



INVESTIGATIONS OF HISTONE H1 GENE-SPECIFIC PROMOTER ELEMENTS

Kym Nicholas Duncliffe, B.Sc. (Hons.)

A thesis submitted for the degree of

Doctor of Philosophy

in

The University of Adelaide

(Faculty of Science)

Department of Biochemistry,
University of Adelaide,
Adelaide, South Australia

June, 1994

Aspart Pris

TABLE OF CONTENTS

Summary		vii
Statement		ix
Note		х
Acknowledgments		xi
Dedication		xiii
w'		
CHAPTER ONE: INTRODUCTION		
1.1 General Introduction 1.2 The Regulation of Eukaryotic Class II Gene Expression 1.2.1 Class II Gene Structure 1.2.2 Transcriptional Regulation 1.2.2 (a) TATA-box 1.2.2 (b) General Promoter Elements 1.2.2 (c) Specific Promoter Elements 1.2.2 (d) Enhancers 1.2.2 (e) Locus Control Regions 1.2.3 Post-Transcriptional Regulation 1.2.3 (a) RNA Splicing 1.2.3 (b) 3' RNA Processing 1.2.3 (c) RNA Stability 1.2.3 (d) mRNA Transport 1.2.3 (e) Translation 1.3 Histones 1.3.1 H1 1.3.2 The Core Histones 1.4 Histones and Chromatin Structure 1.4.1 The Nucleosome 1.4.1 (a) Structure of the Nucleosome Core Particle		1 1 2 2 2 2 3 4 4 5 6 6 7 7 7 8 9 9 10 10 11 11 11
1.4.1 (b) Attachment of H1 to the Nucleosome 1.4.1 (c) Role of the core histone tails 1.4.2 Higher Order Chromatin Structure		12 12 13
 1.4.3 Non histone chromosomal proteins 1.5 Relationship between Histones, Chromatin Structure, and Transcr 1.5.1 Histone Subtypes and Variants 1.5.1 (a) H1 variants 1.5.1 (b) Core histone variants 	ription	14 14 15 16 18
 1.5.2 Histone Modifications 1.5.2 (a) Acetylation 1.5.2 (b) Phosphorylation 1.6 Histone Gene Structure 1.6.1 Structure of the Major Histone Genes 		19 20 21 23 23
1.6.2 Structure of variant histone genes 1.7 The Organisation of Histone Genes 1.7.1 Plants 1.7.2 Fungi		24 25 26 27
1.7.3 Caenorhabditis elegans 1.7.4 Drosophila 1.7.5 Sea Urchin		28 29 31

1.7.6 Trout 1.7.7 Amphibia	3 3	2
1.7.8 Chicken	3.	
1.7.9 Mammals	3	
1.8 Regulation of Histone Gene Expression: Lower Eukaryotes	3	
1.9 Regulation of Histone Gene Expression: Higher Eukaryotes	3	
1.9.1 Replication Independent Expression: Embryonic Uncoupling	3	
1.9.1 (a) Drosophila	3.3	
1.9.1 (b) Sea urchin	4	
1.9.1 (c) <i>Xenopus laevis</i> 1.9.2 Replication Independent Expression: Variant Histone Genes	4	
1.9.3 Replication Dependent Expression: Posttranscriptional Regulation	4	
1.9.3 (a) Histone Transcript Turnover	4	
1.9.3 (b) 3' Processing of Histone Primary Transcripts	4.	5
1.9.4 Replication Dependent Expression: Transcriptional Regulation	4	7
1.9.4 (a) H2A	4	
1.9.4 (b) H2B	4	
1.9.4 (c) H3	50	
1.9.4 (d) H4	5	
1.9.4 (e) H1	5.5	
1.10 The H1 Histone Genes of Chicken	5	b
CHAPTER TWO: MATERIALS AND METHODS		
2.1 Abbreviations	5	
2.2 Materials	59	
2.2.1 Chemicals, Reagents and Kits	5!	
2.2.2 Enzymes	60	
2.2.3 Isotopically Labelled Compounds	6	
2.2.4 Bacterial Strains and Cell Lines	6	
2.2.4 (a) E. coli Strains	6	
2.2.4 (b) Tissue Culture Cell Lines	6	
2.2.5 Media and Buffers	6:	
2.2.5 (a) Media 2.2.5 (b) Buffers	6	
2.2.6 Clones and Vectors	6	
2.2.6 (a) Cloned DNA Sequences	6	
2.2.6 (b) Cloning Vectors	6	4
2.2.6 (c) Reporter Vectors	6-	4
2.2.7 Molecular Weight Markers	6	
2.2.7 (a) DNA Markers	6	
2.2.7 (b) Protein Markers	6	
2.2.8 Oligonucleotides	6	
2.3 Methods	6	
2.3.1 Isolation of Plasmid DNA	6	
2.3.1 (a) General Procedure	6	
2.3.1 (b) Miniscreen Procedure	6	
(i) Alkaline extraction	6	
(ii) STET preps	6	
2.3.1 (c) Large Scale Purification on a CsCl Gradient	6	
2.3.2 Restriction Enzyme Digestion and Analysis of DNA 2.3.2 (a) Restriction Enzyme Digestion	6	
2.3.2 (a) Restriction Elizythe Digestion 2.3.2 (b) Agarose Gel Electrophoresis	6	
2.3.2 (c) Polyacrylamide Gel Electrophoresis	6	
2.3.2 (d) Denaturing polyacrylamide gels	$\ddot{7}$	
2.3.2 (d) Defiationing polyacitylamide gets 2.3.3 Subcloning of DNA fragments into plasmid vectors	Ź	
2.3.3 (a) Isolation of DNA from agarose gels	7	

(i) Geneclean		70
(ii) Spin-columns		71
2.3.3 (b) Isolation of DNA from low gelling temperature agarose gels		71
2.3.3 (c) Preparation of Vector		71
2.3.3 (d) Ligation into plasmid vectors	,	72
2.3.3 (e) Transformation of recombinants into bacteria		72
2.3.3 (f) Plasmid miniscreen procedure		73
2.3.4 Purification of oligonucleotides		73
2.3.5 Labelling of DNA fragments		73
2.3.5 (a) Kinasing of synthetic oligonucleotides		73
2.3.5 (b) Endfilling of restriction fragments		74
2.3.6 Manipulation and Harvesting of tissue culture cells		74
2.3.7 Nuclear extract preparation		75
2.3.7 (a) Tissue culture cells		75 75
		75 75
(i) From Strauss and Varshavsky (1984)		
(ii) From Lee et al (1988)		75
(iii) From Dignam et al (1983)		76
(iv) From Dalton (1987)		76
2.3.7 (b) Small scale nuclear extract preparation from chicken embryos		77
2.3.7 (c) Large scale nuclear extract preparation from chicken embryos		78
2.3.7 (d) Protein concentration determination		78
2.3.8 Detection of DNA binding factors in nuclear extracts		79
2.3.8 (a) Probe preparation		79
2.3.8 (b) Mobility shift assay		79
2.3.8 (c) Polyacrylamide gel electrophoresis of mobility shift assays		79
2.3.9 Chromatographic enrichment of nuclear extract components		80
2.3.9 (a) DEAE- / Heparin-Sepharose chromatography		80
2.3.9 (b) Large scale concentration of partially purified extracts		80
2.3.9 (c) Sephacryl S-300 gel filtration chromatography		80
2.3.9 (d) DNA-affinity microprecipitation		81
(i) Preparation of DNA Ligand		81
(ii) DNA affinity microprecipitation assay		81
(iii) DNA affinity column chromatography		82
2.3.10 SDS Polyacrylamide gel electrophoresis of proteins		83
2.3.10 (a) Gel electrophoresis		83
2.3.10 (b) Silver staining of protein gels		83
2.3.10 (c) Coomassie staining of protein gels		84
2.3.11 Electroblotting of protein to PVDF membranes		84
2.3.12 Association of DNA-binding activity with a protein		85
2.3.12 (a) South-Western transfer		85
2.3.12 (b) Renaturation of binding activity from SDS polyacrylamide gels		85
(i) Guanidine-HCl method		86
(ii) Saturated urea method		86
2.3.13 Transient transfection of tissue culture cells		86
2.3.14 CAT assays		87
2.3.15 Double Stranded Dideoxy Sequencing of DNA		88
2.3.15 (a) Preparation of Template		88
2.3.15 (b) Hybridization		88
2.3.15 (c) Extension and termination		88
2.3.16 Polymerase Chain Reaction (PCR)		89
2.3.17 Autoradiography		89
2.3.17 Autoratiography 2.3.18 Laser Densitometry		90
2.3.19 Phosphorimaging		90
2.3.20 Containment facilities and animal ethics		90
2.3.20 Contaminent facilities and animal eulics		90

CHAPTER THREE: TOWARD PURIFICATION OF AC-BOX-BINDING FACTORS IN CHICKEN

3.1 Introduction	91
3.2 Characterisation of AC-box Binding Activity	92
3.2.1 Source of AC-box Binding Factors	92
3.2.2 Choice and Preparation of Probe	92
3.2.3 Mobility Shift Assays of Chicken Embryo Extracts with the 01-40mer Probe	93
3.2.4 Analysis of AEV ts34 Nuclear Extracts with the 01-40mer Probe	93
3.2.5 Small Scale DEAE- and Heparin-Sepharose Chromatography of ts34 Crude Nuclear Extracts	94
3.2.6 Attempts to Optimise Recovery of 40-MS-e	94
3.2.7 Differences Between H1-SF and 40-MS-e	95
3.2.8 Mobility Shift Assays with the 14mer Probe	96
3.3 Large Scale Preparation and Chromatography of Chicken Embryo Extracts	98
3.3.1 Heparin-Sepharose Chromatography	98
3.3.2 Ultrafiltration of Heparin-Sepharose Extracts	98
3.4 DNA Affinity Chromatography of Partially Purified Extracts	99
3.4.1 Design of a Suitable Oligonucleotide	99
3.4.2 Preparation of the DNA Affinity Ligand	99
3.4.3 DNA Affinity Microprecipitation	100
3.4.4 Optimisation of Microprecipitation Method	100
3.4.5 Scale-up of Microprecipitation Method	102
3.5 Possible Co-purification of H1-Transcription Factors with 14-MS-b	103
3.6 SDS Polyacrylamide Gel Electrophoresis of DNA Affinity Purified Extracts	105
3.7 Direct Amino Acid Sequencing of a DNA Affinity-Purified Protein	105
3.8 2nd-Round DNA Affinity Microprecipitation	106
3.9 Alternative Purification Methods	107
3.9.1 Sephacryl S-300 Gel Filtration Chromatography	107
3.9.2 DNA Affinity Column Chromatography	108
3.10 Renaturation of 14-MS-b	109
3.10.1 Southwestern Analysis	109
3.10.2 Renaturation of Proteins from Gel Slices	109
3.10.3 Analysis of ts34 Extracts with the 01-14mer Probe	110
3.11 Identification and Renaturation of an 01-40mer Binding Factor in DNA Affinity-	
Purified Extracts	111
3.11.1 Characterisation of the 40-MS-1 Band Shift	111
3.11.2 Southwestern Analysis with the 01-40mer Probe	111
3.11.3 Renaturation of 40-MS-1	112
3.12 Use of Intermediate Length Oligonucleotide Probes	113
3.12.1 Design of 25 bp Oligonucleotides	113
3.12.2 Mobility Shift Assays of Chicken Embryo Extracts with the 25mers	113
3.12.3 DNA Affinity Purification with 25mers	114
3.12.4 Characterisation of a Subtype-Specific AC-box-Binding Factor in ts34 Extracts	114
3.12.5 Mobility Shift Assays of Various Cell Lines with the 25mer Probes	115
3.13 Discussion	116
CHAPTER FOUR: FUNCTIONAL ANALYSES OF THE AC-BOX AND	
CHAPTER FOUR: FUNCTIONAL ANALYSES OF THE AC-BOX AND	
A RELATED H1-SPECIFIC PROMOTER ELEMENT	
4.1 Introduction	122
4.2 Effect of the AC-box in a non-S-phase regulated promoter	123
4.2.1 Choice of a suitable reporter system	123
4.2.2 Construction of AC-box/HSVtk promoter fusions in pTKGH	124
4.2.3 Analysis of AC-box/HSVtk promoter activity in p(AC)TKGH series	125
4.2.4 Construction of the p(AC)TKCAT series	126
4.2.5 Analysis of AC-box/HSVtk promoter activity in p(AC)TKCAT.N series	127

4.3 Investigation of a Distal Element in the H1 Histone Promoter	128
4.3.1 Comparison of H1 Histone Promoter Sequences	129
4.3.2 Sequencing of the Chicken 02 H1 Histone Promoter	130
4.3.3 Construction of H1 Promoter / CAT Fusions	131
4.3.3 (a) Generation of p(02)TKCAT vectors	131
4.3.3 (b) Generation of p(02)HCAT vectors	132
4.3.4 Effect of the TG-box on H1 Promoter Activity	132
4.3.5 Effect of an 02 H1 Promoter Fragment on Expression of the HSVtk Promoter	133
4.4 Discussion	134
CHAPTER FIVE: FINAL DISCUSSION	139
BIBLIOGRAPHY	146

SUMMARY

The work presented in this thesis reports investigations of the histone H1 gene-specific promoter element, 5' AAACACA 3', also referred to as the "H1-box" or "AC-box" (Coles and Wells, 1985). The first approach was to isolate and purify a protein factor, H1-SF, from 9 day chicken embryos. This protein had been previously identified in nuclear extracts from AEV ts34 chicken erythroblasts, and had been demonstrated to interact specifically with the AC-box (Dalton and Wells, 1988a).

In mobility-shift assays using a 40 bp oligonucleotide probe (containing the AC-box) derived from the promoter of the chicken histone H1 gene, 01 H1, a candidate AC-box-binding factor, 40-MS-e, was identified in crude nuclear extracts from ts34 cells. Further characterisation revealed anomalies between some properties of this factor and those of H1-SF. In addition, a specific AC-box-binding factor could not be identified in crude chicken embryo extracts. However, a second AC-box-binding factor, 14-MS-b, was identified in chicken embryo extracts using a 14 bp oligonucleotide probe containing only the AC-box.

14-MS-b was purified from chicken embryo extracts on a large scale, using Heparin Sepharose-, Sephacryl S-300-, and finally DNA-affinity chromatography using concatemers of the 14-mer as the ligand. Analysis of DNA affinity-purified extracts on SDS-polyacrylamide gels revealed two enriched proteins of ~86 kD and ~70 kD. To determine whether either of these proteins was 14-MS-b, the MW of 14-MS-b was estimated by renaturation from SDS-polyacrylamide gels. Surprisingly, the renatured 14-MS-b was no greater than 30 kD: much smaller than either of the enriched proteins, or the estimated mass of H1-SF (90 kD - S. Dalton, unpublished results). Thus it was unlikely that 14-MS-b was analogous to H1-SF.

Another factor, 40-MS-1, was identified in mobility shift assays of DNA affinity-purified extracts using the 40 bp oligonucleotide probe. This factor was also renatured from SDS-polyacrylamide gels, but it too was much smaller (42-46 kD) than the estimated molecular mass of H1-SF. Further investigation of the 86 kD and 70 kD proteins indicated that they were non specific DNA binding proteins.

A final series of mobility-shift assays was conducted using four 25 bp oligonucleotides carrying either wild type or mutated AC-boxes derived from the promoters of the chicken histone 01 H1 and 02 H1 genes. In crude nuclear extracts from ts34 cells, a factor (25-MS-1) which specifically bound the wild type 02 H1 oligonucleotide was identified. 25-MS-1 was also detected in extracts from three other chicken cell lines. However, this factor was not detected in 9 day chicken embryo extracts. In addition, no AC-box-specific binding activity analogous to 25-MS-1 was detected with the wild type 01 H1 oligonucleotide.

It had been shown that deletion or mutation of the AC-box both greatly decreased the steady state levels of H1-mRNA and eliminated cell cycle regulation of H1 gene transcription in stably transfected HeLa cell lines (Dalton and Wells, 1988a). To examine the effect of the AC-box on a heterologous promoter, a series of constructs were made in which tandem repeats of the 25 bp AC-box oligonucleotides were cloned immediately 5' to the Herpes Simplex Virus thymidine kinase (HSVtk) promoter of the expression vector pBLCAT.2 (Luckow and Schütz, 1987). It was found that these oligonucleotides, regardless of copy number or orientation, had no significant effect on transcription from the HSVtk promoter in transiently transfected ts34 cells.

Concurrent with the transfection experiments, a computer alignment of the H1 promoters from several species revealed a distal promoter element which was highly conserved in sequence and position. This element (TG-box) was a near perfect inverted repeat of the AC-box, and the spacing between these elements was also well conserved. CAT reporter vectors were constructed which contained fragments of the 02 H1 and 01 H1 promoters. Deletion of a 29 bp region which included the TG-box resulted in a 50% decrease in expression of the 02 H1 promoter in transiently transfected ts34 cells. However, an H1 promoter fragment containing the TG- and AC-boxes, but no proximal H1 promoter elements, had no significant effect on expression of the HSVtk promoter. These data are discussed in relation to the possible role of the TG- and AC-boxes in the regulation of histone H1 gene expression.