

09 PH
A2112



**Characterisation of recombinant
hyaluronidase-1 and -3, and of
hyaluronan turnover in mineralising
osteoblasts**

By

Julian Robert James Adams BSc (Hons)

Thesis submitted for the degree of

Doctor of Philosophy

in

Department of Paediatrics

School of Medicine

Faculty of Health Science

University of Adelaide

December, 2006

CONTENTS

Declaration	i
Acknowledgments	ii
Abbreviations	iii
Abstract	xi
Chapter 1: Introduction	1
1.1 Glycosaminoglycans	2
1.1.1 Glycosaminoglycan structure and function	2
1.1.2 Biosynthesis of glycosaminoglycans	2
1.1.3 Glycosaminoglycan degradation	5
1.2 Hyaluronidases	6
1.2.1 An introduction to hyaluronidases	6
1.2.2 Human hyaluronidases	9
1.2.2.1 <i>PH20</i>	9
1.2.2.2 <i>Hyaluronidase-1</i>	10
1.2.2.3 <i>Hyaluronidase-2</i>	13
1.2.2.4 <i>Hyaluronidase-3 and hyaluronidase-4</i>	14
1.2.2.5 <i>Hyaluronidase pseudogene-1</i>	15
1.2.2.6 <i>Meningioma hyaluronidase</i>	15
1.3 Hyaluronan synthases	18
1.3.1 Introduction to hyaluronan synthases	18
1.3.2 Hyaluronan synthase-1	19
1.3.3 Hyaluronan synthase-2	21
1.3.4 Hyaluronan synthase-3	22
1.3.5 Function of multiple hyaluronidase synthases	22
1.4 Osteoblast biology	23
1.4.1 Introduction to bone	23
1.4.2 Bone formation	23
1.4.3 Matrix mineralisation	24
1.4.4 Role of proteoglycans and glycosaminoglycan in mineralisation	25
1.4.5 Role of small leucine-rich proteoglycans in mineralisation	26
1.4.6 Role of hyaluronan in mineralisation	27
1.5 Aims	28
Chapter 2 Methods	29
2.1 Molecular Biology	30
2.1.1 Synthesis of DNA oligonucleotides	30
2.1.2 Standard cloning PCR protocol	30
2.1.3 Restriction digest	30
2.1.4 Agarose electrophoresis	30
2.1.5 Purification of DNA from agarose gel	31
2.1.6 Preparation of electro-competent cells	31
2.1.7 Bacterial transformation	31
2.1.8 Plasmid preparation	32

2.1.8.1 Mini plasmid preparation	32
2.1.8.2 Midi plasmid preparation	32
2.1.9 TOPO cloning	33
2.1.10 DNA ligation	33
2.1.11 Automated DNA sequencing	33
2.1.11.1 Sequencing primers	34
2.1.12 Site-directed mutagenesis	35
2.1.12.1 Human hyaluronidase-3 His ₆ tagged vector mutagenesis primers	35
2.2 Tissue culture	35
2.2.1 Standard cell culture technique	35
2.2.2 Cell number determination	36
2.2.3 Transient mammalian transfection	36
2.2.4 Fugene transfection	37
2.2.4.1 Stable fugene transfection	37
2.2.5 Immunohistochemical localisation of recombinant protein	38
2.2.6 Intra-cellular localisation of recombinant protein in live cells	38
2.2.6.1 Endoplasmic reticulum staining	39
2.2.6.2 Golgi staining	39
2.2.6.3 Mitochondrial staining	39
2.2.6.4 Lysosomal staining	40
2.2.7 Recombinant protein production	40
2.3 Protein biochemistry methods	40
2.3.1 TNT reaction	40
2.3.2 Talon chelation ion affinity protein purification	41
2.3.3 Bradford protein assay	41
2.3.4 SDS polyacrylamide electrophoresis	42
2.3.4.1 Running gel	42
2.3.4.2 Transfer gel	42
2.3.4.3 Western blot anti-body detection of protein	42
2.3.5 Coomassie-stained SDS-PAGE	43
2.3.6 N-Glycosidase F digestion	43
2.4 Carbohydrate biochemistry methods	44
2.4.1 Hyaluronidase zymography	44
2.4.2 ELISA-based hyaluronidase assay	44
2.4.2.1 Preparation of biotinylated hyaluronan	44
2.4.2.2 Immobilisation of biotinylated hyaluronan onto ELISA plate	45
2.4.2.3 Assay for hyaluronidase activity	45
2.4.3 Substrate gel assay	46
2.4.3.1 Agarose gel analysis of hyaluronidase digests	46
2.4.4 Gradient-PAGE analysis of glycosaminoglycan digests	46
2.4.5 Modified hyaluronidase assay	47
2.4.6 Q-Sepharose anion exchange chromatography	47
2.4.7 Hydroxydiphenyl estimation of uronic acid	48
2.4.8 Extraction of glycosaminoglycan from mouse tissues	48
2.5 Osteoblast methods	49
2.5.1 Mineralising osteoblast tissue culture	49
2.5.2 von Kossa staining	49
2.5.3 Radio-isotope incorporation into glycosaminoglycan	50
2.5.4 Characterisation of glycosaminoglycan type	50
2.6 RNA methods	51
2.6.1 RNA harvest from osteoblasts	51
2.6.2 RNA MOPS gel	52

2.6.3 RNA quantification	53
2.6.4 cDNA synthesis	53
2.6.5. Standard Reverse transcription-polymerisation chain reaction	53
2.6.5.1 Reverse transcription-polymerisation chain reaction primers	53
2.6.6 Real time reverse transcription polymerisation chain reaction	54
2.6.6.1 Real time reverse transcription-polymerisation chain reaction primers	56
Chapter 3 Bioinformatic analysis of the hyaluronidases	56
3.1 Introduction	57
3.2 Hyaluronidase gene and protein sequence comparisons	58
3.3 Analysis of protein sequences	64
3.3.1 Conserved protein motifs	64
3.3.2 Determining molecular mass of hyaluronidases	61
3.3.3 Prediction of signal peptide cleavage of hyaluronidases	61
3.3.4 Prediction of GPI anchor for hyaluronidases	63
3.3.5 Alignments of hyaluronidase	
3.4 Phylogenetic analysis of hyaluronidases	65
3.5 Discussion	66
3.5.1 Conservation of amino acids in the hyaluronidases across species	68
3.5.2 Signal peptide in mammalian hyaluronidases	70
3.5.3 GPI anchors in mammalian hyaluronidases	70
3.5.4 Evolutionary analysis of hyaluronidases	71
3.5.5 Conclusion	71
Chapter 4: Characterisation of recombinant hyaluronidase-1 and -3	73
4.1 Introduction	74
4.2 Construction of recombinant hyaluronidase expression vectors	76
4.2.1 Human hyaluronidase-3 non-tagged vector construction	76
4.2.2 Hyaluronidase-3-His ₆ expression vector	77
4.2.3 Hyaluronidase-1-His ₆ expression vector	78
4.3 Cell-free translation of hyaluronidase-3-His₆ and hyaluronidase-1-His₆	79
4.4 Transient expression of hyaluronidase-3-His₆ and hyaluronidase-1-His₆	80
4.5 Stable expression of recombinant hyaluronidase-3-His₆	85
4.5.1 Isolation of a stable hyaluronidase-3-His ₆ -expressing cell line	85
4.6 Intra-cellular localisation of recombinant hyaluronidase-3-GFP	87
4.7 Stable expression of recombinant hyaluronidase-1-His₆	89
4.8 Production and purification of recombinant hyaluronidase-3-His₆ and hyaluronidase -1-His₆ proteins	89
4.9 Glycosylation of recombinant hyaluronidase-1-His₆ and hyaluronidase-3-His₆	91
4.10 Glycohydrolase activity of recombinant hyaluronidase-1-His₆ and hyaluronidase-3-His₆	94
4.10.1 Activity of recombinant hyaluronidase-3-His ₆ and hyaluronidase-1-His ₆ toward hyaluronan	94
4.10.2 Hyaluronidase activity assay with additives	95
4.10.3 Hyaluronidase activity toward other glycosaminoglycans	97
4.11 Analysis of non-tagged recombinant hyaluronidase-3	99
4.11.1 TNT protein production of non-tagged hyaluronidase-3	99
4.11.2 Transient expression of non-tagged hyaluronidase-3	99
4.11.3 Stable expression of recombinant hyaluronidase-3 non-tagged in COS-7 cells	102

4.12 Analysis of glycosaminoglycans in hyaluronidase-3 knock-out mouse tissues	106
4.13 Discussion	113
4.13.1 Production of recombinant protein	108
4.13.1.1 <i>Cell-free expression of hyaluronidase-1 and hyaluronidase-3</i>	110
4.13.1.2 <i>Transient expression of recombinant hyaluronidase-1 and hyaluronidase-3</i>	110
4.13.1.3 <i>Stable expression of recombinant hyaluronidase-1 and hyaluronidase-3</i>	111
4.13.2 Activity of recombinant hyaluronidase-1.	112
4.13.3 Activity of recombinant hyaluronidase-3	113
4.13.3.1 <i>Protein expression</i>	113
4.13.3.2 <i>Protein purification</i>	115
4.13.3.2.1 His ₆ tag interference	115
4.13.3.2.2 Talon purification column	115
4.13.3.3 <i>Enzymatic assay conditions</i>	116
4.13.4 Glycosaminoglycan accumulation in the hyaluronidase-3 knock-out mouse	117
4.13.5 Conclusions	117
Chapter 5: Hyaluronan metabolism in mineralising osteoblasts	119
5.1 Introduction	120
5.2 Histology of mineralising osteoblasts	121
5.3 RT-PCR in mineralising osteoblasts	121
5.4 Real time reverse transcription-PCR in mineralising osteoblasts	123
5.6 Total glycosaminoglycan synthesis in mineralising MG63 osteoblast cultures	128
5.7 Discussion	129
5.7.1 HAS during mineralisation	131
5.7.2 Hyaluronidase during mineralisation	131
5.7.3 Changes in glycosaminoglycan macromolecules during mineralisation	133
5.7.4 Changes in non-glycosaminoglycan macromolecules during mineralisation	133
5.7.5 Role of hyaluronan in the mineralising matrix	134
5.7.6 Model for the changes in the mineralising osteoblast extra-cellular matrix	134
5.8 Conclusion	135
Chapter 6: Conclusions and future directions	137
6.1 Bioinformatic predictions of recombinant hyaluronidase	138
6.2 Possible functions for hyaluronidase-3	139
6.3 Function of hyaluronidase-1	138
6.4 Glycosaminoglycan turnover in the mineralising matrix of osteoblasts	141
6.5 Conclusions and future directions	143
Chapter 7: References	145
7.1 Publications resulting from this work	146
7.2 References	147
Appendix I: Materials	183
I.i Tissue Culture materials	184
I.ii Molecular biology materials	184
I.iv Carbohydrate biochemistry materials	186
Appendix II: GeneBank sequences	188

Abstract

The mammalian hyaluronidases (HYALs) represent a family of enzymes that can degrade hyaluronan (HA). This thesis examines the properties of hyaluronidase-1 (HYAL-1) and hyaluronidase-3 (HYAL-3), as well as the production of hyaluronan and the expression of HYAL and hyaluronan synthases (HASs) in mineralising osteoblasts.

Recombinant hyaluronidase-1 (rHYAL-1) has a mass of 57 kDa, of which 10 kDa is due to glycosylation and 47 kDa is primary protein translation product. rHYAL-1 was shown to not only degrade HA, but also to function as an endo-glucosaminidase in the degradation of the sulphated gags chondroitin sulphate and dermatan sulphate. ✕

Recombinant hyaluronidase-3 (rHYAL-3) has a mass of 46 kDa, of which 9 kDa is due to glycosylation and 37 kDa is primary protein translation product. Immunofluorescence analysis localised His-tagged rHYAL-3 to the endoplasmic reticulum and lysosomes. *In vitro* activity assays demonstrated that HYAL-3 showed no glycohydrolase activity against any glycosaminoglycan (gag) substrate tested. However, the HYAL-3 knock-out mouse (*hyal-3^{-/-}*) accumulates gag in testis, kidney and muscle, suggesting that HYAL-3 has a highly restrictive substrate specificity. A role for HYAL-3 in the testis is supported by previous data that has shown HYAL-3 is highly expressed in human testis. ✕

HA, the primary substrate of HYAL, has previously been implicated to play an important role in the mineralisation of bone. In this study mRNA expression of the HASs that synthesise HA (HAS-1, HAS-2 and HAS-3), and the HYALs which degrade HA (HYAL-1, HYAL-2, HYAL-3, HYAL-4) were examined in an osteoblast cell line

that could be induced to mineralise *in vitro* and gene expression was compared to the amount of gag production. During mineralisation a 13-fold decrease in HAS-3 expression was observed, as well as a 62-fold increase in HYAL-2 expression, a 13-fold increase in HYAL-3 expression and a 3-fold increase in HYAL-4 expression. These changes in gene expression were coupled to a 5-fold decrease in the production of HA. Therefore, in mineralising osteoblasts, expression of the genes that control HA metabolism are co-ordinated such that a general decrease in the expression of HASs and an increase in HYAL expression corresponds to a decrease in HA. These data implicate a role for HA in the early stages of matrix synthesis and maturation, rather than the later process of mineralisation.