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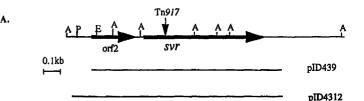
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(54) Title: ANTI-BACTERIAL METHODS AND MATERIALS

(57) Abstract

Staphylococcus aureus virulence genes are identified, thereby allowing the identification of novel anti-bacterial agents that target these virulence genes and their products, and the provision of novel S. aureus mutants useful in vaccines.



B.

MKDEQLYYFEKSPVFKAMMHFSLPMMIGTLLSVIYGILNYIFIGFSEESH MISAISLTLPVFAILMGLGNLFGVGAGTYISRLLGAKDYSKSKFVSSFSI 100 YGGIALGLIVILVTLPFSDQSQQF*

C.



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Inte onal Application No PCT/GB 98/01974

A. CLASSIFICATION OF SUBJECT MATTER IPC 6 C07K14/31 C12Q1/18

A61K39/085

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)

IPC 6 C07K

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

Electronic data base consulted during the international search (name of data base and, where practical, search terms used)

C. DOCUM	ENTS CONSIDERED TO BE RELEVANT	
Category °	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
Х	EP 0 625 575 A (LILLY CO ELI) 23 November 1994 See page 12, line 34-40 see the whole document	1-3,8-14
X	WO 96 17951 A (RPMS TECHNOLOGY LTD ;HOLDEN DAVID WILLIAM (GB)) 13 June 1996 see the whole document	1-3,8-14
Υ	WO 97 11690 A (MICROCIDE PHARMACEUTICALS INC) 3 April 1997 see the whole document	1-3,8-14
	-/	

X Further documents are listed in the continuation of box C.	χ Patent family members are listed in annex.
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Name and mailing address of the ISA European Patent Office, P.B. 5818 Patentlaan 2 NL - 2280 HV Rijswijk Tel. (+31-70) 340-2040, Tx. 31 651 epo ni.	Date of mailing of the international search report 0 3. 03. 1999 Authorized officer
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Inter nal Application No PCT/GB 98/01974

C.(Continu	ation) DOCUMENTS CONSIDERED TO BE RELEVANT	PC1/GB 98/019/4
Category °	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
Y	BERGER-BÄCHI ET AL.: "FemA A HOST MEDIATED FACTOR ESSENTIAL FOR METHICILLIN RESISTANCE IN STAPHYLOCOCCUS AUREUS: MOLECULAR CLONING AND CHARACTERIZATION" MOL.GEN.GENET., vol. 219, 1989, pages 263-269, XP002081494 & DATABASE EMPRO AC:X17688, 100 % identity in 1260 bp overlap with Seq.ID 1 & DATABASE PIR2 AC/ID:S06784, 100 % identity in 419 aa overlap with Seq. ID 2 see abstract	1-3,8-14
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Y	KAMATA ET AL.: "PRIMARY STRUCTURE OF THE ALANINE CARRIER PROTEIN OF THERMOPHILIC BACTERIUM PS3" J.BIOL.CHEM., vol. 267, no. 30, 1992, pages 21650-21665, XP002092572 see the whole document & DATABASE SWISSPROT AC: P30145, 1993 100.0 % identity in 445 aa overlap with Seq.ID 61 see abstract	1,2,8-14

Intel nal Application No PCT/GB 98/01974

C (Continu	ation) DOCUMENTS CONSIDERED TO BE RELEVANT	PC1/GB 98/019/4
Category °	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
A	EBBOLE ET AL.: "CLONING AND CHARACTERIZATION OF A 12-GENE CLUSTER FROM BACILLUS SUBTILIS ENCODING NINE ENZYMES FOR DE NOVO PURINE NUCLEOTIDE SYNTHESIS" J.BIOL.CHEM., vol. 262, no. 17, 1987, pages 8274-8287, XP002092573 & DATABASE SWISSPROT AC: P12042, 1989 54.3 % identity in 738 aa overlap with Seq.ID 30 see abstract	1,2,8-14
A	SAMUELSSON: "A MYCOPLASMA PROTEIN HOMOLOGOUS TO MAMMALIAN SRP54 RECOGNIZES A HIGHLY CONSERVED DOMAIN OF SRP RNA" NUCLEIC ACIDS RESEARCH, vol. 20, no. 21, 1992, pages 5763-5770, XP002092574 & DATABASE SWISSPROT AC: Q01444, 1993 40.7% identity in 427 aa overlap with Seq.ID 74 see abstract	1,2,8-14
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A	RUDNER ET AL.: "THE SPOOK LOCUS OF BACILLUS SUBTILIS IS HOMOLOGOUS TO THE OLIGOPEPTIDE PERMEASE LOCUS AND IS REQUIRED FOR SPORULATION AND COMPETENCE" J.BACTERIOL., vol. 173, no. 4, 1991, pages 1388-1398, XP002092576 & DATABASE SWISSPROT AC:P24136, RUDNER ET AL.: 31.2 % identity in 253 aa overlap with Seq. ID 18 see abstract	1,2,8-14
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Inter Phal Application No
PCT/GB 98/01974

	ation) DOCUMENTS CONSIDERED TO BE RELEVANT	
ategory °	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
1	US 5 587 288 A (CHEUNG AMBROSE ET AL) 24 December 1996 see the whole document	1,2,8-14
1	WO 96 03647 A (ZENECA LTD ;GARMAN ANDREW JOHN (GB); HOLLAND JANET DORA (GB)) 8 February 1996 see the whole document	8-12
1	US 5 283 173 A (FIELDS STANLEY ET AL) 1 February 1994 see the whole document	9-11
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, Y	DATABASE MEDLINE MOL. MICROBIOLOGY, vol. 26, no. 2, pp 399-407, October 1997 MEI ET AL.: "IDENTIFICATION OF STAPHYLOCOCCUS AUREUS VIRULENCE GENES IN A MURINE MODEL OF BACTERAEMIA USING SIGNATURE-TAGGED MUTAGENESIS" XP002092578 see abstract	1-3,8-14
, Ү	DATABASE GCG_GENESEQ_P AC: W28288, BLACK ET AL.: "NOVEL POLYPEPTIDE FROM STAPHYLOCOCCUS AUREUS, ABC TRANSPORTER PROTEIN" XP002092610 94.5 % identity in 181 aa overlap with Seq.ID 20	1,2,8-14
, А	HOLDEN: "DISCOVERING NEW TARGETS FOR ANTI-BACTERIAL AGENTS BY IDENTIFICATION OF GENES ESSENTIAL FOR PATHOGENICITY" BRITISH JOURNAL OF PHARMACOLOGY, vol. 123, March 1998, page 349P XP002092577 see the whole document	1-3,8-14
, А	DATABASE PIR2 AC: F69884, 5 December 1997 KUNST ET AL.: "CONSERVED HYPOTHETICAL PROTEIN ymdA-BACILLUS SUBTILIS" XP002092579 68.5 % in 517 aa overlap with Seq ID 74 see abstract	1,2,8-14

International application No. PCT/GB 98/01974

INTERNATIONAL SEARCH REPORT

Box I Observations where certain claims were found unsearchable (Continuation of item 1 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.: because they are dependent claims and are not drafted in accordance with the second and third sentences of Rule 6.4(a).
Box II Observations where unity of invention is lacking (Continuation of item 2 of first sheet)
This International Searching Authority found multiple inventions in this international application, as follows:
see additional sheet
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 an additional fee, this Authority did not invite payment of any additional fee.
3. X 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.: 1-3,8-14 (all partially). A complete search was done for Seq.ID 1,2,9,10,12,18,20,30,61,66,74 and 80
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.:
Remark on Protest The additional search fees were accompanied by the applicant's protest. X No protest accompanied the payment of additional search fees.

FURTHER INFORMATION CONTINUED FROM PCT/ISA/ 210

This International Searching Authority found multiple (groups of) inventions in this international application, as follows:

1. Claims: 1-3,8-14 (all partially)

Inventions 1-4:

Method of identifying an anti-bacterial agent comprising the steps of assaying potential agents for the ability to interfere with the expression of the virulence gene with Seq. ID 1 or the function of the encoded protein, wherein the protein is involved in cell surface metabolism and a Staphylococcus aureus organism containing a functional mutation in the gene represented by Seq. ID 1 as well as a vaccine composition comprising that mutated S.aureus.

...ibidem for each of sequences represented by Seq. ID 6,60,65.

2. Claims: 1,2,8-14 (all partially)

Inventions 5-67:

Method of identifying an anti-bacterial agent comprising the steps of assaying potential agents for the ability to interfere with the expression of a virulence gene with Seq. ID 2 or the function of the encoded protein and a Staphylococcus aureus organism containing a functional mutation in the gene represented by Seq. ID 2 as well as a vaccine composition comprising that mutated S.aureus.

...ibidem for each of sequences represented by Seq. ID 3,5,7,9,10,12,14,16,18,20,23,24,26,28,30,32,34,36,38,40,42,45,47-53,55-59,61-64,66-74,76,78,80,81,83,85-94

3. Claims: 1,2,4,8-14 (all partiall)

Inventions 68-78

Method of identifying an anti-bacterial agent comprising the steps of assaying potential agents for the ability to interfere with the expression of the virulence gene with Seq. ID 4 or the function of the encoded protein, wherein the protein is involved in cellular biosynthetic pathway and a Staphylococcus aureus organism containing a functional mutation in the gene represented by Seq. ID 4 as well as a vaccine composition comprising that mutated S.aureus.

...ibidem for each of sequences represented by Seq. ID 11,13,22,25,27,29,31,33,35,82

4. Claims: 1,2,7-14 (all partially)

FURTHER INFORMATION CONTINUED FROM PCT/ISA/ 210

Inventions 79-82:

Method of identifying an anti-bacterial agent comprising the steps of assaying potential agents for the ability to interfere with the expression of the virulence gene with Seq. ID 8 or the function of the encoded protein, wherein the protein is involved in cellular regulatory and repair processes, and a Staphylococcus aureus organism containing a functional mutation in the gene represented by Seq. ID 8 as well as a vaccine composition comprising that mutated S.aureus.

...ibidem for each of sequences represented by Seq. ID 21,46,84

5. Claims: 1,2,6,8-14 (all partially)

Inventions 83-89

Method of identifying an anti-bacterial agent comprising the steps of assaying potential agents for the ability to interfere with the expression of the virulence gene with Seq. ID 15 or the function of the encoded protein, wherein the protein is an oligopeptide transport protein and a Staphylococcus aureus organism containing a functional mutation in the gene represented by Seq. ID 15 as well as a vaccine composition comprising that mutated S.aureus.

...ibidem for each of sequences represented by Seq. ID 17,19,54,75,77,79

6. Claims: 1,2,5,8-14 (all partially)

Inventions 90-94:

Method of identifying an anti-bacterial agent comprising the steps of assaying potential agents for the ability to interfere with the expression of the virulence gene with Seq. ID 37 or the function of the encoded protein, wherein the protein is a component of the TCA cycle and a Staphylococcus aureus organism containing a functional mutation in the gene represented by Seq. ID 37 as well as a vaccine composition comprising that mutated S.aureus.

...ibidem for each of sequences represented by Seq. ID 39,41,43,44.

information corpatent family members

Inter nal Application No PCT/GB 98/01974

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