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**Identification and characterisation of an exported immunogenic  
protein of *Mycobacterium avium* subspecies *paratuberculosis***

A thesis presented in partial fulfillment of the requirements for the degree of

**Doctor of Philosophy**

at Massey University, Palmerston North, New Zealand

**Christine Dupont**

**2002**

# MASSEY UNIVERSITY

APPLICATION FOR APPROVAL OF REQUEST TO EMBARGO A THESIS  
(Pursuant to AC98/168 (Revised 2), Approved by Academic Board 16.02.99)

Name of Candidate: Christine Dupont I.D. Number: 99119179

Degree: PhD Dept / Institute / School: IUABS

Thesis Title: Identification and characterisation of an exported immunogenic protein of M. avium subspecies paratuberculosis

Name of Chief Supervisor: Dr Alan Murray Telephone Ext: 7895

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## Abstract

Exported proteins of mycobacteria are available to interact with the immune system at an early stage of infection and are potent inducers of immune responses. Potentially exported proteins of *Mycobacterium avium* subspecies *paratuberculosis* were identified using alkaline phosphatase gene fusion technology. A library of partial gene fusions from a New Zealand clinical isolate of *M. a. paratuberculosis* was constructed in the shuttle vector pJEM11 and expressed in the surrogate hosts *E. coli* and *M. smegmatis*. The DNA inserts from a portion of the resulting clones expressing alkaline phosphatase-positive fusion proteins were partially sequenced to identify the proteins. Eleven proteins not previously described for *M. a. paratuberculosis* were identified as containing signal sequences for export. One of these, a putative lipoprotein named P22 was selected for further study. The full nucleic acid sequence of the *p22* gene was determined and the open reading frame was cloned into the mycobacterial expression vector pMIP12. This enabled P22 to be produced as a polyhistidine-tagged protein in *M. smegmatis* and facilitated purification by chromatography. N-terminal sequencing of the recombinant protein confirmed cleavage of an N-terminal signal sequence. Native P22 was detected in culture supernatants and cell sonicates of *M. a. paratuberculosis* strain 316F using rabbit antibody raised to P22. Investigation of the presence of genes similar to *p22* in other mycobacterial species, revealed *p22* was present in *Mycobacterium avium* subspecies *avium* and similar genes existed in *M. intracellulare* (88.5% identity) and *M. scrofulaceum* (87.7% identity). Database searches showed P22 belonged to the LppX/LprAFG family of mycobacterial lipoproteins also found in *M. leprae* and in members of the *M. tuberculosis* complex. P22 shared less than 75% identity to these proteins. Recombinant P22 was able to elicit significantly increased interferon-gamma secretion in blood from a group of eight sheep vaccinated with a live, attenuated strain of *M. a. paratuberculosis* (strain 316F) compared to a group of five unvaccinated sheep. Antibody to P22 was detected by Western blot analysis in 10 out of 11 vaccinated sheep, in two out of two clinically affected cows and in 11 out of 13 subclinically infected cows.

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Construction of the *M. a. paratuberculosis* PhoA fusion library and characterisation of SodC from this study have been published (Dupont, C. & Murray, A. (2001). Identification, cloning and expression of *sodC* from an alkaline phosphatase gene fusion library of *Mycobacterium avium* subspecies *paratuberculosis*. *Microbios* **106** S1: 7-19).

## Table of Contents

	<b>Page</b>
--	-------------

<b>Abstract.....</b>	<b>i</b>
<b>Acknowledgements.....</b>	<b>ii</b>
<b>List of Figures.....</b>	<b>ix</b>
<b>List of Tables.....</b>	<b>xi</b>
<b>List of Abbreviations.....</b>	<b>xiii</b>
<b>Chapter 1 Literature review</b>	
1.1    History.....	1
1.2    Prevalence and economic impact.....	1
1.3    Host range .....	3
1.3.1    Crohn's disease.....	3
1.4    Classification .....	5
1.5    Clinical signs and transmission.....	6
1.6    Control.....	8
1.6.1    Test and cull.....	8
1.6.2    Vaccination .....	8
1.6.3    Management .....	10
1.7    Detection.....	11
1.7.1    Culture.....	11
1.7.2    Detection of DNA.....	13
1.7.2.1    IS900 insertion element .....	13
1.7.3    Immunological tests.....	15
1.7.3.1    Tests for cellular responses.....	16
1.7.3.2    Tests for humoral responses.....	16
1.7.4    Histopathological detection.....	17
1.8    The mycobacterial envelope and its relationship to pathogenicity and immunology..	18
1.8.1    Lipoarabinomannan .....	19
1.9    Export of proteins in mycobacteria .....	21
1.9.1    Export pathways and signal peptides .....	23
1.9.1.1    Cleavage of signal peptides .....	25
1.9.1.2    Lipidation.....	25
1.9.1.3    C-terminal anchoring .....	26
1.10    Searching for exported proteins of mycobacteria.....	27
1.11    Components isolated from <i>M. a. paratuberculosis</i> .....	29

1.12	Pathogenesis and immune responses .....	32
1.12.1	Pathogenesis .....	32
1.12.2	Immune responses .....	34
1.12.2.1	T-cells.....	35
1.12.2.2	B-cells.....	38
1.13	Summary and aims of the thesis .....	39
<b>Chapter 2 General materials and methods</b>		
2.1	Bacterial strains and plasmids.....	41
2.1.1	Bacterial strains.....	41
2.1.2	Plasmids .....	42
2.2	Bacterial growth and storage conditions.....	43
2.2.1	<i>E. coli</i> .....	43
2.2.2	<i>M. smegmatis</i> .....	43
2.2.3	<i>M. a. paratuberculosis</i> .....	44
2.3	DNA isolations.....	44
2.3.1	Isolation of plasmid DNA from <i>E. coli</i> .....	44
2.3.2	Isolation of genomic DNA from mycobacterial species.....	44
2.4	DNA manipulations and cloning procedures .....	45
2.4.1	Restriction endonuclease digestions.....	45
2.4.2	Electrophoresis .....	45
2.4.3	Extraction from agarose gels.....	46
2.4.4	Southern blotting and hybridisations .....	46
2.4.4.1	DNA probe preparation .....	47
2.4.4.2	Removal of probe from Southern blots.....	47
2.4.5	Polymerase chain reactions .....	47
2.4.5.1	Primer design.....	47
2.4.5.2	PCR conditions.....	48
2.4.6	DNA sequencing .....	49
2.4.7	Ligations .....	49
2.5	Bacterial transformations .....	50
2.5.1	Preparation of electrocompetent <i>M. smegmatis</i> .....	50
2.5.2	Transformation of <i>E. coli</i> .....	50
2.5.3	Transformation of <i>M. smegmatis</i> .....	51
2.6	Protein isolations .....	51
2.6.1	Preparation of cell lysates .....	51

2.6.2	Preparation of culture filtrates.....	52
2.6.2.1	<i>M. smegmatis</i> .....	52
2.6.2.2	<i>M. a. paratuberculosis</i> .....	52
2.6.3	Ni <sup>+2</sup> -affinity chromatography.....	52
2.6.4	Size exclusion chromatography .....	53
2.7	Protein analyses.....	54
2.7.1	Polyacrylamide gel electrophoresis .....	54
2.7.2	Western blotting and immunodetections.....	54
2.7.2.1	Immunodetection of Western blots.....	55
2.7.2.2	Removal of antibody from Western blots .....	55
2.7.3	N-terminal protein sequencing .....	55
2.7.4	Estimation of protein concentration and molecular weight .....	56
2.8	IFN- $\gamma$ assays.....	57
2.9	Rabbit details and immunisations.....	57
2.9.1	Preparation of P22 for immunisation.....	57
2.9.2	Rabbit immunisation protocol .....	58
2.10	Sheep details and immunisations .....	58
2.10.1	Experimental sheep and immunisation protocol.....	58
2.10.2	Naturally infected sheep.....	59
2.11	Naturally infected cattle.....	60
2.12	Bioinformatics .....	60

### **Chapter 3 Identification of *M. a. paratuberculosis* DNA sequences encoding exported proteins**

3.1	Abstract.....	62
3.2	Introduction .....	63
3.3	Materials and methods.....	65
3.3.1	Construction of an <i>M. a. paratuberculosis</i> pJEM11 expression library.....	65
3.3.1.1	Extraction of DNA from <i>M. a. paratuberculosis</i> .....	65
3.3.1.2	Preparation of pJEM11 vector DNA.....	66
3.3.1.3	Partial digestion with <i>Sau</i> 3A of <i>M. a. paratuberculosis</i> DNA.....	66
3.3.1.4	Ligation of <i>M. a. paratuberculosis</i> DNA and pJEM11 and transformation into <i>E. coli</i> .....	66
3.3.1.5	Plasmid isolation from the <i>E. coli</i> recombinant library.....	67
3.3.1.6	Transformation of the recombinant plasmids into <i>M. smegmatis</i> mc <sup>2</sup> 155....	67
3.3.2	Sequencing of DNA inserts encoding putative exported proteins .....	68

<b>3.4</b>	<b>Results.....</b>	<b>69</b>
<b>3.4.1</b>	<b>Construction of an <i>M. a. paratuberculosis</i> pJEM11 expression library.....</b>	<b>69</b>
<b>3.4.1.1</b>	<b>Confirmation of <i>M. a. paratuberculosis</i> DNA for cloning .....</b>	<b>69</b>
<b>3.4.1.2</b>	<b>Subcloning of <i>M. a. paratuberculosis</i> DNA into the vector pJEM11 and expression of the library in <i>E. coli</i> .....</b>	<b>69</b>
<b>3.4.1.3</b>	<b>Expression of the library in <i>M. smegmatis</i> .....</b>	<b>70</b>
<b>3.4.2</b>	<b>Analysis of <i>M. a. paratuberculosis</i> phoA fusions .....</b>	<b>71</b>
<b>3.4.2.1</b>	<b>Sequencing of DNA inserts encoding putative exported proteins .....</b>	<b>71</b>
<b>3.4.2.2</b>	<b>Analysis of phoA fusions.....</b>	<b>71</b>
<b>3.5</b>	<b>Discussion.....</b>	<b>77</b>

**Chapter 4 Cloning, heterologous expression and characterisation of an immunogenic 22 kDa protein from *M. a. paratuberculosis***

<b>4.1</b>	<b>Abstract.....</b>	<b>83</b>
<b>4.2</b>	<b>Introduction .....</b>	<b>84</b>
<b>4.3</b>	<b>Materials and methods.....</b>	<b>86</b>
<b>4.3.1</b>	<b>PCR amplification of the <i>p22</i> gene from <i>M. a. paratuberculosis</i> .....</b>	<b>86</b>
<b>4.3.2</b>	<b>Cloning of the <i>p22</i> open reading frame.....</b>	<b>86</b>
<b>4.3.3</b>	<b>Expression and purification of P22 recombinant protein from <i>M. smegmatis</i> .....</b>	<b>88</b>
<b>4.3.3.1</b>	<b>Western blot analyses of P22 recombinant protein.....</b>	<b>88</b>
<b>4.3.4</b>	<b>Preparation of rabbit antibody raised to P22.....</b>	<b>89</b>
<b>4.3.5</b>	<b>PCR amplification of the <i>p22</i> ORF from genomic DNA.....</b>	<b>89</b>
<b>4.4</b>	<b>Results.....</b>	<b>90</b>
<b>4.4.1</b>	<b>Sequence analysis of plasmid pTB-16 and identification of the <i>p22</i> open reading frame .....</b>	<b>90</b>
<b>4.4.1.1</b>	<b>Sequence analysis of pTB-16 .....</b>	<b>90</b>
<b>4.4.1.2</b>	<b>Identification of the <i>p22</i> ORF.....</b>	<b>91</b>
<b>4.4.2</b>	<b>Sequence analysis of <i>p22</i>.....</b>	<b>91</b>
<b>4.4.2.1</b>	<b>Sequence similarities between P22 and a family of mycobacterial lipoproteins.....</b>	<b>91</b>
<b>4.4.3</b>	<b>Cloning and expression of the <i>p22</i> ORF .....</b>	<b>93</b>
<b>4.4.4</b>	<b>Purification of recombinant P22 from cell lysates.....</b>	<b>94</b>
<b>4.4.5</b>	<b>Analysis of <i>M. smegmatis</i> culture filtrates for the presence of recombinant P22.....</b>	<b>94</b>
<b>4.4.6</b>	<b>Immune responses to P22.....</b>	<b>94</b>

4.4.6.1	Humoral immune responses to P22.....	94
4.4.6.2	Cell-mediated immune responses to P22 .....	96
4.4.7	Localisation of P22 in <i>M. a. paratuberculosis</i> .....	97
4.4.7.1	Detection of P22 with serum from sheep vaccinated with <i>M. a. paratuberculosis</i> strain 316F culture filtrate.....	97
4.4.7.2	Production of rabbit antibody raised to P22 and detection of P22 in cellular fractions.....	98
4.4.8	Species distribution of the <i>p22</i> gene .....	98
4.4.8.1	PCR amplification of the <i>p22</i> gene from <i>Mycobacterium</i> species and strains.....	99
4.4.8.2	Southern blot detection of the <i>p22</i> gene in <i>Mycobacterium</i> species and strains.....	100
4.5	Discussion.....	101
	<b>General discussion and conclusions.....</b>	<b>111</b>
<b>Appendix 1</b>	Commonly used solutions .....	117
<b>Appendix 2</b>	Diagnosis of Johne's disease and results for "Limestone Downs" sheep.	119
<b>Appendix 3</b>	Most significant protein database alignments obtained using the translated DNA segments fused to <i>phoA</i> .....	120
<b>Appendix 4</b>	<i>M. a. paratuberculosis p22</i> and <i>M. bovis lpp-27 (lprG)</i> alignment (Fasta 3).....	129
<b>Appendix 5</b>	Raw data for IFN- $\gamma$ assay Figure 4.13.....	131
<b>Appendix 6</b>	Raw data for IFN- $\gamma$ assay Figure 4.14.....	132
<b>Appendix 7</b>	Alignment of DNA and protein sequences of <i>M. a. paratuberculosis</i> <i>p22</i> with partial sequences of <i>M. intracellulare</i> and <i>M. scrofulaceum</i> ....	133
	<b>References .....</b>	<b>135</b>

## List of Figures

Figure	Page
Figure 1.1. Relationships between <i>Mycobacterium</i> species based on 16S rRNA sequence homology.....	6a
Figure 1.2. General relationship of tests and immune responses to infection with <i>M. a. paratuberculosis</i> over time.....	16a
Figure 1.3. Common features of a signal peptide. ....	24a
Figure 3.1. Schematic of the <i>E. coli/Mycobacterium</i> shuttle vector pJEM11.....	64a
Figure 3.2. Schematic representation of the location of the oligonucleotide primers designed for sequencing the <i>M. a. paratuberculosis</i> inserts in the pJEM11 constructs.....	68a
Figure 3.3. Schematic representation of the construction of the PhoA fusion library.....	69a
Figure 3.4. PCR amplification of <i>M. a. paratuberculosis</i> genetic elements. ....	69b
Figure 3.5. PhoA <sup>+</sup> recombinant <i>E. coli</i> colonies from the <i>M. a. paratuberculosis</i> pJEM11 library.....	69c
Figure 3.6. PhoA <sup>+</sup> recombinant <i>M. smegmatis</i> colonies from the <i>M. a. paratuberculosis</i> pJEM11 library.....	70a
Figure 3.7. Restriction endonuclease analysis of selected PhoA <sup>+</sup> clones.....	71a
Figure 4.1. Schematic of the <i>Mycobacterium</i> expression vector pMIP12 for the production of histidine-tagged recombinant proteins in <i>M. smegmatis</i> . ....	86a
Figure 4.2. Sequence analysis of the p22 ORF.....	91a
Figure 4.3. Amino acid sequence comparison between P22 of <i>M. a. paratuberculosis</i> and database search results.....	91b
Figure 4.4. Kyte-Doolittle plot (top) and signal sequence features (bottom) of the P22 precursor protein.....	92a
Figure 4.5. Comparison of the promoter regions of <i>M. bovis lpp-27</i> and <i>M. a. paratuberculosis p22</i> .....	92b
Figure 4.6. Restriction endonuclease digest of plasmid pMIP-p22. ....	93a
Figure 4.7. Expression of recombinant P22 from <i>M. smegmatis</i> .....	93b

Figure 4.8. Affinity chromatography of recombinant P22.....	94a
Figure 4.9. Detection of recombinant P22 from <i>M. smegmatis</i> culture filtrates.....	94b
Figure 4.10. Detection of antibody to P22 in sheep vaccinated with Neoparasec.....	94c
Figure 4.11. Detection of antibody to P22 in individual sheep from a naturally infected flock.....	95a
Figure 4.12. Detection of antibody to P22 in naturally infected cattle.....	95b
Figure 4.13. IFN- $\gamma$ induction using Ni <sup>+2</sup> -affinity-enriched P22 in Neoparasec-vaccinated sheep blood.....	96a
Figure 4.14. IFN- $\gamma$ induction by purified recombinant P22 in Neoparasec-vaccinated sheep blood.....	97a
Figure 4.15. Detection of antibody to P22 from sheep vaccinated with <i>M. a. paratuberculosis</i> strain 316F culture filtrate.....	97b
Figure 4.16. Western blot detection of rabbit antibody raised to P22.....	98a
Figure 4.17. Detection of native P22 in Western blots of <i>M. a. paratuberculosis</i> strain 316F cell fractions and comparison to recombinant P22 using rabbit antibody raised to P22.....	98b
Figure 4.18. PCR amplification of the <i>p22</i> gene from 13 isolates of <i>M. a. paratuberculosis</i> .....	99a
Figure 4.19. PCR amplification from 22 mycobacterial strains using primers designed to the <i>p22</i> ORF.....	99b
Figure 4.20. Southern blot analyses using a <i>p22</i> probe from genomic DNA of 13 mycobacterial strains.....	100a

## List of Tables

<b>Table</b>	<b>Page</b>
1.1 A summary of the clinical stages of paratuberculosis .....	8
1.2 Components isolated from <i>M. a. paratuberculosis</i> .....	29
2.1 Bacterial strains used in this study .....	41
2.2 Plasmids used in this study .....	42
2.3 Antibiotics and supplements used in microbiological media.....	44
2.4 Sheep treatment groups.....	59
3.1 Oligonucleotide primers designed for analysis of <i>M. a. paratuberculosis</i> DNA....	68
3.2 Identification of selected <i>M. a. paratuberculosis</i> PhoA fusion proteins .....	73
4.1 Summary of results for detection of <i>M. a. paratuberculosis</i> by serum ELISA, faecal culture and P22 Western blot analysis.....	96

## List of Abbreviations

$A_{280\text{ nm}}$	absorbance at 280 nm
ATCC	American type culture collection
Avian PPD	Purified protein derivative from <i>M. a. avium</i>
BCG	bacillus Calmette-Guerin
BCIP	5-bromo-4-chloro-3-indolyl phosphate
BLAST	basic local alignment search tool
ConA	concanavalin A
dTTP	deoxythymidine triphosphate
dUTP	deoxyuridine triphosphate
DIG	digoxigenin
DNA	deoxyribonucleic acid
EDTA	ethylenediamine tetraacetic acid
ELISA	enzyme-linked immunosorbent assay
HPLC	high pressure liquid chromatography
IFN- $\gamma$	interferon-gamma
Johnin PPD	purified protein derivative from <i>M. a. paratuberculosis</i>
kan	kanamycin
kb	kilobase pairs
kDa	kilodalton(s)
LAM	lipooligosaccharide
LB	Luria-Bertani
OD	optical density
ORF	open reading frame
PBS	phosphate-buffered saline
PCR	polymerase chain reaction
PhoA	alkaline phosphatase
POD	peroxidase
PVDF	polyvinylidene difluoride
RBS	ribosome binding site
SDS-PAGE	sodium dodecyl sulphate-polyacrylamide gel electrophoresis
TAE	Tris-acetate, EDTA
UV	ultraviolet

Amino acids	A	alanine	C	cysteine
	D	aspartic acid	E	glutamic acid
	F	phenylalanine	G	glycine
	H	histidine	I	isoleucine
	K	lysine	L	leucine
	M	methionine	N	asparagine
	P	proline	Q	glutamine
	R	arginine	S	serine
	T	threonine	W	tryptophan
	V	valine	Y	tyrosine

Nucleic acids	A	adenosine
	T	thymidine
	C	cytidine
	G	guanosine