Copyright is owned by the Author of the thesis. Permission is given for a copy to be downloaded by an individual for the purpose of research and private study only. The thesis may not be reproduced elsewhere without the permission of the Author.

ISOLATION AND CHARACTERISATION OF HOST DEFENCE PEPTIDES OSTRICACINS FROM OSTRICH HETEROPHILS

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

Masterate of Technology in Biotechnology

at Massey University, Palmerston North, New Zealand

Haryadi Sugiarto 2006

ABSTRACT

Host defence peptides are ubiquitous components of innate immunity within all living organisms. These peptides are small, positively charged and amphiphilic molecules. The biological roles of these peptides are direct antimicrobial activity against pathogens and to induce the innate and adaptive immune response within the host.

The research presented in this thesis was focused on isolating host defence peptides from ostrich blood and characterising their antimicrobial properties. Four ostrich β-defensins, named ostricacins-1-4 (Osp-1-4), were successfully purified from ostrich blood. These peptides contained 36-42 amino acid residues, with the main residues including: arginines, lysines, glycines and cysteines. The molecular weight of these four ostricacins ranged between 4-5 kDa. They displayed antimicrobial activity against Gram-negative bacteria and Gram-positive bacteria with minimum inhibitory concentration (MIC), ranging between 1-12 µg/ml. In addition, Osp-2 displayed antimicrobial activity against yeast, with MIC of 6.2 µg/ml. Osp-1 and Osp-2 were further characterised with the investigation of the effects of cationic ions and temperature changes on their antimicrobial activity against Gram-negative bacteria and Gram-positive bacteria. The antimicrobial activity of both peptides significantly declined with the presence of cationic ions. Both peptides were relatively stable when heated to temperatures between 30-70°C. Finally, an investigation of the mode of action of Osp-1 and Osp-2 against Gram-negative bacteria was carried out. Both peptides were compared with a sheep cathelicidin, SMAP-29, and a human α -defensin, HNP-1. SMAP-29 showed the strongest affinity to LPS and it was the most potent peptide to cause disruption of the outer and cytoplasmic membrane. The two ostricacins showed stronger affinity than HNP-1 and they also indicated partial permeabilisation of the outer membrane and a slight depolarisation of the cytoplasmic membrane. HNP-1 did not indicate disruption of the outer membranes or depolarisation of the cytoplasmic membrane. Further investigation indicated that the partial disruption allowed the ostricacins to pass through the membranes and interact with the intracellular components. However, these peptides could not inhibit the bacterial colony forming potential, and therefore, they were considered bacteriostatic. It is recommended that further research be carried out to investigate the feasibility of ostricacins in adding value to existing topical products.



ACKNOWLEDGEMENT

First and foremost, I would like to express my deepest appreciation to my supervisor, Dr Pak-Lam Yu for providing knowledge, guidance, advice and motivation throughout this research project. I would also like to thank Dr Jon-Paul Powers at the University of British Columbia, Vancouver, Canada. His advice and assistance in the fluorescence assays via email and telephone correspondence was extremely helpful. In addition, I would like to thank Ms Diana Carne and her team at the Protein Microchemistry Facility, University of Otago, Dunedin, which provided the service of mass spectrometry and N-terminal sequencing.

This study was made possible with funding by C. Alma Baker Trust and Massey University Research Fund (MURF). I am also grateful for the staff at Feilding Meat Packers who assist me in collecting the ostrich blood used for these experiments.

Furthermore, I would like to thank the administration and the technical staff of Institute Technology and Engineering (ITE), especially Ms Ann-Marie Jackson, Mr Michael Sahayam, Mr John Edwards, Mr John Sykes, Mr Michael Stevens, Mr Martin Peak, Ms Dannie van Der Linden, Ms Joan Brooks and Ms Gayle Leader. These wonderful people have assisted me in dealing with paper work, ordering media, finding equipment and technical difficulties in the laboratory. I would also like to acknowledge Dr Andy Shilton for lending me the spectrofluorophotometer.

Many thanks to all the other postgraduate students and visiting students who have shared the same or worse problem during my time, including: Antje, Craig, Hans, Jeremy, Nick, Preyas, Rachel, Robert W., Steven.

Finally, I would like to express gratitude for my friends. Christopher, Donny, Fernando, Findy, Herry, Liony, Villy & Kandy, Christian and Lucia are a few names whom I am indebted for their friendship and support. Especially for Felicia, thank you for your patience, companionship and understanding. Last but not least, I would like to dedicate this thesis to my parents. Mum and Dad thank you for your love, support and encouragement through all my life.

TABLE OF CONTENTS

Abstractii
Acknowledgementv
Table of Contentsvii
List of Figuresxi
List of Tablexiii
Chapter 1 Literature Review1
1.1 Introduction
1.2 Animal and Human Immune Systems
1.2.1 Innate Immunity
1.2.2 Adaptive Immunity6
1.3 Antimicrobial Peptides
1.3.1 Classes of Host Defence Peptides
1.3.2 Host Defence Peptides Roles in Immunity
1.3.2.1 Roles in Innate Immunity
1.3.2.2 Roles in Adaptive Immunity
1.4 Mode of Actions
1.4.1 Membrane Disruptive Mechanisms
1.4.2 Non-Membrane Disruptive Mechanisms
1.5 Defensins
1.5.1 Animal and Human Defensins24
1.5.2 Biological roles
1.6 Purification and Characterisation of Avian β-Defensins
1.6.1 Avian Heterophil β-Defensins
1.6.2 Avian Non-Heterophil β-Defensins
1.6.3 Antimicrobial Activity of Avian β-Defensins
1.6.4 Purification and Characterisation Techniques of Avian Heterophil β-Defensins
1.7 Evolutionary of β-defensins

1.8	App	lications of Host Defence Peptides	41
1.8	8.1	Current Developments of Host Defence Peptides into Therapeu Applications	
1.8	8.2	Potential Applications for Ostrich Host Defence Peptides	44
1.9	Con	clusion	45
1.10	Obje	ectives of Research	47
Chapte	er 2 M	aterials and Methods	49
2.1	Mat	erials, Chemicals and Media	49
2.2	Pep	tide Extraction and Purification	51
2.2	2.1	Crude Extraction	51
2.2	2.2	Cation-exchange Chromatography	52
2.2	2.3	RP-HPLC (Analytical Column)	52
2.2	2,4	Mass Spectrometry	53
2.2	2.5	N-Terminal Sequencing	53
2.2	2.6	Radial Diffusion Plate Assay	54
2.2	2.7	Protein Concentration Determination	54
2.2	2.8	Minimum Inhibitory Concentration (MIC)	55
2.3	Ass	essment of Peptide Activity in Variety of Conditions	55
2.3	3.1	RP-HPLC (Semi-prep column)	55
2.3	3.2	MIC using Micro Broth Dilution	56
2.3	3.3	Monovalent cation Effects	57
2.3	3.4	Divalent Cation Effects	57
2.3	3.5	Temperature Effects	58
2,3	3.6	Analysis of Variance (ANOVA)	58
2.4	Stuc	dy of Antimicrobial Peptide Mechanisms of Action	59
2.4	4.1	LPS Binding Assay	59
2.4	4.2	Outer Membrane Permeabilisation Assay	60
2.4	4.3	Cytoplasmic Membrane Depolarisation Assay	61
2.4	4.4	Optical Density and Colony Forming Unit Measurements	62
2.4	4.5	DNA Gel Electrophoresis	62

NOT.	xtraction and Purification of Host Defence Peptides from Ostrich 65
3.1 Intro	duction
3.2 Resu	ilts and Discussion
3.2.1	Peptide Extraction
3.2.2	Peptide Purification with Cationic Exchange
3.2.3	Peptide Purification with RP-HPLC (Analytical Column)
3.2.4	Mass Spectrometry and N-terminal sequence Analysis
3.2.5	Antimicrobial activity of OSP-1-4
3.2.6	Amino Acid Composition Analysis
3.2.7	Evolutionary Analysis of the Ostricacins
3.3 Cone	elusion
- 1	vestigation of Antagonist Factors Affecting Antimicrobial Activity of and 2 on E. coli O157:H7 and S. aureus 1056MRSA83
4.1 Intro	oduction
4.2 Resu	ilts and Discussion
4.2.1	Effect of Monovalent Cation on E. coli O157:H785
4.2.2	Effect of Monovalent Cation on S. aureus 1056MRSA 86
4.2.3	Effect of Divalent Cation on E. coli O157:H789
4.2.4	Effect of Divalent Cation on S. aureus 1056MRSA
4.2.5	Effect of Temperature
4.3 Cone	clusion96
	Investigation of Mechanisms of Action of Ostricacins against ve Bacteria99
5.1 Intro	oduction99
5.2 Resu	alts and Discussion
5.2.1	LPS Binding Assay
5.2.2	Outer Membrane Permeabilisation Assay
5.2.3	Cytoplasmic Membrane Depolarisation Assay
5.2.4	Kill curve
5.2.5	DNA Gel Electrophoresis

5.3	Conclusion	113
Chapte	er 6 Conclusions and Recommendations	115
6.1	Overall Conclusions	115
6.2	Recommendations for Future Work	118
Refere	ences	121
	idix 1 Raw Data and Calculations from Investigation of Antagoniing Osp-1 and 2	
1.1	Example Calculations of MIC	133
1.2	Calculation of the Average and Confidence Intervals of MIC from Data	
1.3	Analysis of Variance (ANOVA) of MIC Data from Different Condit	ions 139
	ndix 2 Replicates Results of the Investigation of Mechanisms of acins	
2.1	Raw Data of LPS Binding Assay and Example Calculations of Fluorescence Inhibited	
2.2	Results of LPS Binding Assay	142
2.3	Results of Outer Membrane Permeabilisation Assay	144
2.4	Results of Cytoplasmic Membrane Depolarisation Assay	146
2.5	Results of Kill Curves on E. coli O111	148
Appen	idix 3 List of Publications	151

LIST OF FIGURES

Figure 1.1 Molecular Components of Innate Immune System
Figure 1.2 The Phagocytosis Mechanisms of a Phagocyte on a Microbe or a Foreign Particle
Figure 1.3 Molecular Components of Adaptive Immune System
Figure 1.4 Adaptive Immune Responses Upon Recognition of Antigens
Figure 1.5 Structural Representations of The Four-Classes Host Defence Peptides 11
Figure 1.6 Host Defence Peptides Roles in Innate and Adaptive Immunity
Figure 1.7 The three models of host defence peptides membrane disruptive mechanisms on Gram-negative membrane.
Figure 1.8 Micellar-aggregate on Gram-negative bacteria
Figure 1.9 Intracellular Targets of Host Defence Peptides in E. coli
Figure 1.10 Schematic processing of two nine-amino-acid segments of mutated α -defensins precursor into θ -defensins.
Figure 1.11 Sequence and Structure of Human Neutrophil Peptide-3 (HNP-3) and Human β-Defensins 2 (HBD-2).
Figure 1.12 Interaction of Defensins with Immune System
Figure 1.13 Evolutionary tree of existing β-defensins. 40
Figure 3.1 Microscopic Examination (400x) of Blood Cells Stained with Diff-Quick 68
Figure 3.2 Ion-Exchange Chromatograph of the Ostrich Crude Extract
Figure 3.3 RP-HPLC chromatograph of Fraction 2 of the Cationic Exchange Column. 71
Figure 3.4 Phylogenetic tree illustrating the homology of Osp-1-4 with other avian and mammalian β -defensins.
Figure 4.1 The effect of monovalent cations, Na ⁺ (A) and K ⁺ (B), on the MIC of Osp-1 and 2 against <i>E. coli</i> O157:H7
Figure 4.2 The effect of monovalent cations, Na ⁺ (A) and K ⁺ (B), on the MIC of Osp-1 and 2 against <i>S. aureus</i> 1056MRSA.
Figure 4.3 The effect of divalent cations, Mg ²⁺ (A) and Ca ²⁺ (B), on the MIC of Osp-1 and 2 against <i>E. coli</i> O157:H7
Figure 4.4 The effect of divalent cations, Mg ²⁺ (A) and Ca ²⁺ (B), on the MIC of Osp-1 and 2 against <i>S. aureus</i> 1056MRSA
Figure 4.5 The effect of temperatures on the MIC of Osp-1 and 2 against <i>E. coli</i> O157:H7 (A) and S. aureus 1056MRSA (B)

Figure 5.1 DPX/LPS binding assay of SMAP-29, HNP-1, OSP-1 and 2 indicating the affinity of the peptides to LPS
Figure 5.2 Kinetics of NPN uptake assay using <i>E. coli</i> UB 1005 that was treated with SMAP-29, HNP-1, OSP-1, OSP-2 and Polymyxin B (PXB)
Figure 5.3 Cytoplasmic membrane depolarisation that induced by SMAP-29, HNP-1 OSP-1, OSP-2 and Gramicidin
Figure 5.4 Measurements of OD _{600nm} taken in parallel with the colonies counting of OSP-1 and 2 against <i>E. coli</i> O111.
Figure 5.5 Interactions between different ratios of bacterial λDNA and the host defence peptide using DNA Gel Electrophoresis
Figure 5.6 Model illustrating the mechanism of ostricacins on Gram-negative bacteria

LIST OF TABLE

Table 1.1 Comparison of Innate and Adaptive Immune Response
Table 1.2 Comparison of Vertebrate Defensins
Table 1.3 Amino acid sequences of existing avian and mammalian β -defensins 30
Table 1.4 Antimicrobial activity of avian β-defensins
Table 2.1 Sequence and Molecular Mass of The Ostrich Antimicrobial Peptide, Osp-1 and 2
Table 2.2 Ingredients for Peptide-DNA Binding Gel Electrophoresis
Table 3.1 Plate Assay of Ion Exchange Column Fractions and Crude Extract70
Table 3.2 Plate Assay of RP-HPLC Peaks
Table 3.3 Mass Spectrometry and N-terminal Sequencing of Osp-1-4. Using FASTA, these Ostricacins were aligned with chicken gallinacins and turkey THPs73
Table 3.4 Minimum Inhibitory Concentrations (μg/ml) of Ostricacins-1-475
Table 3.5 Amino Acid Composition of Ostricacins-1-4 (mole percent)