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The applicant/owner of the patent is registered as Trilochan Kanwaljit Singh Mukkur. They used the patent attorney firm Fisher Adams Kelly Pty Ltd to file this.

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PUBLICATIONS

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CLAIMS

CLAIMS 1. A genetically modified Bordetella strain having a partial or complete loss of 00 function in the endogenous aroQ gene and a lower capacity to propagate in a mammalian host but remaining viable in the host for a period of time sufficient to NO induce an immune response against a pathogenic Bordetella strain. 2. The genetically modified strain of claim 1, wherein the pathogenic Bordetella strain is a natural counterpart of the genetically modified Bordetella strain, or is a 0 10 related pathogenic Bordetella organism. 3. The genetically modified strain of claim 1, wherein the pathogenic strain is selected from Bordetella avium, Bordetella bronchiseptica, Bordetella holmesii, Bordetella parapertussis and Bordetella pertussis. 4. The genetically modified strain of claim 1, wherein the pathogenic Bordetella strain is Bordetella pertussis. The genetically modified strain of claim 1, comprising a disruption in the endogenous aroQ gene. 6. The genetically modified strain of claim 5, wherein the disruption has been introduced into the genome of a pathogenic strain of Bordetella by homologous recombination with a DNA targeting construct such that the targeting construct is stably integrated in the genome, wherein the disruption of the aroQ gene results in a reduced level and/or functional activity of the 3-dehydroquinase. 7. The genetically modified strain of claim 5, comprising an exogenous nucleic acid sequence in its genome, or on an extrachromosomal element, which is capable of abolishing or otherwise reducing the expression of aroQ or the level and/or functional activity of the 3-dehydroquinase encoded by aroQ.-76- 00 8. The genetically modified strain of claim 5, comprising an exogenous nucleic N acid sequence in its

genome, or on an extrachromosomal element, which is capable of abolishing or otherwise reducing the expression of aroQ or the level and/or functional activity of the 3-dehydroquinase encoded by aroQ, wherein the nucleic acid sequence comprises at least a portion of aroQ, in the sense or anti-sense orientation, which is operably linked to a transcriptional control element. 9. The genetically modified strain of claim 1, further having a partial or complete loss of function in at least one other endogenous gene selected from a purgene, another aro gene, a pertussis toxin gene, or any other gene which contributes to survival in the host and/or to bacterial virulence, or a combination thereof. The genetically modified Bordetella strain of claim 1, comprising at least one exogenous gene which is capable for expressing an antigen that is heterologous or foreign to the Bordetella strain. 11. The genetically modified Bordetella strain of claim 10, wherein the heterologous or foreign antigen is derived from a pathogen that is unrelated to the Bordetella strain. 12. The genetically modified Bordetella strain of claim 10, wherein the heterologous or foreign antigen is derived from a pathogen that infects by the mucosal route. 13. An isolated polynucleotide comprising a nucleotide sequence set forth in SEQ ID NO: 1 or 3, or is complementary to the sequence set forth in SEQ ID NO: 1 or 3. 14. An isolated polynucleotide comprising a nucleotide sequence having at least sequence identity to at least a portion of the sequence set forth in SEQ ID NO: 1 or 3, wherein said portion is at least 150 nucleotides in length. 15. The isolated polynucleotide of claim 14 wherein the nucleotide sequence has at least 90% sequence identity at least a portion of the sequence set forth in SEQ ID NO: 1 or 3, wherein said portion is at least 150 nucleotides in length. 16. A nucleic acid construct for disrupting an aroQ gene in a Bordetella cell, comprising: a) a non-homologous replacement portion; b) a first homology region located upstream of the non-homologous replacement portion, the first homology region having a nucleotide sequence with substantial identity to a first aroQ gene sequence; and c) a second homology region located downstream of the non-homologous replacement portion, the second homology region having a nucleotide sequence with substantial identity to a second aroQ gene sequence, the second aroQ gene sequence having a location downstream of the first aroQ gene sequence in a naturally occurring endogenous aroQ gene of the Bordetella cell. 17. The construct of claim 16, wherein the aroQ gene comprises the isolated polynucleotide as set forth in any one of claims 13 to 18. A vector comprising a nucleotide sequence that corresponds or is complementary to the isolated polynucleotide sequence as set forth in any one of claims 13 to 19. The vector of claim 18, wherein the vector is a DNA targeting vector. A host cell containing the construct of claim 16 or claim 17 or the vector of claim 18 or claim 19. 21. A method for producing a genetically modified Bordetella strain, comprising introducing the nucleic acid construct of claim 25 into a Bordetella cell under conditions such that the nucleic acid construct is homologously recombined into the aroQ gene in the genome of that cell to produce a genetically modified Bordetella cell containing a disrupted aroQ gene. 22. The method of claim 21, wherein the genetically modified Bordetella cell containing the homologously recombined nucleic acid construct is further characterised by expressing reduced or undetectable levels of aroQ. 23. The method of claim 21, wherein the genetically modified Bordetella cell lacks the ability to produce a functional 3-dehydroquinase encoded by said aroQ gene. 24. A composition, comprising the genetically modified Bordetella strain of claim 1, together with a pharmaceutically acceptable carrier. The composition of claim 24, further comprising an adjuvant. 26. A composition of matter comprising dendritic cells which have been exposed to the genetically modified Bordetella strain of claim 1 for a time and under conditions sufficient to express a processed or modified antigen derived from the Bordetella strain for presentation to, and modulation of, T cells. 27. The composition of matter of claim 26, which is in the form of an in vitro cell culture. 28. A method for modulating an immune response, comprising administering to a patient in need of such treatment an effective amount of the genetically modified Bordetella strain of claim 1, or the composition of any one of claims 22-25 or the composition of matter of claim 26. 29. A method for the treatment and/or prophylaxis of whooping cough or related condition, comprising administering to a patient in need of such treatment an effective amount of the genetically modified Bordetella strain of claim 1, or the composition of any one of claims 22-25 or the composition of matter of claim 26.

WO 03/060105 WO 03/60105PCT/A1J02/01768 1/8 BvgI (245) PvuII (306) EcoRI (396) SmaI (606) NgoMIV (671) EcoRI KpnI (659) aroQ fragment BssHI (874) FIGURE 1Z TdI7IDHj 099 Tog 009 3 6aa 3D 2*65vl me 465bz htm~~~~qS0 wnb vf3b'e jf1e06frat ona tm BES I i e &Z-6 Tslbv-z6 w o 14e62.ej 2.azi II.6CDI-{ n o-j 0;low ;;)3Be -4a~ b4f6W.(I m :X 0051P 3~m z4.\$mn 4 nnb& ie6bqb' DLLW DZ.. NZ.. Tot :uebG~u One 6d.) onefinodv Qumfied cuebdo .u bdz uwasjdod aucbanad avuEpd0 euebqnode eufou anc6;in.d auabdo *ueBanode, ZNZO tmgBOu ausbd= cuchnndv numbe~jsd 21430 Oaf P 6 .0 :m:>Dvnmb~ 6=16.P m NIPUr m

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03/60105PCT/A1102/01768 3/8 p 998 193 pUSQBord8 S Highlighted insert ligated into pRTP 1 (a s BIS HI
pUSQBord9 898193 Mi Triple Ligation: Inserts frm pUSQBord9 and pUSQBordl ligated together and inserted
into pRTP1 in the same reaction Ea.ndad 11-1s JgOM V Hindlil f9ll BaMI-II PUSQBard1O ,Final Canetuact
FIGURE 3WO 03/060105 WO 03/60105PCT/AU02/01768 4/8 1 23 45 6 2332 bD 2027 bp Inactivated gene size
1 approximately 1150 bp 564 bp 2000 bp 1200 bp aw 800 bp AroQ gene size approximately 500 bp FIGURE
4WO 03/060105 WO 03160105PCT/AU02/01768 5/8 2 4 EmrBa Mew~ -2.0 SD Days Post Inoculation I 1C 12 14
FIGURE c' 6 CL C. 4- 0 2 0 0 2 4 6 8 10 12 14 Days Post Vaccination FIGURE WO 03/060105 WO
03/60105PCT/A1J02/01768 6/8 12.00. 10.00 -I 9.00--i 8.00-- 600-1 4 00 J .001 2 4 0 a 10 12 14 Days Post
Inorulation 8-r Slv M.-s 2.0 SE) DeAe lvM-r FIGURE control mice X4- vaccinated mice 0 2 4 6 8 10 12 14 16
1820 2224 2628 Days Post Challenge FIGURE 6WO 03/060105 WO 03/60105PCT/AU02/01768 7/8 Eont,- mice
vacc.intd mice] 1 10 4 21 Days Post Challenge FIGURE 7A 0 control mice .1 10 14 21 Days Post Challenge
FIGURE 7BWO 03/060105 WO 03/60105PCT/AU02/01768 8/8 200 0 control mice U vaccinated mice E150-____
E 100- E L 0- -1 10 14 21 Days Post Challenge FIGURE 8A 0 control mice U vaccinated mice S150 E 100 z LL.
50 0 -1 10 14 21 Days Post Challenge FIGURE 8BWO 031060105 WO 03160105PCT/AU02/01768 1-
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Edward (US only) Mukkur, Trilochan Kanwaljit Singh Rossetti, Tony Robert (US only) states except US) (US
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CHANGE OF OWNERSHIP

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DRAWINGS



Defective entities and uses therefor

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