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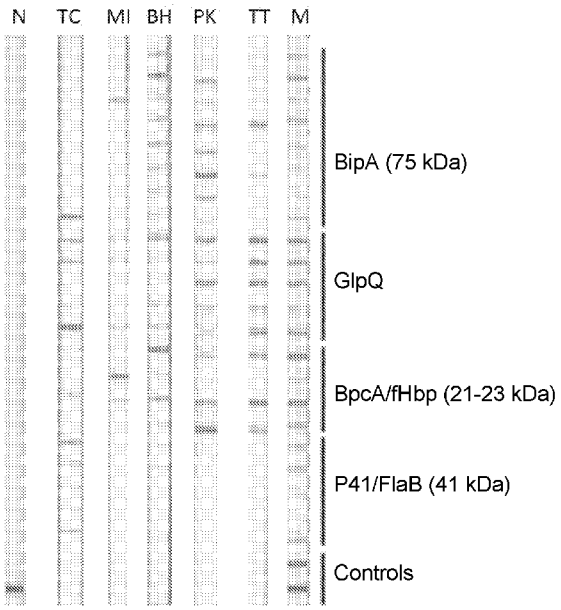
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(54) Title: SPECIES-SPECIFIC ANTIGEN SEQUENCES FOR TICK-BORNE RELAPSING FEVER (TBRF) AND METHODS OF USE

Fig. 1



(57) Abstract: The disclosure, in some aspects, provides antigen-specific amino acid sequences for tick-borne relapsing fever (TBRF) Borrelia species. Specifically, the disclosure provides compositions and methods that may be used to identify antibodies to infection by one or more TBRF Borrelia species in samples from subjects suspected of having TBRF. Further disclosed are compositions that include one or more labeled and/or tagged and/or bound amino acid sequences.

[Continued on next page]

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SPECIES-SPECIFIC ANTIGEN SEQUENCES FOR TICK-BORNE RELAPSING FEVER (TBRF) AND METHODS OF USE

FIELD OF THE INVENTION

Aspects of the present invention provide novel compositions and methods for diagnosing Tick-Borne Relapsing Fever (TBRF) resulting from infection by diverse TBRF *Borrelia* species.

SEQUENCE LISTING

The instant application incorporates by reference the Sequence Listing in the ASCII text file filed January 15, 2021, entitled "ID-FISH 0153-2015US01_ST25.txt", which file was created on January 7, 2021, the size of which file is 27,388 bytes.

BACKGROUND OF THE INVENTION

Borreliosis is caused by two groups of *Borrelia*, the *B. burgdorferi* group and the Tick-Borne Relapsing Fever (TBRF) *Borrelia* group (also referred to herein as RFB (relapsing fever *Borrelia*)). The *B. burgdorferi* group was once thought to be the only group that caused Lyme-like symptoms in infected subjects, but it is now known that TBRF *Borrelia* also causes Lyme-like symptoms. Several *Borrelia* species cause TBRF, and these are usually associated with specific species of ticks. For instance, *B. hermsii* is transmitted by *Ornithodorus hermsi* ticks, *B. parkerii* by *O. parkeri* ticks, and *B. turicatae* by *O. turicata* ticks. Each tick species has a preferred habitat and preferred set of hosts. Typically, hard (*Ixodes*) ticks transmit Lyme disease pathogens whereas soft (*Ornithodorus*) ticks transmit TBRF pathogens, but there are exceptions to this rule: several RFB agents are transmitted only by hard ticks (e.g., *B. miyamotoi*) and one species, *B. recurrentis*, is louse transmitted.

The main symptoms of TBRF are high fever (*e.g.*, 103°F), headache, and muscle and joint aches. Symptoms can reoccur, producing a telltale pattern of fever lasting roughly three days, followed by 7 days without fever, followed by another three days of fever. Without antibiotic treatment, this process can repeat several times.

Currently, the standard for identification is by identification of TBRF spirochetes in blood smears of a subject presenting symptoms consistent with TBRF. After obtaining a blood draw, a sample must be cultured for at least 24 hours to facilitate identification. However, even early in the disease when spirochetes are highest, positive identification is only made about 70% of the time. (*See*, www.cdc.gov/relapsing-fever/clinicians/index.html).

Thus, materials and methods of the current state of the art result in a delay in diagnosis and provide a relatively low level of sensitivity and specificity. New materials and methods are needed that are suitable for the identification of TBRF causative agents with decreased assay time and increased sensitivity and specificity.

SUMMARY

The invention, in part, relates to compositions and methods that may be used to identify antibodies to infection by one or more TBRF *Borrelia* species in samples from subjects suspected of having TBRF. With compositions and methods of the present invention, identification of TBRF *Borrelia* in subject samples may be performed with greater speed, sensitivity, and specificity than with existing compositions and methods. Antigen-specific amino acid sequences of the present invention may be used in diagnostic and scientific assays. Non-limiting examples of suitable assays include immunoblots, ELISA (enzyme-linked immunosorbent assay), etc. Amino acid sequences of the present invention may be used for detecting TBRF *Borrelia* specific T-cells (*e.g.*, the IgXSPOT test; IGeneX, Palo Alto, CA).

According to an aspect of the invention, compositions that include one or more labeled and/or tagged and/or bound amino acid sequences, wherein the one or more labeled and/or tagged and/or bound amino acid sequences comprise amino acid sequences selected from the group consisting of SEQ ID NO: 1, SEQ ID NO: 2, SEQ ID NO: 3, SEQ ID NO: 4, SEQ ID NO: 5, SEQ ID NO: 6, SEQ ID NO: 7, SEQ ID NO: 8, SEQ ID NO: 9, SEQ ID NO: 10, and SEQ ID NO: 11, and variants thereof which retain the immunological binding profile of Fig. 3 are provided. In some embodiments, the composition also includes amino acid sequences that are bound to a substance selected from the group consisting of nitrocellulose, nylon, polyvinylidene difluoride (PVDF), magnetic beads, and agarose. In some embodiments, each of said one or more amino acid sequences are tagged with an antibody with specificity for said amino acid sequence.

According to an aspect of the invention, methods are provided for detecting antibodies resulting from infection by one or more members of Relapsing Fever (TBRF) Borrelia genus, wherein the genus comprises *B. hermsii*, *B. turicatae*, *B. miyamotoi*, *B. duttonii*, and *B. parkeri*, if present in a sample from a subject suspected of having tick-borne relapsing fever (TBRF), the methods including: providing a biological sample obtained from a subject suspected of having TBRF; mixing the biological sample with one or more of the labeled and/or tagged and/or bound amino acid sequences selected from the group consisting

of SEQ ID NO: 1, SEQ ID NO: 2, SEQ ID NO: 3, SEQ ID NO: 4, SEQ ID NO: 5, SEQ ID NO: 6, SEQ ID NO: 7, SEQ ID NO: 8, SEQ ID NO: 9, SEQ ID NO: 10, and SEQ ID NO: 11, and variants thereof which retain the immunological binding profile of Fig. 3; and detecting a positive immunobinding reaction which indicates the presence of TBRF specific antibodies in the sample. In some embodiments, two or more of the labeled and/or tagged and/or bound amino acid sequences are mixed with the biological sample and a sample is considered positive for TBRF if at least two amino acid sequences are detected. In some embodiments, the labeled and/or tagged and/or bound amino acid sequences are detected with anti-human IgG or anti-human IgM antibody linked to a detectable moiety. In some embodiments, the detectable moiety is selected from the group consisting of chromophores, radioactive moieties and enzymes. In some embodiments, the detectable moiety includes alkaline phosphatase. In some embodiments, the detectable moiety includes biotin.

According to an aspect of the invention, a vector including a nucleic acid sequence encoding an amino acid sequence selected from the group consisting of SEQ ID NO: 1, SEQ ID NO: 2, SEQ ID NO: 3, SEQ ID NO: 4, SEQ ID NO: 5, SEQ ID NO: 6, SEQ ID NO: 7, SEQ ID NO: 8, SEQ ID NO: 9, SEQ ID NO: 10, and SEQ ID NO: 11, and variants thereof which retain the immunological binding profile of Fig. 3, and a heterologous nucleic acid sequence is provided. In some embodiments, the nucleic acid sequence is operatively linked to a promoter sequence. In some embodiments, the vector is a prokaryotic vector. In some embodiments, a cell includes the vector. In some embodiments, the cell is a bacterial cell.

BRIEF DESCRIPTION OF THE DRAWINGS

Fig. 1 shows a photomicrographic image of a line immunoblot. TBRF ImmunoBlot strips were tested with rabbit anti-TBRF *Borrelia* serum samples. Antibodies to TBRF *Borrelia* species *B. turcica*, *B. miyamotoi*, *B. hermsii*, *B. parkeri,* and *B. turicatae* were detected by TBRF ImmunoBlot strips prepared from recombinant antigens from all five species, demonstrating that the TBRF ImmunoBlot detected TBRF *Borrelia* group antibodies. N, negative control; TC, *B. turcica*; MI, *B. miyamotoi*; BH, *B. hermsii*; PK, *B. parkeri*, TT, *B. turicatae*; and M, positive control.

Fig. 2 shows a photomicrographic image of TBRF ImmunoBlots of representative patient serum samples. Patient serum samples were tested on Tick-Borne Relapsing Fever (TBRF) IgM and IgG ImmunoBlots. These results illustrate how detection of antibodies to TBRF Borrelia species identified patients with Chronic Lyme Disease symptoms that would

otherwise have been missed by only testing for *Borrelia burgdorferi* species antigens. P, positive control; N, negative control; G, TBRF ImmunoBlot IgG; M, TBRF ImmunoBlot IgM; 1, *B. hermsii*-positive IgM; 2, *B. turicatae*-positive IgG; 3, *B. miyamotoi*-positive IgM; 4, *B. turcica*-positive IgG; 5, *B. parkeri*-positive IgM and IgG.

Fig. 3 presents a table showing representative results of TBRF ImmunoBlot IgM and IgG tests with SEQ ID NOs: 1–11 on patient serum samples. Pos, positive; I, indeterminate; Tt, *B. turicatae*; Pk, *B. parkeri*; Mia, *B. miyamotoi*; Du, *B. duttonii*; Bh, *B. hermsii*;

DESCRIPTION OF THE SEQUENCES

SEQ ID NO: 1 *B. hermsii* Vlp7

MQQPEAGKTGVSGGVNGNLGNSLMELGRSAENAFYAFIELVSDVLGFTAKSDTTKQ
EVGGYFNSLGAKLGEASNDLEQVAVKAETGVDKSDSSKNPIREAVNEAKEVLGTLK
GYVESLGTIGDSNPVGYANNAAGSGTTAADDELRKAFKALQEIVKAATDAGVKALK
IGATTLQANGGADNKEGAKILATSGGNPAAADVAKAAAILSSVSGEEMLSSIVKSGE
NDAQLAAAADGNTSAISFAKGGSDAHLAGANTPKAAAVAGGIALRSLVKTGKLAA
GAADNATGGGKEVQGVGVAAANKLLRAVEDVIKKTVKNVLEKAKEKIDKARGSQE
PVSESSK

SEQ ID NO: 2 *B. turicatae* BipA-2H

MDNVMSGIDNVIQGAGTFATAAMQGVGTVIDVLQDVGTFVISDIQNMGARMLFGT
GENSSVASEGSESVMSLSSNDSSEAKDVTVVLSSDVTSKGNDVAVGLSSDETQVIGR
LEKYLQSAIKINGRSDSDQSNLESGRKKFFNWLKTSDTNASKRKELVQDLQKVFDLI
KEKSSDSTELKHWVQSIVDRIEDKSTIVDIDSDDELNNDKEVDFLIENTLASRDYSGF
AVSLLFQSLADTLYDSDNNRDKSEEQIFQDLRKVFSDSSDKSEGVLGFKSKIEATN

SEQ ID NO: 3 *B. turicatae* BpcA/fHbp

MSETSLLNIETNLLNTLDDNQKQALITFKDLLQDKKHLSILEKQQKSILEDLKANQKN
YNLQDKLKKTLNSEYDKNQLNKLFDELGNIKTKQFLQQLHIILQSIKDGKPTNFASSN
FNNLNQTLEQKKEQALKYIKDKLYTDYYLYINGIQDANYFFERIMSLLE

SEQ ID NO: 4 *B. turicatae* P41/FlaB

MRNNSINAANLSKTQEKLSSGHRINRASDDAAGMGVAGKINAQIRGLSQASRNTSK
AINFIQTTEGNLNEVEKVLVRMKELAVQSGNGTYSDADRGSIQIEIEQLTDEINRIADQ
AQYNQMHMLSNKSAAQNIKTAEELGMQPAKINTPASLAGSQASWTLRVHVGANQD
EAIAVNIYAANVANLFAGEGAQVSPAQEGAQQEGVQAAPAPAAAPAQGGVNSPVN
VTTTIDANMSLSKIENAIRMVSDQRANLGAFQNRLESIKASTEYAIENLKSSYAQIQD
ATMTDEIVASTTNSILTQSAMAMIAQANQVPQYVLSLLR

SEQ ID NO: 5 *B. parkeri* BipA-1H

MDMGSTRDWLTNDDGFVRGTKGFDDSPFRRPDRVDKEVSAGGREIEKAFSRNLGV
AGGQRKGTDDVKNGIAGARESGGVLKEAENAGQRDVDDSGEGIKNDVIQNLGSVG
VQVAVGSENNGDDSGQEAEKGSQNLGDTGTQRVVSTSDLNSHLGVESKGGMSTNK
EGISTNHVTENRNSINSITSTSSGLSTALQIAGTSTRASGYEGEVTTNAQDRSFIDTKTQ
DSKKQYSDFSDQDIRDKILGNVVGGVV

SEQ ID NO: 6 *B. parkeri* BipA-2H

MGNVMSGIDNVIQGAGTFATAAMQGVGTVIDVLQDVGTFVISDIQNMGARMFFGT
GESSSVASEDSESVMSLSSKASSEAKDTTVGLSSDVTSKGNNVAVGLSSDEIQIIGRLE
KYLKSAIKINGRSDSDQSKLESGHKKFFQWLKTSDTNASKRKELVQNLQRVFNLIKE
KSSDSTELKKWMQSIVDDIENKSTIIDINSDDKLNNDKEVDFLIEKTLGSSDYSGFAVS
LLFQALADTLYDSENSRDKSEEQIFKDLRKVFSDKSEGVLEFKSKIEATN

SEQ ID NO: 7 *B. parkeri* GlpQ

MCQNEKMSMTNKKSPLTIAHRGASGYLPEHTLESKAFAYALGADYLEQDIVLTKDN
VPIIMHDPELDTTTNVAKLFPERARENGRYYSVDFTLDELKSLSLSERFDLETRKPIYP
KRFPLNEYNFKIPTLEEEIQFIQGLNKSTGRNVGIYPEIKKPLWHKQQGKDISKIVIEIL
NKYGYKSKEDKIYLQTFDFDELKRIREELGYQGKLIMLVGENDWNEAPTDYEYIKSQ
EGMTEVAKYADGIGPWIPQIIIDGKITDLTSLAHKYNMEVHAYTFRIDSLPSYVKDAN
ELLDLLFNQAKIDGLFTDFTDTVVKFIKQ

SEQ ID NO: 8 *B. parkeri* BpcA/fHbp

MSETSLLNTLDNNQKQALITFKDLLQDKNHRSILEKQQKSILKDLEKHQENSNLQDK
LKKTLNSEYDKTQLNKLFDELGNIKTKQFLQKLHIMLKSINNGTLTSFSSSNFKDSNQ
TLEQKKEQALQYIKGQLYTDYYLYINGIQDANYFFERIMSVLEI

SEQ ID NO: 9 *B. parkeri* P41/FlaB

MRNNSINAANLRKTQEKLSSGHRINRASDDAAGMGVAGKINAQIRGLSQASRNTSK
AINFIQTTEGNLNEVEKVLVRMKELAVQSGNGTYSDADRGSIQIEIEQLTDEINRIADQ
AQYNQMHMLSNKSAAQNIKTAEELGMQPAKINTPASLAGAQASWTLRVHVGANQ
DEAIAVNIYASNVANLFAGEGAQVSPAQEGAQQEGVQAAPAPAAAPAQGGVNSPV
NVTTTVDANMSLSKIENAIRMVSDQRANLGAFQNRLESIKASTEYAIENLKSSYAQIK
DATMTDEIVASTTNSILTQSAMAMIAQANQVPQYVLSLLR

SEQ ID NO: 10 *B. duttonii* GlpQ

MENAKINKKSALIIAHRGASGYLPEHTLEAKAYAHALGADYIEQDIVLTKDDIPIVMH
DPELDTTTNVAKLFPGRARENGKYYSVDFTLAEIKSLSLSERFDPETQQPIYPNRFPAT
EYDFKIPTLEEEIKFIQGLNKSTGKNIGIYPEIKKPLWHKQQGKDISKIVIDILNKYGYK
SKEDKIYLQTFDFDEIKRIREELGYQGKLIMLVGENDWEEAPTDYEYIKSEEGMAEV
AKYADGIGPWIPQIIINGQITGLISLAHKYNMQVHPYTFRIDALPSYVKDPNELLELLFI
KAKVDGLFTDFVDISIKFMQ

SEQ ID NO: 11 *B. miyamotoi* GlpQ

MASMTGGQQMGRGSEMGENKKSPLIIAHRGASGYLPEHTLEAKAYAYALGADYLE
QDIVLTKDNIPVIMHDPEIDTTTNVAQLFPNRARENGRYYATDFTLTELKSLNLSERF
DPENKKPIYPNRFPLNEYNFKIPTLEEEIQFIQGLNKSTGKNVGIYPEIKKPFWHKQQG
KDISKIVIEILNKYGYKSKEDKIYLQTFDFDELKRIRKELGYQGKLIMLVGENDWNEA
PTDYEYIKSEEGIAEVAKYSDGIGPWIPQIIIDGKITELTNLAHKYNIEVHPYTFRTDAL
PSYVKNENELLDLLFNKAKVDGIFTDFTDTVMNFIKK

DETAILED DESCRIPTION OF THE INVENTION

The RFB species best known for causing TBRF in the United States of America (USA) are *B. hermsii, B. miyamotoi, B. parkeri*, and *B. turicatae*. However, other RFB species that cause TBRF continue to be identified. For example, a patient infected with *B. johnsonii*-like species, previously found only in bat ticks, was identified in Wisconsin. Moreover, TBRF has also been reported in Central and South America. *B. hispanica*, *B. persica*, and *B. miyamotoi* are important causes of TBRF in Europe and Asia, and *B. hispanica*, *B. crocidurae*, and *B. duttonii* are important causes of TBRF in Africa. Although most RFB species are transmitted by soft ticks of the genus *Ornithodoros, B. miyamotoi* may be transmitted by the same hard ticks of the genus *Ixodes* that transmit Lyme Disease *Borrelia* (LDB) species. Other RFB species, *B. lonestari,* the recently described *B. turcicia,* and *B. tachyglossi,* are members of a rapidly expanding *Borrelia* clade associated with reptile (*B. turcica*) or echidna (*B. tachyglossi*) hosts and are also transmitted by hard ticks. In order to provide timely treatment, clinicians must be able to rapidly and correctly identify whether a patient presenting with symptoms of tick-borne disease have been exposed to TBRF species.

Aspects of the invention provide a method of quickly and accurately detecting *Borrelia* antisera in a sample from a subject suspected of having TBRF. A subject suspected of having TBRF can be identified as having symptoms such as a high fever (*e.g.*, 103° F), headache, and muscle and joint aches. Symptoms typically reoccur, producing a telltale pattern of fever lasting roughly 3 days, followed by approximately 7 days without fever, followed by another 3 days of fever. Without proper antibiotic treatment, this process can repeat several times. Since the symptoms of TBRF can mimic, for example, viral flu-like symptoms, accurate diagnosis of TBRF is important for providing an effective treatment for the subject. The present invention provides a quick and easy diagnostic test for detecting the presence of antibodies specific for causative *Borrelia* species, thereby satisfying the need for such a test.

Because TBRF can be caused by several RFB species, and because geographic ranges of RFB species may overlap, tests for RFB species need to be inclusive—that is, a test needs to be able to detect antibodies to multiple species concurrently. The present invention provides for antigenic amino acid sequences specific for various TBRF *Borrelia* species. The amino acid sequences of the present invention encode antigenic peptides that have high specificity and/or sensitivity for the indicated genus. The inclusion of antigenic peptides that exhibit cross-reactivity across *Borrelia* species boundaries is also important with respect to the development of inclusive serological, or other immunologically-based assays, wherein the goal is to detect infection, not necessarily to identify a particular species responsible for infection. For example, the disclosure includes multi-panel immunoassays wherein, in the context of a single test screen, multiple RFB species are detectable.

The present invention provides novel compositions and methods for diagnosing infection by TBRF *Borrelia* species. The invention is based, in part, on the discovery of species-specific amino acid sequences encoding antigenic peptides (which may also be referred to in the art as peptide antigens or antigens), as described herein.

The present invention, in one aspect, is a composition comprising one or more labeled and/or tagged and/or bound amino acid sequences, wherein the one or more labeled and/or tagged and/or bound amino acid sequences have at least 90%, 95%, 98%, 99%, 99.5%, or 100% homology to SEQ ID NO: 1, SEQ ID NO: 2, SEQ ID NO: 3, SEQ ID NO: 4, SEQ ID NO: 5, SEQ ID NO: 6, SEQ ID NO: 7, SEQ ID NO: 8, SEQ ID NO: 9, SEQ ID NO: 10, and SEQ ID NO: 11. Sequences with less than 100% homology may be modified with one or more substitutions, deletions, insertions, or other modifications with respect to the amino acid sequences provided herein. Exemplary modifications include, but are not limited to conservative amino acid substitutions, which will produce molecules having functional characteristics similar to those of the molecule from which such modifications are made. Conservative amino acid substitutions are substitutions that do not result in a significant change in the activity or tertiary structure of a selected polypeptide or protein. Such substitutions typically involve replacing a selected amino acid residue with a different residue having similar physico-chemical properties. For example, substitution of Glu for Asp is considered a conservative substitution because both are similarly-sized negatively-charged amino acids. Groupings of amino acids by physico-chemical properties are known to those of skill in the art. The following groups each contain amino acids that are conservative substitutions for one another: 1) Alanine (A), Glycine (G); 2) Aspartic acid (D), Glutamic acid (E); 3) Asparagine (N), Glutamine (Q); 4) Arginine (R), Lysine (K); 5) Isoleucine (I),

Leucine (L), Methionine (M), Valine (V); 6) Phenylalanine (F), Tyrosine (Y), Tryptophan (W); 7) Serine (S), Threonine (T); and 8) Cysteine (C), Methionine (M) (see, e.g., Creighton, Proteins (1984)). One of ordinary skill in the art can determine if sequences with less than 100% homology can bind naturally- or non-naturally-occurring TBRF-related antibodies, as well as the sensitivity and specificity of the antibody to the modified sequences. One of ordinary skill in the art will be able to identify sequences with significant homology to SEQ ID NOs: 1–11 of the present invention that give acceptable or equivalent responses in the methods of the present invention without undue experimentation, in view of the teachings of this specification.

In some embodiments, the present invention is a composition comprising one or more labeled and/or tagged and/or bound amino acid sequences, wherein the one or more labeled and/or tagged and/or bound amino acid sequences comprise amino acid sequences selected from the group consisting of SEQ ID NO: 1, SEQ ID NO: 2, SEQ ID NO: 3, SEQ ID NO: 4, SEQ ID NO: 5, SEQ ID NO: 6, SEQ ID NO: 7, SEQ ID NO: 8, SEQ ID NO: 9, SEQ ID NO: 10, and SEQ ID NO: 11, and variants thereof which retain the immunological binding profile of Fig. 3. As used herein, "consisting of", when used as a claim transition referring to an amino acid sequence, refers to amino acid sequences having 100% homology to the specified amino acid sequence (i.e., SEQ ID NOs: 1–11).

Aspects of the present invention provide antigen-specific amino acid sequences for TBRF *Borrelia* specific species. These novel amino acid sequences may be used in assays to identify TBRF specific *Borrelia* in samples from subjects suspected of having TBRF. With the amino acid sequences of the present invention, identification of TBRF *Borrelia* in subject samples is performed with greater speed, sensitivity and specificity than other current methods. The amino acid sequences of the present invention may be used in diagnostic and scientific assays. Non-limiting examples of suitable assays include immunoblots, line immunoblots, ELISA (enzyme-linked immunosorbent assay), etc. The amino acid sequences of the present invention may be used for the detection of TBRF *Borrelia* specific T-cells, for example, with the IgXSPOT test (IGeneX, Milpitas, CA).

In some embodiments, the invention is a composition comprising one or more labeled and/or tagged and/or bound amino acid sequences, wherein the one or more labeled and/or tagged and/or bound amino acid sequences comprise amino acid sequences selected from the group consisting of SEQ ID NO: 1, SEQ ID NO: 2, SEQ ID NO: 3, SEQ ID NO: 4, SEQ ID NO: 5, SEQ ID NO: 6, SEQ ID NO: 7, SEQ ID NO: 8, SEQ ID NO: 9, SEQ ID NO: 10, and SEQ ID NO: 11, and variants of those sequences. The expression "variants" encompasses

any modification(s) of a specified amino acid sequence (i.e., SEQ ID NOs. 1–11) which retain(s) the immunological binding profile of Fig. 3. Such modifications may include insertions and deletions (internal or from the N- or C- terminus, or both).

Nucleic acid sequences, including polynucleotides and oligonucleotides, encoding the amino acid sequences of the present invention, and portions thereof, may be expressed in cultured cells to provide isolatable quantities of peptides displaying biological (*e.g.*, immunological) properties of the antigenic peptide encoded by the amino acid sequences of the present invention. Because of redundancy of the genetic code, multiple nucleic acid sequences may be suitable for the production of the peptide sequences of the present invention. One of ordinary skill in the art will be able to determine one or more nucleic acid sequences for production of the amino acid sequences of the present invention. A nucleic acid sequence encoding an amino acid sequence of the present invention may be labeled by any suitable label known to one of ordinary skill in the art.

In this regard, nucleic acid sequences suitable for the production of the amino acid sequences of the present invention may be substantially homologous to naturally occurring sequences. Substantial homology of a nucleic acid sequence as used herein means that: (a) there is greater than 65%, 75%, 85%, 95%, 98%, or 99% homology with the naturally occurring sequence, or (b) the homologous nucleic acid sequence will hybridize to the compared sequence or its complementary strand under stringent conditions of the temperature and salt concentration. These stringent conditions will generally be a temperature greater than about 22°C, usually greater than about 30°C and more usually greater than about 45°C, and a salt concentration generally less than about 1 M, usually less than about 500 mM, and preferably less than about 200 mM. The combination of temperature and salt concentration is more important in defining stringency than either the temperature or the salt concentration alone. Other conditions which affect stringency include GC content of the compared sequence, extent of complementarity of the sequences, and length of the sequences involved in the hybridization, as well as the composition of buffer solution(s) used in the hybridization mixture. These and other factors affecting stringency are well described in the scientific and patent literature. One of ordinary skill in the art will be able to determine suitable conditions for determining the homology of the nucleic acid sequences encoding the antigenic peptides of the present invention.

Homologous nucleic acid sequences may be determined based on the nature of a nucleotide substitution in the nucleic acid sequence. For example, synonymous nucleotide substitutions, that is, nucleotide changes within a nucleic acid sequence that do not alter the

encoded amino acid sequence, will be better tolerated and, therefore, may be more numerous in a particular nucleic acid sequence than non-synonymous nucleotide substitutions. One of ordinary skill in the art will be able to determine the suitable number and location of substitutions that may be allowed in a nucleic acid sequence that encodes an amino acid sequence of the present invention without adversely affecting the antigenicity of the encoded antigenic peptide, without undue experimentation.

Labels and Tags

One or more amino acid sequences of the invention may be labeled and/or tagged and/or bound. As used herein, a "label" or "tag" is a detectable moiety that may be attached to an amino acid sequence of the invention. A label or tag may be covalently or non-covalently attached to an amino acid sequence of the invention. Non-limiting examples of such "tags" are natural and synthetic (*i.e.*, non-naturally occurring) nucleic acid and amino acid sequences (*e.g.*, poly-AAA tags), antibodies and detectable moieties such as labels (discussed elsewhere herein). Thus, the definitions of the phrases "labeled" and "tagged" may have overlap in that a tag may also, in some instances, function as a label. Furthermore, tags useful with the present invention may be linked to a label.

The amino acid sequences of the present invention, or any tags attached to an amino acid sequence of the present invention, may be labeled with any suitable label known to one of ordinary skill in the art. Such labels may include, but are not limited to, biotin/streptavidin, enzyme conjugates (*e.g.*, horseradish peroxidase (HRP), alkaline phosphatase (AP), glucose oxidase, and β-galactosidase), fluorescent moieties (*e.g.*, FITC, fluorescein, rhodamine, etc.), biological fluorophores (*e.g.*, green fluorescent protein (GFP), R-phycoerythrin) or other luminescent proteins, etc. Any suitable label known to one of ordinary skill in the art may be used with the present invention.

In some embodiments, amino acid sequences of the invention may be "bound." A "bound" amino acid sequence is an amino acid sequence that has been immobilized in order to permit the use of the amino acid sequence in a biological test such as, for example, an immunoassay. In the context of the present invention, a "bound" amino acid sequence is an amino acid sequence attached (*e.g.*, covalently or non-covalently bound, etc.) directly or indirectly to a non-natural surface or substance. Additionally or alternatively, "bound" amino acid sequences of the present invention may be attached, directly or indirectly, to a natural surface or substance, either of which is not naturally associated with the amino acid sequence. Non-limiting examples of substances to which the amino acid sequences of the present

invention may be bound are nitrocellulose, nylon, polyvinylidene difluoride (PVDF) plastics, metals, magnetic beads and agarose (*e.g.*, beads). Linking agents known to those of ordinary skill in the art may be used to aid or enhance binding of the amino acid sequences of the present invention to a surface or substance.

Production of amino acid sequences

In some embodiments, amino acid sequences of the invention may be non-natural, synthetic sequences, such as sequences produced by recombinant technology or sequences synthesized by protein synthesizing apparatuses. As such, the amino acid sequences of the present invention may be produced by recombinant technology, as is described and enabled in the literature and in commonly referred to manuals such as, *e.g.*, Short Protocols in Molecular Biology, Second Edition, F.M. Ausubel, Ed., all John Wiley & Sons, N.Y., edition as of 2008; and, Sambrook, et al., Molecular Cloning: A Laboratory Manual, 3rd ed., Cold Spring Harbor Laboratory Press, Cold Spring Harbor, 2001, and as is well known to one of ordinary skill in the art. In one embodiment, the amino acid sequences of the present invention are made recombinantly in *E. coli*.

As used herein, the term "vector" refers to a nucleic acid molecule capable of transporting between different genetic environments another nucleic acid to which it has been operatively linked. In addition to the nucleic acid sequence encoding an amino acid sequence selected from the group consisting of SEQ ID NO: 1, SEQ ID NO: 2, SEQ ID NO: 3, SEQ ID NO: 4, SEQ ID NO: 5, SEQ ID NO: 6, SEQ ID NO: 7, SEQ ID NO: 8, SEQ ID NO: 9, SEQ ID NO: 10, and SEQ ID NO: 11, and variants thereof which retain the immunological binding profile of Fig. 3, the vectors of the present invention also include a heterologous nucleic acid sequence. As used herein, heterologous refers to a nucleic acid sequence that does not naturally occur in the organism from which the Markush group sequences are derived. The term "vector" may also refer to a virus or organism that is capable of transporting the nucleic acid molecule. One type of vector is a plasmid, a small, circular, double-stranded, extrachromosomal DNA molecule that is physically separate from and can self-replicate independently from chromosomal DNA. Some useful vectors are those capable of autonomous replication and/or expression of nucleic acids to which they are linked. Vectors capable of directing the expression of nucleic acids to which they are operatively linked are referred to herein as "expression vectors." Other useful vectors, include, but are not limited to bacterial plasmids and bacterial artificial chromosomes (BACs), cosmids, and viruses such as lentiviruses, retroviruses, adenoviruses, and phages.

Vectors useful in methods of the invention may include additional sequences including, but not limited to, one or more signal sequences and/or promoter sequences, or a combination thereof. Promoters that may be used in methods and vectors of the invention include, but are not limited to, cell-specific promoters or general promoters. Non-limiting examples of promoters that can be used in vectors of the invention are: ubiquitous promoters, such as, but not limited to: CMV, CAG, CBA, and EF1a promoters. Methods to select and use suitable promoters are well known in the art.

Vectors useful in methods of the invention may be used to express a fusion protein comprising sequences of the invention in a cell. Expression vectors and methods of their preparation and use are well known in the art. In some embodiments of the invention, a nucleic acid sequence of an expression vector encodes a fusion protein comprising an amino acid sequence of the invention. It is well known in the art how to prepare and utilize fusion proteins that comprise a polypeptide sequence. In some embodiments, a fusion protein comprising an amino acid sequence of the invention may also include an epitope tag that may be used for purification of the fusion protein or in a method of the invention. Non-limiting examples of epitope tags are a FLAG tag, a fluorescent tag (including but not limited to green fluorescent protein (GFP)), a GST tag, a hemagglutinin (HA), a poly-histidine (poly-His) tag, a Myc tag, an MBP tag, or a V5 tag. In some embodiments, a fusion protein comprising an amino acid sequence of the invention may also include a detectable label, as described elsewhere herein.

Subjects and cells

As used herein, a subject may be an animal, such as a mammal or a non-mammal. Non-limiting examples of mammalian subjects include primates (including but not limited to humans), rodents (including but not limited to mice, rats, squirrels, chipmunks, prairie dogs), lagomorphs, deer, canids (including but not limited to dogs, foxes, coyotes, and wolves), felids (including but not limited to domestic cats, bobcats, cougars, and other wild cats), bears, horses, cows, sheep, goats, and pigs. Non-limiting examples of non-mammalian subjects include birds, amphibians, lizards, insects, and arthropods. As used herein, a cell may be a bacterial cell, including but not limited to *E. coli*, or an animal cell, either mammalian or non-mammalian.

Assays and methods of detection

Amino acid sequences of the invention may be tagged with an antibody with specificity for any of said amino acid sequences. Specificity for said amino acid sequence, *i.e.*, antibody specificity, is the property of antibodies which enables them to react preferentially with some antigenic determinants and not with others. Specificity is dependent on chemical composition, physical forces and molecular structure at the binding site. Sensitivity is how strongly the antibody binds to the antigenic determinant. One of ordinary skill in the art can determine specificity and sensitivity of an antibody for a particular amino acid sequence using standard affinity assays, such as immunoblotting, Ouchterlony assays, titer assays, etc. In another aspect, the present invention provides a method of quickly and accurately detecting TBRF *Borrelia* antisera in a sample from a subject suspected of having TBRF. The method of the present invention for detecting TBRF *Borrelia* antisera in a sample from a subject suspected of having TBRF, may comprise, for example, providing a biological sample (including but not limited to blood, saliva) obtained from a subject suspected of having TBRF, mixing the biological sample with one or more of the labeled and/or tagged and/or bound amino acid sequences of the present invention and detecting a positive reaction which indicates the presence of Lyme disease antisera in the sample. The antisera may be detected by, for example, immunoblotting, ELISPOT, ELISA, Western blotting, lateral flow assay, or any other appropriate immunoassay known to one of ordinary skill in the art. These techniques are known to one of ordinary skill in the art and procedures can be found in common technical references. While similar, each of these techniques has its advantages and disadvantages. Other suitable techniques may be known to those of skill in the art and are incorporated herein.

To assess the impact of testing limitations and to determine levels of exposure to TBRF Borrelia, a modified Western blot procedure, the line immunoblot, was developed and employed in aspects of the invention described herein. A line immunoblot uses recombinant antigens from common strains and species of the TBRF *Borrelia* complex for serological identification and diagnosis of TBRF in serum from patients with a suspected tick-borne disease. As described elsewhere herein, the serotype makeup of TBRF *Borrelia* exposure may be more complex than has been previously acknowledged, and infection with more than one TBRF *Borrelia* species is possible.

Western blotting can involve separating proteins by electrophoresis and then transferring to nitrocellulose or other solid media (*e.g.*, polyvinylidene fluoride or PVDF-membrane and nylon membrane), and is described in more detail below. Immunoblotting can also involve applying proteins to a solid media manually or by machine. Preferably, the

proteins are applied in straight lines or spots and dried, binding them to the solid support medium, *e.g.*, nitrocellulose. The proteins used in an immunoblot can be isolated from biological samples or produced by recombinant technology, as is well known by those of ordinary skill in the art. The bound proteins are then exposed to a sample or samples suspected of having antibodies specific for the target proteins. With this procedure, a known antibody can be used to determine if a protein is present in a sample, such as when the proteins of lysed cells are separated by electrophoresis and transferred to the solid medium. Western blotting allows for the identification of proteins by size as well as by specificity for a specific antibody.

Similarly, with a procedure called immunoblotting, known proteins can be bound to the solid medium and samples, such as samples from subjects suspected of having an infection, can be tested for the presence of specific antibodies in the sample by contacting the bound protein with the sample. An antibody that binds the target protein is usually referred to as the primary antibody. A secondary antibody, specific for conserved regions of the primary antibody (for example, a rabbit- anti-human IgG antibody may be used to detect primary human antibodies) is used to detect any bound primary antibodies. The secondary antibody is usually labeled with a detectable moiety for visualization. Non-limiting examples of suitable labels include, for example, chromophores such as biotin, radioactive moieties and enzymes such as alkaline phosphatase, etc. The use of these and other materials for the visualization of antibodies are well known to one of ordinary skill in the art.

The Enzyme-Linked ImmunoSpot (ELISPOT) method can detect human T-cells that respond to Lyme-specific antigens *in vitro*. In an ELISPOT assay, the surfaces of PVDF membrane in a 96- well microtiter plate are coated with capture antibody that binds, for example, anti-Interferon gamma (IFNγ) or other cytokine-specific antibody. During the cell incubation and stimulation step, the T-cells isolated from patient whole blood are seeded into the wells of the plate along with aforementioned sequence(s), and form substantially a monolayer on the membrane surface of the well. Upon stimulation of any antigen-specific cells with one or more of the sequences of the present invention they are activated and they release the IFNγ, which is captured directly on the membrane surface by the immobilized antibody. The IFNγ is thus "captured" in the area directly surrounding the secreting cell, before it has a chance to diffuse into the culture media, or to be degraded by proteases and bound by receptors on bystander cells. Subsequent detection steps visualize the immobilized IFNγ as an ImmunoSpot; essentially the secretory footprint of the activated cell.

For a specific example of an ELISPOT test, each well of the plate is coated with a purified cytokine-specific antibody specific for the test or cell being detected. T-cells are isolated from a subject (for example, a subject suspected of having Lyme Disease or TBRF) and cultured in each well and stimulated with recombinant antigens of one or more sequences of the present invention. Lyme-positive patient cells secrete cytokine in response to stimuli, which is captured by the antibody coated in the well and further detected by ELISA.

ELISA assays may also be used to detect antigens. ELISA assays permit quantification of a specific protein in a mix of proteins (for example, a lysate) or may be used to determine if a peptide is present in a sample. Likewise, ELISA assays may be used to determine if a specific antibody is present by using a specific antigen as a target. As used with the present invention, target amino acid sequence(s) are attached to a surface. Then, if present in the sample being tested, the reactive antibody can bind to the antigen. A secondary antibody linked to an enzyme is added, and, in the final step, a substance containing the enzyme's substrate is added. The subsequent reaction produces a detectable signal, most commonly a color change in the substrate.

Lateral flow assays, also referred to by a variety of other names that include but are not limited to lateral flow tests, lateral flow devices, lateral flow immunoassays, lateral flow immunochromatographic assays, and rapid tests, are simple, versatile, paper-based platforms for detecting and/or quantifying the presence of one or more analytes, such as an antigen, in a mixture, such as a liquid sample. Lateral flow assays may be qualitative or quantitative. In a lateral flow assay, a sample containing one or more analytes of interest is applied to an adsorbent sample pad and is drawn via capillary action through various zones of polymeric test strips to which are attached molecules that can interact with the analyte(s). The sample migrates to the conjugate release pad, which contains molecules that specifically bind to the analyte(s) of interest and are conjugated to fluorescent, colored, or otherwise detectable particles. Finally, the sample, including the bound analyte(s) migrates into the detection zone. Within the porous membrane of the detection zone are biological components such as antibodies or antigens, that are immobilized in lines and that will react with the detectable particles. Lateral flow assays typically have a control line for confirming sample flow through the strip and one or more test lines for detecting the presence of the analyte(s) of interest. The results may be read by eye or with a machine capable of reading and interpreting the results. A lateral flow assay may be designed as a direct or “sandwich” assay, in which the presence of a colored line at the test line position indicates a positive test,

or as a competitive assay, in which the absence of a colored line indicates a positive test. Direct and competitive assays may be multiplexed.

In one aspect of the method of the present invention, a positive result is indicated when two or more of the labeled and/or bound amino acid sequences of the present invention are mixed with the biological sample and when at least two amino acid sequences are detected. In another aspect of the invention, a positive result is indicated when at least one of the labeled and/or bound amino acid sequences of the present invention are mixed with the biological sample and when at least one amino acid sequence is detected.

In the method of the present invention, any primary antibody bound to a peptide encoded by an amino acid sequence of the present invention may be detected with anti-human antibodies, such as IgG or IgM, used as the secondary antibody conjugated to a detectable moiety. As described elsewhere herein, the detectable moiety may be selected from the group consisting of chromophores, radioactivity moieties and enzymes or other detectable moiety known to one of ordinary skill in the art. In one embodiment, the detectable moiety comprises alkaline phosphatase. In another embodiment the detectable moiety comprises biotin.

In another aspect of the invention, a method is provided for detecting and distinguishing various species of *Borrelia* in a sample. The sample may be from a subject suspected of having Lyme disease. The method may comprise, for example, providing a sample, for example, a biological sample obtained from a subject suspected of having TBRF and mixing or contacting the biological sample with one or more of the labeled and/or bound amino acid sequences of the present invention. Amino acids may be labeled to confirm their presence if positive results are not obtained in the assay.

In some embodiments of the invention, a sample may be considered positive in a multi-species panel assay for TBRF *Borrelia* if at least one amino acid sequence is detected. In some embodiments of the invention, a sample may be considered positive in a multi-species panel assay for TBRF *Borrelia* if at least two amino acid sequences are detected. In some embodiments of the invention, a sample may be considered positive for a specific species of TBRF *Borrelia* if at least two amino acid sequences identified with a species are detected.

Equivalents

Although several embodiments of the present invention have been described and illustrated herein, those of ordinary skill in the art will readily envision a variety of other

means and/or structures for performing the functions and/or obtaining the results and/or one or more of the advantages described herein, and each of such variations and/or modifications is deemed to be within the scope of the present invention. More generally, those skilled in the art will readily appreciate that all parameters, dimensions, materials, and configurations described herein are meant to be exemplary and that the actual parameters, dimensions, materials, and/or configurations will depend upon the specific application or applications for which the teachings of the present invention is/are used. Those skilled in the art will recognize, or be able to ascertain using no more than routine experimentation, many equivalents to the specific embodiments of the invention described herein. It is, therefore, to be understood that the foregoing embodiments are presented by way of example only and that, within the scope of the appended claims and equivalents thereto; the invention may be practiced otherwise than as specifically described and claimed. The present invention is directed to each individual feature, system, article, material, and/or method described herein. In addition, any combination of two or more such features, systems, articles, materials, and/or methods, if such features, systems, articles, materials, and/or methods are not mutually inconsistent, is included within the scope of the present invention.

All definitions, as defined and used herein, should be understood to control over dictionary definitions, definitions in documents incorporated by reference, and/or ordinary meanings of the defined terms.

The indefinite articles "a" and "an," as used herein in the specification and in the claims, unless clearly indicated to the contrary, should be understood to mean "at least one." The phrase "and/or," as used herein in the specification and in the claims, should be understood to mean "either or both" of the elements so conjoined, i.e., elements that are conjunctively present in some cases and disjunctively present in other cases. Other elements may optionally be present other than the elements specifically identified by the "and/or" clause, whether related or unrelated to those elements specifically identified, unless clearly indicated to the contrary.

All references, patents and patent applications and publications that are cited or referred to in this application are incorporated by reference in their entirety herein.

EXAMPLES

Example 1. TBRF ImmunoBlot antigen inclusivity study

TBRF ImmunoBlot IgG and IgM tests, qualitative immunoblot assays, were designed and performed to detect IgG and IgM antibodies directed against TBRF *Borrelia* species in

serum samples suspected of having Relapsing Fever or Lyme-like symptoms. Recombinant *TBRF Borrelia* antigens were sprayed as straight lines onto nitrocellulose strips, which were then used in the TBRF ImmunoBlot Test. Experiments were undertaken to determine the specificity of the TBRF ImmunoBlot Test.

Methods

Antigen preparation

Recombinant target proteins were obtained by cloning hybrid gene constructs or portions of genes into pET vectors, expressing the gene products in *Escherichia coli* (GenScript, Piscataway, NJ), then isolating the proteins to > 90% purity, as previously described [Liu et al., *Healthcare* (2018) 6(3) pii: E99; Shah et al., *Healthcare* (2019) 7:121]. Recombinant proteins from several TBRF *Borrelia* species (*B. hermsii*, *B. miyamotoi*, *B. turicatae* and *B. turcica*) were derived for the detection of four target antigens: BipA, GlpQ, BpcA/fHbp and P41/FlaB, as previously described [Shah et al., *Healthcare* (2019) 7:121].

Preparation of antigen strips

Antigen strips for TBRF immunoblots were prepared as previously described [Liu et al., *Healthcare* (2018) 6(3) pii: E99; Shah et al., *Healthcare* (2019) 7:121]. Purified proteins and control proteins were diluted (7–19 ng protein/line) and sprayed in straight lines on nitrocellulose sheets (Cytiva, Marlborough, MA) using a BioDot liquid dispenser (BioDot, Irvine, CA). The sheets were then blocked with 5% non-fat dry milk and sliced into 3 mm wide strips.

Detection of antibody reactivity

Serological immunoblot testing was performed at IGeneX Reference Laboratory, a high-complexity testing facility with Clinical Laboratory Improvement Amendments (CLIA) certification.

Reactivity between TBRF *Borrelia*-specific antibodies from test sera and TBRF *Borrelia* antigens on immunoblots was detected as previously described [Liu et al., *Healthcare* (2018) 6(3) pii: E99; Shah et al., *Healthcare* (2019) 7:121]. Strips were labeled, soaked in diluent (100 mM Tris, 0.9% NaCl, 0.1% Tween-20, and 1% non-fat dry milk) for 5 minutes (min) in a trough; then, a 10 µL aliquot of either test serum or control serum was added to the strip. Strips with sera were incubated at room temperature for one hour, washed three times with wash buffer (KPL, Gaithersburg, MD) at room temperature, and the

final wash solution was then aspirated. To detect IgG and IgM reactivity, strips were incubated with alkaline phosphatase-conjugated goat anti-human IgG at 1:10,000 dilution or IgM at 1:6,000 dilution, respectively (KPL, Gaithersburg, MD), for one hour, and then washed three times. To visualize bands of antibody/antigen reactivity, the strips were reacted with a chromogenic substrate, 5-bromo-4-chloro-3-indolylphosphatenitro-blue tetrazolium (BCIP/NBT, KPL, Gaithersburg, MD), and the reaction was terminated by washing with distilled water after the calibration control produced a visible band at 39 kDa. Bands demonstrating an intensity lower than that of the calibration control were reported as negative. Human sera from patients with confirmed *Borrelia* infection were used as positive controls and sera from uninfected persons were used as negative controls. All immunoblot testing of patient samples was performed with simultaneous testing of positive and negative control serum samples.

Scoring of Immunoblots

For TBRF *Borrelia* ImmunoBlots, detection of either IgG or IgM antibodies against P41/FlaB (41kDa), as well as any two out of the four antigens BipA (75kDa), GlpQ, BpcA/fHbp (21-23kDa) gave the best specificity for TBRF species.

Detection of either IgG or IgM antibodies to two or more proteins within each antigen type was regarded as a positive reaction for that antigen type. Applying the same criteria for reactivity of either IgM or IgG antibodies to the four TBRF *Borrelia* antigens led to optimum sensitivity of detection, as previously described [Shah et al., *Healthcare* (2019) 7:121].

Results

As shown in Fig. 1, antibodies to TBRF *Borrelia* species *B. turicica*, *B. miyamotoi*, *B. hermsii*, *B. parkeri*, and *B. turicatae* species were detected by the TBRF ImmunoBlot strip prepared from recombinant antigens from all five species, indicating that TBRF ImmunoBlots detected TBRF *Borrelia* group antibodies.

Example 2. TBRF ImmunoBlot Clinical Study of Patients with Suspected Tick-borne Disease

A study of patients who met the case definition of Chronic Lyme Disease (CLD) was performed to assess the impact of testing limitations and to determine levels of exposure to TBRF *Borrelia*. The results also revealed that all had exposure to *B. burgdorferi* species.

Methods

Methods used were as described in Example 1 and elsewhere herein.

Antigen preparation and preparation of antigen strips

Additional recombinant proteins from several TBRF *Borrelia* species (*B. hermsii*, *B. miyamotoi*, *B. turicatae*, *B. parkeri*, and *B. duttonii*) were derived as described elsewhere herein for the detection of target antigens BipA, GlpQ, BpcA/fHbp, P41/FlaB, and Vlp7. These recombinant proteins, comprising the amino acid sequences of SEQ ID NOs: 1-11, were included on TBRF ImmunoBlot strips.

Clinical testing

A total of 265 patient serum samples were tested by TBRF ImmunoBlots IgM and IgG, including 90 patients with CLD symptoms and 212 patient serum samples that were negative for TBRF.

Chronic Lyme Disease patient cohort (n=90)

A patient cohort of 90 patients with CLD was recruited from a medical practice located in San Francisco, CA, specializing in the diagnosis and treatment of tick-borne diseases. The Western Institutional Review Board (WIRB), Puyallup, WA approved the anonymous retrospective data collection protocol and consent form. Patients of either sex qualified for the study provided they were at least 18 years of age, had a medical history of musculoskeletal, neuropsychiatric and/or cardiac symptoms consistent with LD, and gave written informed consent for data collection. Subjects were included in the study if they met the case definition of untreated or previously treated CLD with symptoms lasting more than six months, as described in detail elsewhere [Cameron et al., *Expert Rev Anti Infect Ther*. (2014) 12:1103–1135; Stricker et al., *Am J Infect Dis* (2018) 14:1-44]. Patients were not required to have had a documented tick bite or erythema migrans rash for participation in the study because serological testing was used to detect exposure rather than active infection. De-identified patient samples were coded according to the patient's place of residence. Blood was drawn and serum was separated at independent laboratories including BioReference®, LabCorp®, and AnyLabTestNow®, and serum samples were transported to the reference laboratory for immunoblot testing.

Control Patients (n=175)

A total of 175 human sera expected to be negative for *B. burgdorferi* species and TBRF *Borrelia* species were obtained from the Centers for Disease Control and Prevention (CDC, Atlanta, GA), College of American Pathologists, New York State Department of Health, New York Biologics (Southampton, NY, USA) and IGeneX Reference Laboratory (Milpitas, CA, USA). The IGeneX samples were leftover sera received for routine testing for tick-borne diseases that would otherwise have been discarded. ImmunoBlot IgM and IgG testing of patient sera and control sera was performed by laboratory personnel in a blinded fashion in the same manner as testing of clinical samples from TBRF patients. Results are shown in Tables 1-2 and Figs. 2-3.

Results

Immunoblot reactivity for TBRF *Borrelia* in representative patient serum samples is shown in Fig. 2, demonstrating that TBRF ImmunoBlots successfully detect TBRF *Borrelia* group antibodies in serum samples.

The results obtained with the 175 control sera that were expected to be negative for *B. burgdorferi* species and RFB yielded a false positive rate of 2.3% (4/175 samples) for the *B. burgdorferi* species immunoblot and 2.9% (5/175 samples) for the RFB immunoblot (Table 1). False positive tests for RFB were seen with an allergy patient serum (one control), multiple sclerosis (one control), viral infection (one control), and syphilis (two controls).

Using the TBRF ImmunoBlot test described in Example 1 and including SEQ ID NOs: 1–11, the study of patients who met the CLD case definition revealed that all had exposure to *B. burgdorferi* species, and 62% had exposure to TBRF species. In the CLD cohort (Table 2), ImmunoBlot testing revealed that out of the 90 subjects with suspected LD, a total of 48 patients (53%) were seropositive for TBRF alone, and 8 patients (9%) had mixed infection with TBRF *Borrelia* species and *B. burgdorferi* species. Fig. 2 shows TBRF ImmunoBlots of representative patient serum samples. Positive immunoblots were further characterized for the presence of antibodies to one or more RFB species, and instances of infection with more than one RFB species were identified (Table 2). Immunoblot testing of control sera demonstrated excellent specificity of 97.1% for the TBRF assay (Table 2). As shown in Fig. 3, various combinations of SEQ ID NOs: 1–11 were identified in patient serum samples, demonstrating that antigens from multiple species are required for an inclusive test.

These results illustrated how detection of antibodies to TBRF Borrelia species identified patients with Chronic Lyme Disease symptoms that would otherwise be missed by

only testing for *Borrelia burgdorferi* antigens. These results also confirmed that the serotype makeup of TBRF *Borrelia* exposure is complex, thereby showing the power and importance of inclusivity in TBRF testing, that is, simultaneous testing for antigens from multiple TBRF species.

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Table 1. Reference human sera for determining specificity of *B. Burgdorferi* species and RFB species immunoblots

Source	Characteristic	Total No. of Sera	BbsI Immunoblots (+)	RFB Immunoblots (+)
CDC Reference Set (n = 25)	Fibromyalgia	2	0	0
	Healthy endemic	7	1	0
	Healthy non-endemic	6	0	0
	Mononucleosis	2	0	0
	Multiple sclerosis	2	0	1
	Rheumatoid arthritis	2	0	0
	Severe Periodontitis	2	0	0
	Syphilis	2	0	1
CAP and NYSHD (n = 42) Autoimmunity and Allergy	**Autoimmune**			
	ANA (+)	3	0	0
	ANA (−)	2	0	0
	DNA (+)	1	0	0
	Rheumatoid factor (+)	9	0	0
	Rheumatoid factor (−)	8	0	0
	Allergy (n = 19)			
	IgG (+)	13	0	1
	Spec. IgE (+)	4	0	0
	Spec. IgE (−)	2	0	0
NYB (n = 21) Viruses and RPR (+)	Epstein–Barr virus (EBV)	4	1	0
	Human immunodeficiency virus (HIV)	4	0	0
	Cytomegalovirus (CMV)	5	0	1
	Hepatitis C virus (HCV)	0	0	0
	RPR (+)	8	2	1
IGeneX (n = 87)	*Bartonella henselae* infection	7	0	0
	Human granulocytic anaplasmosis	16	0	0
	Babesia microti infection	14	0	0
	Babesia duncani infection	41	0	0
	Human monocytic ehrlichiosis	5	0	0
	Negative controls	4	0	0
False Positives		**175**	4	5
Specificity			97.7%	97.1%

ANA—anti-nuclear antibodies; CAP—College of American Pathologists; CDC—Center for Disease Control; NYB—New York Biologics, Southampton, NY; NYSH—New York State Department of Health; RF—rheumatoid factor; RPR—rapid plasma regain test for syphilis.

Table 2. Summary of seroreactivity for subjects in Group 1 (*B. Burgdorferi* species), Group 2 (RFB species), or both

Immuno Blot	Total
Group 1 Bbsl Positive	**42**
Group 2 RFB Positive	**56**
Dual Group 1 and 2 Positive	**8**
Group 2. RFB Positive Samples	**48 (53%)**
RFB alone	25
B. hermsii alone	7
B. miyamotoi alone	4
B. turicatae alone	8
B. turcica alone	2
B. hermsii + B. turcica	1
B. hermsii + B. miyamotoi	1
Dual Group 1 and 2	**8 (9%)**
Bb + RFB	2
Bb + B. hermsii	1
B. californiensis + RFB	2
B. spielmanii + RFB	1
B. afzelii/garinii + RFB	1
B. afzelii/garinii + B. turicatae	1
Total cases	**56**

Discussion for Examples 1-2

TBRF are found worldwide and are a significant cause of morbidity and mortality, particularly with respect to unexplained fever cases. Advances in molecular testing have led to widespread identification of *Borrelia* species in human specimens, thus challenging prevailing thought regarding geographic distribution and prevalence of RFB infection. Although RFB is a growing concern in the Western United States, Central America, and South America, TBRF is not reportable nationally in the United States, and there is no standard case definition. In 2011, TBRF was reportable in 12 Western states, yielding 504 cases, with 70% of the cases reported in three states: California (33%), Washington (25%), and Colorado (11%). Disease-causing species in the United States include *B. miyamotoi*, *B. hermsii*, *B. lonestari*, *B. parkeri*, *B. turicatae*, and *B. mazzotii*. Most cases in the USA are caused by *B. hermsii*, transmitted by *Ornithodoros hermsi* ticks, but *B. miyamotoi*, *B.*

hermsii, and *B. parkeri* human infections were reported in California, and *B. coriaceae* was detected in ticks, although human infection was not confirmed [Fesler et al. Healthcare 8(2): 97-112 (2020)].

The genetic diversity of *Borrelia* spirochetes, as well as the symptoms of infection that are as diverse as the organisms causing them, makes it challenging to diagnose *Borrelia*-associated disease. In summary, exposure to TBRF *Borrelia* is a cause for concern, and TBRF *Borrelia* may explain Lyme Disease symptoms in patients who are seronegative for *B. burgodorferi sensu lato* species. As shown herein, some patients may demonstrate dual exposure to both *B. burgodorferi* and TBRF species, further complicating diagnosis and treatment. Immunoblot testing for RFB species using methods and compositions of the invention allows the detection of a diverse group of TBRF *Borrelia* serotypes and provides a greater understanding of human exposure to pathogenic *Borrelia*.

Equivalents

Although several embodiments of the present invention have been described and illustrated herein, those of ordinary skill in the art will readily envision a variety of other means and/or structures for performing the functions and/or obtaining the results and/or one or more of the advantages described herein, and each of such variations and/or modifications is deemed to be within the scope of the present invention. More generally, those skilled in the art will readily appreciate that all parameters, dimensions, materials, and configurations described herein are meant to be exemplary and that the actual parameters, dimensions, materials, and/or configurations will depend upon the specific application or applications for which the teachings of the present invention is/are used. Those skilled in the art will recognize, or be able to ascertain using no more than routine experimentation, many equivalents to the specific embodiments of the invention described herein. It is, therefore, to be understood that the foregoing embodiments are presented by way of example only and that, within the scope of the appended claims and equivalents thereto; the invention may be practiced otherwise than as specifically described and claimed. The present invention is directed to each individual feature, system, article, material, and/or method described herein. In addition, any combination of two or more such features, systems, articles, materials, and/or methods, if such features, systems, articles, materials, and/or methods are not mutually inconsistent, is included within the scope of the present invention.

All definitions, as defined and used herein, should be understood to control over dictionary definitions, definitions in documents incorporated by reference, and/or ordinary meanings of the defined terms.

The indefinite articles "a" and "an," as used herein in the specification and in the claims, unless clearly indicated to the contrary, should be understood to mean "at least one." The phrase "and/or," as used herein in the specification and in the claims, should be understood to mean "either or both" of the elements so conjoined, i.e., elements that are conjunctively present in some cases and disjunctively present in other cases. Other elements may optionally be present other than the elements specifically identified by the "and/or" clause, whether related or unrelated to those elements specifically identified, unless clearly indicated to the contrary.

All references, patents and patent applications and publications that are cited or referred to in this application are incorporated by reference in their entirety herein.

What is claimed is:

CLAIMS

1. A composition comprising one or more labeled and/or tagged and/or bound amino acid sequences, wherein the one or more labeled and/or tagged and/or bound amino acid sequences comprise amino acid sequences selected from the group consisting of SEQ ID NO: 1, SEQ ID NO: 2, SEQ ID NO: 3, SEQ ID NO: 4, SEQ ID NO: 5, SEQ ID NO: 6, SEQ ID NO: 7, SEQ ID NO: 8, SEQ ID NO: 9, SEQ ID NO: 10, and SEQ ID NO: 11, and variants thereof which retain the immunological binding profile of Fig. 3.

2. The composition of Claim 1, wherein said amino acid sequences are bound to a substance selected from the group consisting of nitrocellulose, nylon, polyvinylidene difluoride (PVDF), magnetic beads, and agarose.

3. The composition of Claim 1, wherein each of said one or more amino acid sequences are tagged with an antibody with specificity for said amino acid sequence.

4. A method for detecting antibodies resulting from infection by one or more members of Relapsing Fever (TBRF) Borrelia genus, wherein the genus comprises *B. hermsii*, *B. turicatae*, *B. miyamotoi*, *B. duttonii*, and *B. parkeri*, if present in a sample from a subject suspected of having tick-borne relapsing fever (TBRF), said method comprising:

providing a biological sample obtained from a subject suspected of having TBRF;

mixing the biological sample with one or more of the labeled and/or tagged and/or bound amino acid sequences of Claim 1; and

detecting a positive immunobinding reaction which indicates the presence of TBRF specific antibodies in the sample.

5. The method of Claim 4, wherein two or more of the labeled and/or tagged and/or bound amino acid sequences are mixed with the biological sample and a sample is considered positive for TBRF if at least two amino acid sequences are detected.

6. The method of Claim 4, wherein the labeled and/or tagged and/or bound amino acid sequences are detected with anti-human IgG or anti-human IgM antibody linked to a detectable moiety.

7. The method of Claim 6, wherein said detectable moiety is selected from the group consisting of chromophores, radioactive moieties and enzymes.

8. The method of claim 6, wherein said detectable moiety comprises alkaline phosphatase.

9. The method of Claim 6, wherein said detectable moiety comprises biotin.

10. A vector comprising a nucleic acid sequence encoding an amino acid sequence selected from the group consisting of SEQ ID NO: 1, SEQ ID NO: 2, SEQ ID NO: 3, SEQ ID NO: 4, SEQ ID NO: 5, SEQ ID NO: 6, SEQ ID NO: 7, SEQ ID NO: 8, SEQ ID NO: 9, SEQ ID NO: 10, and SEQ ID NO: 11, and variants thereof which retain the immunological binding profile of Fig. 3, and a heterologous nucleic acid sequence.

11. The vector of claim 10, wherein the nucleic acid sequence is operatively linked to a promoter sequence.

12. The vector of claim 10 or claim 11, wherein the vector is a prokaryotic vector.

13. A cell comprising the vector of any one of claims 10-12.

14. The cell of claim 13, wherein the cell is a bacterial cell.

Fig. 1

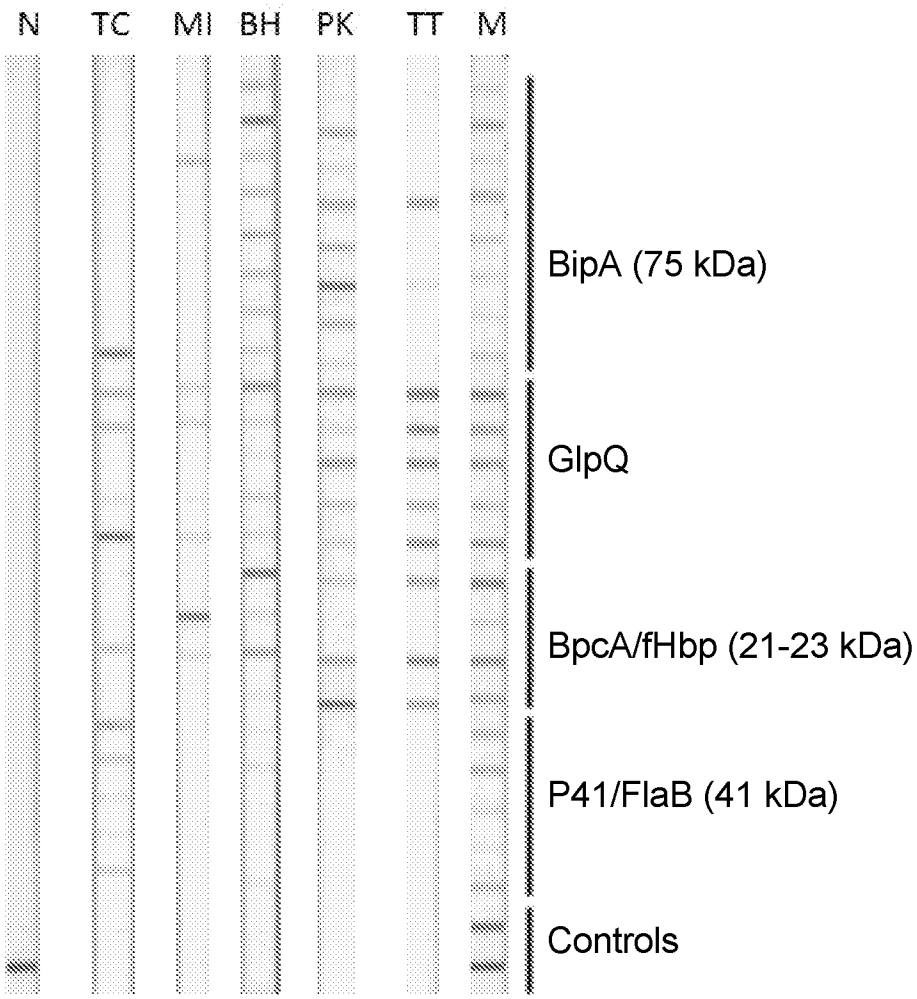

N
TC
MI
BH
PK
TT
M
BipA (75 kDa)
GlpQ
BpcA/fHbp (21-23 kDa)
P41/FlaB (41 kDa)
Controls

Fig. 2

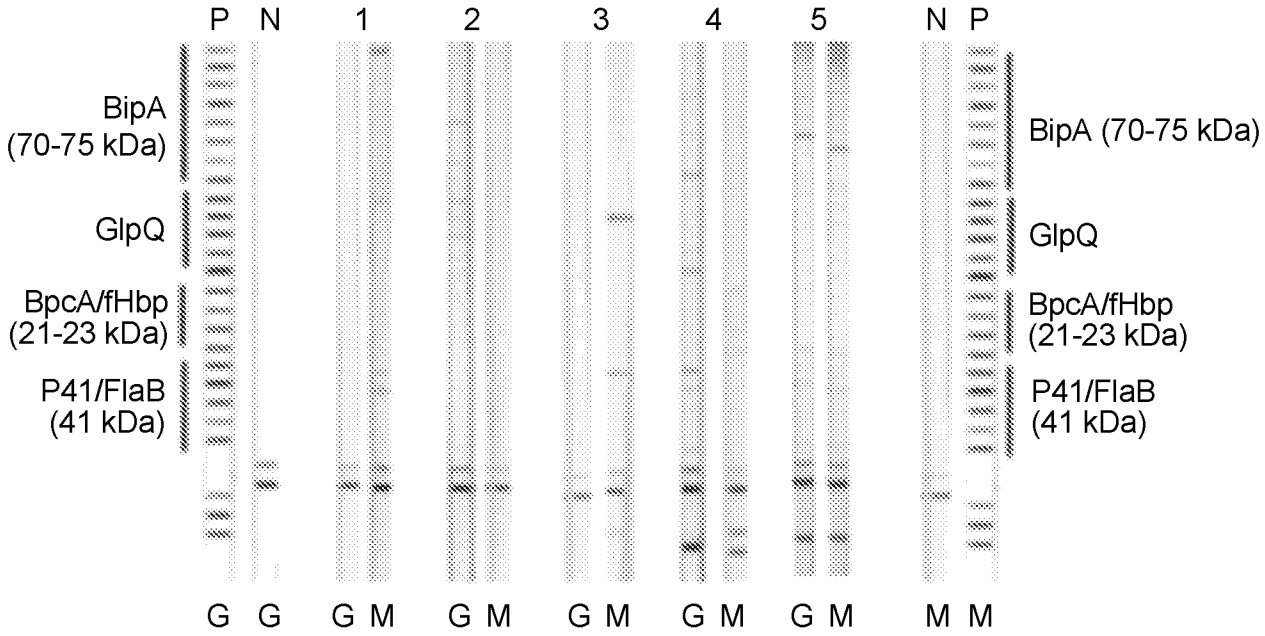

P N 1 2 3 4 5 N P
BipA (70-75 kDa)
GlpQ
BpcA/fHbp (21-23 kDa)
P41/FlaB (41 kDa)
BipA (70-75 kDa)
GlpQ
BpcA/fHbp (21-23 kDa)
P41/FlaB (41 kDa)
G G G M G M G M G M G M M M

Fig. 3

TBRF IgM and IgG Blots - Performance on Patient Serum Samples												
TBRF ImmunoBlot IgM												Result
Antigens	Bip A (70-75kDa)			GlpQ			BpcA/fHbp		P41/FlaB		Vlp7	
SEQ ID NO	2	5	6	11	7	10	3	8	4	9	1	
Sample #	Tt-2	Pk-1	Pk-2	Mia	Pk	Du	Tt	Pk	Tt	Pk	Bh	
1												Neg
2	Pos	Pos	Pos			Pos						Pos
3				I								Neg
4											Pos	Neg
5	Pos	Pos	Pos	Pos	Pos		Pos		Pos	I	Pos	Pos
6						I						Neg
7						Pos		Pos		Pos		Pos
8												Neg
9												Neg
10												Neg
TBRF ImmunoBlot IgG												Result
Antigens	Bip A (70-75kDa)			GlpQ			BpcA/fHbp		P41/FlaB		Vlp7	
SEQ ID NO	2	5	6	11	7	10	3	8	4	9	1	
Sample#	Tt-2	Pk-1	Pk-2	Mia	Pk	Du	Tt	Pk	Tt	Pk	Bh	
1									I	I		Neg
2												Neg
3		Pos										Neg
4				I							I	Neg
5	I					Pos	Pos		I	I		Pos
6												Neg
7					Pos							Neg
8	Pos						Pos					Pos
9								Pos		Pos		Pos
10			Pos					I				Neg

INTERNATIONAL SEARCH REPORT

International application No.
PCT/US 22/12229

A. CLASSIFICATION OF SUBJECT MATTER

IPC - G01N 33/00; G01N 33/53; G01N 33/554 (see supplemental sheet)

CPC - C07K 16/18; G01N 33/5436; G01N 33/543; G01N 33/53; G01N 33/68; G01N 33/58; G01N 33/56911; C07K 2319/40; C07K 2319/00; C07K 14/195; C12N 15/00; C07K 2319/60; C07H 21/00; G01N 33/6893

According to International Patent Classification (IPC) or to both national classification and IPC

B. FIELDS SEARCHED

Minimum documentation searched (classification system followed by classification symbols)
See Search History document

Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched
See Search History document

Electronic data base consulted during the international search (name of data base and, where practicable, search terms used)
See Search History document

C. DOCUMENTS CONSIDERED TO BE RELEVANT

Category*	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
X	US 2019/0277847 A1 (ID-FISH TECHNOLOGIES, INC.) 12 September 2019 (12.09.2019), para [0002], [0005], [0010], [0011], [0012], [0013], [0014], [0015], [0016], [0046], and SEQ ID NO: 1	1-9
A	FESLER et al., Lyme Disease: Diversity of Borrelia Species in California and Mexico Detected Using a Novel Immunoblot Assay. Healthcare (Basel). 2020, Vol. 8(2): 97. Epub 2020 Apr 14. PDF File: pg 1-16. Entire documentation especially Abstract; pg 3, last para; pg 4, para 2, and last para; and pg 9-12, Table 1	1-9
A	LIU et al., Pilot Study of Immunoblots with Recombinant Borrelia burgdorferi Antigens for Laboratory Diagnosis of Lyme Disease. Healthcare (Basel). 2018, Vol. 6(3): 99. PDF File: pg 1-15. Entire documentation especially Abstract; pg 2, para 3; and pg 8, Table 2	1-9
A	US 2020/0255889 A1 (THE TRUSTEES OF COLUMBIA UNIVERSITY IN THE CITY OF NEW YORK) 13 August 2020 (13.08.2020), para [0004], [0006], [0010], [0012], and [0019]	1-9
A	SHAH et al., Line Immunoblot Assay for Tick-Borne Relapsing Fever and Findings in Patient Sera from Australia, Ukraine and the USA. Healthcare (Basel). 2019, Vol. 7(4): 121. PDF File: pg 1-17. Entire documentation especially Abstract	1-9

☐ Further documents are listed in the continuation of Box C. ☐ See patent family annex.

* Special categories of cited documents:
"A" document defining the general state of the art which is not considered to be of particular relevance
"D" document cited by the applicant in the international application
"E" earlier application or patent but published on or after the international filing date
"L" document which may throw doubts on priority claim(s) or which is cited to establish the publication date of another citation or other special reason (as specified)
"O" document referring to an oral disclosure, use, exhibition or other means
"P" document published prior to the international filing date but later than the priority date claimed
"T" later document published after the international filing date or priority date and not in conflict with the application but cited to understand the principle or theory underlying the invention
"X" document of particular relevance; the claimed invention cannot be considered novel or cannot be considered to involve an inventive step when the document is taken alone
"Y" document of particular relevance; the claimed invention cannot be considered to involve an inventive step when the document is combined with one or more other such documents, such combination being obvious to a person skilled in the art
"&" document member of the same patent family

Date of the actual completion of the international search	Date of mailing of the international search report
08 April 2022	MAY 11 2022
Name and mailing address of the ISA/US Mail Stop PCT, Attn: ISA/US, Commissioner for Patents P.O. Box 1450, Alexandria, Virginia 22313-1450 Facsimile No. 571-273-8300	Authorized officer Kari Rodriquez Telephone No. PCT Helpdesk: 571-272-4300

INTERNATIONAL SEARCH REPORT

International application No.
PCT/US 22/12229

Box No. I Nucleotide and/or amino acid sequence(s) (Continuation of item 1.c of the first sheet)

1. With regard to any nucleotide and/or amino acid sequence disclosed in the international application, the international search was carried out on the basis of a sequence listing:

   a. [X] forming part of the international application as filed:

      [X] in the form of an Annex C/ST.25 text file.

      [ ] on paper or in the form of an image file.

   b. [ ] furnished together with the international application under PCT Rule 13*ter*.1(a) for the purposes of international search only in the form of an Annex C/ST.25 text file.

   c. [ ] furnished subsequent to the international filing date for the purposes of international search only:

      [ ] in the form of an Annex C/ST.25 text file (Rule 13*ter*.1(a)).

      [ ] on paper or in the form of an image file (Rule 13*ter*.1(b) and Administrative Instructions, Section 713).

2. [ ] In addition, in the case that more than one version or copy of a sequence listing has been filed or furnished, the required statements that the information in the subsequent or additional copies is identical to that forming part of the application as filed or does not go beyond the application as filed, as appropriate, were furnished.

3. Additional comments:

INTERNATIONAL SEARCH REPORT

International application No.
PCT/US 22/12229

Box No. II Observations where certain claims were found unsearchable (Continuation of item 2 of first sheet)

This international search report has not been established in respect of certain claims under Article 17(2)(a) for the following reasons:

1. ☐ Claims Nos.:
because they relate to subject matter not required to be searched by this Authority, namely:

2. ☐ Claims Nos.:
because they relate to parts of the international application that do not comply with the prescribed requirements to such an extent that no meaningful international search can be carried out, specifically:

3. ☒ Claims Nos.: 13-14
because they are dependent claims and are not drafted in accordance with the second and third sentences of Rule 6.4(a).

Box No. III Observations where unity of invention is lacking (Continuation of item 3 of first sheet)

This International Searching Authority found multiple inventions in this international application, as follows:
This application contains the following inventions or groups of inventions which are not so linked as to form a single general inventive concept under PCT Rule 13.1. In order for all inventions to be searched, the appropriate additional search fees must be paid.

Groups I+, claims 1-9, directed to a composition comprising one or more labeled and/or tagged and/or bound amino acid sequences, wherein the one or more labeled and/or tagged and/or bound amino acid sequences
comprise amino acid sequences; or a method for detecting antibodies resulting from infection by one or more members of Relapsing Fever (TBRF) Borrelia genus, comprising mixing the biological sample with one or more of the labeled and/or tagged and/or bound amino acid sequences. The one or more of the labeled and/or tagged and/or bound amino acid sequences will be searched to the extent that the one or more of the labeled and/or tagged and/or bound amino acid sequences encompasses SEQ ID NO: 1. It is believed that claims 1 (in part) and 2-9 encompass this first named invention, and thus these claims will be searched without fee to the extent that they encompass the one or more of the labeled and/or tagged and/or bound amino acid sequences encompasses SEQ ID NO: 1.

**********Continued in the extra sheet**********

1. ☐ As all required additional search fees were timely paid by the applicant, this international search report covers all searchable claims.

2. ☐ As all searchable claims could be searched without effort justifying additional fees, this Authority did not invite payment of additional fees.

3. ☐ As only some of the required additional search fees were timely paid by the applicant, this international search report covers only those claims for which fees were paid, specifically claims Nos.:

4. ☒ No required additional search fees were timely paid by the applicant. Consequently, this international search report is restricted to the invention first mentioned in the claims; it is covered by claims Nos.:
1 (in part) and 2-9, limited to SEQ ID NO: 1

Remark on Protest

☐ The additional search fees were accompanied by the applicant's protest and, where applicable, the payment of a protest fee.

☐ The additional search fees were accompanied by the applicant's protest but the applicable protest fee was not paid within the time limit specified in the invitation.

☐ No protest accompanied the payment of additional search fees.

INTERNATIONAL SEARCH REPORT

International application No.

PCT/US 22/12229

Continuation of:
Box No III (unity of invention is lacking)

(Continuation of Groups I+)
Additional the one or more labeled and/or tagged and/or bound amino acid sequences will be searched upon the payment of additional fees. Applicants must specify the claims that encompass any additionally elected the one or more labeled and/or tagged and/or bound amino acid sequences. Applicants must further indicate, if applicable, the claims which encompass the first named invention, if different than what was indicated above for this group. Failure to clearly identify how any paid additional invention fees are to be applied to the "+" group(s) will result in only the first claimed invention to be searched. An exemplary election would be the one or more labeled and/or tagged and/or bound amino acid sequences encompasses SEQ ID NO: 2 [claims 1 (in part), (2-9)].

Group II, claims 10-12, directed to a vector.

The inventions listed as Groups I+ and II do not relate to a single general inventive concept under PCT Rule 13.1 because, under PCT Rule 13.2, they lack the same or corresponding special technical features for the following reasons:

Special Technical Feature
Groups I+ includes the special technical feature of one or more labeled and/or tagged and/or bound amino acid sequences, not required by Group II.

Group II includes the special technical feature of a vector comprising a nucleic acid sequence, not required by Groups I+.

Among Groups I+, each SEQ ID NO represents a structurally different protein sequence.

Common Technical Features
The inventions of Groups I+ and Group II share the technical feature of an amino acid sequence selected from the group consisting of SEQ ID NO: 1; and
--the inventions of Groups I+ further share the technical feature of a composition comprising one or more labeled and/or tagged and/or bound amino acid sequences, wherein the one or more labeled and/or tagged and/or bound amino acid sequences comprise amino acid sequences specified (part of claim 1); and
--a method for detecting antibodies resulting from infection by one or more members of Relapsing Fever (TBRF) Borrelia genus, wherein the genus comprises B. hermsii, B. Turicatae, B. miyamotoi, B. duttonii, and B. parker', if present in a sample from a subject suspected of having tick-borne relapsing fever (TBRF), said method comprising: providing a biological sample obtained from a subject suspected of having TBRF; mixing the biological sample with one or more of the labeled and/or tagged and/or bound amino acid sequences of Claim 1; and detecting a positive immunobinding reaction which indicates the presence of TBRF specific antibodies in the sample (claim 4; for claims 4-9 of Groups 1+).

Therefore, the common technical features shared by inventions of Groups I+ and Group II can be covered by:

A composition comprising one or more labeled and/or tagged and/or bound amino acid sequences, wherein the one or more labeled and/or tagged and/or bound amino acid sequences comprise amino acid sequences selected from the group consisting of SEQ ID NO: 1 (part of claim 1); and
--a method for detecting antibodies resulting from infection by one or more members of Relapsing Fever (TBRF) Borrelia genus, wherein the genus comprises B. hermsii, B. Turicatae, B. miyamotoi, B. duttonii, and B. parker', if present in a sample from a subject suspected of having tick-borne relapsing fever (TBRF), said method comprising: providing a biological sample obtained from a subject suspected of having TBRF; mixing the biological sample with one or more of the labeled and/or tagged and/or bound amino acid sequences; and detecting a positive immunobinding reaction which indicates the presence of TBRF specific antibodies in the sample (claim 4).

However, these shared technical features do not represent a contribution over prior art as being anticipated by US 2019/0277847 A1 to ID-FISH TECHNOLOGIES, INC. (hereinafter 'ID-Fish') as follows:

ID-Fish discloses a composition comprising one or more labeled and/or tagged and/or bound amino acid sequences (para [0010] - 'a composition comprising one or more labeled and/or tagged and/or bound amino acid sequences'),
--wherein the one or more labeled and/or tagged and/or bound amino acid sequences comprise amino acid sequences selected from the group consisting of SEQ ID NO: 1 (para [0010] - 'a composition comprising one or more labeled and/or tagged and/or bound amino acid sequences, said amino acid sequences consisting of SEQ ID NO: 1', wherein SEQ ID NO: 1 is 100% identical to the claimed SEQ ID NO: 1).

ID-Fish further discloses a method for detecting antibodies resulting from infection by one or more members of Relapsing Fever (TBRF) Borrelia genus (para [0013] - 'detecting Borrelia antisera in a sample from a subject suspected of having tick-borne relapsing fever (TBRF)', wherein 'antisera' are 'antibodies'; Abstract - 'the detection and identification of Tick-Borne Relapsing Fever Borrelia sp. antibodies'),
--wherein the genus is selected from the group consisting of comprises B. hermsii, B. turicatae, B. miyamotoi, and B. parkeri, if present in a sample from a subject suspected of having tick-borne relapsing fever (TBRF) (para [0013] - 'detecting Borrelia antisera in a sample from a subject suspected of having tick-borne relapsing fever (TBRF)'; para [0002] - 'Borrelia bacteria that cause TBRF are typically transmitted to humans through the bite of infected "soft ticks" of the genus Ornithodorus'; para [0005] - 'Borrelia species that cause TBRF,... specific species of ticks... B. Hermsii ...B. parkerii by O. parkeri ticks, and B. Turicatae ... Each tick species has a preferred habitat and preferred set of hosts'; para [0016] - 'detecting and distinguishing various species of Borrelia in a sample from a subject suspected of having tick-borne relapsing fever (TBRF),... B. Hermsii, ... B. miyamotoi in the subject; ...detection of B. turicatae in the sample'), said method comprising:

**********Continued in the next extra sheet**********

Continuation of:
The previous extra sheet - Box No III (unity of invention is lacking)

--providing a biological sample obtained from a subject suspected of having TBRF (para [0013] - 'detecting Borrelia antisera in a sample from a subject suspected of having tick-borne relapsing fever (TBRF), ...providing a biological sample obtained from a subject suspected of having TBRF'; para [0016] - 'detecting and distinguishing various species of Borrelia in a sample from a subject suspected of having tick-borne relapsing fever (TBRF), ... providing a biological sample obtained from a subject suspected of having TBRF');
--mixing the biological sample with one or more of the labeled and/or tagged and/or bound amino acid sequences (para [0013] - 'detecting Borrelia antisera in a sample from a subject suspected of having tick-borne relapsing fever (TBRF), ... providing a biological sample obtained from a subject ... mixing the biological sample with one or more of the labeled, tagged and/or bound amino acid sequences of the present invention'; para [0016] - 'detecting and distinguishing various species of Borrelia in a sample from a subject suspected of having tick-borne relapsing fever (TBRF), ... providing a biological sample ... mixing the biological sample with one or more of the labeled and/or tagged and/or bound amino acid sequences of the present invention'); and
--detecting a positive immunobinding reaction which indicates the presence of TBRF specific antibodies in the sample (para [0013] - 'detecting Borrelia antisera in a sample from a subject suspected of having tick-borne relapsing fever (TBRF), ...providing a biological sample ... mixing the biological sample with one or more of the labeled, tagged and/or bound amino acid sequences ... detecting a positive immunobinding reaction which indicates the presence of TBRF antisera in the sample'; para [0014] - 'a sample is considered positive for TBRF if at least one amino acid sequences is detected'; para [0016] - 'detecting and distinguishing various species of Borrelia in a sample from a subject suspected of having tick-borne relapsing fever (TBRF), ... providing a biological sample obtained from a subject suspected of having TBRF, mixing the biological sample with one or more of the labeled and/or tagged and/or bound amino acid sequences of the present invention and detecting a positive immunobinding reaction which indicates the presence of Borrelia in the subject, wherein detection of SEQ ID NO: 1, ... indicates the presence of B. hermsii, in the subject').

Without a shared special technical feature, the inventions lack unity with one another.

Groups I+ and II therefore lack unity under PCT Rule 13 because they do not share a same or corresponding special technical feature.

Continuation of item 4: Claims 13-14 are not drafted in accordance with the second and third sentences of Rule 6.4 (a). These claims are improper multiple dependent claims.

Note:

I) Each of Claims 1 and 10 is partially omnibus type claim, and is not drafted in accordance with PCT Rule 6.2(a). Each of the claims is partially indefinite as it is unclear what is included or excluded with the limitation "variants thereof which retain the immunological binding profile of Fig. 3", because Fig 3 shows each different SEQ ID NOs is associated with different profiles, and it is impossible for this ISR to test each sequence with unknown serum samples. For the purposes of this ISR, the limitation "and variants thereof which retain the immunological binding profile of Fig. 3" is removed from each of claims 1 and 10.

II) Claim 4 is using an improper Markush group. For the purposes of this ISR, the limitation "the genus comprises B. hermsii, B. turicatae, B. miyamotoi, B. duttonii, and B. parkeri" in claim 4 is assumed to read as "the genus is selected from the group consisting of B. hermsii, B. turicatae, B. miyamotoi, B. duttonii, and B. Parkeri".

INTERNATIONAL SEARCH REPORT

International application No.

PCT/US 22/12229

Continuation of:

A. CLASSIFICATION OF SUBJECT MATTER

C07K 14/00; C12N 15/00; C07K 16/00; C07H 21/04 (2022.01)