virus, notably a bacteriophage, i.e. lambda-gt11 phage (Galan J E *et al.*, Infect Immun, 1987, 55(12): 3181-3187). Another approach is to use fused genes for the production of chimeric proteins between a *Streptococcus* protein and a *Escherichia coli* protein, notably lipoprotein (Cullen P

A *et al.*, Plasmid, 2003, 49(1): 18-29), or with GST (Zhao G *et al.*, Protein Expr Purif, 1999, 16(2): 331-339). The Streptococcal insert can be linked to promoter, i.e. bacteriophage T7 promoter (Yamamoto M *et al.*, FEMS Microbiol Lett, 1995, 132(3): 209-213), signal peptide sequence, i.e. *Borrelia burgdorferi* outer surface protein A signal peptide (De B K *et al.*, Vaccine, 2000, 18(17): 1811-1821), an *Escherichia coli* major outer membrane lipoprotein signal sequence (Cullen P A *et al.*, Plasmid, 2003, 49(1): 18-29).

For *Lactobacillus*, different strains can be used, notably *Lactobacillus brevis*, *Lactobacillus plantarum*, *Lactobacillus casei* (Seegers J F, Trends Biotechnol., 2002, 20(12): 508-515). Promoters and regulatory genes involved in production of sakacin P are suitable for establishing inducible high-level gene expression in *Lactobacillus* (Mathiesen G *et al.*, Lett Appl Microbiol., 2004, 39(2): 137-143), or lactose operon promoter (Oliveira M L *et al.*, FEMS Microbiol Lett., 2003, 227(1): 25-31). Another approach is to use fused genes for the production of chimeric proteins between a *Lactobacillus* protein and a *Streptococcus* protein (Hung J *et al.*, FEMS Microbiol Lett., 2002, 211(1): 71-75).

For *in vitro* expression in insect cells, vectors are advantageously viruses, i.e. baculoviruses (*see, e.g.*, U.S. Patent No. 4,745,051; Vialard J. *et al.*, J. Virol., 1990 64 (1), 37-50; Verne A., Virology, 1988, 167, 56-71; Oker-Blom C *et al.*, Brief Funct Genomic Proteomic. 2003, 2(3), 244-253), e.g. *Autographa californica* Nuclear Polyhedrosis Virus AcNPV, and insect cells are Sf9 *Spodoptera frugiperda* cells (ATCC CRL 1711; *see also* U.S. Patents Nos. 6,228,846, 6,103,526). Protein production can take place by the transfection of mammalian cells by plasmids, by replication or expression without productive replication of viral vectors on mammal cells or avian cells. Mammalian cells which can be used are advantageously hamster cells (e.g. CHO or BHK-21) or monkey cells (e.g. COS or VERO) or bovine cells (e.g. MDBK), i.e. culture of EHV-1 vectors in MDBK cells (Ibrahim el S M *et al.*, Microbiol. Immunol., 2004, 48(11): 831-842) or in Vero cells (US-A-4,110,433), or culture of VEEV replicons in BHK cells (Lee J S *et al.*, Infect. Immun., 2001, 69(9): 5709-5715).

It is understood to one of skill in the art that conditions for culturing *in vitro* a host cell varies according to the particular gene and that routine experimentation is necessary at times to determine the optimal conditions for culturing a protein depending on the host cell. A "host cell" denotes a prokaryotic or eukaryotic cell that has been genetically altered, or is capable of being genetically altered by administration of an exogenous polynucleotide, such as a recombinant plasmid or vector. When referring to genetically altered cells, the term refers both to the originally altered cell and to the progeny thereof.

The *in vitro* expression vectors can be introduced into a suitable host cell for replication and amplification. The vectors containing the polynucleotides of interest can be introduced into the host cell by any of a number of appropriate means, including direct uptake, endocytosis, transfection, f-mating, electroporation, transfection employing calcium chloride, rubidium chloride, calcium phosphate, DEAE-dextran, or other substances;
5 microprojectile bombardment; lipofection; and infection (where the vector is infectious, for instance, a retroviral vector). The choice of introducing vectors or polynucleotides will often depend on features of the host cell.

The expressed proteins can be harvested in or from the culture supernatant after, or
10 not after secretion (if there is no secretion a cell lysis typically occurs or is performed), optionally concentrated by concentration methods such as ultrafiltration and/or purified by purification means, such as affinity, ion exchange or gel filtration-type chromatography methods, notably by affinity chromatography, i.e. using Ni NTA Superflow (Qiagen) or Ni Sepharose fastflow (Amersham), or by gel-filtration, i.e. using Sephacryl® or Superdex®
15 (Amersham GE Healthcare).

The polypeptides and fragments thereof can also be obtained by extraction, notably acid extraction or mutanolysin extraction (Boschwitz J S *et al.*, Cornell Vet., 1991, 81(1): 25-36), and purification, notably immunoprecipitation (Erickson E D *et al.*, Can J Comp Med., 1975, 39(2): 110-115), from crude culture of streptococcal bacteria, notably of *Streptococcus equi*.
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The polypeptides and fragments thereof can also be synthesised chemically (Luo Y *et al.*, Vaccine 1999, 17(7-8): 821-31).

By "subunit vaccine composition" is meant a composition containing at least one immunogenic polypeptide or a polynucleotide able to express at least one, but not all, antigen
25 derived from or homologous to an antigen from a pathogen of interest. Such a composition is substantially free of intact pathogen cells or particles, or the lysate of such cells or particles. Thus, a "subunit vaccine composition" is prepared from at least partially purified (preferably substantially purified) immunogenic polypeptides from the pathogen, or recombinant analogs thereof. A subunit vaccine composition can comprise the subunit antigen or antigens of
30 interest substantially free of other antigens or polypeptides from the pathogen.

An object of the invention is a subunit immunogenic composition or subunit vaccine comprising at least one polypeptide, analog thereof or fragment thereof according to the invention, and a pharmaceutically acceptable excipient, diluent or vehicle, and optionally an adjuvant and/or a stabilizer.

The subunit immunogenic or vaccine composition can comprise at least one polypeptide selected from the group consisting of sequences SEQ ID NOS: 18, 20, 22, 24, 26, 48, 50 and 52, or analogs thereof, or fragments thereof. The subunit immunogenic or vaccine composition can advantageously comprise two or three or four or five polypeptides selected from the group consisting of sequences SEQ ID NOS: 18, 20, 22, 24, 26, 48, 50 and 52, or analogs thereof, or fragments thereof.

The subunit immunogenic or vaccine composition can comprise at least one polypeptide selected from the group consisting of sequences SEQ ID NOS: 32, 34, 36, 38, 40, 54, 56 and 58, or analogs thereof.

The subunit immunogenic or vaccine composition can advantageously comprise two or three or four or five polypeptides selected from the group consisting of sequences SEQ ID NOS: 32, 34, 36, 38, 40, 54, 56 and 58, or analogs thereof.

Another object of the invention is a recombinant immunogenic composition or vaccine comprising at least one recombinant *in vivo* expression vector, and a pharmaceutically acceptable excipient, diluent or vehicle, and optionally an adjuvant and/or a stabilizer. The recombinant *in vivo* expression vectors are vectors comprising polynucleotides or their fragments inserted hereinto and able to express *in vivo* in the targeted mammal the polypeptide encoded by this polynucleotide or fragment thereof. The vectors can be a polynucleotide vectors or plasmids (EP-A2-1001025; Chaudhuri P Res. Vet. Sci. 2001, 70(3), 255-6) for DNA immunogenic compositions and DNA vaccines, or can be viruses (e.g. herpesvirus such as equine herpesvirus type 1 (Trapp S *et al.*, J. Virol. 2005, 79(9): 5445-5454), equine herpesvirus type 2, equine herpesvirus type 4; poxvirus virus such as vaccinia virus or avipox virus, like fowlpox (US-A-5,174,993 US-A-5,505,941 and US-A-5,766,599) or canarypox (US-A-5,756,103); adenovirus such as human adenovirus (Chroboczek J *et al.*, Virol. 1992, 186: 280-285); encephalitis virus such as venezuelean equine encephalitis virus (Pushko P *et al.* Virol. 1997, 239: 389-401; Davis N L *et al.* J. Virol. 2000, 74: 371-378)) for recombinant viral vector immunogenic compositions and recombinant viral vector vaccines. In a further embodiment, the polynucleotides or their fragments may be inserted into recombinant *in vivo* expression bacterial vectors, notably into *Salmonella*, to produce live recombinant bacterial immunogenic compositions or vaccines.

The term plasmid covers any DNA transcription unit comprising a polynucleotide according to the invention and the elements necessary for its *in vivo* expression in a cell or cells of the desired host or target; and, in this regard, it is noted that a supercoiled or non-supercoiled, circular plasmid, as well as a linear form, are intended to be within the scope of

the invention. In a specific, non-limiting example, the pVR1020 or 1012 plasmid (VICAL Inc.; Luke C. et al., Journal of Infectious Diseases, 1997, 175, 91-97; Hartikka J. et al., Human Gene Therapy, 1996, 7, 1205-1217) can be utilized as a vector for the insertion of a polynucleotide sequence. The pVR1020 plasmid is derived from pVR1012 and contains the human tPA signal sequence. Each plasmid comprises or contains or consists essentially of, in addition to the polynucleotide or analog thereof or fragment thereof, operably linked to a promoter or under the control of a promoter or dependent upon a promoter. In general, it is advantageous to employ a strong promoter functional in eukaryotic cells. The preferred strong promoter is the immediate early cytomegalovirus promoter (CMV-IE) of human or murine origin, or optionally having another origin such as the rat or guinea pig. The CMV-IE promoter can comprise the actual promoter part, which may or may not be associated with the enhancer part. Reference can be made to EP-A-260 148, EP-A-323 597, U.S. Patents Nos. 5,168,062, 5,385,839, and 4,968,615, as well as to PCT Application No WO-A-87/03905. The CMV-IE promoter is advantageously a human CMV-IE (Boshart M. et al., Cell, 1985, 41, 521-530) or murine CMV-IE. In more general terms, the promoter has either a viral or a cellular origin. A strong viral promoter other than CMV-IE that may be usefully employed in the practice of the invention is the early/late promoter of the SV40 virus or the LTR promoter of the Rous sarcoma virus. A strong cellular promoter that may be usefully employed in the practice of the invention is the promoter of a gene of the cytoskeleton, such as e.g. the desmin promoter (Kwissa M. et al., Vaccine, 2000, 18, 2337-2344), or the actin promoter (Miyazaki J. et al., Gene, 1989, 79, 269-277). Functional sub fragments of these promoters, i.e., portions of these promoters that maintain an adequate promoting activity, are included within the present invention, e.g. truncated CMV-IE promoters according to PCT Application No. WO-A-98/00166 or U.S. Patent No. 6,156,567 can be used in the practice of the invention. A promoter in the practice of the invention consequently includes derivatives and sub fragments of a full-length promoter that maintain an adequate promoting activity and hence function as a promoter, preferably promoting activity substantially similar to that of the actual or full-length promoter from which the derivative or sub fragment is derived, e.g., akin to the activity of the truncated CMV-IE promoters of U.S. Patent No. 6,156,567 to the activity of full-length CMV-IE promoters. Thus, a CMV-IE promoter in the practice of the invention can comprise or consist essentially of or consist of the promoter portion of the full-length promoter and/or the enhancer portion of the full-length promoter, as well as derivatives and sub fragments. Advantageously, the plasmids comprise or consist essentially of other expression control elements. It is particularly advantageous to incorporate stabilizing

sequence(s), e.g., intron sequence(s), preferably the first intron of the hCMV-IE (PCT Application No. WO-A-89/01036), the intron II of the rabbit β -globin gene (van Ooyen et al., Science, 1979, 206, 337-344). As to the polyadenylation signal (polyA) for the plasmids and viral vectors other than poxviruses, use can more be made of the poly(A) signal of the bovine growth hormone (bGH) gene (*see* U.S. Patent No. 5,122,458), or the poly(A) signal of the rabbit β -globin gene or the poly(A) signal of the SV40 virus.

For recombinant vectors based on equine herpesvirus, in particular attenuated strains (i.e. EHV-1 KyA strains (Zhang Y *et al.*, Virology, 2000, 268(2): 482-492)) or attenuated by mutation or deletion of genes involve in pathogenicity (Kirisawa R *et al.*, Vet. Microbiol., 2003, 95(3): 159-174) or by serial culture passages (US-A-4,110,433) can be used. For EHV-1, the insertion can be made in the intergenic region between ORF 62 and ORF 63 (Ibrahim el S M *et al.*, Microbiol. Immunol., 2004, 48(11): 831-842; Csellner H *et al.*, Arch. Virol., 1998, 143(11): 2215-2231), or in the genes, eventually after a partial or complete deletion of the genes, i.e. in thymidine kinase (TK) gene, in genes 1 or 71 (Kirisawa R *et al.*, Vet. Microbiol., 2003, 95(3): 159-174), in glycoprotein I gene and glycoprotein E gene (Matsumura T. *et al.*, Virology, 1998, 242(1): 68-79), in IR6 protein gene (Osterrieder N *et al.*, Virology, 1996, 217(2): 442-451), in gene 15 (EP-B1-0,668,355). For EHV-4, the insertion can be made in glycoprotein I gene and glycoprotein E gene (Damiani A M *et al.*, Virus Res., 2000, 67(2): 189-202), in US2 or TK or glycoprotein E gene (US-A-5,741,696; US-A-5,731,188), eventually after a partial or complete deletion of the genes. In one embodiment the polynucleotide to be expressed is inserted under the control of a promoter functional in eukaryotic cells, advantageously a CMV-IE promoter (murine or human). A poly(A) sequence and terminator sequence can be inserted downstream the polynucleotide to be expressed, e.g. bovine growth hormone or a rabbit β -globin gene polyadenylation signal.

For recombinant vector based on poxvirus vector, a vaccinia virus or an attenuated vaccinia virus, (for instance, MVA, a modified Ankara strain obtained after more than 570 passages of the Ankara vaccine strain on chicken embryo fibroblasts; *see* Stickl & Hochstein-Mintzel, Munch. Med. Wschr., 1971, 113, 1149-1153; Sutter *et al.*, Proc. Natl. Acad. Sci. U.S.A., 1992, 89, 10847-10851; available as ATCC VR-1508; or NYVAC, *see* U.S. Patent No. 5,494,807, for instance, Examples 1 to 6 and *et seq* of U.S. Patent No. 5,494,807 which discuss the construction of NYVAC, as well as variations of NYVAC with additional ORFs deleted from the Copenhagen strain vaccinia virus genome, as well as the insertion of heterologous coding nucleic acid molecules into sites of this recombinant, and also, the use of matched promoters; *see* also WO-A-96/40241), an avipox virus or an attenuated avipox virus

(e.g., canarypox, fowlpox, dovepox, pigeonpox, quailpox, ALVAC or TROVAC; see, e.g., U.S. Patent No. 5,505,941, 5,494,807) can be used. Attenuated canarypox viruses are described in U.S. Patent No. 5,756,103 (ALVAC) and WO-A-01/05934. Reference is also made to U.S. Patent No. 5,766,599 which pertains to the attenuated fowlpox strain TROVAC.

5 Reference is made to the canarypox available from the ATCC under access number VR-111. Numerous fowlpox virus vaccination strains are also available, e.g. the DIFTOSECT strain marketed by MERIAL and the NOBILIS VARIOLE vaccine marketed by INTERVET. For information on the method to generate recombinants thereof and how to administer recombinants thereof, the skilled artisan can refer documents cited herein and to WO-A-

10 90/12882, e.g., as to vaccinia virus mention is made of U.S. Patents Nos. 4,769,330, 4,722,848, 4,603,112, 5,110,587, 5,494,807, and 5,762,938 *inter alia*; as to fowlpox, mention is made of U.S. Patents Nos. 5,174,993, 5,505,941 and 5,766,599 *inter alia*; as to canarypox mention is made of U.S. Patent No. 5,756,103 *inter alia*. When the expression vector is a vaccinia virus, insertion site or sites for the polynucleotide or polynucleotides to be expressed

15 are advantageously at the thymidine kinase (TK) gene or insertion site, the hemagglutinin (HA) gene or insertion site, the region encoding the inclusion body of the A type (ATI); see also documents cited herein, especially those pertaining to vaccinia virus. In the case of canarypox, advantageously the insertion site or sites are ORF(s) C3, C5 and/or C6; see also documents cited herein, especially those pertaining to canarypox virus. In the case of

20 fowlpox, advantageously the insertion site or sites are ORFs F7 and/or F8; see also documents cited herein, especially those pertaining to fowlpox virus. The insertion site or sites for MVA virus are advantageously as in various publications, including Carroll M. W. et al., Vaccine, 1997, 15 (4), 387-394; Stittelaar K. J. et al., J. Virol., 2000, 74 (9), 4236-4243; Sutter G. et al., 1994, Vaccine, 12 (11), 1032-1040; and, in this regard it is also noted

25 that the complete MVA genome is described in Antoine G., Virology, 1998, 244, 365-396, which enables the skilled artisan to use other insertion sites or other promoters. Advantageously, the polynucleotide to be expressed is inserted under the control of a specific poxvirus promoter, e.g., the vaccinia promoter 7.5 kDa (Cochran et al., J. Virology, 1985, 54, 30-35), the vaccinia promoter I3L (Riviere et al., J. Virology, 1992, 66, 3424-3434), the

30 vaccinia promoter HA (Shida, Virology, 1986, 150, 451-457), the cowpox promoter ATI (Funahashi et al., J. Gen. Virol., 1988, 69, 35-47), the vaccinia promoter H6 (Taylor J. et al., Vaccine, 1988, 6, 504-508; Guo P. et al. J. Virol., 1989, 63, 4189-4198; Perkus M. et al., J. Virol., 1989, 63, 3829-3836), *inter alia*.

For recombinant vector based on adenovirus vector, a human adenovirus (HAV), advantageously, a human adenovirus serotype 5 (Ad5) vector, an E1-deleted and/or disrupted adenovirus, an E3-deleted and/or disrupted adenovirus or an E1- and E3-deleted and/or disrupted adenovirus can be used. Optionally, E4 may be deleted and/or disrupted from any of the adenoviruses described above. For example, the human Ad5 vectors described in Yarosh et al. and Lutze-Wallace et al. can be used (see, e.g., Yarosh et al., Vaccine. 1996 Sep;14(13):1257-64 and Lutze-Wallace et al., Biologicals. 1995 Dec;23(4):271-7). In one embodiment the viral vector is a human adenovirus, in particular a serotype 5 adenovirus, rendered incompetent for replication by a deletion in the E1 region of the viral genome. The deleted adenovirus is propagated in E1-expressing 293 cells or PER cells, in particular PER.C6 (F. Falloux et al Human Gene Therapy 1998, 9, 1909-1917). The human adenovirus can be deleted in the E3 region eventually in combination with a deletion in the E1 region (see, e.g. J. Shriver et al. Nature, 2002, 415, 331-335, F. Graham et al Methods in Molecular Biology Vol .7: Gene Transfer and Expression Protocols Edited by E. Murray, The Human Press Inc, 1991, p 109-128; Y. Ilan et al Proc. Natl. Acad. Sci. 1997, 94, 2587-2592; S. Tripathy et al Proc. Natl. Acad. Sci. 1994, 91, 11557-11561; B. Tapnell Adv. Drug Deliv. Rev.1993, 12, 185-199;X. Danthinne et al Gene Thrapy 2000, 7, 1707-1714; K. Berkner Bio Techniques 1988, 6, 616-629; K. Berkner et al Nucl. Acid Res. 1983, 11, 6003-6020; C. Chavier et al J. Virol. 1996, 70, 4805-4810). The insertion sites can be the E1 and/or E3 loci eventually after a partial or complete deletion of the E1 and/or E3 regions. Advantageously, when the expression vector is an adenovirus, the polynucleotide to be expressed is inserted under the control of a promoter functional in eukaryotic cells, such as a strong promoter, preferably a cytomegalovirus immediate-early gene promoter (CMV-IE promoter). The CMV-IE promoter is advantageously of murine or human origin. The promoter of the elongation factor 1 α can also be used. A muscle specific promoter can also be used (X. Li et al Nat. Biotechnol. 1999, 17, 241-245). Strong promoters are also discussed herein in relation to plasmid vectors. A poly(A) sequence and terminator sequence can be inserted downstream of the polynucleotide to be expressed, e.g. a bovine growth hormone gene or a rabbit β -globin gene polyadenylation signal.

For recombinant vector based on Encephalitis virus, a venezuelean equine encephalitis virus (VEEV) can be used (Nelson E L *et al.*, Breast Cancer Res. Treat., 2003, 82(3): 169-183), in particular as a replicon, that is to say as a self-replicating RNA containing all of the VEEV non-structural genes and a multiple-cloning site in place of the VEEV structural genes (Lee J S *et al.*, Infect. Immun., 2003, 71(3): 1491-1496; Velders M P et al.,

Cancer Res., 2001, 61(21): 7861-7867). The polynucleotides to be expressed are inserted into this multiple-cloning site, optionally linked to the nucleotide sequence encoding a secretory sequence or a tissue plasminogen activator secretory sequence. The polynucleotides to be expressed can also be inserted downstream of the subgenomic 26S promoter in place of the viral VEEV structural genes (Lee J S *et al.*, Infect. Immun., 2001, 69(9): 5709-5715; Pushko P *et al.*, Vaccine, 2000, 19(1): 142-153).

For recombinant vectors based on bacteria, *Salmonella*, notably *Salmonella typhimurium* (Yang X L *et al.*, Biomed. Environ. Sci., 2005, 18(6): 411-418; Dunstan S J *et al.*, FEMS Immunol. Med. Microbiol., 2003, 37(2-3): 111-119), *Salmonella typhi* (Santiago-Machuca A E *et al.*, Plasmid., 2002, 47(2): 108-119) can be used. The polynucleotides to be expressed can be inserted into the flagellin gene of *Salmonella* (Chauhan N *et al.*, Mol Cell Biochem., 2005, 276(1-2): 1-6), or into the *aroC* gene (Santiago-Machuca A E *et al.*, Plasmid., 2002, 47(2): 108-119). The polynucleotides to be expressed can also be inserted under the control of the anaerobically inducible *nirB* promoter (Santiago-Machuca A E *et al.*, Plasmid., 2002, 47(2): 108-119).

The recombinant immunogenic or vaccine composition can contain at least one recombinant expression vector comprising at least one polynucleotide encoding a polypeptide selected from the group consisting of sequences SEQ ID NOS: 18, 20, 22, 24, 26, 48, 50 and 52, or analogs thereof or fragments thereof. The recombinant immunogenic or vaccine composition can contain at least one recombinant expression vector comprising at least one polynucleotide encoding a polypeptide selected from the group consisting of sequences SEQ ID NOS: 32, 34, 36, 38, 40, 54, 56 and 58, or analogs thereof. The recombinant immunogenic or vaccine composition can advantageously contain at least one recombinant expression vector comprising at least one polynucleotide encoding two or three or four or five polypeptides selected from the group consisting of sequences SEQ ID NOS: 32, 34, 36, 38, 40, 54, 56 and 58, or analogs thereof.

The vectors of the invention may further comprise at least one heterologous polynucleotide. This is useful for reproducing or replicating heterologous polynucleotides and/or for expression of heterologous polynucleotides, either *in vivo* or *in vitro*. Such vectors are also useful for preparing multivalent immunogenic or vaccine compositions, notably multivalent DNA immunogenic compositions or vaccines and multivalent recombinant viral vector immunogenic compositions or vaccines. The heterologous nucleic acid sequence advantageously codes for an immunogen, antigen or epitope from a pathogenic viral, parasitic or bacterial agent, such bacterial agent is different from the streptococcal bacteria at the

origin of the polynucleotides encoding the polypeptides according to the invention. This heterologous sequence may encode an immunogen, antigen or epitope from western equine encephalitis virus (WEEV), eastern equine encephalitis virus (EEEV), venezuelean equine encephalitis virus (VEEV), equine influenza virus, equine herpesvirus type 1 (EHV-1),
5 equine herpesvirus type 2 (EHV-2), equine herpesvirus type 4 (EHV-4), Equine Artheritis virus (EAV), West Nile virus (WNV), tetanus, rhodococcus. In a particular embodiment, the heterologous sequence may encode an immunogen, antigen or epitope from equine influenza virus and from EHV-1 and/or EHV-4. An immunogen or antigen is a protein or polypeptide able to induce an immune response against the pathogenic agent or a secreted antigen of the
10 pathogenic agent, and contains one or more epitopes; an epitope is a peptide or polypeptide which is able to induce an immune response against the pathogenic agent or a secreted antigen of the pathogenic agent.

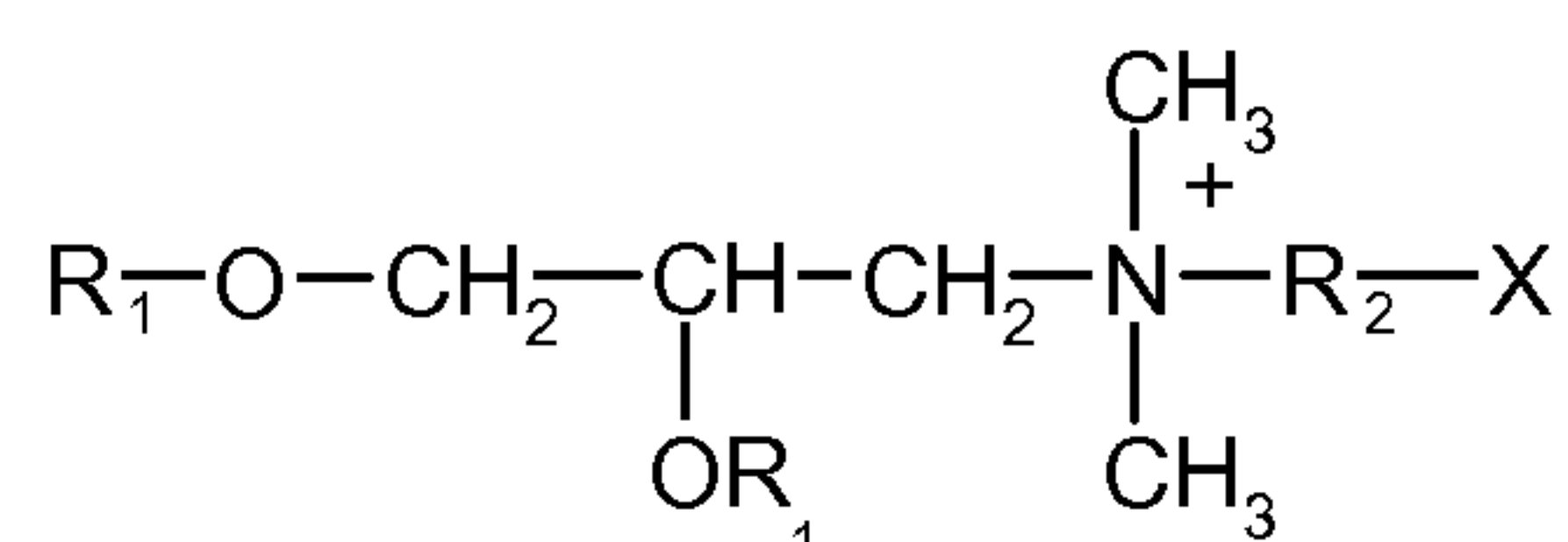
Optionally, the subunit immunogenic composition or vaccine of the invention can be combined with one or more immunogens, antigens or epitopes selected from other pathogenic
15 micro-organisms or viruses to form a multivalent subunit immunogenic composition or vaccine. For the equine, such a multivalent subunit immunogenic composition or vaccine may comprises at least one polypeptide according to the present invention and at least one immunogen, antigen or epitope from WEEV, EEEV, VEEV, equine influenza virus, EHV-1, EHV-4, EAV, WNV, tetanus, rhodococcus. In a particular embodiment, such a multivalent
20 subunit immunogenic composition or vaccine may comprises at least one polypeptide according to the present invention and at least one immunogen, antigen or epitope from equine influenza virus and from EHV-1 and/or EHV-4.

The pharmaceutically or veterinary acceptable excipient, diluent or vehicle may be water or saline, buffer but it may, for example, also comprise Streptococcus culture medium.

25 The pharmaceutically or veterinarily acceptable carriers or vehicles or excipients are well known to the one skilled in the art. For example, a pharmaceutically or veterinarily acceptable carrier or vehicle or excipient can be a 0.9% NaCl (e.g., saline) solution or a phosphate buffer. Other pharmaceutically or veterinarily acceptable carrier or vehicle or excipients that can be used for methods of this invention include, but are not limited to, poly-
30 (L-glutamate) or polyvinylpyrrolidone. The pharmaceutically or veterinarily acceptable carrier or vehicle or excipients may be any compound or combination of compounds facilitating the administration of the vector (or protein expressed from an inventive vector *in vitro*); advantageously, the carrier, vehicle or excipient may facilitate transfection and/or improve preservation of the vector (or protein). Doses and dose volumes are herein discussed

in the general description and can also be determined by the skilled artisan from this disclosure read in conjunction with the knowledge in the art, without any undue experimentation.

The cationic lipids containing a quaternary ammonium salt which are advantageously but not exclusively suitable for plasmids, are advantageously those having the following formula:



in which R₁ is a saturated or unsaturated straight-chain aliphatic radical having 12 to 18 carbon atoms, R₂ is another aliphatic radical containing 2 or 3 carbon atoms and X is an amine or hydroxyl group, e.g. the DMRIE. In another embodiment the cationic lipid can be associated with a neutral lipid, e.g. the DOPE.

Among these cationic lipids, preference is given to DMRIE (N-(2-hydroxyethyl)-N,N-dimethyl-2,3-bis(tetradecyloxy)-1-propane ammonium; WO96/34109), advantageously associated with a neutral lipid, advantageously DOPE (dioleoyl-phosphatidyl-ethanol amine; Behr J. P., 1994, Bioconjugate Chemistry, 5, 382-389), to form DMRIE-DOPE.

Advantageously, the plasmid mixture with the adjuvant is formed extemporaneously and advantageously contemporaneously with administration of the preparation or shortly before administration of the preparation; for instance, shortly before or prior to administration, the plasmid-adjuvant mixture is formed, advantageously so as to give enough time prior to administration for the mixture to form a complex, e.g. between about 10 and about 60 minutes prior to administration, such as approximately 30 minutes prior to administration.

When DOPE is present, the DMRIE:DOPE molar ratio is advantageously about 95: about 5 to about 5: about 95, more advantageously about 1: about 1, e.g., 1:1.

The DMRIE or DMRIE-DOPE adjuvant:plasmid weight ratio can be between about 50: about 1 and about 1: about 10, such as about 10: about 1 and about 1: about 5, and advantageously about 1: about 1 and about 1: about 2, e.g., 1:1 and 1:2.

The immunogenic compositions and vaccines according to the invention may comprise or consist essentially of one or more adjuvants. Suitable adjuvants for use in the practice of the present invention are (1) polymers of acrylic or methacrylic acid, maleic anhydride and alkenyl derivative polymers, (2) immunostimulating sequences (ISS), such as oligodeoxyribonucleotide sequences having one or more non-methylated CpG units

(Klinman et al., Proc. Natl. Acad. Sci., USA, 1996, 93, 2879-2883; WO98/16247), (3) an oil in water emulsion, such as the SPT emulsion described on p 147 of "Vaccine Design, The Subunit and Adjuvant Approach" published by M. Powell, M. Newman, Plenum Press 1995, and the emulsion MF59 described on p 183 of the same work, (4) cation lipids containing a quaternary ammonium salt, e.g., DDA (5) cytokines, (6) aluminum hydroxide or aluminum phosphate, (7) saponin or (8) other adjuvants discussed in any document cited and incorporated by reference into the instant application, or (9) any combinations or mixtures thereof.

The oil in water emulsion (3), which is especially appropriate for viral vectors, can be based on: light liquid paraffin oil (European pharmacopoeia type), isoprenoid oil such as squalane, squalene, oil resulting from the oligomerization of alkenes, e.g. isobutene or decene, esters of acids or alcohols having a straight-chain alkyl group, such as vegetable oils, ethyl oleate, propylene glycol, di(caprylate/caprata), glycerol tri(caprylate/caprata) and propylene glycol dioleate, or esters of branched, fatty alcohols or acids, especially isostearic acid esters.

The oil is used in combination with emulsifiers to form an emulsion. The emulsifiers may be nonionic surfactants, such as: esters of on the one hand sorbitan, mannide (e.g. anhydromannitol oleate), glycerol, polyglycerol or propylene glycol and on the other hand oleic, isostearic, ricinoleic or hydroxystearic acids, said esters being optionally ethoxylated, or polyoxypropylene-polyoxyethylene copolymer blocks, such as Pluronic, e.g., L121.

Among the type (1) adjuvant polymers, preference is given to polymers of crosslinked acrylic or methacrylic acid, especially crosslinked by polyalkenyl ethers of sugars or polyalcohols.

These compounds are known under the name carbomer (Pharmeuropa, vol. 8, no. 2, June 1996). One skilled in the art can also refer to U.S. Patent No. 2,909,462, which provides such

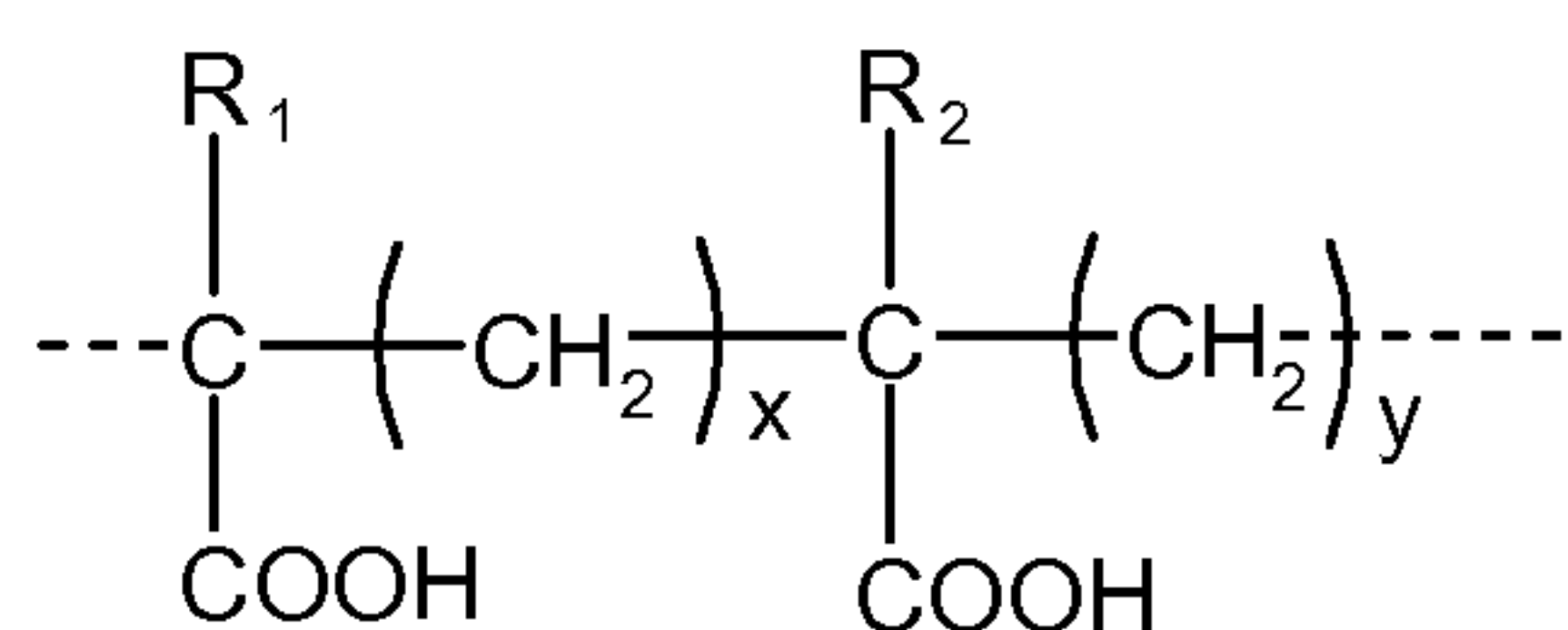
acrylic polymers crosslinked by a polyhydroxyl compound having at least three hydroxyl groups, preferably no more than eight such groups, the hydrogen atoms of at least three hydroxyl groups being replaced by unsaturated, aliphatic radicals having at least two carbon atoms. The preferred radicals are those containing 2 to 4 carbon atoms, e.g. vinyls, allyls and other ethylenically unsaturated groups. The unsaturated radicals can also contain other

substituents, such as methyl. Products sold under the name Carbopol (BF Goodrich, Ohio, USA) are especially suitable. They are crosslinked by allyl saccharose or by allyl pentaerythritol. Among them, reference is made to Carbopol 974P, 934P and 971P.

As to the maleic anhydride-alkenyl derivative copolymers, preference is given to EMA (Monsanto), which are straight-chain or crosslinked ethylene-maleic anhydride copolymers

and they are, for example, crosslinked by divinyl ether. Reference is also made to J. Fields et al., Nature 186: 778-780, June 4, 1960.

With regard to structure, the acrylic or methacrylic acid polymers and EMA are preferably formed by basic units having the following formula:



5 in which:

- R₁ and R₂, which can be the same or different, represent H or CH₃
- x = 0 or 1, preferably x = 1
- y = 1 or 2, with x + y = 2.

For EMA, x = 0 and y = 2 and for carbomers x = y = 1.

10 These polymers are soluble in water or physiological salt solution (20 g/l NaCl) and the pH can be adjusted to 7.3 to 7.4, e.g., by soda (NaOH), to provide the adjuvant solution in which the expression vector(s) can be incorporated. The polymer concentration in the final vaccine composition can range between 0.01 and 1.5% w/v, advantageously 0.05 to 1% w/v and preferably 0.1 to 0.4% w/v.

15 The cytokine or cytokines (5) can be in protein form in the immunogenic or vaccine composition, or can be co-expressed in the host with the immunogen or immunogens or epitope(s) thereof. Preference is given to the co-expression of the cytokine or cytokines, either by the same vector as that expressing the immunogen or immunogens or epitope(s) thereof, or by a separate vector therefor.

20 The invention comprehends preparing such combination compositions; for instance by admixing the active components, advantageously together and with an adjuvant, carrier, cytokine, and/or diluent.

Cytokines that may be used in the present invention include, but are not limited to, granulocyte colony stimulating factor (G-CSF), granulocyte/macrophage colony stimulating factor (GM-CSF), interferon α (IFN α), interferon β (IFN β), interferon γ, (IFN γ), interleukin-1α (IL-1 α), interleukin-1 β (IL-1 β), interleukin-2 (IL-2), interleukin-3 (IL-3), interleukin-4 (IL-4), interleukin-5 (IL-5), interleukin-6 (IL-6), interleukin-7 (IL-7), interleukin-8 (IL-8), interleukin-9 (IL-9), interleukin-10 (IL-10), interleukin-11 (IL-11), interleukin-12 (IL-12), tumor necrosis factor α (TNF α), tumor necrosis factor β (TNF β), and transforming growth factor β (TGF β). It is understood that cytokines can be co-administered and/or sequentially administered with the immunogenic or vaccine composition

of the present invention. Thus, for instance, the vaccine of the instant invention can also contain an exogenous nucleic acid molecule that expresses in vivo a suitable cytokine, e.g., a cytokine matched to this host to be vaccinated or in which an immunological response is to be elicited (for instance, a canine cytokine for preparations to be administered to dogs).

5 Advantageously, the pharmaceutical and/or therapeutic compositions and/or formulations according to the invention comprise or consist essentially of or consist of an effective quantity to elicit a therapeutic response of one or more expression vectors and/or polypeptides as discussed herein; and, an effective quantity can be determined from this disclosure, including the documents incorporated herein, and the knowledge in the art,
10 without undue experimentation.

The recombinant viral vector immunogenic compositions and the recombinant viral vector vaccines according to the invention may be freeze-dried advantageously with a stabiliser. Freeze-drying can be done according to well-known standard freeze-drying procedures. The pharmaceutically or veterinary acceptable stabilisers may be carbohydrates
15 (e.g. sorbitol, mannitol, lactose, sucrose, glucose, dextran, trehalose), sodium glutamate (Tsvetkov T et al., *Cryobiology* 1983, 20(3): 318-23; Israeli E et al., *Cryobiology* 1993, 30(5): 519-23), proteins such as peptone, albumin, lactalbumin or casein, protein containing agents such as skimmed milk (Mills C K *et al.*, *Cryobiology* 1988, 25(2): 148-52; Wolff E *et al.*, *Cryobiology* 1990, 27(5): 569-75), and buffers (e.g. phosphate buffer, alkaline metal
20 phosphate buffer). An adjuvant may be used to make soluble the freeze-dried preparations.

Further the present invention concerns the use of at least one polypeptide having an amino acid sequence as shown in SEQ ID NOS: 18, 20, 22, 24, 26, 48, 50 and 52, or analogs thereof or fragments thereof, for the treatment and/or vaccination of mammals against streptococcal infection, notably of equine, canine and human species. In one embodiment,
25 these fragments are selected from the group consisting of sequences SEQ ID NOS: 32, 34, 36, 38, 40, 54, 56 and 58, or analogs thereof. A particular embodiment concerns the use of at least one polypeptide selected from the group consisting of sequences SEQ ID NOS: 18, 20, 22, 24, 26, 32, 34, 36, 38, 40, 48, 50, 52, 54, 56 and 58, or analogs thereof, for the treatment and/or vaccination of equines against strangles disease. A preferred embodiment concerns the
30 use of at least one polypeptide selected from the group consisting of sequences SEQ ID NOS: 32, 34, 36, 38, 40, 54, 56 and 58, or analogs thereof, for the treatment and/or vaccination of equines against strangles disease.

A further embodiment concerns the use of at least one polypeptide having an amino acid sequence as shown in SEQ ID NOS: 18, 20, 22, 24, 26, 48, 50 and 52, or analogs thereof

or fragments thereof, for the preparation of a subunit vaccine protecting equine against *Streptococcus equi* infection. This further embodiment concerns preferably the use of at least one polypeptide having an amino acid sequence as shown in SEQ ID NOS: 32, 34, 36, 38, 40, 54, 56 and 58, or analogs thereof, for the preparation of a subunit vaccine protecting equine
5 against *Streptococcus equi* infection.

Another further embodiment concerns the use of at least one recombinant vector and of at least one polynucleotide inserted therein coding for a polypeptide having an amino acid sequence as shown in SEQ ID NOS: 18, 20, 22, 24, 26, 48, 50 and 52, or an analog thereof or a fragment thereof, and said vector is able to express *in vivo* this polypeptide in a mammal
10 susceptible to streptococcal infection, for the preparation of a recombinant vaccine protecting equine against *Streptococcus equi* infection. This further embodiment concerns preferably the use of at least one recombinant vector and of at least one polynucleotide inserted therein coding for a polypeptide having an amino acid sequence as shown in SEQ ID NOS: 32, 34, 36, 38, 40, 54, 56 and 58, or an analog thereof, and said vector is able to express *in vivo* this
15 polypeptide in a mammal susceptible to streptococcal infection, for the preparation of a recombinant vaccine protecting equine against *Streptococcus equi* infection.

An additional embodiment concerns the use of at least one recombinant vector and at least one polynucleotide inserted therein, wherein said polynucleotide has a nucleotide sequence as shown in SEQ ID NOS: 17, 19, 21, 23, 25, 31, 33, 35, 37, 39, 47, 49, 51, 53, 55
20 and 57, or an analog thereof, and said vector is able to express *in vivo* the polypeptide encoded by said polynucleotide in a mammal susceptible to streptococcal infection, for the preparation of a recombinant vaccine protecting equine against *Streptococcus equi* infection.

In particular, combinations of polypeptides to be used for the treatment and/or vaccination of equine against strangles disease are combination of at least one polypeptide
25 selected from a first group of polypeptides consisting of SEQ ID NOS: 18, 20, 22, 24, 26, 32, 34, 36, 38, 40, 48, 50, 52, 54, 56 and 58, or analogs thereof or combinations thereof, and at least one another *Streptococcus equi* immunogen, which is not present in this first group, notably at least one immunogen selected from a second group of polypeptides consisting of FNZ protein, EAG protein, SFS protein, SEC protein, SFSC1 fragment, FNZN fragment, SEC2.16 fragment, SEC1.18 fragment, SclC1 fragment (WO-A-2004/032957) and SEQ ID
30 NOS: 28, 30 or analogs thereof or fragments thereof, or combinations thereof. In preferred combinations of polypeptides, this first group consists of SEQ ID NOS: 32, 34, 36, 38, 40, 54, 56 and 58, or analogs thereof or combinations thereof. Further the present invention also concerns the use of these combinations of peptides for the preparation of subunit vaccines

protecting equines against *Streptococcus equi* infection. In these embodiments, the use of polypeptides may be replaced by the use of recombinant expression vectors according to the present invention comprising at least one polynucleotide encoding said polypeptides. In particular, at least one recombinant expression vector comprising at least one polynucleotide encoding at least one polypeptide selected from a first group of polypeptides consisting of SEQ ID NOS: 18, 20, 22, 24, 26, 32, 34, 36, 38, 40, 48, 50, 52, 54, 56 and 58, or analogs thereof, and at least one polynucleotide encoding at least one another *Streptococcus equi* immunogen, which is not present in this first group, notably at least one immunogen selected from a second group of polypeptides consisting of FNZ protein, EAG protein, SFS protein, SEC protein, SFSC1 fragment, FNZN fragment, SEC2.16 fragment, SEC1.18 fragment, SclC1 fragment (WO-A-2004/032957) and SEQ ID NOS: 28, 30 or analogs thereof or fragments thereof, or combinations thereof, are used for the treatment and/or vaccination of equines against strangles disease. In a preferred embodiment, this first group consists of SEQ ID NOS: 32, 34, 36, 38, 40, 54, 56 and 58, or analogs thereof or combinations thereof.

Further the present invention also concerns the use of these recombinant expression vectors for the preparation of recombinant vaccines protecting equines against *Streptococcus equi* infection.

In a particular embodiment, the combinations of polypeptides to be used for the treatment and/or vaccination of equine against strangles disease are combination of polypeptides SEQ ID NOS: 52, 48, 22 and 26, or analogs thereof, or fragments thereof. In a more preferred embodiment, the combinations of polypeptides to be used for the treatment and/or vaccination of equine against strangles disease are combination of polypeptides SEQ ID NOS: 58, 54, 36 and 40, or analogs thereof.

In a particular embodiment, the combinations of polypeptides to be used for the treatment and/or vaccination of equine against strangles disease are combination of polypeptides SEQ ID NOS: 52, 48, 22, 26 and 50, or analogs thereof, or fragments thereof. In a more preferred embodiment, the combinations of polypeptides to be used for the treatment and/or vaccination of equine against strangles disease are combination of polypeptides SEQ ID NOS: 58, 54, 36, 40 and 56, or analogs thereof.

In a particular embodiment, the combinations of polypeptides to be used for the treatment and/or vaccination of equine against strangles disease are combination of polypeptides SEQ ID NOS: 52, 48, 22, 26 and 20, or analogs thereof, or fragments thereof. In a more preferred embodiment, the combinations of polypeptides to be used for the treatment

and/or vaccination of equine against strangles disease are combination of polypeptides SEQ ID NOS: 58, 54, 36, 40 and 34, or analogs thereof.

In a particular embodiment, the combinations of polypeptides to be used for the treatment and/or vaccination of equine against strangles disease are combination of polypeptides SEQ ID NOS: 52, 48, 22, 26 and 18, or analogs thereof, or fragments thereof. In a more preferred embodiment, the combinations of polypeptides to be used for the treatment and/or vaccination of equine against strangles disease are combination of polypeptides SEQ ID NOS: 58, 54, 36, 40 and 32, or analogs thereof.

Further the present invention relates to methods to immunise against or to prevent streptococcal infection in mammals, preferably in equine, canine and in human species. According to these methods, (1) a subunit immunogenic composition or vaccine of the present invention, or (2) a recombinant immunogenic composition or vaccine of the present invention, or their combinations, are administered. Of course, embodiments of the invention may be employed with other vaccines or immunogenic compositions that are not of the invention, e.g., in prime-boost processes, such as where a vaccine or immunogenic composition of the invention is administered first and a different vaccine or immunogenic composition is administered thereafter, or *vice versa*. Particular prime-boost processes may be that a subunit vaccine or immunogenic composition of the invention is administered first and a recombinant vaccine or immunogenic composition of the invention is administered thereafter, or *vice versa*.

The administration may be notably made by intramuscular (IM), intradermal (ID), subcutaneous (SC) or transdermal injection or via intranasal, intratracheal, oral administration, or through the ears or the lip. The immunogenic composition or the vaccine according to the invention is administered by syringe, a syringe with a microneedle (i.e. BD™ Intradermal Delivery System of Becton, Dickinson and Company, Franklin Lakes, NJ, USA), needleless apparatus (like for example Pigjet, Avijet, Dermojet or Biojector (Bioject, Oregon, USA), see US-A-2006/0034867) or a spray. The administration is preferably made by IM injection with a syringe, or by transdermal injection with a needleless apparatus or with a syringe with a microneedle (i.e. BD™ Intradermal Delivery System), or by intranasal or oral administration with a spray, i.e. a liquid nebulisation of a vaccine of the invention, or by oral or nasal administration of a micronized powder of a freeze-dried vaccine according to the invention.

The quantity of immunogenic compositions or vaccines can be determined and optimised by the skilled person, without undue experimentation from this disclosure and the

knowledge in the art. Generally an animal (including a human) may be administered approximately 10^4 - 10^9 CFUs, advantageously approximately 10^5 - 10^8 CFUs and more advantageously approximately 10^6 - 10^8 CFUs in a single dosage unit of recombinant viral immunogenic compositions or vaccines of the present invention; approximately 10ng-1mg, advantageously approximately 100ng-500 μ g and more advantageously approximately 1 μ g-250 μ g per plasmid type in a single dosage unit of recombinant DNA immunogenic compositions or vaccines of the present invention; approximately 5 μ g-1mg, advantageously approximately 50 μ g-500 μ g and more advantageously approximately 100 μ g-200 μ g in a single dosage unit of subunit immunogenic compositions or vaccines of the present invention.

In the case of therapeutic and/or pharmaceutical compositions based on a plasmid vector, a dose can comprise, consist essentially of or consist of, in general terms, about in 1 μ g to about 2000 μ g, advantageously about 50 μ g to about 1000 μ g and more advantageously from about 100 μ g to about 800 μ g of plasmid expressing the antigen, epitope, immunogen, peptide or polypeptide of interest. When the therapeutic and/or pharmaceutical compositions based on a plasmid vector is administered with electroporation the dose of plasmid is generally between about 0.1 μ g and 1mg, advantageously between about 1 μ g and 100 μ g, advantageously between about 2 μ g and 50 μ g. The dose volumes can be between about 0.1 and about 2 ml, advantageously between about 0.2 and about 1 ml. These doses and dose volumes are suitable for the treatment mammalian target species.

The volume of one single dosage unit by syringe can be between about 0.2 ml and about 5.0 ml and advantageously between about 0.5 ml and about 2.0 ml and more advantageously about 1.0 ml. The volume of one single dosage unit by needleless apparatus can be between about 0.1 ml and about 1.0 ml and advantageously between about 0.2 ml and about 0.5 ml. The volume of one single dosage unit by liquid spray can be between about 2.0 ml and about 10.0 ml and advantageously about 5.0 ml (for powder spray, the quantities administered are corresponding to the equivalent volumes).

In a particular method, foals, that is to say horses of 2 to 6 months old and preferably 3 to 4 months old, are vaccinated with subunit vaccines of the present invention, adjuvanted with CTB and/or CTA and/or Carbopol®, via the intranasal or oral route.

In another particular method, foals are vaccinated with recombinant viral vaccines of the present invention, adjuvanted or not with Carbopol®, via the oral route with a spray and liquid nebulisation. Preferably, the viral vectors of these recombinant viral vaccines are EHV-4 or EHV-2, or EHV-1.

In another particular method, foals are vaccinated with recombinant viral vaccines of the present invention, via the lips with a syringe. Preferably, the viral vectors of these recombinant viral vaccines are EHV-4 or EHV-2 or EHV-1.

Preferably for the vaccination of horses and mares, two administrations of subunit
5 vaccines of the present invention, adjuvanted with Carbopol® and/or CpG, and or emulsion are made by IM injection with a syringe. Boost administrations may be injected every 6 months or annually. For pregnant mares, a boost administration may be injected 2 to 4 weeks before the expected foaling date.

Polypeptides of the invention and fragments thereof may also be used in therapy.

10 The polypeptides and fragments may also be used as reagents in antibody-antigen reactions. Accordingly, another aspect of the invention is thus a diagnostic method and/or kit for detecting infection by the streptococcal bacterium. Kits, e.g. ELISA, can include at least one polypeptide or fragment according to the invention (e.g., at least one polypeptide identified by sequence herein or a fragment thereof as herein discussed).

15 Antibodies against the herein polypeptides or fragments (e.g., polypeptides identified by sequence herein or fragments thereof as herein discussed) can be used as a diagnostic reagent or in passive immunization or vaccination or in therapy. The amounts of antibody administered in passive immunization can be the same as or analogous to amounts used in the art, such that from the knowledge in the art, the skilled artisan can practice passive
20 immunization without undue experimentation.

Proteins encoded by the novel viruses of the present invention, or their fragments, can be used to produce antibodies, both polyclonal and monoclonal. If polyclonal antibodies are desired, a selected mammal, (e.g., mouse, rabbit, goat, horse, etc.) is immunized with an antigen of the present invention, or its fragment, or a mutated antigen. Serum from the
25 immunized animal is collected and treated according to known procedures. See, e.g., Jurgens et al. (1985) J. Chrom. 348:363-370. If serum containing polyclonal antibodies is used, the polyclonal antibodies can be purified by immunoaffinity chromatography, using known procedures.

Another aspect of the invention is an antibody preparation comprising an antibody
30 specific to a polypeptide or a fragment according to the invention and methods of diagnosis using the same. With respect to an antibody specific to a polypeptide, it is meant that the antibody binds preferentially to the polypeptide, e.g., the antibody binds to the polypeptide and not to other polypeptides or has a specificity to the polypeptide that is acceptably particular to the polypeptide such that the antibody can be used to isolate the polypeptide

from a sample or detect its presence in a sample with no more than 5% false positives, using techniques known in the art or discussed in documents cited herein, including Sambrook, *infra*.

Antibodies can be polyclonal or monoclonal.

5 If polyclonal antibodies are desired, a selected animal (e.g. mouse, rabbit, goat, horse, etc.) is immunized with a polypeptide or a fragment. Serum from the immunized animal is collected and treated according to known procedures and possibly purified. See, e.g. Jurgens et al. J. Chrom., 1985, 348: 363-370.

10 Monoclonal antibodies to the proteins and to the fragments thereof, can also be readily produced by one skilled in the art. The general methodology for making monoclonal antibodies by using hybridoma technology is well known. Immortal antibody-producing cell lines can be created by cell fusion, and also by other techniques such as direct transformation of B lymphocytes with oncogenic DNA, or transfection with Epstein-Barr virus. See, e.g., M. Schreier et al., *Hybridoma Techniques* (1980); Hammerling et al., *Monoclonal Antibodies and T-cell Hybridomas* (1981); Kennett et al., *Monoclonal Antibodies* (1980); see also U.S. Pat. Nos. 4,341,761; 4,399,121; 4,427,783; 4,444,887; 4,452,570; 4,466,917; 4,472,500, 4,491,632; and 4,493,890. Panels of monoclonal antibodies produced against the desired protein, or fragment thereof, can be screened for various properties; i.e., for isotype, epitope, affinity, etc. Monoclonal antibodies are useful in purification, using immunoaffinity
15 20 techniques, of the individual antigens which they are directed against. Both polyclonal and monoclonal antibodies can also be used for passive immunization or can be combined with subunit vaccine preparations to enhance the immune response. Polyclonal and monoclonal antibodies are also useful for diagnostic purposes.

One embodiment of the invention is a method of eliciting an immune response against
25 the antigen, epitope, immunogen, peptide or polypeptide of interest in an animal, comprising administering a formulation for delivery and expression of a recombinant vaccine in an effective amount for eliciting an immune response. Still another embodiment of the invention is a method of inducing an immunological or protective response in an animal, comprising administering to the animal an effective amount of a formulation for delivery and
30 expression of an antigen, epitope, immunogen, peptide or polypeptide of interest wherein the formulation comprises a recombinant vaccine and a pharmaceutically or veterinarily acceptable carrier, vehicle or excipient.

The invention relates to a method to elicit, induce or stimulate the immune response of an animal, advantageously a mammal or a vertebrate.

Another embodiment of the invention is a kit for performing a method of inducing an immunological or protective response against an antigen, epitope, immunogen, peptide or polypeptide of interest in an animal comprising a recombinant vaccine and instructions for performing the method of delivery in an effective amount for eliciting an immune response in the animal.

The invention will now be further described by way of the following non-limiting examples.

EXAMPLES

Example 1: ELISA study of *Streptococcus equi* polypeptides

The IgG antibody responses directed towards *Streptococcus equi* cell surface proteins in sera taken from ponies intranasally challenged with 4 doses of *Streptococcus equi* strain 4047 were determined by ELISA. These *Streptococcus equi* proteins were obtained by *in vitro* expression in *Escherichia coli*. The antibody responses in sera from the six ponies receiving the higher doses of 3×10^7 and 3×10^5 cfu were averaged and compared to those from ponies challenged with 3000 and 30 cfu of *Streptococcus equi* 4047. Ponies receiving the higher doses developed strangles. Those receiving the low doses remained healthy throughout the study. On the basis of the results of this study, polypeptides of *Streptococcus equi* were selected for expression in *Escherichia coli* and for test of efficacy in mice and in ponies.

Example 2: Expression of recombinant *Streptococcus equi* polypeptides in *Escherichia coli*

Polynucleotides encoding fragments of Se50, Se1459 and Se358 were produced using Vent DNA polymerase (New England Biolabs® Inc.) as per the manufacturers recommended conditions with a 5 minute denaturation step followed by 35 cycles using 30 seconds annealing at 50-55°C, 1 minute extension at 72°C, and 30 second denaturing at 95°C.

Products were finished off with a 4-minute 72°C extension. The template in each case was *Streptococcus equi* strain 4047 chromosomal DNA. The fragments of Se50, Se1459 and Se358 are those designated as SEQ ID NOS: 32, 34 and 40, respectively.

Polynucleotides encoding fragments of Se528 and Se595 were produced using the pfu polymerase with same conditions as above apart from the extension time extended to two minutes at 72°C. The fragments of Se528 and Se595 are those designated as SEQ ID NOS: 38 and 36, respectively.

Table 1: Primers used to clone and sequence recombinant subunits.

Primer name	Sequence	Function	SEQ ID NO:
5'50GST	AAAGGATCCTACAGCCAGCAGCACTAAAA	Se50 cloning	1
3'50GST	AAAGAATTCTTGGGCAGCTTCTTCTA	Se50 cloning	2
5'1459GST	TTTGGATCCTATCGGAACCCAATCCATAT	Se1459 cloning	3
3'1459GST	AAAGAATTCATTTTAAAGTTCAAACCTC	Se1459 cloning	4
5'595GST	AAAGGATCCTAGCTACCCTCATCACAGGA	Se595 cloning	5
3'595GST	TTTGAATTCTTGTTTTTTTAGGTGTTGC	Se595 cloning	6
5'528 LONG	TTTGGATCCTAGAGGTAGTTGAAGTTTGGCCTAATGGG	Se528 cloning	7
3'528 LONG	AAAGAATTCTTTTTCTGTCTTGTTGAAGTAATCTGCCC	Se528 cloning	8
ASW49	GACGGATCCCCTCTAATGGTGGAAACAAAGGAAGC	Se358 cloning	9
ASW50*	GACTGAATTCAAACAAAGCACAAACACCAG	Se358 cloning	10
5'pGEX	GGGCTGCAAGCCACGTTTGGTG	Sequencing	11
3'pGEX	CCGGGAGCTGCATGTGTCAGAGG	Sequencing	12
595 seq1	AAAGAGCTAATCTGGCAG	Sequencing Se595	13
595 seq2	CTCAATTGGAGGCTACAG	Sequencing Se595	14
595 seq3	TGACGCTTCCAAAGCAAG	Sequencing Se595	15
595 seq4	CTCCGGAAACACCAAAGG	Sequencing Se595	16

* The recombinant Se358 ends at the actual stop codon for Se358 as the ASW50 primer site is located down stream from the gene.

All PCR products were purified using the Qiaquick PCR purification kit (Qiagen) with a 32µl elution volume, then digested with the addition of EcoRI buffer (3.5µl) (New England Biolabs® Inc.), BamHI (0.75µl) and EcoRI (0.75µl) for 2 hours at 37°C. 1µg of pGEX-3X plasmid (GE Healthcare) was also digested with EcoRI and BamHI as described above. The pGEX-3X plasmids have a multiple cloning site, a tac promoter, glutathione S-transferase gene (GST) and stop codons in all three frames, for the production of GST fusion proteins and their isolation and purification.

All digestion products were run on 0.7% TAE agarose gels, the desired restriction product excised and purified using the Qiaquick gel extraction kit (Qiagen) with a 35µl final elution.

Ligation reactions were set up using 16.5µl of purified digested PCR product and 1µl of pGEX-3X plasmid with 2µl of ligase buffer and 0.5µl of ligase, and incubated overnight at room temperature (rtp).

The ligation reactions were transformed into *E.coli* DH10B strain and plated onto L-agar plates supplemented with 50µg/ml ampicillin and incubated overnight at 37°C.

Transformants were tested for the presence of insert by colony PCR. A single colony was resuspended in 4µl of water and 2µl of this was added to a Taq DNA polymerase PCR reaction containing the appropriate cloning primers for the polynucleotide of interest and a 35 cycle extension was performed as described above (1 minute for Se50, Se1459 and Se358 and 2 minutes for Se595 and Se528). The presence of the desired insert was confirmed by running the PCR reaction on 0.7% TAE agarose gels.

Plasmids of each of the clones were prepared and sequenced using the 5' and 3' pGEX-3X PCR primers for all of the clones (Table 1). Due to its size, Se595 was also sequenced using the additional primers listed above (Table 1). All of the cloned polynucleotides were confirmed to have the same sequence as the assembled *Streptococcus equi* strain 4047 genome sequence (http://www.sanger.ac.uk/Projects/S_equi/).

Clones of Se1459, Se595, Se528, Se50 and Se358 were induced for expression using the following protocol. A single colony was inoculated into 50ml 2 x YT culture medium containing 50µg/ml ampicillin and was grown overnight with shaking at 220 rpm at 37°C. This culture was then added to 500ml of pre-warmed (37°C) and pre-gassed 2 x YT and incubated for 1 hour at 37°C.

For all cultures, after 1 hour, 200µl was removed and the cells were collected for analysis (this being the pre-induction sample), IPTG (isopropyl-beta-D-thiogalactopyranoside) was added to a final concentration of 0.1mM, and the temperature was lowered to 28°C. The culture was induced for a further four hours and a 100µl sample was removed for analysis, after which the cells were harvested by centrifugation and stored at –70°C.

All samples were analysed using 10% SDS-PAGE.

For Se1459, Se595, Se528, Se50 and Se358, the pellets were resuspended in 20ml of ice cold PBS (Oxoid Limited, catalog # BR0014G), PMSF (Phenylmethyl sulphonyl fluoride) was added to a final concentration of 0.1mM and the cells were lysed by sonication on ice

with 6x10 second pulses with a 10 second rest between each pulse. Triton X-100 was added to 0.1% (v/v) and the insoluble and intact cells were removed by centrifugation (11,000g, 5 min 4°C). Samples of the soluble bacterial lysate and insoluble material were analysed by SDS-PAGE. DTT (dithiothreitol; Cleland's reagent) (5mM) was added and 1.3ml of
5 glutathione sepharose 4B was added (equates to 1ml of beads, these were pre-washed three times with 20 ml PBS). The mixture was incubated at room temperature for 1 hour and then the beads were collected by centrifugation (800g, 2 min, rtp) and washed three times with 50 ml of ice cold PBST (PBS with 0.1% v/v Triton X-100). Samples of each wash (washes 1-3) were taken for analysis by SDS-PAGE as was a sample of the fusion protein attached to the
10 beads. All samples were analysed using 10% SDS-PAGE.

Removal of the GST carrier was done for all fusion proteins by protease cleavage.

For Se1459, Se595, Se528, Se50 and Se358, cleavage reactions were performed with the fusion protein attached to the glutathione sepharose beads. The beads were first equilibrated once with 20ml of wash buffer (Tris-HCL, 50 mM, pH 7.5, NaCl, 150 mM) and
15 then twice with 20ml cleavage buffer (wash buffer with CaCl₂, 1.0mM) and finally resuspended in 1ml of cleavage buffer. Factor Xa (New England Biolabs® Inc.) (50 units) was added and incubated on a rotary mixer for 16 hours at room temperature. The released protein was recovered by pelleting the glutathione beads (800g, 2 min, rtp) and retaining the supernatant. The beads were washed twice with 1ml of wash buffer and the supernatant was
20 retained each time. Aliquots of each supernatant and the remaining glutathione beads were analysed by SDS-PAGE. The beads were washed three times with wash buffer 2 (Tris-HCL, 50 mM, pH8.0) and eluted with 3ml of wash buffer 2 containing 5.0mM reduced glutathione for 2 min at room temperature followed by pelleting the beads (800g, 2 min, rtp) and retaining the supernatant. This elution was repeated twice and the purity of each fraction was
25 analysed by SDS-PAGE.

All the proteins were dialysed twice in 5 litres of 20mM sodium phosphate buffer pH7.4 at 4°C overnight.

After dialysis, any insoluble material was removed by centrifugation (10000g, 2 min, 4°C), and then it was stored at -70°C.

30 **Example 3: Protection in mice after subunit vaccine administration and *Streptococcus equi* challenge**

The objective of this study was to determine if vaccination with *Streptococcus equi* subunit vaccine protected mice from subsequent intranasal *Streptococcus equi* challenge.

Five *Streptococcus equi* subunit vaccines were used, Se595, Se358, Se1459, Se50 and Se528. All these vaccines used the *Streptococcus equi* proteins obtained in Example 2. The Se595, Se358, Se1459, Se50 and Se528 recombinant *Streptococcus equi* proteins were purified by buffer exchanged into 20mM Sodium Phosphate pH7.4 and concentrated to give a concentration equivalent to the molar concentration of Se358 at 10µg/ml where possible. Proteins, Se595 and Se528 were insoluble at the desired concentration. The actual protein concentrations used are shown in Table 2.

Table 2: Actual doses of recombinant subunits administered

Protein	Dose administered	Dose equivalent to Se358
Se595	11.2µg	5.3µg
Se358	10µg	10µg
Se1459	1.7µg	10µg
Se50	1.9µg	9.6µg
Se528	4.2µg	3.4µg

Purified proteins in 20mM Sodium Phosphate pH 7.4, were diluted 1:3 in TS6 adjuvant (example 1 of US-A-2005/0079185) prior to administration.

Groups of 20 3-4 week old female BalbC mice were vaccinated on day 0 (D0) with each of five *Streptococcus equi* subunit vaccines. Vaccinations were given by intramuscular injection in the thigh and intranasally. These mice were given a booster vaccination after 4 weeks (D28) and challenged with *Streptococcus equi* after a further 2 weeks (D42). An additional control group receiving no vaccination was also included, and was challenged on D42. A group of 5 control mice was not vaccinated or challenged with *Streptococcus equi* (see Table 3).

The group size is sufficient to detect 50% protection of vaccinated mice with 80% infection of controls. Blood was taken before vaccination, before boost, before challenge and on euthanasia to determine the level of antibody response to vaccination and challenge.

Streptococcus equi strain 4047 was grown for challenge in Todd Hewitt broth (THB) supplemented with 10% foetal calf serum (FCS). An overnight culture was diluted 1:20 to inoculate fresh pre-warmed and gassed THB+10% FCS and the culture grown until an OD_{600nm} of 0.3 was reached. At this point, dilutions of the culture were made. Following preparation of the required dose of *Streptococcus equi* strain 4047, mice were sedated with 100mg/kg of Ketaset by intramuscular injection and challenged by intranasal administration of 10µl of challenge inocula containing approximately 1x10⁴ cfu *S. equi*.

Immediately after challenge, the dose administered was quantified by back titration and was found to contain 0.9x10⁴ cfu.

All the mice were carefully monitored to ensure complete recovery from sedation and checked regularly for the onset of clinical signs of disease. These included malaise, shivering or staring coat and were taken as a sign of moderate effect with the onset of more severe effects possible. Mice were finally euthanased.

5 Clinical observations were recorded daily throughout the study and mice were weighed pre vaccination, pre boost, pre challenge and then daily post challenge. Nasal dabs were taken from mice post challenge to quantify the level of *Streptococcus equi* shedding. A post mortem examination was carried out on all mice.

Table 3: Summary of study protocol

# of mice	Age of mice	Vaccine Subunit	Dose	Booster	Challenge dose (cfu)	Route
20	3-4 weeks	Se595 + adjuvant	20µl IN + IM	Yes	1x10 ⁴	10µl IN
20	3-4 weeks	Se358 + adjuvant	20µl IN + IM	Yes	1x10 ⁴	10µl IN
20	3-4 weeks	Se1459 + Adjuvant	20µl IN + IM	Yes	1x10 ⁴	10µl IN
20	3-4 weeks	Se50 + adjuvant	20µl IN + IM	Yes	1x10 ⁴	10µl IN
20	3-4 weeks	Se528 + adjuvant	20µl IN + IM	Yes	1x10 ⁴	10µl IN
60	3-4 weeks	No vaccine	No vaccine	No	1x10 ⁴	10µl IN
5	3-4 weeks	None	None	No	none	10µl IN

10 The vaccination phase proceeded without incident, all groups gained weight in line with controls. No adverse injection site reactions were identified. All mice seroconverted with respect to the vaccination administered.

Following challenge, Groups vaccinated with Se595, Se358, Se1459, Se528 and Se50 generally had reduced clinical signs of disease, reduced weight loss, more weight gain, reduced nasal colonisation and improved survival when compared with controls. Each mouse was examined on the day of challenge and on each of the following 7 days for the occurrence of symptoms associated with *Streptococcus equi* (lethargy, weight loss, sneezing, piloerection, hunched up, inner ear imbalance). Mice were examined at least 4 times a day. The following recording and scoring method was applied (Table 4)

20 Table 4: Strangles scoring method

Observation	Score
Normal	0
Slight piloerection	1
Piloerection	3

Observation	Score
Quiet	3
Marked sneezing	3
Hunched	5
Rapid breathing	5
Ear imbalance	5
Immobile	10

For example: A mouse with piloerection that is quiet, hunched and immobile would score $3 + 3 + 5 + 10 = 21$.

Weights of mice were analysed on a spreadsheet each day to determine %weight loss relative to the day of challenge.

5 Following challenge, the nose of each mouse was touched onto a clearly labelled Streptococcal selective agar plate each day (bioMerieux). Plates were streaked to determine if viable *S. equi* bacteria were present. The level of *S. equi* shedding was graded according to the following 4 point scoring system; 0 = no growth, 1 = 1-10 colonies present, 2 = 11-100 colonies present, 3 = greater than 100 colonies and 4 = confluent growth.

10 Blood samples were taken from the thigh vein. Serum was prepared and stored frozen at -20°C or below until use.

Clinical signs of disease were observed from D44, 2 days post challenge. The vaccinated groups generally had lower clinical signs than the control groups. Mice vaccinated with Se50 and Se358 had the lowest level of clinical signs of disease. Those vaccinated with
15 Se595 had clinical signs of disease for a reduced period compared with controls (FIG. 1). On examining the mean percentage change in mass following challenge, relative to the day of challenge (D42), it was apparent that Groups vaccinated with Se528, Se1459 and Se50 had a slightly reduced weight loss. However, Se358 and Se595 vaccinated mice appeared best protected (FIG. 2).

20 As expected, the mean number of β -haemolytic bacteria isolated from nasal dabs increased following challenge, peaking at D45 (3 days post challenge). On D45, the lowest nasal dabs scores were seen in groups vaccinated with Se358, Se595, Se50 and Se1459. The vaccinated groups generally had lower scores than the control groups. The exception being Se358 vaccinated mice, which peaked on D46 (FIG. 3).

25 **Table 5: IgG Serological responses**

	Sera dilution in ELISA	Seroconversion
Se595	1:80000*	All mice seroconverted
Se358	1:160000*	All mice seroconverted
Se1459	1:100**	All mice seroconverted
Se50	1:160000*	All mice seroconverted
Se528	1:3000*	All mice seroconverted

*Sera dilution to achieve an ELISA reading of around 1.5 in preliminary studies.

**Sera dilution to achieve an ELISA reading of around 0.3 in preliminary studies.

Example 4: Expression of recombinant *Streptococcus equi* Se1631, Se1681 and Sla polypeptides in *Escherichia coli*

Polynucleotides encoding fragments of Se1631, Se1681 and Sla were produced using Vent DNA polymerase (New England Biolabs® Inc.) as per the manufacturers recommended conditions with a 5 minute denaturation step followed by 35 cycles using 30 seconds annealing at 50-55°C, 1 minute extension at 72°C, and 30 second denaturing at 95°C.

Products were finished off with a 4-minute 72°C extension. The template in each case was *Streptococcus equi* strain 4047 chromosomal DNA.

The fragments of Se1631, Se1681 and Sla are those designated as SEQ ID NOS: 53, 55 and 57, respectively.

Table 6: Primers used to clone and sequence recombinant subunits.

Primer name	Sequence	Function	SEQ ID NO:
5'1631GST	AAAGGATCCTGCAGTCTACAAATGACAATAC	Se1631 cloning	41
3'1631GST	AAAGAATTCTTATTTTTCTGACTTAGATTTAGAAG	Se1631 cloning	42
5'1681GST	AAAGGATCCTGGCGACTACCCTAGCAGGACAAAC	Se1681 cloning	43
3'1681GST	AAAGAATTCTTATGTATCTTGACAGTGCTTAAG	Se1681 cloning	44
5'SLAGST	CCCGGATCCTAGAAGGGATAAATGATAAAATAG	Sla cloning	45
3'SLAGST	TTTGAATTCTTACAAAACATCTACTACTGGCAAC	Sla cloning	46

All PCR products were purified using the Qiaquick PCR purification kit (Qiagen) with a 32µl elution volume, then digested with the addition of EcoRI buffer (3.5µl) (New England Biolabs® Inc.), BamHI (0.75µl) and EcoRI (0.75µl) for 2 hours at 37°C. 1µg of pGEX-3X plasmid (GE Healthcare) was also digested with EcoRI and BamHI as described above. The pGEX-3X plasmids have a multiple cloning site, a tac promoter, glutathione S-transferase gene (GST) and stop codons in all three frames, for the production of GST fusion proteins and their isolation and purification.

All digestion products were run on 0.7% TAE agarose gels, the desired restriction product excised and purified using the Qiaquick gel extraction kit (Qiagen) with a 35µl final elution.

Ligation reactions were set up using 16.5µl of purified digested PCR product and 1µl of pGEX-3X plasmid with 2µl of ligase buffer and 0.5µl of ligase, and incubated overnight at room temperature (rtp).

The ligation reactions were transformed into *E.coli* DH10B strain and plated onto L-agar plates supplemented with 50µg/ml ampicillin and incubated overnight at 37°C.

Transformants were tested for the presence of insert by colony PCR. A single colony was resuspended in 4µl of water and 2µl of this was added to a Taq DNA polymerase PCR reaction containing the appropriate cloning primers for the polynucleotide of interest and a 35 cycle extension was performed as described above. The presence of the desired insert was confirmed by running the PCR reaction on 0.7% TAE agarose gels.

Plasmids of each of the clones were prepared and sequenced using the 5' and 3' pGEX-3X PCR primers for all of the clones (Table 6). All of the cloned polynucleotides were confirmed to have the same sequence as the assembled *Streptococcus equi* strain 4047 genome sequence (http://www.sanger.ac.uk/Projects/S_equi/).

All clones were induced for expression using the following protocol. A single colony was inoculated into 50ml 2 x YT culture medium containing 50µg/ml ampicillin and was grown overnight with shaking at 220 rpm at 37°C. This culture was then added to 500ml of pre-warmed (37°C) and pre-gassed 2 x YT and incubated for 1 hour at 37°C. After 1 hour, 200µl was removed and the cells were collected for analysis (this being the pre-induction sample), IPTG (isopropyl-beta-D-thiogalactopyranoside) was added to a final concentration of 0.1mM, and the temperature was lowered to 28°C. The culture was induced for a further four hours and a 100µl sample was removed for analysis, after which the cells were harvested by centrifugation and stored at -70°C.

All samples were analysed using 10% SDS-PAGE.

The pellets were resuspended in 20ml of ice cold PBS (Oxoid Limited, catalog # BR0014G), PMSF (Phenylmethyl sulphonyl fluoride) was added to a final concentration of 0.1mM and the cells were lysed by sonication on ice with 6x10 second pulses with a 10 second rest between each pulse. Triton X-100 was added to 0.1% (v/v) and the insoluble and intact cells were removed by centrifugation (11,000g, 5 min 4°C). Samples of the soluble bacterial lysate and insoluble material were analysed by SDS-PAGE. DTT (dithiothreitol; Cleland's reagent) (5mM) was added and 1.3ml of glutathione sepharose 4B was added (equates to 1ml of beads, these were pre-washed three times with 20 ml PBS). The mixture was incubated at room temperature for 1 hour and then the beads were collected by centrifugation (800g, 2 min, rtp) and washed three times with 50 ml of ice cold PBST (PBS with 0.1% v/v Triton X-100). Samples of each wash (washes 1-3) were taken for analysis by SDS-PAGE as was a sample of the fusion protein attached to the beads. All samples were analysed using 10% SDS-PAGE.

Removal of the GST carrier was done for all fusion proteins by protease cleavage.

Cleavage reactions were performed with the fusion protein attached to the glutathione sepharose beads. The beads were first equilibrated once with 20ml of wash buffer (Tris-HCL, 50 mM, pH 7.5, NaCl, 150 mM) and then twice with 20ml cleavage buffer (wash buffer with CaCl₂, 1.0mM) and finally resuspended in 1ml of cleavage buffer. Factor Xa
 5 (New England Biolabs® Inc.) (50 units) was added and incubated on a rotary mixer for 16 hours at room temperature. The released protein was recovered by pelleting the glutathione beads (800g, 2 min, rtp) and retaining the supernatant. The beads were washed twice with 1ml of wash buffer and the supernatant was retained each time. Aliquots of each supernatant and the remaining glutathione beads were analysed by SDS-PAGE. The beads were washed
 10 three times with wash buffer 2 (Tris-HCL, 50 mM, pH8.0) and eluted with 3ml of wash buffer 2 containing 5.0mM reduced glutathione for 2 min at room temperature followed by pelleting the beads (800g, 2 min, rtp) and retaining the supernatant. This elution was repeated twice and the purity of each fraction was analysed by SDS-PAGE.

The beads were washed three times with wash buffer 2 (Tris-HCL, 50 mM, pH8.0)
 15 and eluted with 3ml of wash buffer 2 containing 5.0 mM reduced glutathione for 2 min at room temperature followed by pelleting the beads (800g, 2 min, rtp) and retaining the supernatant. This elution was repeated twice and the purity of each fraction was analysed by SDS-PAGE.

All the proteins were dialysed twice in 5 litres of 20mM sodium phosphate buffer
 20 pH7.4 at 4°C overnight.

After dialysis, any insoluble material was removed by centrifugation (10000g, 2 min, 4°C), and then it was stored at -70°C.

Example 5: Protection in mice after Se1631, Se1681 or Sla subunit vaccine administration and *Streptococcus equi* challenge

25 Groups of 20 3/4 week old female BalbC mice were vaccinated with each of Se1631, Se 1681 and Sla *S. equi* peptides, and a GST + adjuvant control group. This group size is sufficient to detect 50% protection of mice with 80% infection of controls. An additional control group of 5 mice receiving no vaccination or challenge was included (see Table 7).

30 Three *Streptococcus equi* subunit vaccines were used, Se1631, Se1681358 and Sla. All these vaccines used the *Streptococcus equi* proteins obtained in Example 4.

Recombinant proteins were prepared for vaccination by centricon concentration in 0.02M phosphate buffer pH 7.4 to 1.5 mg/ml and adding 2 parts of TS6 adjuvant (example 1 of US-A-2005/0079185).

Mice were vaccinated on Day 0 and boosted on Day 28. IM vaccinations were administered into the left hind leg on Day 0 and the right hind leg on Day 28 according to the following table. IN vaccinations will be administered via a pipette onto the nares of mice.

- 5 Blood was taken before vaccination, before boost and before challenge to determine the level of antibody response to vaccination.

Mice were challenged with *S. equi* after a further 2 weeks. *S. equi* strain 4047 were prepared as described in example 3. Following dilution to the required dose, 10 μ l were administered to sedated mice by the intranasal route in a contained environment. Mice
10 were carefully monitored to ensure complete recovery from sedation and checked regularly for the onset of clinical signs of disease. These include malaise, shivering or staring coat and were taken as a sign of moderate effect with the onset of more severe effects possible.

Table 7: Summary of study protocol

# of mice	Age of mice	Vaccine Subunit	Dose	Booster	Challenge dose (cfu)	Route
20	3-4 weeks	Se1631 + adjuvant	20 μ l IN + IM	Yes	1x10 ⁴	10 μ l IN
20	3-4 weeks	Se1681 + adjuvant	20 μ l IN + IM	Yes	1x10 ⁴	10 μ l IN
20	3-4 weeks	Sla + adjuvant	20 μ l IN + IM	Yes	1x10 ⁴	10 μ l IN
20	3-4 weeks	GST + adjuvant	20 μ l IN + IM	Yes	1x10 ⁴	10 μ l IN
5	3-4 weeks	None	None	No	none	10 μ l IN

- Clinical observations were recorded daily throughout the study (for the scoring
15 method see Table 4) and mice were weighed pre vaccination, pre boost, pre challenge and then daily post challenge. Nasal dabs were taken from mice post challenge to quantify the level of *S. equi* shedding.

Clinical signs of disease were observed from D44, 2 days post challenge. The vaccinated groups generally had lower clinical signs than the control groups. Mice
20 vaccinated with Se1631 had the lowest level of clinical signs of disease. Those vaccinated with Se1681 and Sla had clinical signs of disease for a reduced period compared with controls (FIG. 4).

On examining the mean percentage change in mass following challenge, relative to the day of challenge (D42), it was apparent that Groups vaccinated with Se1681, Se1631
25 and Sla had a slightly reduced weight loss. On day D49, Sla vaccinated mice appeared less protected (FIG. 5).

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As expected, the mean number of β -haemolytic bacteria isolated from nasal dabs increased following challenge, peaking for the control group at D49 (7 days post challenge). On D49, the lowest nasal dabs scores were seen in groups vaccinated with Sel631 and Sel681. The vaccinated groups generally had lower scores than the control groups. The exception being Sla vaccinated mice, which peaked on D48 (FIG. 6). On examining percentage of mice surviving and not losing more than 7% weight (FIG. 7), it was apparent that Groups vaccinated with Sel631, Sel681 and Sla had a reduced weight loss and a better surviving rate than the control groups.

* * *

10 Having thus described in detail preferred embodiments of the present invention, it is to be understood that the invention defined by the above paragraphs is not to be limited to particular details set forth in the above description as many apparent variations thereof are possible without departing from the scope of the present invention.

SEQUENCE LISTING IN ELECTRONIC FORM

In accordance with Section 111(1) of the Patent Rules, this description contains a sequence listing in electronic form in ASCII text format (file: 51440-108 Seq 17-DEC-08 v1.txt).

A copy of the sequence listing in electronic form is available from the Canadian Intellectual Property Office.

The sequences in the sequence listing in electronic form are reproduced in the following table.

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Leu Gly Gly Ile Asp Lys Lys Val Glu Glu Ala Ala Gln Lys Pro Gly
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Ile Pro Gly Pro Thr Gly Pro Gln Gly Pro Lys Gly Asp Lys Gly Asp
100 105 110
Pro Gly Ala Pro Gly Glu Arg Gly Pro Ala Gly Pro Lys Gly Asp Thr
115 120 125
Gly Glu Ala Gly Pro Arg Gly Glu Gln Gly Pro Ala Gly Gln Ala Gly
130 135 140
Glu Arg Gly Pro Lys Gly Asp Pro Gly Ala Pro Gly Pro Lys Gly Glu
145 150 155 160
Lys Gly Asp Thr Gly Ala Val Gly Pro Lys Gly Glu Lys Gly Asp Thr
165 170 175
Gly Ala Thr Gly Pro Lys Gly Asp Lys Gly Glu Arg Gly Glu Lys Gly
180 185 190
Glu Gln Gly Gln Arg Gly Glu Lys Gly Glu Gln Gly Gln Arg Gly Glu
195 200 205
Lys Gly Glu Gln Lys Pro Lys Gly Asp Gln Gly Lys Asp Thr Lys Pro
210 215 220
Ser Ala Pro Lys Ala Pro Glu Lys Ala Pro Ala Pro Lys Ala Pro Lys
225 230 235 240
Ala Ser Glu Gln Ser Ser Asn Pro Lys Ala Pro Ala Pro Lys Ser Ala
245 250 255
Pro Ser Lys Ser Ala Ala Pro Thr Gly Gln Lys Ala Ala Leu Pro Ala
260 265 270
Thr Gly Glu Ile Asn His Pro Phe Phe Thr Leu Ala Ala Leu Ser Val
275 280 285
Ile Ala Ser Val Gly Val Leu Thr Leu Lys Gly Lys Lys Asp
290 295 300

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<210> 19
 <211> 1092
 <212> DNA
 <213> Streptococcus equi

<400> 19
 atgatgaatc gtaaaaagca taccaaactc attcgtcact acagtatctg ctcagcagtg 60
 gctgtcctag cagcagttag cttgggaaca ggtcaggacg tacaggcttc ggaacccaat 120
 ccatatccag atgtgaggcg tttccttgat gagaagtacg atggagatgt ggataaatta 180
 tctaaacaac ttcaaggtta ttttggtagt ttaagagagt atatagagtt tgaacttaaa 240
 aatggcaaac aaggtcctgc tggccctcaa ggcccaaagg gtgataaagg agaccccgga 300
 gagccaggtc ctcaagggtc agtggggccc gctgggtccaa aggggtgatcg tggtttagac 360
 ggacgacgtg gccagcagg tcagcaaggg gaagccggcc cagttgggtcc tcagggtcct 420
 caaggcgagc aagggccaaa aggcgaccgt ggcgaacaag gaccaaaggg tgaacgcggt 480
 gagcaaggac caaagggtga acgcggtgag caagggccaa aaggcgaccg tggcgaacaa 540
 ggaccaaagg gtgaacgcgg tgagcaagga ccaaagggtg aacgcggtga gcaagggccaa 600
 aaaggcgatc gtggcgagca aggaccaaag ggtgaacgcg gtgagcaagg gccaaaaggc 660
 gatcgtggcg agcaaggacc aaagggtgaa cgcggtgagc aagggccaaa aggcgatcgt 720
 ggcgagcaag gaccaaaggg tgatcgtggc gagcaaggac caaaagggtga tcgtggcgaa 780
 caaggaccaa aggggtgatcg tggcgagcaa ggaccaaagg gtgaacgcgg tgagcaagga 840
 ccaaaggcg atcgtggcga gcaagggtcca aagaaggagc cagaaaaaga caccaagcca 900
 tcagtgccaa aggcaccaga taatatggct gcaccaaagg ctccggagaa gaagaaccct 960
 aaagcaccgg ctcttaagtc agctccaaag gcattcattgc catcaacagg agacactagc 1020
 caatcatttg ttgcagcagc ccttggtctt atcgctagcg cgggcttgct agtcttcaaa 1080
 agaaagaaaa ac 1092

<210> 20
 <211> 364
 <212> PRT
 <213> Streptococcus equi

<400> 20
 Met Met Asn Arg Lys Lys His Thr Lys Leu Ile Arg His Tyr Ser Ile
 1 5 10 15
 Cys Ser Ala Val Ala Val Leu Ala Ala Val Ser Leu Gly Thr Gly Gln
 20 25 30
 Asp Val Gln Ala Ser Glu Pro Asn Pro Tyr Pro Asp Val Arg Arg Phe
 35 40 45
 Leu Asp Glu Lys Tyr Asp Gly Asp Val Asp Lys Leu Ser Lys Gln Leu
 50 55 60
 Gln Gly Tyr Phe Gly Ser Leu Arg Glu Tyr Ile Glu Phe Glu Leu Lys
 65 70 75 80
 Asn Gly Lys Gln Gly Pro Ala Gly Pro Gln Gly Pro Lys Gly Asp Lys
 85 90 95
 Gly Asp Pro Gly Glu Pro Gly Pro Gln Gly Pro Val Gly Pro Ala Gly
 100 105 110
 Pro Lys Gly Asp Arg Gly Leu Asp Gly Arg Arg Gly Pro Ala Gly Gln
 115 120 125
 Gln Gly Glu Ala Gly Pro Val Gly Pro Gln Gly Pro Gln Gly Glu Gln
 130 135 140
 Gly Pro Lys Gly Asp Arg Gly Glu Gln Gly Pro Lys Gly Glu Arg Gly
 145 150 155 160
 Glu Gln Gly Pro Lys Gly Glu Arg Gly Glu Gln Gly Pro Lys Gly Asp
 165 170 175
 Arg Gly Glu Gln Gly Pro Lys Gly Glu Arg Gly Glu Gln Gly Pro Lys
 180 185 190
 Gly Glu Arg Gly Glu Gln Gly Pro Lys Gly Asp Arg Gly Glu Gln Gly
 195 200 205
 Pro Lys Gly Glu Arg Gly Glu Gln Gly Pro Lys Gly Asp Arg Gly Glu
 210 215 220
 Gln Gly Pro Lys Gly Glu Arg Gly Glu Gln Gly Pro Lys Gly Asp Arg
 225 230 235 240

Gly Glu Gln Gly Pro Lys Gly Asp Arg Gly Glu Gln Gly Pro Lys Gly
 245 250 255
 Asp Arg Gly Glu Gln Gly Pro Lys Gly Asp Arg Gly Glu Gln Gly Pro
 260 265 270
 Lys Gly Glu Arg Gly Glu Gln Gly Pro Lys Gly Asp Arg Gly Glu Gln
 275 280 285
 Gly Pro Lys Lys Glu Pro Glu Lys Asp Thr Lys Pro Ser Val Pro Lys
 290 295 300
 Ala Pro Asp Asn Met Ala Ala Pro Lys Ala Pro Glu Lys Lys Asn Pro
 305 310 315 320
 Lys Ala Pro Ala Pro Lys Ser Ala Pro Lys Ala Ser Leu Pro Ser Thr
 325 330 335
 Gly Asp Thr Ser Gln Ser Phe Val Ala Ala Ala Leu Gly Phe Ile Ala
 340 345 350
 Ser Ala Gly Leu Leu Val Phe Lys Arg Lys Lys Asn
 355 360

<210> 21
 <211> 2019
 <212> DNA
 <213> Streptococcus equi

<400> 21
 atgataacta aaaaacagag aaaactaaaa agacacacaa caatcagacg cctgtgtcat 60
 caagccctag cagctaccct catcacagga agcctgatgg gaacaggact acttggaat 120
 aaagtcgtgt atggggatat gaaaagcgat gccaaaaagg tattagcaac attgcaaaaa 180
 ggtaatcata aacaagagtt aaaaaaagtt cttagaacta ttgggtgatga tagtttgcta 240
 ttattatggg tacttgtatc ggggtgattcc aatttagatg aaagtaataa aagttttgat 300
 tatttatcaa tttggaatac tggatctaga gttggtaaag tgatatctcc tactgaatac 360
 tttaaacaag aatcaaaaaa cactgaaaat aggcaagctg tttatagtga atttaaaca 420
 cgcgtagaag aacgatctaa aaaagccaag gaagccactg aagcactaaa ccaaaaagct 480
 caattggagg ctacagttaa aaatataaac gaagaacttg aaaaaactag agaaggtttt 540
 aaagttgtga gtgaaaactc aaacaaatta gaaaaacaat taatggctga gaaaataaaa 600
 acacgtacag ctgaagaaac tgccaaacaa gctaaaaccg ataaagaaag agcagaggct 660
 gaagctaaaa aagcgaaaga ggaagctaag actgcggaag gaaaagttaa acaagcagaa 720
 actgagaaaa ggaatgcgga agctaaggct agaaccgctg aagaagaagc taaacaagct 780
 acggctgata aagaaaaggc agagactgaa gctaaaaaag cgaaagaaga agctaagacc 840
 gctaaagaag ctgctcatca agaacaagaa aaagctaaac aattagaaca agctaataca 900
 caggctaacc aaagagctaa tctggcagaa aaatctaaaa aagatttaga aactcaaaaa 960
 gaaaaacttg aacaagagat taaagaagct acagaagcta aaaacaaagc cgagcaaaaa 1020
 ttaaaagacc tacaagattc agctagtcaa ggcagtgaat tgtctaaaca gctgttaaaa 1080
 gaaaaagaag aattaacaac aaaacttcaa gagcttcaaa aacaagctga ggaaaaaaca 1140
 actgaaattg aaaagctaaa acaagagctt gaagctaata aacaaaacag tggtcagcta 1200
 ggtcaacagg aacaaaagct tcaagagcaa ttgaataagg ttcaaaaaga gctgaaacaa 1260
 aaggaaatgg agttaaaca agctcaagaa cagctgaaac aagaacaaaa acctcacgaa 1320
 ggtggcggag attctgacgc ttccaaagca agaactactg agctagaaaa acaagttcaa 1380
 aactaacca aggaaaaggc agatctgtca tcaacccttg agtcaacaaa agctcaattg 1440
 tcagagactc aagcaagact atcagaagct caaaaacaat tgactgctgc tcaagaaaag 1500
 ttaacaacac ttgaagccga gaaaacagcg cttcaacatc aagttgagac catttcaaaa 1560
 cagctatcag aaacacgtga ccttagcgaa aaggaaaaag cagcactcca agagcaaatac 1620
 aataagctaa aggctgagat tgagcaaaaag accaaggaaa ttgaagcctt aaagcaaggc 1680
 atgcaatcac atcaaggtea agaaaaacca aaggatccaa aaactccgga aacaccaaaag 1740
 gatccaaaga ctcttgagaa aaacgatcag ccacaagcgc ctgaaaaaac tagtgccct 1800
 tggactgctc ttacaccagc taagccaatt gacactacca aagctcctaa gagctcagca 1860
 ccaagcccac agactggggc agcaaacact aaaaaacaat taccagcaac tggtgataca 1920
 gcgacaccta gcttctttac cgctgctgct atggcagtat tggctagtgc ctgtgcccta 1980
 accctctcac ctagacgcaa aaagaatcaa aacaaccgt 2019

<210> 22
 <211> 673
 <212> PRT
 <213> Streptococcus equi

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<400> 22
Met Ile Thr Lys Lys Gln Arg Lys Leu Lys Arg His Thr Thr Ile Arg
1      5      10      15
Arg Leu Cys His Gln Ala Leu Ala Ala Thr Leu Ile Thr Gly Ser Leu
20     25     30
Met Gly Thr Gly Leu Leu Gly Asn Lys Val Val Tyr Gly Asp Met Lys
35     40     45
Ser Asp Ala Lys Lys Val Leu Ala Thr Leu Gln Lys Gly Asn His Lys
50     55     60
Gln Glu Leu Lys Lys Val Leu Arg Thr Ile Gly Asp Asp Ser Leu Leu
65     70     75     80
Leu Leu Trp Leu Leu Val Ser Gly Asp Ser Asn Leu Asp Glu Ser Asn
85     90     95
Lys Ser Phe Asp Tyr Leu Ser Ile Trp Asn Thr Gly Ser Arg Val Gly
100    105    110
Lys Val Ile Ser Pro Thr Glu Tyr Phe Lys Gln Glu Ser Lys Asn Thr
115    120    125
Glu Asn Arg Gln Ala Val Tyr Ser Glu Phe Lys Gln Arg Val Glu Glu
130    135    140
Arg Ser Lys Lys Ala Lys Glu Ala Thr Glu Ala Leu Asn Gln Lys Ala
145    150    155    160
Gln Leu Glu Ala Thr Val Lys Asn Ile Asn Glu Glu Leu Glu Lys Thr
165    170    175
Arg Glu Gly Phe Lys Val Val Ser Glu Asn Ser Asn Lys Leu Glu Lys
180    185    190
Gln Leu Met Ala Glu Lys Ile Lys Thr Arg Thr Ala Glu Glu Thr Ala
195    200    205
Lys Gln Ala Lys Thr Asp Lys Glu Arg Ala Glu Ala Glu Ala Lys Lys
210    215    220
Ala Lys Glu Glu Ala Lys Thr Ala Glu Gly Lys Val Lys Gln Ala Glu
225    230    235    240
Thr Glu Lys Arg Asn Ala Glu Ala Lys Ala Arg Thr Ala Glu Glu Glu
245    250    255
Ala Lys Gln Ala Thr Ala Asp Lys Glu Lys Ala Glu Thr Glu Ala Lys
260    265    270
Lys Ala Lys Glu Glu Ala Lys Thr Ala Lys Glu Ala Ala His Gln Glu
275    280    285
Gln Glu Lys Ala Lys Gln Leu Glu Gln Ala Asn Gln Gln Ala Asn Gln
290    295    300
Arg Ala Asn Leu Ala Glu Lys Ser Lys Lys Asp Leu Glu Thr Gln Lys
305    310    315    320
Glu Lys Leu Glu Gln Glu Ile Lys Glu Ala Thr Glu Ala Lys Asn Lys
325    330    335
Ala Glu Gln Lys Leu Lys Asp Leu Gln Asp Ser Ala Ser Gln Gly Ser
340    345    350
Glu Leu Ser Lys Gln Leu Leu Lys Glu Lys Glu Glu Leu Thr Thr Lys
355    360    365
Leu Gln Glu Leu Gln Lys Gln Ala Glu Glu Lys Thr Thr Glu Ile Glu
370    375    380
Lys Leu Lys Gln Glu Leu Glu Ala Asn Lys Gln Asn Ser Gly Gln Leu
385    390    395    400
Gly Gln Gln Glu Gln Lys Leu Gln Glu Gln Leu Asn Lys Val Gln Lys
405    410    415
Glu Leu Lys Gln Lys Glu Met Glu Leu Lys Gln Ala Gln Glu Gln Leu
420    425    430
Lys Gln Glu Gln Lys Pro His Glu Gly Gly Gly Asp Ser Asp Ala Ser
435    440    445
Lys Ala Arg Ile Thr Glu Leu Glu Lys Gln Val Gln Thr Leu Thr Lys
450    455    460
Glu Lys Ala Asp Leu Ser Ser Thr Leu Glu Ser Thr Lys Ala Gln Leu
465    470    475    480
Ser Glu Thr Gln Ala Arg Leu Ser Glu Ala Gln Lys Gln Leu Thr Ala
485    490    495

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Ala Gln Glu Lys Leu Thr Thr Leu Glu Ala Glu Lys Thr Ala Leu Gln
500 505 510
His Gln Val Glu Thr Ile Ser Lys Gln Leu Ser Glu Thr Arg Asp Leu
515 520 525
Ser Glu Lys Glu Lys Ala Ala Leu Gln Glu Gln Ile Asn Lys Leu Lys
530 535 540
Ala Glu Ile Glu Gln Lys Thr Lys Glu Ile Glu Ala Leu Lys Gln Gly
545 550 555 560
Met Gln Ser His Gln Gly Gln Glu Lys Pro Lys Asp Pro Lys Thr Pro
565 570 575
Glu Thr Pro Lys Asp Pro Lys Thr Pro Glu Lys Asn Asp Gln Pro Gln
580 585 590
Ala Pro Glu Lys Arg Ser Val Pro Trp Thr Ala Leu Thr Pro Ala Lys
595 600 605
Pro Ile Asp Thr Thr Lys Ala Pro Lys Ser Ser Ala Pro Ser Pro Gln
610 615 620
Thr Gly Ala Ala Thr Pro Lys Lys Gln Leu Pro Ala Thr Gly Asp Thr
625 630 635 640
Ala Thr Pro Ser Phe Phe Thr Ala Ala Ala Met Ala Val Leu Ala Ser
645 650 655
Ala Cys Ala Leu Thr Leu Ser Pro Arg Arg Lys Lys Asn Gln Asn Asn
660 665 670
Arg

<210> 23
<211> 1155
<212> DNA
<213> Streptococcus equi

<400> 23
atgatgaaaa aacaatcatt cacacactca cgtaaacctta aattcggtat gagaaaatta 60
tctattggcc ttgcctcatg tatgctagga atgatgttcc taacaacagg acatgtttct 120
ggtgaggttag ttgaagtttg gcctaattggg caaaatccta atggtaaaat agaaattcta 180
agtcaaactg agcactctga gcattttacag aaattacgcg atattgaaga tttccaagct 240
caaaagcaag ctgatcatgt tcgttacact aaatgggttag atggggtaac tgttgatgag 300
catgaattca gaaaaatcaa ggaatatgac acagaatatt atgtaacacc tcttttaagt 360
ggtaaagggtt actatgatat caataaagat ttcaatcaag atagtataa atgtgctgcc 420
gctgtagcgg ctaatatggt ccattattgg tttgatagaa atagagacag tattaatcgt 480
ttcttaagtc aaagtccagg tgaaaatggt gttattaaac ttgaaaatga aaaaacaata 540
gaagtatcaa aattttttaga aacttaccgt agtgatggtg attatcttga taaaagtccg 600
ttttttgacc ttatcagtaa cagcttttaa ggtcctgttt gggcaaataa gctattggat 660
gcttacatta acggctatgg ttatatccat aaatttgcta aaaatactcc acattctaaa 720
aataataata gtaaatttaa tttcttttaa aaagtatttg atggtaatct cttgacagat 780
attcaccaaa tttttgatta taacactttt tcagataaat taagtgaggc tctctatact 840
ggtaaagcca ttggattggc ctacggacct ggagacttgc gtcgttcact aggtcatatt 900
atcttctgtct ggggagctga tcttgacgat cagaatcgcg tggtagctat ttatgtaact 960
gattctgatg ataaaaagtt aactatagga aatgagagag ttggtttgaa gcgatataaa 1020
gtatctagcg atgatcaagg tcgtgctcgt ctgacgactc gtgataaaga taacacaggt 1080
ggtgaaattc gatctattga aacattagat atgggtacac aagagtgggc agattacttc 1140
aacaagacag aaaaa 1155

<210> 24
<211> 385
<212> PRT
<213> Streptococcus equi

<400> 24
Met Met Lys Lys Gln Ser Phe Thr His Ser Arg Lys Pro Lys Phe Gly
1 5 10 15
Met Arg Lys Leu Ser Ile Gly Leu Ala Ser Cys Met Leu Gly Met Met
20 25 30

Phe Leu Thr Thr Gly His Val Ser Gly Glu Val Val Glu Val Trp Pro
 35 40 45
 Asn Gly Gln Asn Pro Asn Gly Lys Ile Glu Ile Leu Ser Gln Thr Glu
 50 55 60
 His Ser Glu His Leu Gln Lys Leu Arg Asp Ile Glu Asp Phe Gln Ala
 65 70 75 80
 Gln Lys Gln Ala Asp His Val Arg Tyr Thr Lys Trp Leu Asp Gly Val
 85 90 95
 Thr Val Asp Glu His Glu Phe Arg Lys Ile Lys Glu Tyr Asp Thr Glu
 100 105 110
 Tyr Tyr Val Thr Pro Leu Leu Ser Gly Lys Gly Tyr Tyr Asp Ile Asn
 115 120 125
 Lys Asp Phe Asn Gln Asp Ser Asp Lys Cys Ala Ala Ala Val Ala Ala
 130 135 140
 Asn Met Phe His Tyr Trp Phe Asp Arg Asn Arg Asp Ser Ile Asn Arg
 145 150 155 160
 Phe Leu Ser Gln Ser Pro Gly Glu Asn Gly Val Ile Lys Leu Glu Asn
 165 170 175
 Glu Lys Thr Ile Glu Val Ser Lys Phe Leu Glu Thr Tyr Arg Ser Asp
 180 185 190
 Gly Asp Tyr Leu Asp Lys Ser Pro Phe Phe Asp Leu Ile Ser Asn Ser
 195 200 205
 Phe Lys Gly Pro Val Trp Ala Asn Lys Leu Leu Asp Ala Tyr Ile Asn
 210 215 220
 Gly Tyr Gly Tyr Ile His Lys Phe Ala Lys Asn Thr Pro His Ser Lys
 225 230 235 240
 Asn Asn Asn Ser Lys Phe Asn Phe Phe Lys Lys Val Phe Asp Gly Asn
 245 250 255
 Leu Leu Thr Asp Ile His Gln Ile Phe Asp Tyr Asn Thr Phe Ser Asp
 260 265 270
 Lys Leu Ser Glu Ala Leu Tyr Thr Gly Lys Ala Ile Gly Leu Ala Tyr
 275 280 285
 Gly Pro Gly Asp Leu Arg Arg Ser Leu Gly His Ile Ile Ser Val Trp
 290 295 300
 Gly Ala Asp Leu Asp Asp Gln Asn Arg Val Val Ala Ile Tyr Val Thr
 305 310 315 320
 Asp Ser Asp Asp Lys Lys Leu Thr Ile Gly Asn Glu Arg Val Gly Leu
 325 330 335
 Lys Arg Tyr Lys Val Ser Ser Asp Asp Gln Gly Arg Ala Arg Leu Thr
 340 345 350
 Thr Arg Asp Lys Asp Asn Thr Gly Gly Glu Ile Arg Ser Ile Glu Thr
 355 360 365
 Leu Asp Met Gly Thr Gln Glu Trp Ala Asp Tyr Phe Asn Lys Thr Glu
 370 375 380
 Lys
 385

<210> 25
 <211> 930
 <212> DNA
 <213> Streptococcus equi

<400> 25
 atgaaaaaga aacttagcct ggccttgagt gcttttttgg ctgttttcat gctggttgcc 60
 tgctctaattg gtggaacaa ggaagctaag agtgacaagc tcaaggttgt gggtaccaat 120
 tccattattg ctgacatgac taaaaatatt gcaggtgaca agattgactt gcacagcatt 180
 gtgccgattg gtcaagatcc gcatgagtat gagccattgc cagaagacgt tgagaagacc 240
 agcaatgccg atcttattct ctataacggt attaaccttg aagacggcgg acaggcttgg 300
 tttacaaagc ttgtaaaaaa tgctaagaag gaaaaggaca aggactactt cgctgtatct 360
 gatggtattg atgtgattta cctagaaggt gaaagtgaag aaggcaagga agaccctcat 420
 gcttggttga atcttgaaaa tggggttatc tattcaaaaa atattgcaaa gcaattgatc 480
 gctaaagacc ctaaaaacaa ggaaacctat gaaaacaacc taaaagctta cgttgagaag 540
 ctagagaagc ttgataagga agcaaaatca aaatttgatg ctattgctga caataaaaag 600

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ctaatacgta caagtgaagg ctgcttcaaa tactttctcaa aagcttacgg tgtcccatca 660
gcctacatct gggaaatcaa cactgaagaa gaaggaactc cggatcaaat ttcattcattg 720
atcgaaaaat tgaaagacgt taagccatca ggcgtttttg ttgagtcaag tgttgacaga 780
cgtccaatgg agactgtttc taaggacagt ggtattccga tctatgcaga aatctttact 840
gactctgtag ccaaaaaagg ccaagatgga gacagctact atgccatgat gaaatggaac 900
cttgataaaa tttctgaagg cttagcaaaa 930

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<210> 26
<211> 310
<212> PRT
<213> Streptococcus equi

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<400> 26
Met Lys Lys Lys Leu Ser Leu Ala Leu Ser Ala Phe Leu Ala Val Phe
1      5      10      15
Met Leu Val Ala Cys Ser Asn Gly Gly Asn Lys Glu Ala Lys Ser Asp
20      25      30
Lys Leu Lys Val Val Val Thr Asn Ser Ile Ile Ala Asp Met Thr Lys
35      40      45
Asn Ile Ala Gly Asp Lys Ile Asp Leu His Ser Ile Val Pro Ile Gly
50      55      60
Gln Asp Pro His Glu Tyr Glu Pro Leu Pro Glu Asp Val Glu Lys Thr
65      70      75      80
Ser Asn Ala Asp Leu Ile Leu Tyr Asn Gly Ile Asn Leu Glu Asp Gly
85      90      95
Gly Gln Ala Trp Phe Thr Lys Leu Val Lys Asn Ala Lys Lys Glu Lys
100     105     110
Asp Lys Asp Tyr Phe Ala Val Ser Asp Gly Ile Asp Val Ile Tyr Leu
115     120     125
Glu Gly Glu Ser Glu Lys Gly Lys Glu Asp Pro His Ala Trp Leu Asn
130     135     140
Leu Glu Asn Gly Val Ile Tyr Ser Lys Asn Ile Ala Lys Gln Leu Ile
145     150     155     160
Ala Lys Asp Pro Lys Asn Lys Glu Thr Tyr Glu Asn Asn Leu Lys Ala
165     170     175
Tyr Val Glu Lys Leu Glu Lys Leu Asp Lys Glu Ala Lys Ser Lys Phe
180     185     190
Asp Ala Ile Ala Asp Asn Lys Lys Leu Ile Val Thr Ser Glu Gly Cys
195     200     205
Phe Lys Tyr Phe Ser Lys Ala Tyr Gly Val Pro Ser Ala Tyr Ile Trp
210     215     220
Glu Ile Asn Thr Glu Glu Glu Gly Thr Pro Asp Gln Ile Ser Ser Leu
225     230     235     240
Ile Glu Lys Leu Lys Asp Val Lys Pro Ser Ala Leu Phe Val Glu Ser
245     250     255
Ser Val Asp Arg Arg Pro Met Glu Thr Val Ser Lys Asp Ser Gly Ile
260     265     270
Pro Ile Tyr Ala Glu Ile Phe Thr Asp Ser Val Ala Lys Lys Gly Gln
275     280     285
Asp Gly Asp Ser Tyr Tyr Ala Met Met Lys Trp Asn Leu Asp Lys Ile
290     295     300
Ser Glu Gly Leu Ala Lys
305      310

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<210> 27
<211> 1287
<212> DNA
<213> Streptococcus equi

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<400> 27
atggaaaaaa acaaaaaggt atcctacttt ttacgtcaat cagctgttgg tttagcttca 60
gtgtcggcag cttttctggt tggaacgact tcagtgggtg ctttagacgc agcaacagtg 120

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ttagagccta caacagcctt cattagagaa gctggttaggg aaatcaatca gctgagtgat 180
gactacgctg acaatcaaga gcttcaggct gttcttgcta atgctggagt tgaggcactt 240
gctgcagata ctgttgatca ggctaaagca gctcttgaca aagcaaaggc agctgttgct 300
ggtgttcagc ttgatgaagc aagacgtgag gcttacagaa caatcaatgc ctttaagtga 360
cagcacaaaa gcgatcaaaa ggttcagcta gctctagttg ctgcagcagc taaggtggca 420
gatgctgctt cagttgatca agtgaatgca gccattaatg atgctcatac agctattgcg 480
gacattacag gagcagcctt gttggaggct aaagaagctg ctatcaatga actaaagcag 540
tatggcatta gtgattacta tgtgacctta atcaacaaag ccaaaactgt tgaaggtgtc 600
aatgcgctta aggcagagat tttatcagct ctaccgagtt ctgaggtcat tgacgcagca 660
gaactaacac cagccttgac tagctataag cttgtcatca agggagcaac tttttcaggt 720
gaaacagcta ctaaggcagt agacgcagct gtagctgagc agaccttcag agactatgct 780
aataaaaaatg gtgtagacgg cgtttgggct tatgatgctg ccacaaagac atttacagtc 840
actgaacagc ctgtagctga gactattgag gcggcagaat taacaccagc tttgaccacc 900
tataggcttg ttattaaggg ggttaccttc tcaggtgaaa cagctaccaa ggcagtagac 960
gcagccacag ccgagcagac cttcaggcaa tatgccaatg acaatggcgt tactggtgaa 1020
tgggcttatg acgctgctac aaagacattt acagtcactg aggcctttaga ggagagtcca 1080
gctgatccag aaaagccatc agcttctctt ccgttgacac cgcttacacc agcaactaag 1140
acagcaccag ctaagcaaaa agataagcaa aaggctaaga ctcttccaac gactggtgag 1200
aaggctaatc cattctttac agcagcggct cttgccatta tggcgagtgc aggtgcttta 1260
gcagtgactt caaagcgtca gcaggac 1287

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<210> 28
<211> 429
<212> PRT
<213> Streptococcus equi

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<400> 28
Met Glu Lys Asn Lys Lys Val Ser Tyr Phe Leu Arg Gln Ser Ala Val
1          5          10          15
Gly Leu Ala Ser Val Ser Ala Ala Phe Leu Val Gly Thr Thr Ser Val
          20          25          30
Gly Ala Leu Asp Ala Ala Thr Val Leu Glu Pro Thr Thr Ala Phe Ile
          35          40          45
Arg Glu Ala Val Arg Glu Ile Asn Gln Leu Ser Asp Asp Tyr Ala Asp
          50          55          60
Asn Gln Glu Leu Gln Ala Val Leu Ala Asn Ala Gly Val Glu Ala Leu
65          70          75          80
Ala Ala Asp Thr Val Asp Gln Ala Lys Ala Ala Leu Asp Lys Ala Lys
          85          90          95
Ala Ala Val Ala Gly Val Gln Leu Asp Glu Ala Arg Arg Glu Ala Tyr
          100          105          110
Arg Thr Ile Asn Ala Leu Ser Asp Gln His Lys Ser Asp Gln Lys Val
          115          120          125
Gln Leu Ala Leu Val Ala Ala Ala Ala Lys Val Ala Asp Ala Ala Ser
          130          135          140
Val Asp Gln Val Asn Ala Ala Ile Asn Asp Ala His Thr Ala Ile Ala
145          150          155          160
Asp Ile Thr Gly Ala Ala Leu Leu Glu Ala Lys Glu Ala Ala Ile Asn
          165          170          175
Glu Leu Lys Gln Tyr Gly Ile Ser Asp Tyr Tyr Val Thr Leu Ile Asn
          180          185          190
Lys Ala Lys Thr Val Glu Gly Val Asn Ala Leu Lys Ala Glu Ile Leu
          195          200          205
Ser Ala Leu Pro Ser Ser Glu Val Ile Asp Ala Ala Glu Leu Thr Pro
          210          215          220
Ala Leu Thr Ser Tyr Lys Leu Val Ile Lys Gly Ala Thr Phe Ser Gly
225          230          235          240
Glu Thr Ala Thr Lys Ala Val Asp Ala Ala Val Ala Glu Gln Thr Phe
          245          250          255
Arg Asp Tyr Ala Asn Lys Asn Gly Val Asp Gly Val Trp Ala Tyr Asp
          260          265          270
Ala Ala Thr Lys Thr Phe Thr Val Thr Glu Gln Pro Val Ala Glu Thr
          275          280          285

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Ile Glu Ala Ala Glu Leu Thr Pro Ala Leu Thr Thr Tyr Arg Leu Val
 290                      295                      300
Ile Lys Gly Val Thr Phe Ser Gly Glu Thr Ala Thr Lys Ala Val Asp
305                      310                      315                      320
Ala Ala Thr Ala Glu Gln Thr Phe Arg Gln Tyr Ala Asn Asp Asn Gly
                      325                      330                      335
Val Thr Gly Glu Trp Ala Tyr Asp Ala Ala Thr Lys Thr Phe Thr Val
                      340                      345                      350
Thr Glu Ala Leu Glu Glu Ser Pro Ala Asp Pro Glu Lys Pro Ser Ala
                      355                      360                      365
Ser Leu Pro Leu Thr Pro Leu Thr Pro Ala Thr Lys Thr Ala Pro Ala
                      370                      375                      380
Lys Gln Lys Asp Lys Gln Lys Ala Lys Thr Leu Pro Thr Thr Gly Glu
385                      390                      395                      400
Lys Ala Asn Pro Phe Phe Thr Ala Ala Ala Leu Ala Ile Met Ala Ser
                      405                      410                      415
Ala Gly Ala Leu Ala Val Thr Ser Lys Arg Gln Gln Asp
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<210> 29
<211> 2004
<212> DNA
<213> Streptococcus equi

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<400> 29
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cttagtgaca acatcacatc attgacggtt gcttcttcat cactccgaga tggagagaga 180
acgacggtaa aggttgcggt tgatgacaaa aaacagaaaa tcaaggcagg ggatacgata 240
gaggtcacct ggcctacaag tggtaatgtc tacattcagg gctttaataa aaccataaccg 300
cttaataatta gaggggtaga tgttggtacc ttggagggtca cgctagacaa ggctgttttc 360
acattcaatc aaaatattga aacaatgcat gatgtctctg gttggggaga gtttgatatt 420
actgtagaaa atgtgacaca aaccaccgct gaaacatcag gaacgaccac agtaaaggta 480
ggcaatcgca ctgctactat cactgttact aagcctgagg caggcactgg taccagctca 540
ttttattata agactggtga tatgcagccc aatgatactg agcgtgtgag atgggttcctg 600
ctgattaaca acaacaagga atgggtggcc aatactgtta cagtcgaaga cgatattcaa 660
ggtggtcaaaa ccttgatat gagcagcttt gacatcaccc tatctggtta tcgtaacgag 720
cgcttcggtt gggaaaacgc tctgacagag ttctatacaa catttccaaa ttctgtcatt 780
acggcaacag ataatacat tagtgtgcgg ttagatcaat atgatgcctc acaaaacact 840
gtcaacattg cttataagac aaagataacg gactttgacc aaaaagaatt tgccaacaac 900
agtaaaatct ggtaccagat tttatacaag gatcagggtat cgggtcaaga gtcaaaccac 960
caagtagcca atatcaatgc taacggcggg gttgatggca gtcgctatac cagctttact 1020
gtcaagaaaa tttggaatga caaggaaaat caagacggta agcgtccaaa gactattact 1080
gttcagcttt acgccaatga tcagaaagtt aatgataaga ccattgaatt gagtgatact 1140
aatagctggc aagcaagttt tggtaagctg gataagtatg acagtcagaa ccaaaaaatt 1200
acctacagtg tcaaggaagt gatggttcct gttggctacc aatcgcaggt tgagggggat 1260
agtggagtag gatttaccat taccaacacc tatacaccag aggtcattag cattaccggt 1320
caaaaaactt gggacgacag ggaaaaccaa gacggtaaac gtcctaagga gattacggtt 1380
cgtttatttg caaatgacgc tgcaactgac aaggtagcaa ctgcttcaga gcaaaccggc 1440
tggaagtata catttaccaa tctaccgaaa tacaagatg gtaaacagat cacctacacg 1500
atccaagagg accctgtggc agattacacc acaaccattc agggatttga tattaccaat 1560
catcatgagg tagccttgac cagcctaaag gtcacaaagg tttggaatga taaggacgat 1620
tattaccata aacgtcccaa ggagattacc attttgctaa aggcagatgg caaggtgatt 1680
cgtgaacatc agatgacacc ggatcagcaa ggaaaatggg aatacacctt tgaccagctg 1740
ccggtctatc agacaggcaa gaaaatcagc tacagcattg aggaaaaaca ggttgctggc 1800
tatcaagccc ctgtctatga ggttgatgaa ggcttgaagc aggtcactgt aaccaacacc 1860
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2004

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<210> 30
<211> 668

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<212> PRT

<213> Streptococcus equi

<400> 30

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Lys	Ile	Val	Ser	Val	Val	Leu	Leu	Leu	Val	Phe	Thr	Leu	Ser	Ala	Ser	20	25	30	
Leu	His	Lys	Val	Arg	Ala	Thr	Asn	Leu	Ser	Asp	Asn	Ile	Thr	Ser	Leu	35	40	45	
Thr	Val	Ala	Ser	Ser	Ser	Leu	Arg	Asp	Gly	Glu	Arg	Thr	Thr	Val	Lys	50	55	60	
Val	Ala	Phe	Asp	Asp	Lys	Lys	Gln	Lys	Ile	Lys	Ala	Gly	Asp	Thr	Ile	65	70	75	80
Glu	Val	Thr	Trp	Pro	Thr	Ser	Gly	Asn	Val	Tyr	Ile	Gln	Gly	Phe	Asn	85	90	95	
Lys	Thr	Ile	Pro	Leu	Asn	Ile	Arg	Gly	Val	Asp	Val	Gly	Thr	Leu	Glu	100	105	110	
Val	Thr	Leu	Asp	Lys	Ala	Val	Phe	Thr	Phe	Asn	Gln	Asn	Ile	Glu	Thr	115	120	125	
Met	His	Asp	Val	Ser	Gly	Trp	Gly	Glu	Phe	Asp	Ile	Thr	Val	Arg	Asn	130	135	140	
Val	Thr	Gln	Thr	Thr	Ala	Glu	Thr	Ser	Gly	Thr	Thr	Thr	Val	Lys	Val	145	150	155	160
Gly	Asn	Arg	Thr	Ala	Thr	Ile	Thr	Val	Thr	Lys	Pro	Glu	Ala	Gly	Thr	165	170	175	
Gly	Thr	Ser	Ser	Phe	Tyr	Tyr	Lys	Thr	Gly	Asp	Met	Gln	Pro	Asn	Asp	180	185	190	
Thr	Glu	Arg	Val	Arg	Trp	Phe	Leu	Leu	Ile	Asn	Asn	Asn	Lys	Glu	Trp	195	200	205	
Val	Ala	Asn	Thr	Val	Thr	Val	Glu	Asp	Asp	Ile	Gln	Gly	Gly	Gln	Thr	210	215	220	
Leu	Asp	Met	Ser	Ser	Phe	Asp	Ile	Thr	Val	Ser	Gly	Tyr	Arg	Asn	Glu	225	230	235	240
Arg	Phe	Val	Gly	Glu	Asn	Ala	Leu	Thr	Glu	Phe	His	Thr	Thr	Phe	Pro	245	250	255	
Asn	Ser	Val	Ile	Thr	Ala	Thr	Asp	Asn	His	Ile	Ser	Val	Arg	Leu	Asp	260	265	270	
Gln	Tyr	Asp	Ala	Ser	Gln	Asn	Thr	Val	Asn	Ile	Ala	Tyr	Lys	Thr	Lys	275	280	285	
Ile	Thr	Asp	Phe	Asp	Gln	Lys	Glu	Phe	Ala	Asn	Asn	Ser	Lys	Ile	Trp	290	295	300	
Tyr	Gln	Ile	Leu	Tyr	Lys	Asp	Gln	Val	Ser	Gly	Gln	Glu	Ser	Asn	His	305	310	315	320
Gln	Val	Ala	Asn	Ile	Asn	Ala	Asn	Gly	Gly	Val	Asp	Gly	Ser	Arg	Tyr	325	330	335	
Thr	Ser	Phe	Thr	Val	Lys	Lys	Ile	Trp	Asn	Asp	Lys	Glu	Asn	Gln	Asp	340	345	350	
Gly	Lys	Arg	Pro	Lys	Thr	Ile	Thr	Val	Gln	Leu	Tyr	Ala	Asn	Asp	Gln	355	360	365	
Lys	Val	Asn	Asp	Lys	Thr	Ile	Glu	Leu	Ser	Asp	Thr	Asn	Ser	Trp	Gln	370	375	380	
Ala	Ser	Phe	Gly	Lys	Leu	Asp	Lys	Tyr	Asp	Ser	Gln	Asn	Gln	Lys	Ile	385	390	395	400
Thr	Tyr	Ser	Val	Lys	Glu	Val	Met	Val	Pro	Val	Gly	Tyr	Gln	Ser	Gln	405	410	415	
Val	Glu	Gly	Asp	Ser	Gly	Val	Gly	Phe	Thr	Ile	Thr	Asn	Thr	Tyr	Thr	420	425	430	
Pro	Glu	Val	Ile	Ser	Ile	Thr	Gly	Gln	Lys	Thr	Trp	Asp	Asp	Arg	Glu	435	440	445	
Asn	Gln	Asp	Gly	Lys	Arg	Pro	Lys	Glu	Ile	Thr	Val	Arg	Leu	Leu	Ala	450	455	460	
Asn	Asp	Ala	Ala	Thr	Asp	Lys	Val	Ala	Thr	Ala	Ser	Glu	Gln	Thr	Gly	465	470	475	480

Trp Lys Tyr Thr Phe Thr Asn Leu Pro Lys Tyr Lys Asp Gly Lys Gln
 485 490 495
 Ile Thr Tyr Thr Ile Gln Glu Asp Pro Val Ala Asp Tyr Thr Thr Thr
 500 505 510
 Ile Gln Gly Phe Asp Ile Thr Asn His His Glu Val Ala Leu Thr Ser
 515 520 525
 Leu Lys Val Ile Lys Val Trp Asn Asp Lys Asp Asp Tyr Tyr His Lys
 530 535 540
 Arg Pro Lys Glu Ile Thr Ile Leu Leu Lys Ala Asp Gly Lys Val Ile
 545 550 555 560
 Arg Glu His Gln Met Thr Pro Asp Gln Gln Gly Lys Trp Glu Tyr Thr
 565 570 575
 Phe Asp Gln Leu Pro Val Tyr Gln Thr Gly Lys Lys Ile Ser Tyr Ser
 580 585 590
 Ile Glu Glu Lys Gln Val Ala Gly Tyr Gln Ala Pro Val Tyr Glu Val
 595 600 605
 Asp Glu Gly Leu Lys Gln Val Thr Val Thr Asn Thr Leu Asn Pro Ser
 610 615 620
 Tyr Lys Leu Pro Asp Thr Gly Gly Gln Gly Val Lys Trp Tyr Leu Leu
 625 630 635 640
 Ile Gly Gly Gly Phe Ile Ile Val Ala Ile Leu Val Leu Ile Ser Leu
 645 650 655
 Tyr Gln Lys His Lys Arg His Asn Met Ser Lys Pro
 660 665

<210> 31
 <211> 165
 <212> DNA
 <213> Streptococcus equi

<400> 31
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 gatgttattt atgatgagga tataaaaaga tatttttgagg atttagaagc ctatttaaca 120
 gctagacttg gtgggattga taaaaaagta gaagaagctg cccaa 165

<210> 32
 <211> 55
 <212> PRT
 <213> Streptococcus equi

<400> 32
 Gln Pro Ala Ala Leu Lys Tyr Pro Glu Pro Arg Asp Tyr Phe Leu His
 1 5 10 15
 Thr Arg Glu Gly Asp Val Ile Tyr Asp Glu Asp Ile Lys Arg Tyr Phe
 20 25 30
 Glu Asp Leu Glu Ala Tyr Leu Thr Ala Arg Leu Gly Gly Ile Asp Lys
 35 40 45
 Lys Val Glu Glu Ala Ala Gln
 50 55

<210> 33
 <211> 135
 <212> DNA
 <213> Streptococcus equi

<400> 33
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 gtggataaat tatctaaaca acttcaaggt tatttttggt gtttaagaga gtatatagag 120
 tttgaactta aaaat 135

<210> 34
 <211> 45
 <212> PRT
 <213> Streptococcus equi

<400> 34
 Ser Glu Pro Asn Pro Tyr Pro Asp Val Arg Arg Phe Leu Asp Glu Lys
 1 5 10 15
 Tyr Asp Gly Asp Val Asp Lys Leu Ser Lys Gln Leu Gln Gly Tyr Phe
 20 25 30
 Gly Ser Leu Arg Glu Tyr Ile Glu Phe Glu Leu Lys Asn
 35 40 45

<210> 35
 <211> 1827
 <212> DNA
 <213> Streptococcus equi

<400> 35
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 caagagttaa aaaaagttct tagaactatt ggtgatgata gtttgctatt attatgggta 180
 cttgtatcgg gtgattccaa tttagatgaa agtaataaaa gttttgatta tttatcaatt 240
 tggaataactg gatctagagt tggtaaagtg atatctccta ctgaataactt taaacaagaa 300
 tcaaaaaaca ctgaaaatag gcaagctggt tatagtgaat ttaaacaacg cgtagaagaa 360
 cgatctaaaa aagccaagga agccactgaa gcactaaacc aaaaagctca attggaggct 420
 acagttaaaa atataaacga agaacttgaa aaaactagag aagggttttaa agttgtgagt 480
 gaaaactcaa acaaattaga aaaacaatta atggctgaga aaataaaaac acgtacagct 540
 gaagaaactg ccaaacaagc taaaaccgat aaagaaagag cagaggctga agctaaaaaa 600
 gcgaaagagg aagctaagac tgcggaagga aaagttaaac aagcagaaac tgagaaaagg 660
 aatgcggaag ctaaggctag aaccgctgaa gaagaagcta aacaagctac ggctgataaa 720
 gaaaaggcag agactgaagc taaaaaagcg aaagaagaag ctaagaccgc taaagaagct 780
 gctcatcaag aacaagaaaa agctaaacaa ttagaacaag ctaatcaaca ggctaaccaa 840
 agagctaatac tggcagaaaa atctaaaaaa gatttagaaa ctcaaaaaga aaaacttgaa 900
 caagagatta aagaagctac agaagctaaa aacaaagccg agcaaaaatt aaaagaccta 960
 caagattcag ctagtcaagg cagtgaattg tctaaacagc tgttaaaaga aaaagaagaa 1020
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 aagctaaaaac aagagcttga agctaataaa caaacagtg gtcagctagg tcaacaggaa 1140
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 gaaaaggcag atctgtcatc aacccttgag tcaacaaaag ctcaattgtc agagactcaa 1380
 gcaagactat cagaagctca aaaacaattg actgctgctc aagaaaagtt aacaacactt 1440
 gaagccgaga aaacagcgct tcaacatcaa gttgagacca tttcaaaaca gctatcagaa 1500
 acacgtgacc ttagcgaaaa ggaaaaagca gcactccaag agcaaataca taagctaaag 1560
 gctgagattg agcaaaagac caaggaaatt gaagccttaa agcaaggcat gcaatcacat 1620
 caaggtcaag aaaaacccaaa ggatccaaaa actccggaaa caccaaagga tccaaagact 1680
 cctgagaaaa acgatcagcc acaagcgcct gaaaaacgta gtgtcccttg gactgctctt 1740
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 actggggcag caacacctaa aaaacaa 1827

<210> 36
 <211> 609
 <212> PRT
 <213> Streptococcus equi

<400> 36
 Ala Thr Leu Ile Thr Gly Ser Leu Met Gly Thr Gly Leu Leu Gly Asn
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 Lys Val Val Tyr Gly Asp Met Lys Ser Asp Ala Lys Lys Val Leu Ala
 20 25 30

Thr	Leu	Gln	Lys	Gly	Asn	His	Lys	Gln	Glu	Leu	Lys	Lys	Val	Leu	Arg
		35					40					45			
Thr	Ile	Gly	Asp	Asp	Ser	Leu	Leu	Leu	Leu	Trp	Leu	Leu	Val	Ser	Gly
	50					55					60				
Asp	Ser	Asn	Leu	Asp	Glu	Ser	Asn	Lys	Ser	Phe	Asp	Tyr	Leu	Ser	Ile
65					70					75					80
Trp	Asn	Thr	Gly	Ser	Arg	Val	Gly	Lys	Val	Ile	Ser	Pro	Thr	Glu	Tyr
				85					90					95	
Phe	Lys	Gln	Glu	Ser	Lys	Asn	Thr	Glu	Asn	Arg	Gln	Ala	Val	Tyr	Ser
			100					105					110		
Glu	Phe	Lys	Gln	Arg	Val	Glu	Glu	Arg	Ser	Lys	Lys	Ala	Lys	Glu	Ala
		115					120					125			
Thr	Glu	Ala	Leu	Asn	Gln	Lys	Ala	Gln	Leu	Glu	Ala	Thr	Val	Lys	Asn
	130					135						140			
Ile	Asn	Glu	Glu	Leu	Glu	Lys	Thr	Arg	Glu	Gly	Phe	Lys	Val	Val	Ser
145					150					155					160
Glu	Asn	Ser	Asn	Lys	Leu	Glu	Lys	Gln	Leu	Met	Ala	Glu	Lys	Ile	Lys
				165					170					175	
Thr	Arg	Thr	Ala	Glu	Glu	Thr	Ala	Lys	Gln	Ala	Lys	Thr	Asp	Lys	Glu
			180					185					190		
Arg	Ala	Glu	Ala	Glu	Ala	Lys	Lys	Ala	Lys	Glu	Glu	Ala	Lys	Thr	Ala
	195						200					205			
Glu	Gly	Lys	Val	Lys	Gln	Ala	Glu	Thr	Glu	Lys	Arg	Asn	Ala	Glu	Ala
	210					215					220				
Lys	Ala	Arg	Thr	Ala	Glu	Glu	Glu	Ala	Lys	Gln	Ala	Thr	Ala	Asp	Lys
225					230					235					240
Glu	Lys	Ala	Glu	Thr	Glu	Ala	Lys	Lys	Ala	Lys	Glu	Glu	Ala	Lys	Thr
			245					250					255		
Ala	Lys	Glu	Ala	Ala	His	Gln	Glu	Gln	Glu	Lys	Ala	Lys	Gln	Leu	Glu
			260					265					270		
Gln	Ala	Asn	Gln	Gln	Ala	Asn	Gln	Arg	Ala	Asn	Leu	Ala	Glu	Lys	Ser
		275					280					285			
Lys	Lys	Asp	Leu	Glu	Thr	Gln	Lys	Glu	Lys	Leu	Glu	Gln	Glu	Ile	Lys
	290					295					300				
Glu	Ala	Thr	Glu	Ala	Lys	Asn	Lys	Ala	Glu	Gln	Lys	Leu	Lys	Asp	Leu
305					310					315					320
Gln	Asp	Ser	Ala	Ser	Gln	Gly	Ser	Glu	Leu	Ser	Lys	Gln	Leu	Leu	Lys
			325					330					335		
Glu	Lys	Glu	Glu	Leu	Thr	Thr	Lys	Leu	Gln	Glu	Leu	Gln	Lys	Gln	Ala
			340					345					350		
Glu	Glu	Lys	Thr	Thr	Glu	Ile	Glu	Lys	Leu	Lys	Gln	Glu	Leu	Glu	Ala
	355					360						365			
Asn	Lys	Gln	Asn	Ser	Gly	Gln	Leu	Gly	Gln	Gln	Glu	Gln	Lys	Leu	Gln
	370					375					380				
Glu	Gln	Leu	Asn	Lys	Val	Gln	Lys	Glu	Leu	Lys	Gln	Lys	Glu	Met	Glu
385					390					395					400
Leu	Lys	Gln	Ala	Gln	Glu	Gln	Leu	Lys	Gln	Glu	Gln	Lys	Pro	His	Glu
			405					410					415		
Gly	Gly	Gly	Asp	Ser	Asp	Ala	Ser	Lys	Ala	Arg	Ile	Thr	Glu	Leu	Glu
			420					425					430		
Lys	Gln	Val	Gln	Thr	Leu	Thr	Lys	Glu	Lys	Ala	Asp	Leu	Ser	Ser	Thr
	435						440					445			
Leu	Glu	Ser	Thr	Lys	Ala	Gln	Leu	Ser	Glu	Thr	Gln	Ala	Arg	Leu	Ser
	450					455					460				
Glu	Ala	Gln	Lys	Gln	Leu	Thr	Ala	Ala	Gln	Glu	Lys	Leu	Thr	Thr	Leu
465					470					475					480
Glu	Ala	Glu	Lys	Thr	Ala	Leu	Gln	His	Gln	Val	Glu	Thr	Ile	Ser	Lys
			485					490					495		
Gln	Leu	Ser	Glu	Thr	Arg	Asp	Leu	Ser	Glu	Lys	Glu	Lys	Ala	Ala	Leu
			500					505					510		
Gln	Glu	Gln	Ile	Asn	Lys	Leu	Lys	Ala	Glu	Ile	Glu	Gln	Lys	Thr	Lys
	515						520					525			
Glu	Ile	Glu	Ala	Leu	Lys	Gln	Gly	Met	Gln	Ser	His	Gln	Gly	Gln	Glu
	530					535					540				

Lys Pro Lys Asp Pro Lys Thr Pro Glu Thr Pro Lys Asp Pro Lys Thr
 545 550 555 560
 Pro Glu Lys Asn Asp Gln Pro Gln Ala Pro Glu Lys Arg Ser Val Pro
 565 570 575
 Trp Thr Ala Leu Thr Pro Ala Lys Pro Ile Asp Thr Thr Lys Ala Pro
 580 585 590
 Lys Ser Ser Ala Pro Ser Pro Gln Thr Gly Ala Ala Thr Pro Lys Lys
 595 600 605
 Gln

<210> 37
 <211> 1032
 <212> DNA
 <213> Streptococcus equi

<400> 37
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 aagcaagctg atcatgttcg ttacactaaa tggttagatg gggtaactgt tgatgagcat 180
 gaattcagaa aaatcaagga atatgacaca gaatattatg taacacctct ttttaagtgg 240
 aaagggtact atgatatcaa taaagatttc aatcaagata gtgataaatg tgctgccgct 300
 gtagcggcta atatgttcca ttattgggtt gatagaaata gagacagtat taatcgtttc 360
 ttaagtcaaa gtccagggtga aaatgggtgt attaaacttg aaaatgaaaa aacaatagaa 420
 gtatcaaaat ttttagaaac ttaccgtagt gatgggtgatt atcttgataa aagtcctgtt 480
 tttgacctta tcagtaacag ctttaaagggt cctgtttggg caaataagct attggatgct 540
 tacattaacg gctatgggtta tatccataaa tttgctaaaa atactccaca ttctaaaaat 600
 aataatagta aatttaattt ctttaaaaaa gtatttgatg gtaatctctt gacagatatt 660
 caccaaattt ttgattataa cactttttca gataaattaa gtgaggctct ctatactgg 720
 aaagccattg gattggccta cggacctgga gacttgctgc gttcactagg tcatattatt 780
 tctgtctggg gagctgatct tgacgatcag aatcgcgtgg tagctattta tgtaactgat 840
 tctgatgata aaaagttaac tataggaaat gagagagttg gtttgaagcg atataaagta 900
 tctagcgatg atcaaggctg tgctcgtctg acgactcgtg ataaagataa cacagggtgg 960
 gaaattcgat ctattgaaac attagatatg ggtacacaag agtgggcaga ttacttcaac 1020
 aagacagaaa aa 1032

<210> 38
 <211> 344
 <212> PRT
 <213> Streptococcus equi

<400> 38
 Glu Val Val Glu Val Trp Pro Asn Gly Gln Asn Pro Asn Gly Lys Ile
 1 5 10 15
 Glu Ile Leu Ser Gln Thr Glu His Ser Glu His Leu Gln Lys Leu Arg
 20 25 30
 Asp Ile Glu Asp Phe Gln Ala Gln Lys Gln Ala Asp His Val Arg Tyr
 35 40 45
 Thr Lys Trp Leu Asp Gly Val Thr Val Asp Glu His Glu Phe Arg Lys
 50 55 60
 Ile Lys Glu Tyr Asp Thr Glu Tyr Tyr Val Thr Pro Leu Leu Ser Gly
 65 70 75 80
 Lys Gly Tyr Tyr Asp Ile Asn Lys Asp Phe Asn Gln Asp Ser Asp Lys
 85 90 95
 Cys Ala Ala Ala Val Ala Ala Asn Met Phe His Tyr Trp Phe Asp Arg
 100 105 110
 Asn Arg Asp Ser Ile Asn Arg Phe Leu Ser Gln Ser Pro Gly Glu Asn
 115 120 125
 Gly Val Ile Lys Leu Glu Asn Glu Lys Thr Ile Glu Val Ser Lys Phe
 130 135 140
 Leu Glu Thr Tyr Arg Ser Asp Gly Asp Tyr Leu Asp Lys Ser Pro Phe
 145 150 155 160

Phe Asp Leu Ile Ser Asn Ser Phe Lys Gly Pro Val Trp Ala Asn Lys
 165 170 175
 Leu Leu Asp Ala Tyr Ile Asn Gly Tyr Gly Tyr Ile His Lys Phe Ala
 180 185 190
 Lys Asn Thr Pro His Ser Lys Asn Asn Asn Ser Lys Phe Asn Phe Phe
 195 200 205
 Lys Lys Val Phe Asp Gly Asn Leu Leu Thr Asp Ile His Gln Ile Phe
 210 215 220
 Asp Tyr Asn Thr Phe Ser Asp Lys Leu Ser Glu Ala Leu Tyr Thr Gly
 225 230 235 240
 Lys Ala Ile Gly Leu Ala Tyr Gly Pro Gly Asp Leu Arg Arg Ser Leu
 245 250 255
 Gly His Ile Ile Ser Val Trp Gly Ala Asp Leu Asp Asp Gln Asn Arg
 260 265 270
 Val Val Ala Ile Tyr Val Thr Asp Ser Asp Asp Lys Lys Leu Thr Ile
 275 280 285
 Gly Asn Glu Arg Val Gly Leu Lys Arg Tyr Lys Val Ser Ser Asp Asp
 290 295 300
 Gln Gly Arg Ala Arg Leu Thr Thr Arg Asp Lys Asp Asn Thr Gly Gly
 305 310 315 320
 Glu Ile Arg Ser Ile Glu Thr Leu Asp Met Gly Thr Gln Glu Trp Ala
 325 330 335
 Asp Tyr Phe Asn Lys Thr Glu Lys
 340

<210> 39
 <211> 867
 <212> DNA
 <213> Streptococcus equi

<400> 39
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 attattgctg acatgactaa aaatattgca ggtgacaaga ttgacttgca cagcattgtg 120
 ccgattgggtc aagatccgca tgagtatgag ccattgccag aagacgttga gaagaccagc 180
 aatgccgatc ttattctcta taacgggtatt aaccttgaag acggcggaca ggcttggttt 240
 acaaagcttg taaaaaatgc taagaaggaa aaggacaagg actacttcgc tgtatctgat 300
 ggtattgatg tgattttacct agaaggtgaa agtgaaaaag gcaaggaaga ccctcatgct 360
 tggttgaatc ttgaaaatgg gggttatctat tcaaaaaata ttgcaaagca attgatcgct 420
 aaagacccta aaaacaagga aacctatgaa aacaaccta aagcttacgt tgagaagcta 480
 gagaagcttg ataaggaagc aaaatcaaaa tttgatgcta ttgctgacaa taaaaagcta 540
 atcgtcacaa gtgaaggctg cttcaaatac ttctcaaaag cttacgggtgt cccatcagcc 600
 tacatctggg aaatcaacac tgaagaagaa ggaactccgg atcaaatttc atcattgatc 660
 gaaaaattga aagacgttaa gccatcagcg ctttttgttg agtcaagtgt tgacagacgt 720
 ccaatggaga ctgtttctaa ggacagtggg attccgatct atgcagaaat ctttactgac 780
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 gataaaattt ctgaaggctt agcaaaa 867

<210> 40
 <211> 289
 <212> PRT
 <213> Streptococcus equi

<400> 40
 Ser Asn Gly Gly Asn Lys Glu Ala Lys Ser Asp Lys Leu Lys Val Val
 1 5 10 15
 Val Thr Asn Ser Ile Ile Ala Asp Met Thr Lys Asn Ile Ala Gly Asp
 20 25 30
 Lys Ile Asp Leu His Ser Ile Val Pro Ile Gly Gln Asp Pro His Glu
 35 40 45
 Tyr Glu Pro Leu Pro Glu Asp Val Glu Lys Thr Ser Asn Ala Asp Leu
 50 55 60

Ile Leu Tyr Asn Gly Ile Asn Leu Glu Asp Gly Gly Gln Ala Trp Phe
 65 70 75 80
 Thr Lys Leu Val Lys Asn Ala Lys Lys Glu Lys Asp Lys Asp Tyr Phe
 85 90 95
 Ala Val Ser Asp Gly Ile Asp Val Ile Tyr Leu Glu Gly Glu Ser Glu
 100 105 110
 Lys Gly Lys Glu Asp Pro His Ala Trp Leu Asn Leu Glu Asn Gly Val
 115 120 125
 Ile Tyr Ser Lys Asn Ile Ala Lys Gln Leu Ile Ala Lys Asp Pro Lys
 130 135 140
 Asn Lys Glu Thr Tyr Glu Asn Asn Leu Lys Ala Tyr Val Glu Lys Leu
 145 150 155 160
 Glu Lys Leu Asp Lys Glu Ala Lys Ser Lys Phe Asp Ala Ile Ala Asp
 165 170 175
 Asn Lys Lys Leu Ile Val Thr Ser Glu Gly Cys Phe Lys Tyr Phe Ser
 180 185 190
 Lys Ala Tyr Gly Val Pro Ser Ala Tyr Ile Trp Glu Ile Asn Thr Glu
 195 200 205
 Glu Glu Gly Thr Pro Asp Gln Ile Ser Ser Leu Ile Glu Lys Leu Lys
 210 215 220
 Asp Val Lys Pro Ser Ala Leu Phe Val Glu Ser Ser Val Asp Arg Arg
 225 230 235 240
 Pro Met Glu Thr Val Ser Lys Asp Ser Gly Ile Pro Ile Tyr Ala Glu
 245 250 255
 Ile Phe Thr Asp Ser Val Ala Lys Lys Gly Gln Asp Gly Asp Ser Tyr
 260 265 270
 Tyr Ala Met Met Lys Trp Asn Leu Asp Lys Ile Ser Glu Gly Leu Ala
 275 280 285
 Lys

<210> 41
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 <213> Artificial

<220>
 <223> Oligonucleotides as PCR primer

<400> 41
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31

<210> 42
 <211> 35
 <212> DNA
 <213> Artificial

<220>
 <223> Oligonucleotides as PCR primer

<400> 42
 aaagaattct tatttttctg acttagattt agaag

35

<210> 43
 <211> 34
 <212> DNA
 <213> Artificial

<220>
 <223> Oligonucleotides as PCR primer

<400> 43
 aaaggatcct ggcgactacc ctagcaggac aaac 34

<210> 44
 <211> 33
 <212> DNA
 <213> Artificial

<220>
 <223> Oligonucleotides as PCR primer

<400> 44
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<210> 45
 <211> 33
 <212> DNA
 <213> Artificial

<220>
 <223> Oligonucleotides as PCR primer

<400> 45
 cccggatcct agaagggata aatgataaaa tag 33

<210> 46
 <211> 34
 <212> DNA
 <213> Artificial

<220>
 <223> Oligonucleotides as PCR primer

<400> 46
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<210> 47
 <211> 1002
 <212> DNA
 <213> Streptococcus equi

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 agtgtttagtg atttttacaa tgaaacaaaa aatacagaga tttctcaaag agcaatgcta 180
 aaccttgtgg ttagtcgtgt ttttgaggac caatacggta aaaaggtttc taagaaaaga 240
 acggaagaag cttacaataa atcagctgag caatacggtg cgtcattctc tgcagccctt 300
 gcgcagtctg gcttgacaac agatacctac aagcgtcaaa ttcgctcagc catgctgggt 360
 gaatatgctg ttaaagaagc agctaaaaaa gagctgacag atgctgatta caaaaaagcc 420
 tatgagtcac acacaccaga aatgactact caggtcacta ctctagacaa tgaagaaaca 480
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 gctgaaaaga cacaaaacat tgattttcaa aataagggtca ttgcaaaggc cttagacaag 840
 gcaaatgtta agatcaaaga ccaagcattt gctaatatct tggcacagta tgccaatact 900
 gataaaaaag caagcaaggc gaacacaagc aagtcagatc agaaatcatc ttcagactca 960
 agcaaggata gtcaatcttc taaatctaag tcagaaaaat ag 1002

<210> 48
 <211> 333
 <212> PRT
 <213> Streptococcus equi

<400> 48
 Met Lys Lys Ser Thr Lys Leu Leu Ala Gly Ile Val Thr Leu Ala Ser
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 20 25 30
 Ile Thr Met Lys Gly Asp Thr Ile Ser Val Ser Asp Phe Tyr Asn Glu
 35 40 45
 Thr Lys Asn Thr Glu Ile Ser Gln Arg Ala Met Leu Asn Leu Val Val
 50 55 60
 Ser Arg Val Phe Glu Asp Gln Tyr Gly Lys Lys Val Ser Lys Lys Arg
 65 70 75 80
 Thr Glu Glu Ala Tyr Asn Lys Ser Ala Glu Gln Tyr Gly Ala Ser Phe
 85 90 95
 Ser Ala Ala Leu Ala Gln Ser Gly Leu Thr Thr Asp Thr Tyr Lys Arg
 100 105 110
 Gln Ile Arg Ser Ala Met Leu Val Glu Tyr Ala Val Lys Glu Ala Ala
 115 120 125
 Lys Lys Glu Leu Thr Asp Ala Asp Tyr Lys Lys Ala Tyr Glu Ser Tyr
 130 135 140
 Thr Pro Glu Met Thr Thr Gln Val Thr Thr Leu Asp Asn Glu Glu Thr
 145 150 155 160
 Ala Lys Ala Val Leu Gly Glu Val Lys Ala Glu Gly Ala Asp Phe Ala
 165 170 175
 Ala Ile Ala Lys Glu Lys Thr Thr Ala Ala Asp Lys Lys Val Asp Tyr
 180 185 190
 Lys Phe Asp Ser Gly Asp Thr Lys Leu Pro Ala Asp Val Ile Lys Ala
 195 200 205
 Ala Ser Gly Leu Lys Glu Gly Asp Ile Ser Glu Val Val Ser Val Leu
 210 215 220
 Asp Pro Ala Thr Tyr Gln Asn Lys Phe Tyr Ile Val Lys Val Thr Lys
 225 230 235 240
 Lys Ala Glu Lys Ala Ser Asp Trp Lys Lys Tyr Lys Lys Arg Leu Lys
 245 250 255
 Glu Ile Val Leu Ala Glu Lys Thr Gln Asn Ile Asp Phe Gln Asn Lys
 260 265 270
 Val Ile Ala Lys Ala Leu Asp Lys Ala Asn Val Lys Ile Lys Asp Gln
 275 280 285
 Ala Phe Ala Asn Ile Leu Ala Gln Tyr Ala Asn Thr Asp Lys Lys Ala
 290 295 300
 Ser Lys Ala Asn Thr Ser Lys Ser Asp Gln Lys Ser Ser Ser Asp Ser
 305 310 315 320
 Ser Lys Asp Ser Gln Ser Ser Lys Ser Lys Ser Glu Lys
 325 330

<210> 49
 <211> 1179
 <212> DNA
 <213> Streptococcus equi

<400> 49
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 atgaacaaaa aatcagcaag acgcaggcgt aagaatctta ttacgaagct tgcgatgaca 120
 agtgccttaa ccctgggtgt aggcgcgact accctagcag gacaaacaga agtacgggct 180
 gataatatct tacgcttaga tatgacagat aaagaagcag ttgaaaaatt cgctaacgag 240
 cttaaaaaatg aagtccataa aaactatcgt ggtagtaata cttggcaaaa gcttaccctt 300
 atacttaatg gttatcaaaa ccttagagaa caaatagaga ccgagctaaa aaatagtgaa 360
 caaaaagtaa aagagcttaa tgataagggt aatagtgaaa ctcaaggaaa acaagagtta 420
 cagaatcagc ttgagaaaga aaaagaagag ttagaaacac taaaaaaga gcttgaagct 480

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gagaaggcta aaggaactgg agaaacagag aagcttcaaa aggaaattga agcaaaaaat 540
gcaatgattt ctgacctaca aaaacagctt gaggaacta agcaaagggt tcaagagttt 600
gaagctgaag taggtaaatt aatggccgaa aaggcagacc tacaaacaaa attaaatgaa 660
caagagcagc ttaacgctaa gcttcaaaaa gaaattgaag acttaaaggc tcagattgaa 720
aagcttaagc actgtcaaga tacacctaag ccagagccta agccagagcc taagccagag 780
cctaagccag agcctaagcc agagcctaag ccagagccta agccagagcc taagccagag 840
cctaagccag ggcctaagcc agagcctaag ccagagccta agccagggcc taagccagag 900
cctaagccag agcctaagcc agggcctaag ccagggccta agccagagcc taagccaggg 960
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gaacaaccta aaccaatgac taaaccagga gctaagaagc ctgagcaatc acttccatca 1080
actggtgaca tcagaaatcc attcttcacg cctgcagcta ttgctattat gatcgagca 1140
ggtaccattg ccattccaaa acgcaaggaa gaagattaa 1179

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<210> 50
<211> 393
<212> PRT
<213> Streptococcus equi

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<400> 50
Met Thr Val Leu Lys Tyr Leu Gln Ser Lys Leu Ile Cys Leu Ser Leu
1      5      10      15
Lys Glu Arg Glu Met Asn Lys Lys Ser Ala Arg Arg Arg Arg Lys Asn
20      25      30
Leu Ile Thr Lys Leu Ala Met Thr Ser Ala Leu Thr Leu Gly Val Gly
35      40      45
Ala Ala Thr Thr Leu Ala Gly Gln Thr Glu Val Arg Ala Asp Asn Ile
50      55      60
Leu Arg Leu Asp Met Thr Asp Lys Glu Ala Val Glu Lys Phe Ala Asn
65      70      75      80
Glu Leu Lys Asn Glu Val His Lys Asn Tyr Arg Gly Ser Asn Thr Trp
85      90      95
Gln Lys Leu Thr Leu Ile Leu Asn Gly Tyr Gln Asn Leu Arg Glu Gln
100     105     110
Ile Glu Thr Glu Leu Lys Asn Ser Glu Gln Lys Val Lys Glu Leu Asn
115     120     125
Asp Lys Val Asn Ser Glu Thr Gln Gly Lys Gln Glu Leu Gln Asn Gln
130     135     140
Leu Glu Lys Glu Lys Glu Glu Leu Glu Thr Leu Lys Lys Glu Leu Glu
145     150     155     160
Ala Glu Lys Ala Lys Gly Thr Gly Glu Thr Glu Lys Leu Gln Lys Glu
165     170     175
Ile Glu Ala Lys Asn Ala Met Ile Ser Asp Leu Gln Lys Gln Leu Glu
180     185     190
Glu Thr Lys Gln Arg Val Gln Glu Phe Glu Ala Glu Val Gly Lys Leu
195     200     205
Met Ala Glu Lys Ala Asp Leu Gln Thr Lys Leu Asn Glu Gln Glu Gln
210     215     220
Leu Asn Ala Lys Leu Gln Lys Glu Ile Glu Asp Leu Lys Ala Gln Ile
225     230     235     240
Glu Lys Leu Lys His Cys Gln Asp Thr Pro Lys Pro Glu Pro Lys Pro
245     250     255
Glu Pro Lys Pro Glu Pro Lys Pro Glu Pro Lys Pro Glu Pro Lys Pro
260     265     270
Glu Pro Lys Pro Glu Pro Lys Pro Glu Pro Lys Pro Gly Pro Lys Pro
275     280     285
Glu Pro Lys Pro Glu Pro Lys Pro Gly Pro Lys Pro Glu Pro Lys Pro
290     295     300
Glu Pro Lys Pro Gly Pro Lys Pro Gly Pro Lys Pro Glu Pro Lys Pro
305     310     315     320
Gly Pro Lys Pro Glu Pro Lys Pro Glu Pro Lys Pro Glu Pro Lys Pro
325     330     335
Glu Ala Lys Lys Pro Glu Gln Pro Lys Pro Met Thr Lys Pro Gly Ala
340     345     350

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Lys Lys Pro Glu Gln Ser Leu Pro Ser Thr Gly Asp Ile Arg Asn Pro
 355 360 365
 Phe Phe Thr Pro Ala Ala Ile Ala Ile Met Ile Ala Ala Gly Thr Ile
 370 375 380
 Ala Ile Pro Lys Arg Lys Glu Glu Asp
 385 390

<210> 51
 <211> 573
 <212> DNA
 <213> Streptococcus equi

<400> 51
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 ttactacaa atgtattagc tgaagggata aatgataaaa tagaaaatgg cactgaaagc 120
 gatattagct tccaaaatgg tgaactccta aaaaattatc ttatcctaga aggtgaacga 180
 gtatactttg attatgatag agcaactcaa gataaagtat cagatgatgt tctagagatg 240
 ggaatgtag ttgaagctat aagtaaggat tattctgaga agacattcac cccagataaa 300
 tatttttaaag ctagttggcc tatccatggt aactattgtg gaccagggtca taatgggaat 360
 aactttacgt tgccagtagt agatgttttg gatcaagggt gccaaaacca cgatagttgc 420
 tataagtggg gtgccggtat tgggtgctaat tgtgaatgta accgtcagct cgtaattac 480
 ataaaagttt atagacgatg gatgccagcg aatgtccttg gcgtagccga tgctattaga 540
 gtatatatttg agacagtagg ttctatagga tgt 573

<210> 52
 <211> 191
 <212> PRT
 <213> Streptococcus equi

<400> 52
 Met Lys Lys Val Ile Asn Thr Ile Leu Leu Ala Ala Cys Phe Leu Phe
 1 5 10 15
 Phe Leu Gly Asn Phe Thr Thr Asn Val Leu Ala Glu Gly Ile Asn Asp
 20 25 30
 Lys Ile Glu Asn Gly Thr Glu Ser Asp Ile Ser Phe Gln Asn Gly Glu
 35 40 45
 Leu Leu Lys Asn Tyr Leu Ile Leu Glu Gly Glu Arg Val Tyr Phe Asp
 50 55 60
 Tyr Asp Arg Ala Thr Gln Asp Lys Val Ser Asp Asp Val Leu Glu Met
 65 70 75 80
 Gly Met Leu Val Glu Ala Ile Ser Lys Asp Tyr Ser Glu Lys Thr Phe
 85 90 95
 Thr Pro Asp Lys Tyr Phe Lys Ala Ser Trp Pro Ile His Gly Asn Tyr
 100 105 110
 Cys Gly Pro Gly His Asn Gly Asn Asn Phe Thr Leu Pro Val Val Asp
 115 120 125
 Val Leu Asp Gln Gly Cys Gln Asn His Asp Ser Cys Tyr Lys Trp Gly
 130 135 140
 Ala Gly Ile Gly Ala Asn Cys Glu Cys Asn Arg Gln Leu Val Asn Tyr
 145 150 155 160
 Ile Lys Val Tyr Arg Arg Trp Met Pro Ala Asn Val Leu Gly Val Ala
 165 170 175
 Asp Ala Ile Arg Val Tyr Phe Glu Thr Val Gly Ser Ile Gly Cys
 180 185 190

<210> 53
 <211> 930
 <212> DNA
 <213> Streptococcus equi

<400> 53
cagtctacaa atgacaatac aagtgtcatt acgatgaagg gcgacactat cagtgttagt 60
gattttttaca atgaaacaaa aaatacagag attttctcaa gagcaatgct aaaccttgtg 120
gttagtcgtg tttttgagga ccaatacggg aaaaagggtt ctaagaaaag aacggaagaa 180
gcttacaata aatcagctga gcaatacggg gcgtcattct ctgcagccct tgcgcagtct 240
ggcttgacaa cagataccta caagcgtcaa attcgtcag ccatgctggg tgaatatgct 300
gttaaagaag cagctaaaaa agagctgaca gatgctgatt acaaaaaagc ctatgagtca 360
tacacaccag aaatgactac tcaggtcact actctagaca atgaagaaac agcaaaggct 420
gttttaggtg aggttaaggc tgagggtgct gactttgctg ctattgctaa ggaaaagaca 480
acagcagcag acaagaaggc agactataag tttgactcag gagacactaa gttaccagca 540
gatgtgatca aggccgcctc aggattaaaa gagggtgata tttcagaggg ggtttctgtc 600
ttagatccgg ctacttatca aaacaagttc tatattgtta aggtaaccaa aaaagccgaa 660
aaggcttctg attggaagaa atataagaaa cgtctaaaag aaattgtctt ggctgaaaag 720
acacaaaaca ttgatttcca aaataaggctc attgcaaagg ccttagacaa ggcaaagtgt 780
aagatcaaag accaagcatt tgctaataatc ttggcacagt atgccaatat tgataaaaaa 840
gcaagcaagg cgaacacaag caagtcagat cagaaatcat cttcagactc aagcaaggat 900
agtcaatctt ctaaattctaa gtcagaaaaa 930

<210> 54
<211> 310
<212> PRT
<213> Streptococcus equi

<400> 54
Gln Ser Thr Asn Asp Asn Thr Ser Val Ile Thr Met Lys Gly Asp Thr
1 5 10 15
Ile Ser Val Ser Asp Phe Tyr Asn Glu Thr Lys Asn Thr Glu Ile Ser
20 25 30
Gln Arg Ala Met Leu Asn Leu Val Val Ser Arg Val Phe Glu Asp Gln
35 40 45
Tyr Gly Lys Lys Val Ser Lys Lys Arg Thr Glu Glu Ala Tyr Asn Lys
50 55 60
Ser Ala Glu Gln Tyr Gly Ala Ser Phe Ser Ala Ala Leu Ala Gln Ser
65 70 75 80
Gly Leu Thr Thr Asp Thr Tyr Lys Arg Gln Ile Arg Ser Ala Met Leu
85 90 95
Val Glu Tyr Ala Val Lys Glu Ala Ala Lys Lys Glu Leu Thr Asp Ala
100 105 110
Asp Tyr Lys Lys Ala Tyr Glu Ser Tyr Thr Pro Glu Met Thr Thr Gln
115 120 125
Val Thr Thr Leu Asp Asn Glu Glu Thr Ala Lys Ala Val Leu Gly Glu
130 135 140
Val Lys Ala Glu Gly Ala Asp Phe Ala Ala Ile Ala Lys Glu Lys Thr
145 150 155 160
Thr Ala Ala Asp Lys Lys Val Asp Tyr Lys Phe Asp Ser Gly Asp Thr
165 170 175
Lys Leu Pro Ala Asp Val Ile Lys Ala Ala Ser Gly Leu Lys Glu Gly
180 185 190
Asp Ile Ser Glu Val Val Ser Val Leu Asp Pro Ala Thr Tyr Gln Asn
195 200 205
Lys Phe Tyr Ile Val Lys Val Thr Lys Lys Ala Glu Lys Ala Ser Asp
210 215 220
Trp Lys Lys Tyr Lys Lys Arg Leu Lys Glu Ile Val Leu Ala Glu Lys
225 230 235 240
Thr Gln Asn Ile Asp Phe Gln Asn Lys Val Ile Ala Lys Ala Leu Asp
245 250 255
Lys Ala Asn Val Lys Ile Lys Asp Gln Ala Phe Ala Asn Ile Leu Ala
260 265 270
Gln Tyr Ala Asn Thr Asp Lys Lys Ala Ser Lys Ala Asn Thr Ser Lys
275 280 285

Ser Asp Gln Lys Ser Ser Ser Asp Ser Ser Lys Asp Ser Gln Ser Ser
 290 295 300
 Lys Ser Lys Ser Glu Lys
 305 310

<210> 55
 <211> 600
 <212> DNA
 <213> Streptococcus equi

<400> 55
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 acagataaag aagcagttga aaaattcgct aacgagctta aaaatgaagt ccataaaaac 120
 tatcgtggta gtaataacttg gcaaaagctt acccttatac ttaatgggta tcaaaacctt 180
 agagaacaaa tagagaccga gctaaaaaat agtgaacaaa aagtaaaaga gcttaatgat 240
 aagggttaata gtgaaactca aggaaaacaa gagttacaga atcagcttga gaaagaaaaa 300
 gaagagtttag aaacactaaa aaaagagctt gaagctgaga aggctaaagg aactggagaa 360
 acagagaagc ttcaaaagga aattgaagca aaaaatgcaa tgatttctga cctacaaaaa 420
 cagcttgagg aaactaagca aagggttcaa gagtttgaag ctgaagtagg taaattaatg 480
 gccgaaaagg cagacctaca aacaaaatta aatgaacaag agcagcttaa cgctaagctt 540
 caaaaagaaa ttgaagactt aaaggctcag attgaaaagc ttaagcactg tcaagataca 600

<210> 56
 <211> 200
 <212> PRT
 <213> Streptococcus equi

<400> 56
 Ala Thr Thr Leu Ala Gly Gln Thr Glu Val Arg Ala Asp Asn Ile Leu
 1 5 10 15
 Arg Leu Asp Met Thr Asp Lys Glu Ala Val Glu Lys Phe Ala Asn Glu
 20 25 30
 Leu Lys Asn Glu Val His Lys Asn Tyr Arg Gly Ser Asn Thr Trp Gln
 35 40 45
 Lys Leu Thr Leu Ile Leu Asn Gly Tyr Gln Asn Leu Arg Glu Gln Ile
 50 55 60
 Glu Thr Glu Leu Lys Asn Ser Glu Gln Lys Val Lys Glu Leu Asn Asp
 65 70 75 80
 Lys Val Asn Ser Glu Thr Gln Gly Lys Gln Glu Leu Gln Asn Gln Leu
 85 90 95
 Glu Lys Glu Lys Glu Glu Leu Glu Thr Leu Lys Lys Glu Leu Glu Ala
 100 105 110
 Glu Lys Ala Lys Gly Thr Gly Glu Thr Glu Lys Leu Gln Lys Glu Ile
 115 120 125
 Glu Ala Lys Asn Ala Met Ile Ser Asp Leu Gln Lys Gln Leu Glu Glu
 130 135 140
 Thr Lys Gln Arg Val Gln Glu Phe Glu Ala Glu Val Gly Lys Leu Met
 145 150 155 160
 Ala Glu Lys Ala Asp Leu Gln Thr Lys Leu Asn Glu Gln Glu Gln Leu
 165 170 175
 Asn Ala Lys Leu Gln Lys Glu Ile Glu Asp Leu Lys Ala Gln Ile Glu
 180 185 190
 Lys Leu Lys His Cys Gln Asp Thr
 195 200

<210> 57
 <211> 309
 <212> DNA
 <213> Streptococcus equi

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<400> 57
gaagggataa atgataaaat agaaaatggc actgaaagcg atattagctt ccaaaatggt 60
gaactcctaa aaaattatct taccctagaa ggtgaacgag tatactttga ttatgataga 120
gcaactcaag ataaagtatc agatgatggt ctagagatgg gaatgttagt tgaagctata 180
agtaaggatt attctgagaa gacattcacc ccagataaat attttaaagc tagttggcct 240
atccatggta actattgtgg accaggatcat aatgggaata actttacgtt gccagtagta 300
gatgttttg 309

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<210> 58
<211> 103
<212> PRT
<213> Streptococcus equi

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<400> 58
Glu Gly Ile Asn Asp Lys Ile Glu Asn Gly Thr Glu Ser Asp Ile Ser
1          5          10          15
Phe Gln Asn Gly Glu Leu Leu Lys Asn Tyr Leu Ile Leu Glu Gly Glu
20          25          30
Arg Val Tyr Phe Asp Tyr Asp Arg Ala Thr Gln Asp Lys Val Ser Asp
35          40          45
Asp Val Leu Glu Met Gly Met Leu Val Glu Ala Ile Ser Lys Asp Tyr
50          55          60
Ser Glu Lys Thr Phe Thr Pro Asp Lys Tyr Phe Lys Ala Ser Trp Pro
65          70          75          80
Ile His Gly Asn Tyr Cys Gly Pro Gly His Asn Gly Asn Asn Phe Thr
85          90          95
Leu Pro Val Val Asp Val Leu
100

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CLAIMS:

1. An immunogenic composition comprising at least one polypeptide, wherein said polypeptide is a fragment of Se50 protein, said fragment consisting of the amino acid sequence as shown in SEQ ID NO: 32, and a pharmaceutically acceptable excipient, diluent or
5 vehicle.
2. An immunogenic composition comprising at least one recombinant vector and at least one polynucleotide inserted therein coding for a polypeptide, wherein said polypeptide is a fragment of Se50 protein, said fragment consisting of the amino acid sequence as shown in SEQ ID NO: 32, and a pharmaceutically acceptable excipient, diluent or vehicle, and said
10 vector is able to express *in vivo* the polypeptide in a mammal susceptible to streptococcal infection caused by *Streptococcus equi*.
3. An immunogenic composition comprising at least one recombinant vector and at least one polynucleotide inserted therein, wherein said polynucleotide consists of the nucleotide sequence as shown in SEQ ID NO: 31 or said polynucleotide consists of a
15 nucleotide sequences having a sequence identity which is equal or more than about 95% with SEQ ID NO: 31, and a pharmaceutically acceptable excipient, diluent or vehicle, and said vector is able to express *in vivo* the polypeptide encoded by said polynucleotide in a mammal susceptible to streptococcal infection caused by *Streptococcus equi*, and said encoded peptide elicits an immunogenic response against *Streptococcus equi*.
- 20 4. The immunogenic composition according to claim 3 wherein said polynucleotide has a sequence identity which is equal or more than about 98% with SEQ ID NO: 31.
5. The immunogenic composition of claim 3 or 4, wherein the vector is an expression vector.
- 25 6. The immunogenic composition of claim 5, wherein the vector is a plasmid or a virus.

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7. The immunogenic composition of claim 6, wherein the virus is a poxvirus, an equine herpesvirus (EHV), a human adenovirus, or an equine encephalitis virus.

8. Use of at least one polypeptide which is a fragment of Se50 protein, said fragment consisting of the amino acid sequence as shown in SEQ ID NO: 32, in the
5 preparation of a medicament for eliciting an immunogenic response in equine against *Streptococcus equi* infection.

9. Use of at least one immunogenic composition as defined in any one of claims 2 to 7 in the preparation of a medicament for eliciting an immunogenic response in equine against *Streptococcus equi* infection.

10 10. An antibody preparation comprising at least one antibody specific for at least one polypeptide, wherein said polypeptide is a fragment of Se50 protein, said fragment consisting of the amino acid sequence as shown in SEQ ID NO: 32, and which antibody is polyclonal or monoclonal, or a fragment of said antibody that is specific for said polypeptide, and a pharmaceutically acceptable excipient, diluent or vehicle.

15 11. A diagnostic kit for detecting infection by a streptococcal bacterium comprising at least one polypeptide, wherein said polypeptide is a fragment of Se50 protein, said fragment consisting of the amino acid sequence as shown in SEQ ID NO: 32, or the antibody preparation of claim 10, and a pharmaceutically acceptable excipient, diluent or vehicle.

20 12. A diagnostic method for detecting infection by a streptococcal bacterium, comprising using the diagnostic kit of claim 11 and detecting in a sample the polypeptide or an antibody specific to that polypeptide.

13. Use of the antibody preparation of claim 10 in the preparation of a medicament for passive immunization wherein said antibody preparation is formulated for administration
25 to an equine susceptible to infection by *Streptococcus equi* or infected by *Streptococcus equi*.

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14. The antibody preparation of claim 10 for use in passive immunization wherein said antibody preparation is formulated for administration to an equine susceptible to infection by *Streptococcus equi* or infected by *Streptococcus equi*.

15. A polypeptide that is a fragment of Se50 protein, said fragment consisting of
5 the amino acid sequence as shown in SEQ ID NO: 32 for use in eliciting an immunogenic response in equine against *Streptococcus equi* infection.

16. An immunogenic composition as defined in any one of claims 2 to 7 for use in eliciting an immunogenic response in equine against *Streptococcus equi* infection.

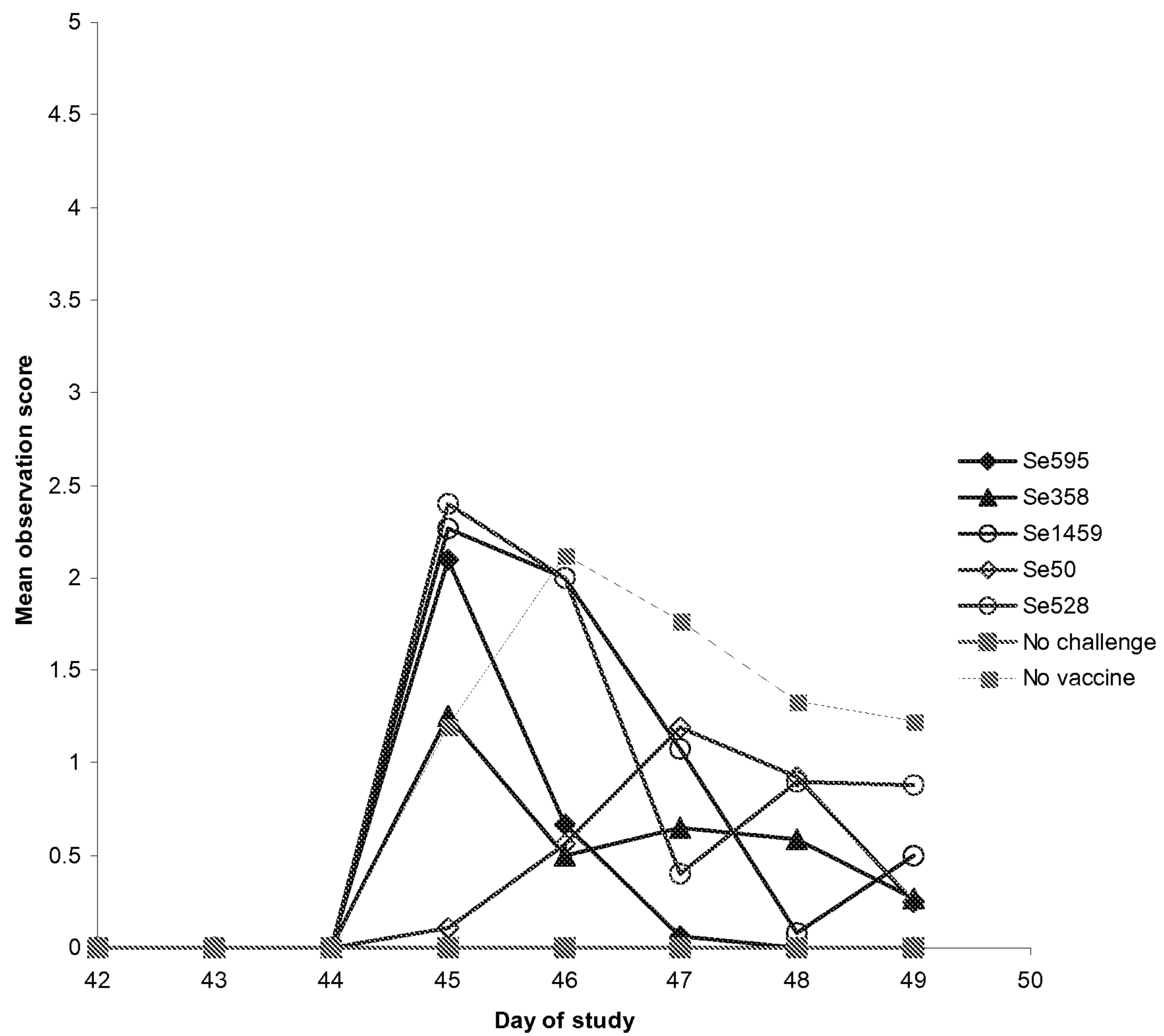


FIG. 1

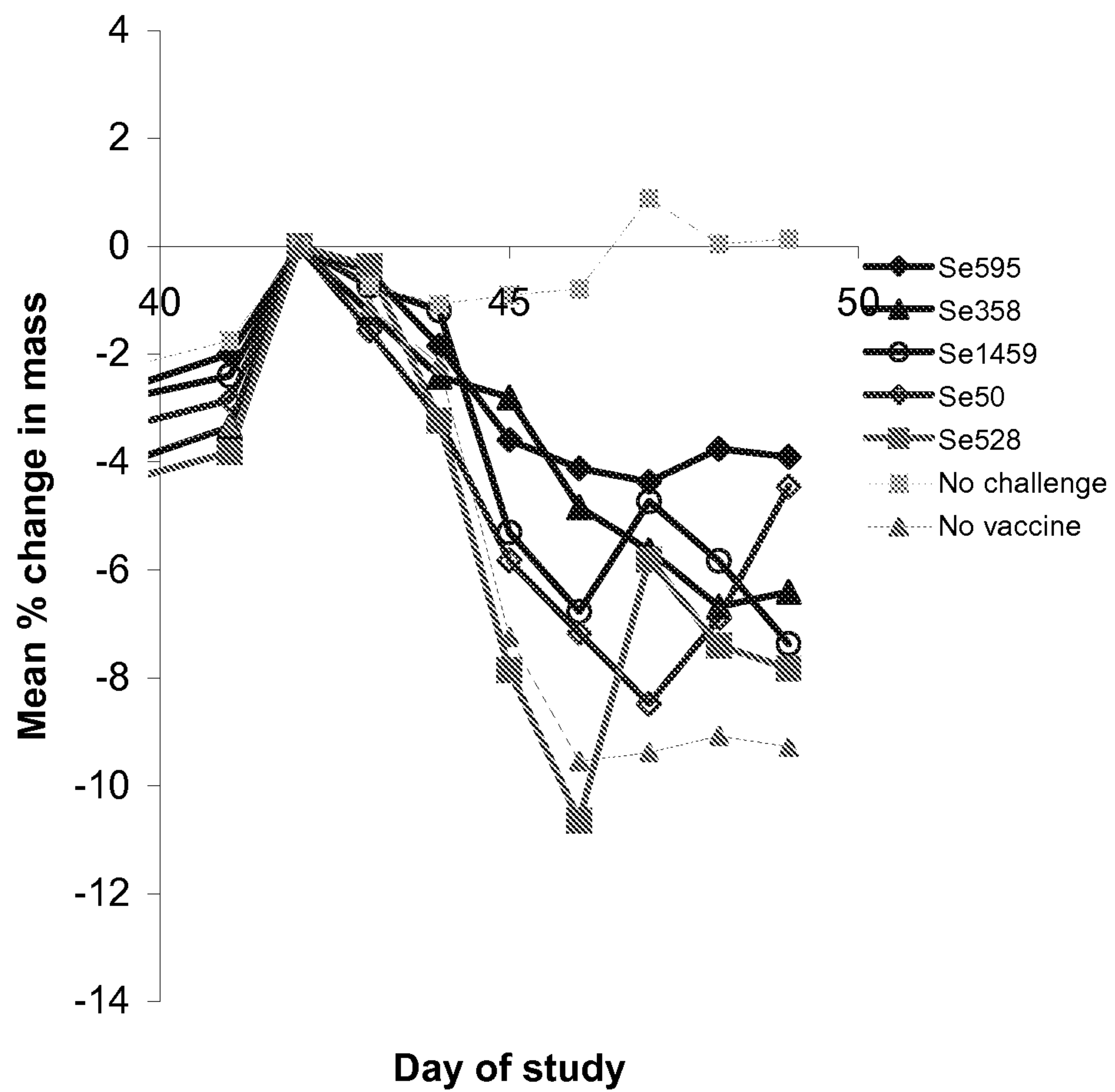


FIG. 2

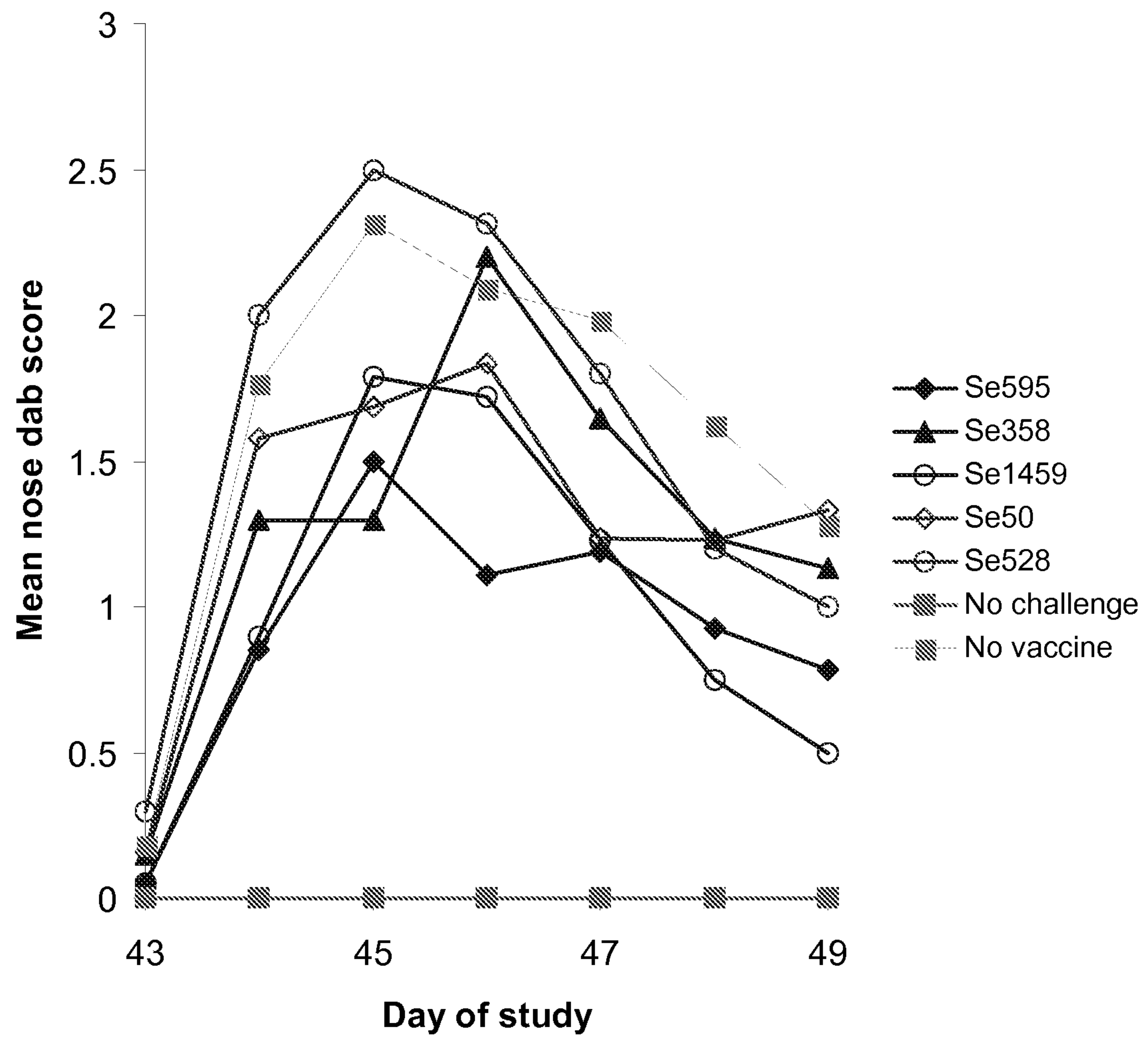


FIG. 3

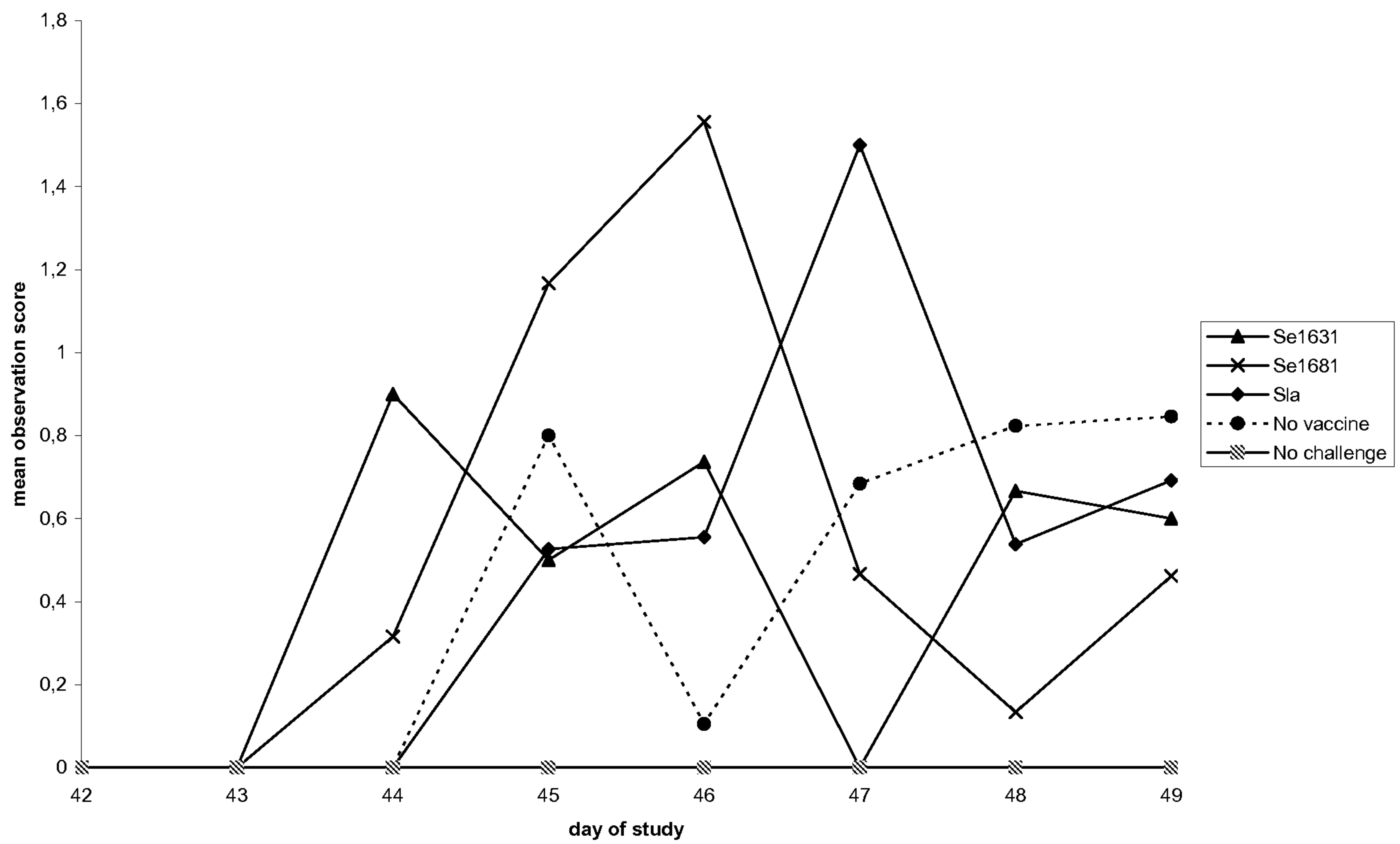


FIG. 4

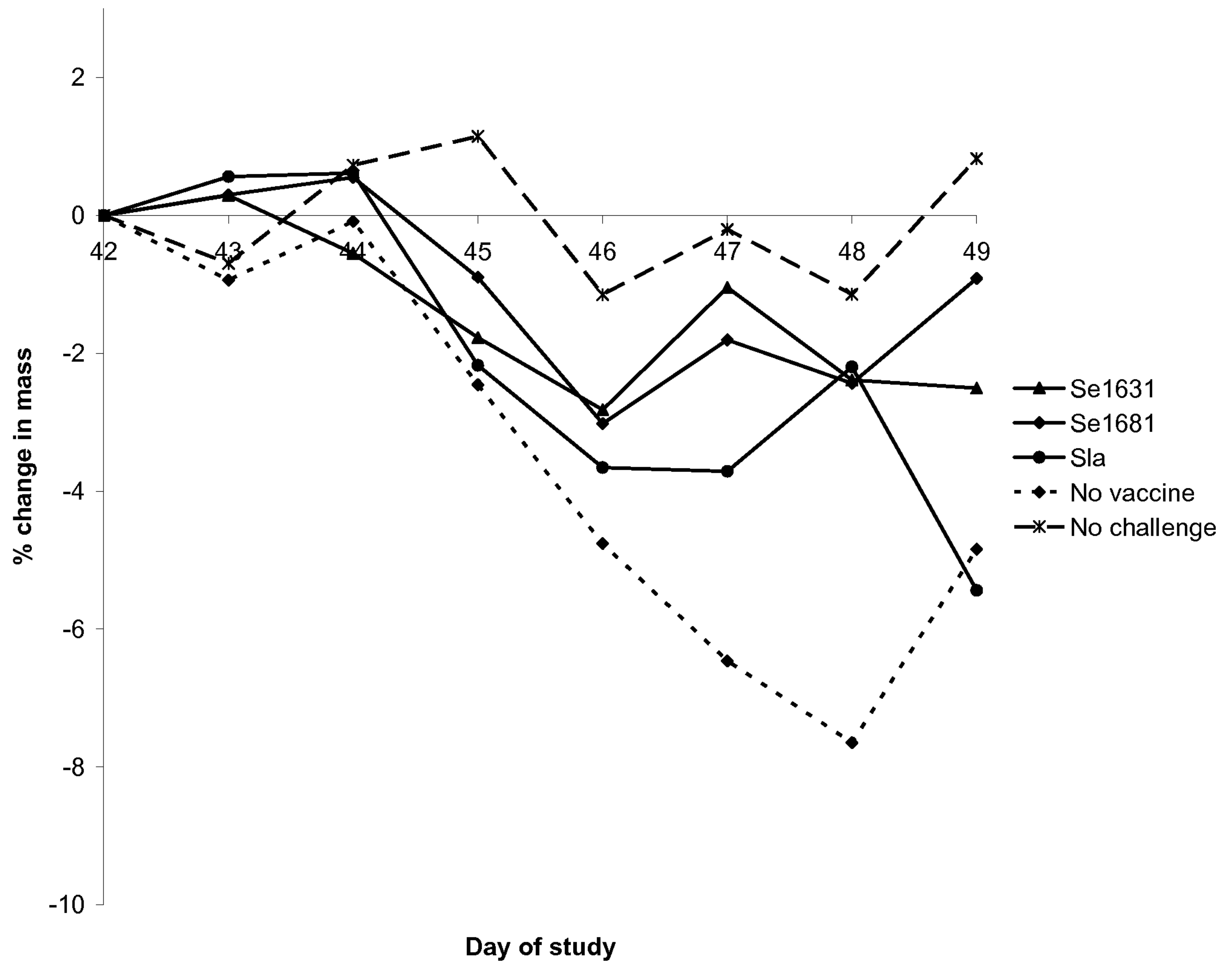


FIG. 5

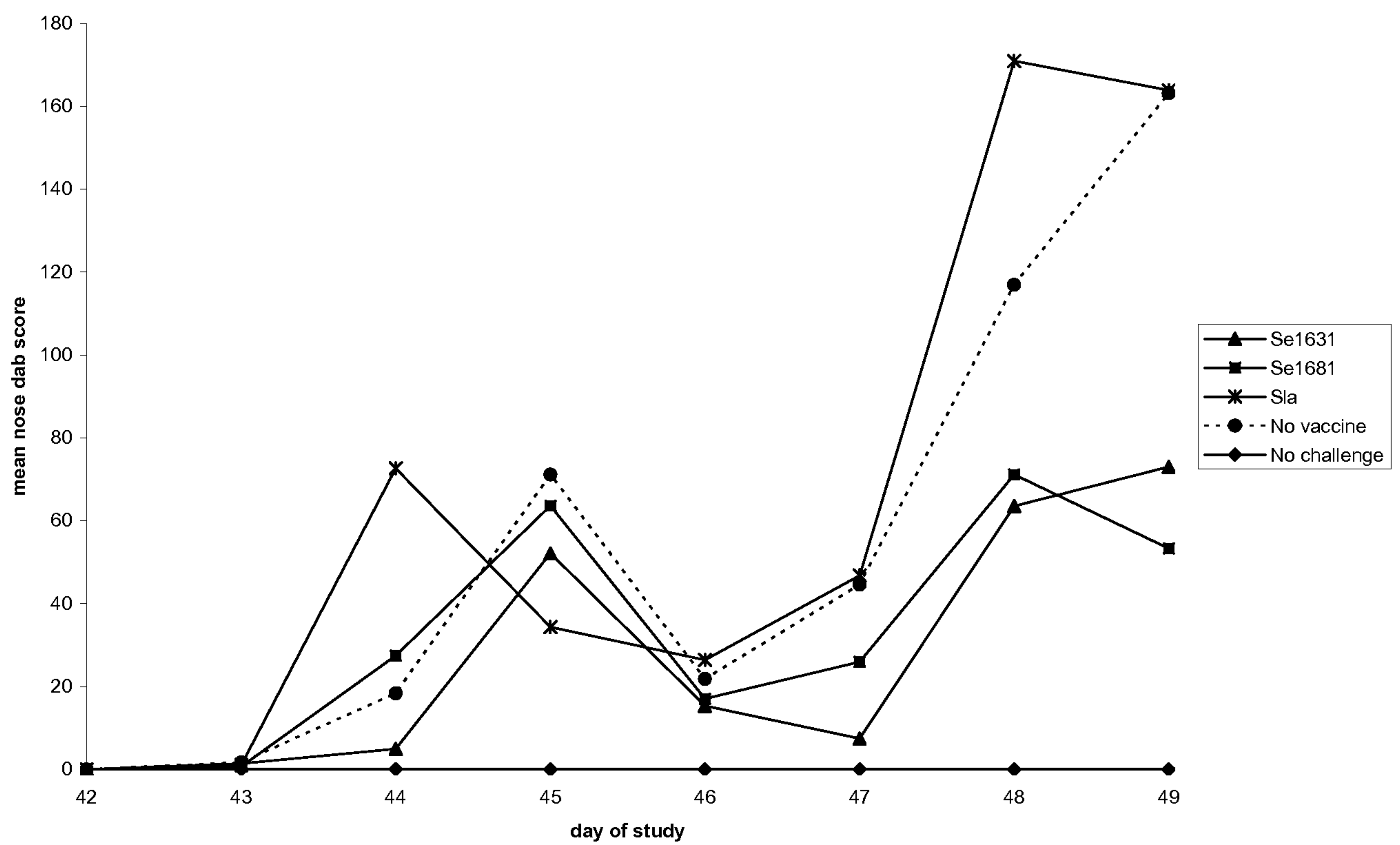


FIG. 6

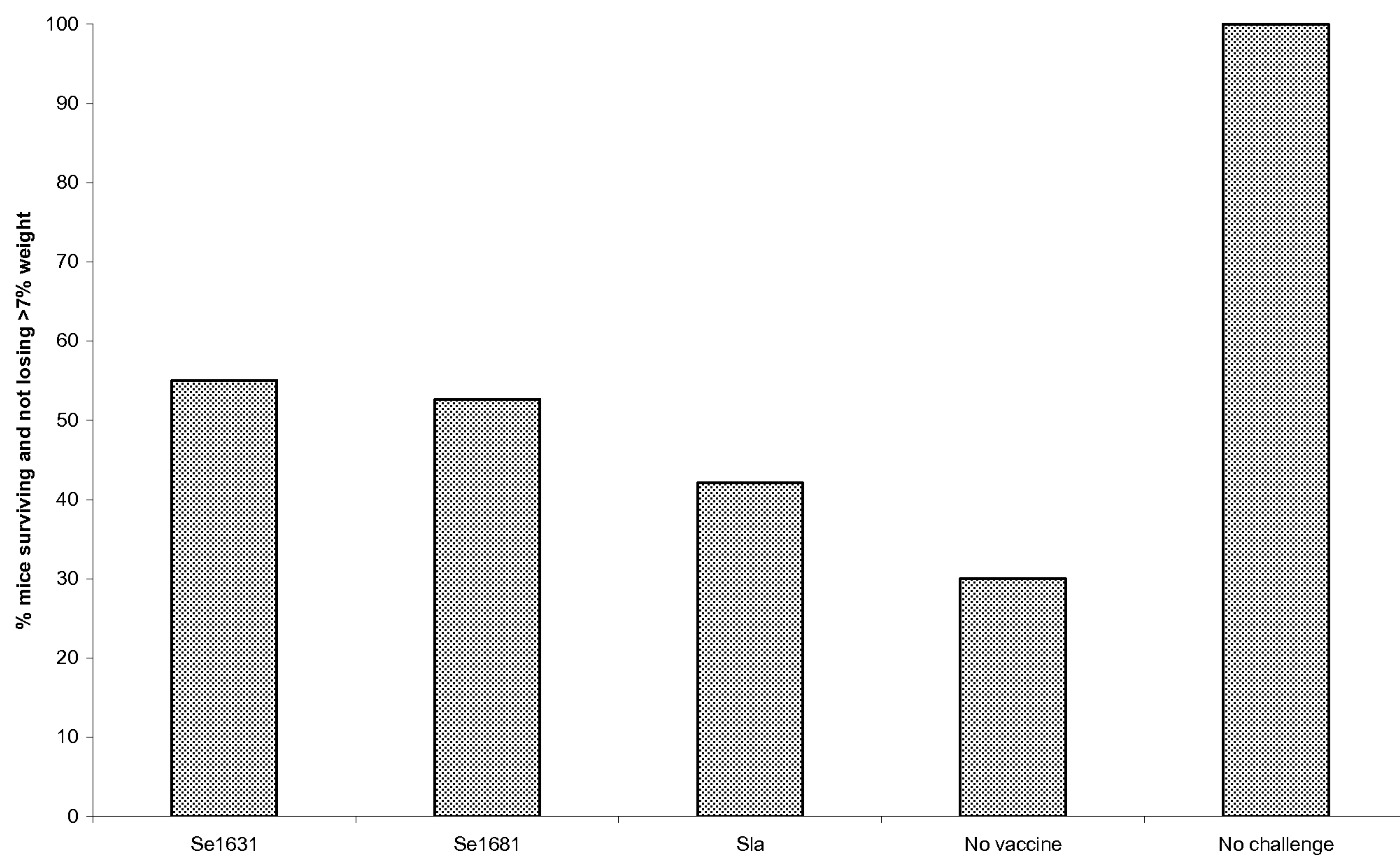


FIG. 7

