



THE COMPLEX CHITINOLYTIC SYSTEM OF

ASPERGILLUS FUMIGATUS

by

MARIAM TAIB

Submitted in accordance with the requirements for the degree of Doctor of Philosophy

The University of Leeds School of Biochemistry and Microbiology

May 2005

The candidate confirms that the work submitted is her own and that appropriate credit has been given where reference has been made to the work of others

This copy has been supplied on the understanding that it is copyright material and that no quotation from the thesis may be published without proper acknowledgement

1100042644

Acknowledgements

I am extremely grateful to my supervisor, Dr. David Adams for the opportunity to undertake this study and for his continuous advice, support and encouragement. I have the pleasure of thanking Dr. Alex Jaques for his invaluable help and advice throughout my study. This has been a great experience that I shall never forget.

I would like to express my gratitude to the staff of the School of Biochemistry and Microbiology, especially Ms Diana Hall for the technical assistance. My gratitude extends to Dr Kenny McDowall, his lab members Jonathan Stead and Jane Towle, Prof. Steve Baldwin and his lab members, for their help with parts of the study.

I would like to thank the Government of Malaysia for the financial support.

My heartfelt thanks go to my mother and father, mother and father in-law, and my families, for their love, support and patience. I would also like to thank all my friends for their support, especially Azna, Farid and Erin.

Special thanks and love to my husband Zaba, whose love shines through his patience, support and understanding.

i

Abstract

Aspergillus fumigatus is one of the most important fungal pathogens of humans and there is an urgent need for new drugs to counter infections caused by A. fumigatus and other pathogenic species. Enzymes of chitin metabolism, and their regulators present novel targets for antifungal agents. During the work described here, the patterns of expression of the chiA1 and chiB1 chitinase genes of A. fumigatus during batch culture were investigated using real-time, reversetranscription PCR. The chiA1 gene, encoding the fungal/plant chitinase ChiA1, was expressed at significant levels throughout the six days of culture. However, the level of expression of *chiB1*, encoding the fungal/bacterial chitinase ChiB1, was only just detectable on day one but had been induced 1280-fold, to a level similar to that detected for chiA1 expression, by day 6. The results suggest markedly different roles for these enzymes. The gene encoding the transcription factor CreA was cloned and expressed, as a glutathione S-transferase (GST) fusion protein, in Escherichia coli. In electrophoretic mobility shift assays purified GST-CreA, or an A. fumigatus cell extract, bound specifically to putative CreA binding sites upstream of the chiB1 gene. CreA may therefore have a role in the regulation of chitinase activity in A. fumigatus. The effects of a range of compounds on A. fumigatus chitinase activity were determined. The cyclopentapeptides, argadin and argifin (each at 0.6 µM), were potent inhibitors of enzyme activity. The cyclic dipeptides, D-Leu-D-Pro, cyclo-(D-Leu-D-Pro) and cyclo-(L-His-L-Pro) (each at 300 µM), did not inhibit chitinase activity, while the methylxanthines, pentoxyfylline and theophylline, caused significant inhibition at concentrations of 75 µM and 300 µM, respectively. In preliminary expression studies, ChiA1 was fused with GST or maltose-binding protein (MBP) and expressed in E. coli. In addition, ChiA1-His₆ peptide was expressed in Pichia pastoris. These constructs will be used in future work which will further explore the complex chitinolytic system of A. fumigatus and which may lead to the exploitation of this system as a target for antifungals.

Contents

	Page
Acknowledgements	i
Abstract	ii
Contents	ili
Tables and Figures	xviii
Abbreviations	xxiii
Chapter 1: General Introduction	1
1.1 Human Mycoses	2
1.2 Aspergillus and Aspergillosis	3
1.3 Putative Virulence Factors of A. fumigatus	6
1.4 Fungal cell wall	8
1.4.1 Architecture of the fungal cell wall	9
1.4.2 Biosynthesis of the fungal cell wall	12
1.5 Roles for chitinases	15
1.5.1 Roles for chitinases in yeast	19
1.5.2 Roles for chitinases in filamentous fungi	20
1.6 Regulation of chitinase activity	27
1.6.1 The role of transcription factors in the regulation	30
of chitinase expression	
1.7 Antifungal Agents	33
1.8 Aims of Study	36

iii

Chapter 2: General Methods	37
2.1 Materials	38
2.2 Growth and maintenance of A. fumigatus	38
2.3 Preparation of mycelial pads of A. fumigatus	38
2.4 Agarose gel electrophoresis	39
2.5 Isolation of genomic DNA from A. fumigatus	39
2.6 General PCR reaction	40
2.7 Growth and maintenance of <i>E. coli</i>	40
2.8 Cloning of PCR products	41
2.9 Ligation	41
2.10 Transformation	41
2.11 Isolation of plasmid DNA	42
2.12 Restriction digests	42
2.13 Gel-purification of DNA	42
2.14 Protein estimation	42
2.15 Sodium dodecyl sulphate polyacrylamide gel electrophoresis	43
(SDS-PAGE)	
2.16 Coomassie blue staining of polyacrylamide gels	43
2.17 Preparation of DIG-labelled probes	44
2.18 Southern Blotting	44
2.19 Western blotting	45
2.20 Micro-fluorescence assay for chitinase activity	46
2.21 Glycol chitin gel electrophoresis	47

iv

Chapter 3. Expression of Chitinase Genes of A. fumigatus	49
3.1 Introduction	50
3.2 Materials and Methods	54
3.2.1 Real-time reverse transcription PCR analysis of	54
A. fumigatus chitinase gene expression	
3.2.1.1 Induction of chitinase expression under conditions	54
of starvation	
3.2.1.2 Total RNA isolation	54
3.2.1.3 DNase I treatment	54
3.2.1.4 Quality control of total RNA by standard PCR	55
amplification and reverse transcription PCR	
amplification (RT-PCR)	
3.2.1.5 Reverse transcription of total RNA	56
3.2.1.6 Quantitative assay of chitinase expression by	56
Real-time PCR	
3.2.1.7 Determination of dry weight of A. fumigatus	57
and microfluorescence chitinase assay	
3.2.2 Green Fluorescent Protein as a Reporter for Chitinase	58
Gene Expression in A. fumigatus	
3.2.2.1 Amplification of the promoter and N-terminal	58
domains of chiA1 or chiB1 by polymerase chain	
reaction (PCR)	
3.2.2.2 Ligation and transformation of competent E. coli	60
JM109 cells	

v

vi	
3.2.2.3 Sub-cloning of the <i>chiA1</i> or <i>chiB1</i> insert into the	60
pUCGH expression vector	
3.2.2.4 Transformation of A. fumigatus with	61
recombinant plasmids pUCGH-chiA1	
or pUCGH- <i>chiB1</i>	
3.2.2.4.1 Protoplast preparation	61
3.2.2.4.2 Transformation using protoplasts	62
3.2.2.5 Southern blotting and hybridization analysis	63
using DIG-labelled DNA probes	
3.2.2.5.1 Preparation of DIG-labelled	63
DNA probes	
3.2.2.5.2 Isolation of chromosomal DNA from	63
A. fumigatus transformants and	
digestion with restriction enzymes	
3.2.2.5.3 Southern blotting and hybridization	64
with DIG-labelled probes	
3.2.2.6 Expression of the chitinase-green fluorescent	64
protein construct and fluorescence microscopy	
analyses	
3.2.2.6.1 Induction of chitinase-GFP expression	64
under conditions of starvation	
3.2.2.6.2 Fluorescence and light microscopy	65
analyses	

3.3 Results	66
3.3.1 Real-time reverse transcription PCR analysis of	66
A. fumigatus chitinase gene expression	
3.3.1.1 Determining the quality of A. fumigatus RNA	66
preparations prior to real-time RT-PCR analysis	
3.3.1.2 Establishing the reaction conditions for real-time	68
PCR analysis	
3.3.1.3 Analysis and quantification of <i>chiA1</i> and <i>chiB1</i>	69
mRNA transcript levels during conditions	
of starvation	
3.3.2 Green Fluorescent Protein as a Reporter for Chitinase	77
Gene Expression in A. fumigatus	
3.3.2.1 Amplification of the chiA1 or chiB1	77
gene fragments	
3.3.2.2 Ligation of the PCR product into pGEM-T and	77
transformation of <i>E. coli</i> JM109	
3.3.2.3 Sub-cloning of the chiA1 or chiB1 insert into	79
pUCGH, transformation of E. coli JM109 and	
transformation of A. fumigatus	
3.3.2.4 Amplification of DIG-labelled DNA probes	79
3.3.2.5 Southern blot analysis of A. fumigatus	79
transformants	
3.3.2.6 Fluorescence and light microscopy analyses of	82
chiB-egfp gene expression under conditions of	
starvation	
	96
3.4 Discussion	86

vii

Chapter 4.0: Heterologous expression of the gene encoding the 90 A. fumigatus chitinase ChiA1

4.1 Introduction		91
4.2 Materials and Methods		95
4.2.1 Expression of A	. <i>fumigatus chiA1</i> gene in	95
Pichia pastoris		
4.2.1.1 Recomb	inant P. pastoris strains	95
4.2.1.2 Induction	n of heterologous gene expression	95
4.2.1.3 Analysis	of heterologously-expressed gene	96
products	3	
4.2.2 Expression of A	<i>fumigatus chiA1</i> gene in	96
Escherichia coli		
4.2.2.1 Express	on of <i>A. fumigatus</i> ChiA1 as	96
MBP-fus	sion protein	
4.2.2.1.1	Polymerase chain reaction	96
4.2.2.1.2	Cloning of the PCR product using	97
	pGEM-T, and transformation of	
	competent <i>E. coli</i> JM109 cells	
4.2.2.1.3	Sub-cloning of the cloned PCR	98
	product using pMAL-c2x	
	and pMAL-p2x expression vectors	
	4.2.2.1.3.1 Preparation of insert	98
	4.2.2.1.3.2 Preparation of vector	98
	4.2.2.1.3.3 Ligation of isolated insert	99

viii

			into pMAL expression	
			vectors, transformation of	
			E. coli JM109 and	
			screening of colonies	
	4.2.2.1.4	Expression	of the MBP-ChiA1 fusion	99
		protein in E	E. coli BL21(DE3)pLysS	
		4.2.2.1.4.1	Induction of heterologous	99
			expression	
		4.2.2.1.4.2	Analysis of recombinant	100
			MBP-ChiA1 protein	
			expression	
		4.2.2.1.4.3	Purification of recombinant	101
			MBP-ChiA1 fusion protein	
		4.2.2.1.4.4	Cleavage of MBP-ChiA1	102
			fusion protein by Factor Xa	
			protease	
	4.2.2.1.5	Expression	of the MBP-ChiA1	103
		fusion prote	ein in <i>E. coli</i>	
		BL21-Code	onPlus(DE3)-RP	
	4.2.2.1.6	Expression	of the MBP-ChiA1	103
		fusion prote	ein in <i>E. coli</i>	
		Origami B((DE3)pLysS	
4.2.2.2	Expression	on of A. fumi	<i>igatus</i> ChiA1 as	104
	GST-fusio	on protein		
	4.2.2.2.1	Polymerase	chain reaction	104
	4.2.2.2.2	Cloning of th	ne PCR product	105

ix

using pGEM-T, subcloning using	
pGEX-4T-3 expression vector,	
and transformation of E. coli	
JM109 competent cells	
4.2.2.2.3 Expression of the GST-ChiA1fusion	106
protein in <i>E. coii</i> Origami B(DE3)pLysS	
4.2.2.3.1 Analysis of recombinant	106
GST-ChiA1 protein	
expression	
4.2.2.3.2 Purification of recombinant	107
GST-ChiA1 fusion protein	
4.3 Results	108
4.3.1 Expression of A. fumigatus chiA1 gene in P. pastoris	108
4.3.2 Expression of A. fumigatus chiA1 gene in E. coli	108
4.3.2.1 Expression of A. fumigatus ChiA1 as	108
MBP-fusion protein	
4.3.2.1.1 Amplification of the chiA1 gene	108
4.3.2.1.2 Ligation of the <i>chiA1</i> PCR product	111
into pGEM-T, and transformation	
of <i>E. coii</i> JM109	
4.3.2.1.3 Sub-cloning of <i>chiA1</i> gene using	111
pMAL-c2x or pMAL-p2X, and	
transformation of E. coii JM109	
4.3.2.1.4 Expression of MBP-ChiA1	114
fusion protein in <i>E. coli</i> BL21	
(DE3)pLysS	

х

4.3.2.1.4.1 Purification of MBP-ChiA1	
fusion protein	114
4.3.2.1.4.2 Cleavage of MBP-ChiA1	117
fusion protein by	
Factor Xa protease	
4.3.2.1.5 Expression of chiA1 in E. coli	117
BL21-CodonPlus(DE3)-RP	
4.3.2.1.5.1 Purification of MBP-ChiA1	120
fusion protein	
4.3.2.1.5.2 Cleavage of MBP-ChiA1	120
fusion protein by	
Factor Xa protease	
4.3.2.1.6 Expression of chiA1 in E. coli	123
Origami B(DE3)pLysS	
4.3.2.1.6.1 Purification of MBP-ChiA1	123
fusion protein	
4.3.2.2 Expression of A. fumigatus ChiA1 as	126
GST-fusion protein	
4.3.2.2.1 Amplification of the chiA1 gene	126
4.3.2.2.2 Ligation of the chiA1 gene	126
PCR product into pGEM-T,	
and transformation of E. coli JM109	
4.3.2.2.3 Sub-cloning of chiA1 gene using	127
pGEX-4T-3, and transformation	
of <i>E. coli</i> JM109	
4.3.2.2.4 Expression of GST-ChiA1 fusion	127

xi

xii	
protein in <i>E. coii</i> Origami	
B(DE3) pLysS	
4.3.2.2.5 Purification of GST-ChiA1	129
fusion protein	
4.4 Discussion	132
ter 5.0: The Transcriptional Regulator, CreA	136
5.1 Introduction	137
5.2 Materials and Methods	14
5.2.1 Cloning and heterologous expression of	14
the A. fumigatus transcriptional regulator CreA	
5.2.1.1 Expression of A. fumigatus creA gene	14
in <i>Pichia pastoris</i>	
5.2.1.1.1 Polymerase chain reaction	141
5.2.1.1.2 Cloning of the PCR product	142
using pGEM-T, and transformation	
of E. coli JM109 competent cells	
5.2.1.1.3 Sub-cloning of the cloned	142
PCR product using pPICZ A	
expression vector	
5.2.1.1.3.1 Preparation of insert	142
5.2.1.1.3.2 Preparation of expression	143
vector	
5.2.1.1.3.3 Ligation of isolated insert	143
into pPICZ A expression	

<i>E. coli</i> JM109 and	
screening of colonies	
5.2.1.1.4 Transformation of <i>P. pastoris</i>	144
5.2.1.1.4.1 Preparation of	144
recombinant plasmid	
pPICZ A/creA	
5.2.1.1.4.2 Preparation of P. pastoris	144
strain X-33	
5.2.1.1.4.3 Transformation of	145
P. pastoris with	
recombinant plasmid DNA	
5.2.1.1.5 Isolation of genomic DNA from	146
transformed P. pastoris cells and	
PCR analysis	
5.2.1.1.6 Induction of heterologous gene	146
expression	
5.2.1.1.7 Analysis of recombinant	146
creA expression	
5.2.1.2 Expression of A. fumigatus creA gene as	147
GST-CreA protein in Escherichia coli	
5.2.1.2.1 Polymerase chain reaction	147
5.2.1.2.2 Cloning of the PCR product using	147
pGEM-T, sub-cloning using	
pGEX-4T-3 expression vector,	
and transformation of E. coli JM109	
competent cells	

xiii

xiv	
5.2.1.2.3 Expression of the GST-CreA fusion	148
protein in <i>E. coii</i> BL21 (DE3)plysS	
5.2.1.2.4 Analysis and purification of GST-CreA	148
protein	
5.2.2 Binding of native and recombinant CreA to sites	149
upstream of A. fumigatus chitinase genes	
5.2.2.1 Amplification of regions upstream regions of	149
the chiA1 and chiB1 genes using the	
polymerase chain reaction (PCR)	
5.2.2.1.1 Design of primers	149
5.2.2.1.2 PCR and cloning of the	149
reaction products	
5.2.2.2 Putative CreA binding sites upstream of chiB1	150
5.2.2.3 Preparation of A. fumigatus cytosolic extract	151
5.2.2.4 Electrophoretic Mobility Shift Assay (EMSA)	152
5.2.2.4.1 DIG Gel Shift	152
5.2.2.4.2 Fluorescein Method	153
5.2.3 Polymerase chain reaction and cloning of the	154
A. fumigatus transcriptional regulator ACE2 gene	
5.2.3.1 PCR amplification	154
5.2.3.2 Cloning of the PCR product using pGEM-T	155
and transformation of <i>E. coli</i> JM109	
competent cells	
3 Results	156
5.3.1 Cloning and heterologous expression of the	156
A. fumigatus creA gene	

5.

	XV	
5	3.1.1 Expression of A. fumigatus creA gene	156
	in Pichia pastoris	
	5.3.1.1.1 Polymerase chain reaction	156
	5.3.1.1.2 Cloning of the PCR product using	156
	pGEM-T, and transformation of	
	E. coli JM109 competent cells	
	5.3.1.1.3 Sub-cloning of the creA gene using	159
	pPICZ A expression vector	
	5.3.1.1.4 Isolation of genomic DNA from	159
	transformed P. pastoris cells and	
	PCR analysis	
	5.3.1.1.5 Expression of CreA in <i>P. pastoris</i>	163
5.	3.1.2 Expression of <i>A. fumigatus creA</i> gene as	163
	GST-CreA fusion protein in E. coli	
	5.3.1.2.1 Polymerase chain reaction	163
	5.3.1.2.2 Cloning of the PCR product using	163
	pGEM-T and transformation of	
	E. coli BL21 (DE3)plysS	
	competent cells	
	5.3.1.2.3 Sub-cloning of creA gene using	166
	pGEX-4T-3 and transformation of	
	E. coli JM109 competent cells	
	5.3.1.2.4 Expression of GST-CreA fusion	166
	protein in <i>E. coli</i> BL21 (DE3)plysS	

XVI	
5.3.2 Binding of native and recombinant CreA to sites	170
upstream of A. fumigatus chitinase gene	
5.3.2.1 PCR for <i>chiA1</i> and <i>chiB1</i> upstream regions,	170
and cloning of the reaction products	
5.3.2.2 Electrophoretic Mobility Shift Assay (EMSA)	173
5.3.2.2.1 DIG Gel Shift	173
5.3.2.2.1.1 EMSA with putative CreA	173
binding sites and cytosolic	
extract	
5.3.2.2.1.2 EMSA with putative	173
CreA binding sites and	
recombinant CreA	
5.3.2.2.2 Fluorescein Method	176
5.3.2.2.1 EMSA with putative CreA	176
binding sites and cytosolic	
extract	
5.3.2.2.2 EMSA with putative CreA	176
binding sites and	
recombinant CreA	
5.3.3 Amplification and cloning of an apparent ACE2 gene	177
homologue from A. fumigatus	
5.3.3.1 PCR Amplification	177
5.3.3.2 Cloning of the PCR product into pGEM-T	177
and transformation of E. coli JM109	
5.4 Discussion	183

XVII	
Chapter 6: Inhibition of Extracellular Chitinase Activity	186
6.1 Introduction	187
6.2 Materials and Methods	190
6.2.1 Test compounds	190
6.2.2 NCCLS Microdilution Broth Method	190
6.2.2.1 Preparation of medium	190
6.2.2.2 Preparation and dilution of stock solutions	190
6.2.2.3 Preparation of A. fumigatus inoculum	191
6.2.2.4 Inoculation of the microtitre plates and	191
plate reading	
6.2.3 Micro-fluorescence assay for chitinase activity	192
6.3 Results	193
6.3.1 Microdilution Broth Method	193
6.3.2 Microfluorescence Assay	193
6.3.2.1 Enzyme kinetics	193
6.3.2.2 Effect of test compounds on A. fumigatus	193
chitinase activity	
6.4 Discussion	199
Chapter 7: Final Discussion	201
Chapter 8: References	208

Tables and Figures

xviii

	Page
Tables	
Table 3.1 The primer pairs used for PCR amplification of a 180-bp fragment of	54
chiA1 and chiB1	
Table 5.1 CreA binding-sites located upstream of the A. nidulans creA	150
regulatory domain and the putative CreA binding sites located upstream of the	
A. <i>fumigatus chiB1</i> chitinase gene	

Figures

Figure 1.1 Sporulating conidiophores of <i>A. fumigatus</i>	
Figure 1.2 Schematic diagram of the architecture of the fungal cell wall	10
Figure 1.3 Comparison of chemical structures of the fungal polysaccharide cell	13
wall core	
Figure 1.4 Putative successive enzymatic events involved in the biosynthesis of	14
the fungal wall	
Figure 1.5 Mechanisms of action of exochitinase and endochitinase	16
Figure 1.6 Conserved amino acids (in bold) of (A) Family 18 chitinases and (B)	18
Family 19 chitinases	
Figure 1.7 Multiple alignment of the apparent active site from the 14 (2 known	23
and 12 hypothetical) chitinase sequences	
Figure 1.8 Phylogenetic tree generated from a multiple alignment of 209	24
residues surrounding the apparent active site in the genes found	
Figure 1.9 Comparison of the domains of the ChiB1 fungal/bacterial chitinase	26

and ChiA1 fungal/plant chitinase of A. fumigatus	
Figure 1.10 Mechanisms of action of caspofungin, fluconazole and voriconazole	34
Figure 2.1 Standard curve generated for micro-fluorescence assay for chitinase	47
activity	
Figure 3.1 Amplification plots of IL-4 plasmid cDNA	51
Figure 3.2 Expression plasmid pUCGH	59
Figure 3.3 Monitoring the quality of total A. fumigatus RNA	67
Figure 3.4 Establishing the reaction conditions for real time PCR analysis for	70
real-time PCR analysis of the chiA1 and chiB1 genes	
Figure 3.5 Fluorescence profiles of the chiA1 and chiB1 amplicons	72
Figure 3.6 Fluorescence profiles of the pGEMT chiA1 and pGEMT chiB1	73
amplicons	
Figure 3.7 Standard curves for pGEMT chiA1 and pGEMT chiB1	75
Figure 3.8 Expression profiles for <i>chiA1</i> and <i>chiB1</i> in <i>A. fumigatus</i> during batch	76
culture	
Figure 3.9 Amplification of the promoter and N-terminal domains of	78
A. fumigatus	
Figure 3.10 Restriction digests for positive clones containing the 760 bp	80
fragment of <i>chiA1</i> or <i>chiB1</i> in pUCGH	
Figure 3.11 Amplification of DIG-labelled 760 bp fragment of	81
A. fumigatus chiA1 and chiB1	
Figure 3.12 Identification of an A. fumigatus transformant carrying the PchiB1-	83
egfp fusion integrated in a single copy at the chiB1 gene locus	
Figure 3.13 Bright light (a, e, i, c, g, k) and fluorescence (b, f, j, d, h, l)	85
microscopy for wild-type A. fumigatus and apparent A. fumigatus PchiB1-egfp	
transformant (lane 3, Figure 3.12) grown under batch culture conditions	

xix

Figure 4.1 Glycol chitin zymography analysis of ChiA1 (full length protein with	109
histidine tag) expressed in <i>P. pastoris</i>	
Figure 4.2 Amplification of the A. fumifatus chiA1 gene	110
Figure 4.3 Restriction digest of the pGEM-T Easy vector from a 'positive' colony	112
of <i>E. coli</i> JM109	
Figure 4.4 Restriction digests for positive clones, containing the 2.5 kb PCR	113
product in (a) pMAL-c2x and (b) pMAL-p2x, to determine the orientation of the	
insert in the plasmid	
Figure 4.5 Analysis of expression of the MBP-ChiA1 fusion protein in	115
E. coli BL21(DE3)pLysS cells	
Figure 4.6 SDS-PAGE analysis following fractionation of the E. coli	116
BL21(DE3)pLysS cell lysate using amylose affinity chromatography	
Figure 4.7 SDS-PAGE analysis of MBP-ChiA1 after incubation with Factor Xa	118
Figure 4.8 Analysis of expression of the MBP-ChiA1 fusion protein in	119
E. coli BL21-CodonPlus(DE3)-RP	
Figure 4.9 SDS-PAGE analysis following fractionation of the E. coli BL21-	121
CodonPlus(DE3)-RP cell lysate using amylose affinity chromatography	
Figure 4.10 SDS-PAGE analysis of MBP-ChiA1 after incubation with Factor Xa	122
Figure 4.11 Analysis of expression of the MBP-ChiA1 fusion protein in	124
E. coli Origami B(DE3)pLysS	
Figure 4.12 SDS-PAGE analysis following fractionation of the E. coli Origami	125
B(DE3)pLysS cell lysate using amylose affinity chromatography	
Figure 4.13 Restriction digests for positive clones (lanes 2-6), containing the 2.5	128
kb PCR product in pGEX-4T-3 to determine the orientation of the insert in the	
plasmid	
Figure 4.14 Analysis of expression of the GST-ChiA1 fusion protein in E. coli	130

XX

Origami B(DE3)pLysS	
Figure 4.15 SDS-PAGE and glycol-chitin SDS-PAGE zymography analyses	131
following purification of the E. coli Origami B cell lysate using Glutathione	
Sepharose 4B affinity chromatography	
Figure 5.1 Alignment of the amino acid sequence of A. fumigatus CreA with the	ə 139
amino acid sequence of A. nidulans CreA	
Figure 5.2 Amplification of the creA gene for expression in P. pastoris	157
Figure 5.3 Restriction digests for positive clones of the apparent creA PCR	158
product in pGEM-T	
Figure 5.4 Restriction digests for positive clones containing the 1.3 kb PCR	160
product in pPICZ A	
Figure 5.5 Restriction digests for positive clones containing the 1.3 kb PCR	161
product in pPICZ A to determine the orientation of the insert in the plasmid	
Figure 5.6 Agarose gel electrophoresis following the integration of creA into the	e 162
P. pastoris genome	
Figure 5.7 Amplification of the creA gene for expression in E. coii	164
BL21(DE3)pLysS	
Figure 5.8 Restriction digests for positive clones of the PCR product of the creation digests for positive clones of the posit	A 165
gene	
Figure 5.9 Restriction digests for positive clones containing the 1.3 kb PCR	167
product in pGEX-4T-3	
Figure 5.10 Restriction digests for positive clones containing the 1.3 kb PCR	168
product in pGEX-4T-3 to determine the orientation of the insert in the plasmid	
Figure 5.11 SDS-PAGE analysis of the GST-CreA fusion protein following	169
expression in E. coli BL21 and purification using Glutathione Sepharose 4B	
affinity chromatography	

xxi

Figure 5.12 Amplification of the regions upstream of A. fumigatus chiA1 and	171
chiB1 genes	
Figure 5.13 Restriction digests for positive clones of PCR products of chitinase	172
gene upstream regions	
Figure 5.14 EMSA with putative CreA binding sites and cytosolic extract using	174
DIG gel shift kit	
Figure 5.15 EMSA with putative CreA binding site 2 upstream of the	175
A. fumigatus chiB1 gene, and recombinant GST-CreA using the DIG gel shift kit	
Figure 5.16 EMSA with A. fumigatus cytosolic extract	178
Figure 5.17 EMSAs with recombinant A. fumigatus creA: titration of binding shift	179
using increasing amounts of unlabelled specific and non-specific competitors	
Figure 5.18 Amplification of the apparent ACE2 gene	181
Figure 5.19 Restriction digests for positive clones containing the apparent	182
ACE2 PCR product in pGEM-T	
Figure 6.1 Allosamidin	187
Figure 6.2 Cyclopentapeptides	188
Figure 6.3 Cyclic dipeptides	189
Figure 6.4 Methylxanthines	189
Figure 6.5 A. fumigatus extracellular chitinase activity with time	194
Figure 6.6 Effect of cyclopentapeptides on chitinase activity	195
Figure 6.7 Effect of allosamidin on chitinase activity	196
Figure 6.8 Effect of methylxanthines on chitinase activity	198

xxii

Abbreviations

The following abbreviations are used:

ATCC	American Type Culture Collection
ddH₂O	Double distilled water
DIG	Digoxygenin
DMSO	Dimethylsulfoxide
DTT	Dithiothreitol
EDTA	Diaminoethane tetraacetic acid, disodium salt
EMSA	Electrophoretic Mobility Shift Assay
GIcNAc	N-acetylglucosamine
GST	Glutathione S-transferase
HEPES	N-[2-Hydroxyethyl]piperazine-N'-(2-ethylsulfonic acid)
IPTG	Isopropyl β-D thiogalactoside
LB	Luria-Bertani
MBP	Maltose-binding protein
MMS	Minimal Medium with 1 M sucrose
MOPS	3-(N-morpholino)propanesulfonic acid
MU	Methylumbelliferone
NBT/BCIP	Nitro blue tetrazolium chloride / 5-Bromo-4-chloro-3-indoyl phosphate
NCCLS	The U.S National Committee of Clinical Lab Standard
NTP	Nucleoside Triphosphates
PAGE	Del com demide del electronic
	Polyacrylamide gel electrophoresis
PBS	Polyacrylamide gel electrophoresis Phosphate-buffered saline

xxiii

xxiv

ŝ

PMSF	Phenylmethylsulphonylfluoride
rpm	Revolutions per minute
SDS	Sodium dodecyl sulphate
TBS	Tris-buffered saline
Tris	Tris(hydroxymethyl)aminoethane
TTBS	Tween Tris-buffered saline
YNB	Yeast Nitrogen Base with ammonium sulfate without amino acids
YPD	Yeast Peptone Dextrose

1.1 Harrish Mycones

Chapter 1

General Introduction

-

1.1 Human Mycoses

Fungi can cause a variety of diseases of humans and other animals ranging from minor superficial skin and mucous membrane infections to life-threatening, systemic involvement of the internal organs. There are three major types of disease caused by fungi: allergies, poisonings and fungal infections (Schaechter *et al.*, 1998). Allergic reactions to fungi are caused by sensitivity to fungal proteins, such as those present in inhaled fungal spores, while poisonings result from the ingestion of fungal toxins in contaminated food or poisonous mushrooms. Fungal allergies and poisonings are important concerns in agriculture and other industries where fungal contamination is common. Fungal infections, or mycoses, result from the invasion of living tissue by a fungus and they represent the most common form of fungal disease (http://www.mercksharpedohme.com).

There are more than 100,000 recognised species of fungi, but only about 400 are known to infect humans (de Hoog & Guarro, 1995). Some species of fungi, called primary pathogens, cause disease regardless of the individual's state of health. Other species that infect individuals with a weakened immune system are described opportunistic pathogens. However, as among patients with severe immunosuppression, almost any type of fungus can exhibit pathogenic potential which may have devastating consequences for the host (Dunn, 2000). The most common fungal infections in immunocompromised hosts are candidiasis, aspergillosis, cryptococcosis, mucormycoses (zygomycoses) and Pneumocystis carinii pneumonia (http://merckmedicus.com).

2