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# Involvement of the Osteoblast in Paget's Disease of Bone

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# ABSTRACT

Paget's disease is characterised by focal regions of accelerated bone turnover. The aetiology is unknown, but genetic and environmental factors have been implicated. Pagetic lesions contain increased numbers of osteoclasts with abnormal morphology, so an osteoclast defect has been considered central to the pathogenesis. However, given osteoblasts regulate osteoclast differentiation and activity; osteoblast abnormalities may be important in the disease. This study aimed to identify features of pagetic osteoblasts that could clarify their role in Paget's disease.

Gene expression in osteoblasts and bone marrow cultured from pagetic lesions of 23 patients was compared to cells from unaffected tissue using both microarrays and real time RT-PCR. The results indicated global changes in gene expression in pagetic osteoblasts. A number of genes that can stimulate osteoclastogenesis, including interleukins 6 and 1 $\beta$ , and monocyte chemotactic factor 1 were up-regulated, but the *RANKL/OPG* ratio tended to be decreased. Genes involved in osteoblast differentiation were down-regulated, including the transcription factors *RUNX2*, *DLX5* and *SATB2*, the osteogenic factor *BMP2*, and the matrix proteins osteocalcin and bone sialoprotein. Markers of less mature osteoblastic cells, alkaline phosphatase and matrix gla protein were up-regulated. The intermediate filament, keratin 18, was very significantly up-regulated in pagetic cells. Over-expression of this protein in osteoblasts using an adenoviral vector produced some changes in gene expression, but did not produce an overtly pagetic phenotype. Over-expression of *SQSTM1* mutants found in some patients with Paget's disease also produced only minor changes in osteoblast phenotype. The RNA from the primary cell cultures was also used to investigate the presence of measles virus and somatic mutations in *SQSTM1* in the disease, but neither were identified in any of the patients.

These results suggest that there are important changes in pagetic osteoblasts that are maintained when the cells are removed from the affected bone microenvironment. These include enhanced production of factors to stimulate osteoclastogenesis, while osteoblast differentiation and activity may be impaired. We were unable to identify genetic or environmental factors that could trigger these changes. The pagetic osteoblast is distinct from control cells, and is likely to contribute to the development of Paget's disease.

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# TABLE OF CONTENTS

ABSTRACT .....	i
ACKNOWLEDGEMENTS .....	ii
LIST OF TABLES .....	viii
LIST OF FIGURES.....	ix
ABBREVIATIONS.....	xiii
CHAPTER 1: INTRODUCTION .....	1
Part A. BONE BIOLOGY.....	1
1.1 Macrostructure of bone .....	1
1.1.1 Cortical bone.....	1
1.1.2 Trabecular bone.....	2
1.2 Bone formation and modelling.....	2
1.2.1 Intramembranous ossification .....	3
1.2.2 Endochondral ossification.....	4
1.3 Bone remodelling .....	4
1.4 Bone mass .....	6
1.5 Bone matrix .....	6
1.5.1 Mineralisation .....	8
1.6 Bone cells .....	8
1.6.1 Osteocytes .....	8
1.6.2 Osteoblasts .....	10
1.6.3 Osteoclasts .....	16
1.7 Hormones and growth factors involved in bone metabolism.....	19
1.7.1 Parathyroid hormone and parathyroid hormone related protein .....	19
1.7.2 Vitamin D.....	20
1.7.3 Oestrogen and sex hormones .....	21
1.7.4 Glucocorticoids .....	22
1.7.5 Transforming growth factor $\beta$ .....	22
1.7.6 Other growth factors .....	23
1.8 Common disorders of bone metabolism.....	24
1.8.1 Osteoporosis.....	24
1.8.2 Hyperparathyroidism.....	25
1.8.3 Cancer and bone.....	25

Part B. PAGET'S DISEASE OF BONE .....	27
1.9 Epidemiology .....	27
1.10 Clinical features.....	29
1.11 Diagnosis and treatment .....	30
1.12 Pathophysiology .....	34
1.13 Aetiology .....	37
1.13.1 Genetics.....	37
1.13.2 Paramyxoviruses and Paget's disease .....	41
1.14 Genetic disorders similar to Paget's disease .....	42
1.14.1 Familial expansile osteolysis and other diseases caused by <i>RANK</i> mutations..	42
1.14.2 Idiopathic hyperphosphatasia.....	44
1.14.3 Inclusion body myopathy, Paget's disease and frontotemporal dementia .....	45
Part C. AIM .....	46
1.15 Aims of this study .....	46
CHAPTER 2: METHODS .....	47
2.1 Materials.....	47
2.1.1 Cell culture .....	47
2.1.2 Molecular biology .....	48
2.1.3 Protein detection .....	48
2.1.4 Solutions.....	48
2.2 Cell culture .....	49
2.2.1 Human osteoblasts .....	49
2.2.2 Human bone marrow cells .....	50
2.2.3 Primary rat osteoblasts .....	51
2.2.4 Cell lines .....	51
2.2.5 Proliferation assay .....	51
2.2.6 Mineralisation assay.....	52
2.2.7 Mouse bone marrow osteoclastogenesis culture .....	53
2.2.8 Luciferase assay .....	53
2.2.9 Cultures in three dimensional scaffolds .....	54
2.3 Molecular biology .....	55
2.3.1 RNA extraction .....	55
2.3.2 RT-PCR.....	55
2.3.3 Sequencing .....	58

2.3.4 Real time RT-PCR .....	58
2.3.5 Microarrays .....	61
2.3.6 Cloning.....	62
2.3.7 Adenoviral vectors .....	63
2.4 Protein detection.....	65
2.4.1 ELISA .....	65
2.4.2 Western blotting.....	65
2.4.3 Luminex .....	66
2.5 Statistical analysis and graphs.....	66
<b>CHAPTER 3: GLOBAL ANALYSIS OF GENE EXPRESSION IN PAGETIC OSTEOPBLASTS .....</b>	<b>67</b>
3.1 Introduction .....	67
3.2 Methods .....	68
3.2.1 Sample details .....	68
3.2.2 Cell characteristics .....	70
3.3 Microarray analysis results.....	71
3.3.1 Overview of Affymetrix microarrays.....	71
3.3.2 Quality control and normalisation.....	71
3.3.3 Global analysis of gene expression .....	73
3.3.4 Expression of individual genes .....	77
3.3.5 Pathway analysis .....	81
3.4 Discussion .....	84
<b>CHAPTER 4: DIFFERENTIAL GENE EXPRESSION IN PAGETIC CELLS .....</b>	<b>87</b>
4.1 Introduction .....	87
4.2 Methods .....	87
4.3 Results – Differential gene expression in pagetic cultures .....	89
4.3.1 Local regulators of bone metabolism.....	89
4.3.2 Osteoblast matrix proteins .....	92
4.3.3 Osteoblast transcription factors.....	95
4.3.4 Wnt signalling molecules.....	95
4.3.5 Other genes with known roles in bone biology.....	97
4.3.6 Genes with unexplored roles in osteoblast biology .....	97
4.3.7 Overall changes in gene expression in pagetic and non-pagetic samples.....	100
4.3.8 Effects of patient age and sex on gene expression.....	104

4.3.9 Correlation of gene expression for different genes .....	105
4.4 Results – Gene expression in differentiating osteoblasts .....	106
4.4.1 Osteoblast matrix proteins .....	108
4.4.2 Local regulators of bone turnover .....	112
4.4.3 Osteoblast transcription factors.....	113
4.4.4 Wnt signalling molecules.....	113
4.4.5 Other genes with known roles in bone biology.....	114
4.4.6 Genes with unknown roles in bone biology.....	116
4.4.7 Summary of gene expression in differentiating osteoblasts.....	118
4.5 Discussion .....	119
4.5.1 Agreement between real time PCR and microarray results .....	121
4.5.2 RANKL/OPG.....	122
4.5.3 Interleukins.....	123
4.5.4 CCL2 .....	123
4.5.5 BMP2 and osteoblast transcription factors .....	124
4.5.6 Osteoblast matrix proteins .....	125
4.5.7 Wnt signalling molecules.....	126
4.5.8 FGF signalling.....	127
4.5.9 Prostaglandin D synthase .....	127
4.5.10 Genes with unexplored roles in osteoblast biology.....	128
4.5.11 Differences between osteoblasts and bone marrow .....	130
4.5.12 Paired osteoblast samples.....	130
4.5.13 Conclusion .....	131
CHAPTER 5: EFFECTS OF DICKKOPF 1 AND KERATIN 18 ON BONE CELLS.....	132
5.1 Introduction .....	132
5.1.1 Dickkopf 1.....	132
5.1.2 Keratin 18.....	132
5.2 Results – Dickkopf 1 .....	133
5.3 Results – Keratin 18 .....	134
5.3.1 Optimising the transduction conditions .....	135
5.3.2 Effects of keratin 18 over-expression on cell proliferation.....	138
5.3.3 Effects of keratin 18 over-expression on osteoblast gene expression .....	141
5.3.4 Other effects of keratin 18 in osteoblasts.....	146
5.4 Discussion .....	149

CHAPTER 6: EFFECT OF WILD-TYPE AND MUTANT SQSTM1 IN OSTEOBLASTS	154
6.1 Introduction .....	154
6.2 Methods .....	155
6.2.1 Vector construction .....	155
6.2.2 Site-directed mutagenesis.....	156
6.2.3 Transfection and protein expression .....	157
6.3 Results .....	157
6.3.1 Basal <i>SQSTM1</i> expression in osteoblasts.....	157
6.3.2 Over-expression of SQSTM1 in transfected cells.....	159
6.3.3 Effect of wild-type and mutant SQSTM1 expression on proliferation.....	160
6.3.4 Effect of SQSTM1 over-expression on NF $\kappa$ B signalling in osteoblasts.....	160
6.3.5 Effects of SQSTM1 over-expression on cell signalling pathways.....	164
6.3.6 Effects of SQSTM1 over-expression on gene expression.....	166
6.4 Discussion .....	168
CHAPTER 7: DETECTION OF MEASLES VIRUS RNA IN PAGETIC CELLS	172
7.1 Introduction .....	172
7.2 Methods .....	172
7.3 Further characterisation of bone marrow cells.....	172
7.4 Measles virus detection .....	174
7.5 Discussion .....	174
CHAPTER 8: DETECTION OF SOMATIC <i>SQSTM1</i> MUTATIONS IN PAGETIC CELLS	177
8.1 Introduction .....	177
8.2 Sequence analysis of exons 7 and 8 of SQSTM1.....	177
8.3 Allelic discrimination.....	178
8.3.1 Development and validation of methodology .....	178
8.3.2 Testing of pagetic samples .....	180
8.4 Discussion .....	181
CHAPTER 9: GENERAL DISCUSSION AND CONCLUSIONS	184
9.1 Discussion .....	184
9.2 Future directions.....	192
9.3 Conclusions .....	194
REFERENCES.....	196

# LIST OF TABLES

Table 2.1: PCR primers used in this thesis.....	57
Table 2.2: Genes included in the human low density arrays containing 24 primer-probesets.	59
Table 2.3: Genes included in the human low density arrays containing 48 primer-probesets.	60
Table 3.1: Patient and sample details for the patients with Paget's disease.....	69
Table 3.2: Summary of patient details and samples collected from patients with Paget's disease and controls.....	69
Table 3.3: Patient details for the 12 RNA samples run on the microarrays.....	73
Table 3.4: Genes showing statistically significant changes in expression in microarray analysis of pagetic and non-pagetic osteoblast RNA samples .....	77
Table 3.5: Top 10 up-regulated and down-regulated genes in the microarray analysis of pagetic and non-pagetic osteoblast RNA samples ranked according to fold change.....	77
Table 3.6: Top differentially regulated genes in the three paired samples .....	80
Table 3.7: Pathway analysis showing KEGG pathways that had statistically significant changes using more than one method of analysis .....	83
Table 3.8: Gene Ontology analysis showing the most significantly changed categories between pagetic and non-pagetic samples .....	84
Table 4.1: Microarray results for genes that have been examined using real time PCR .....	88
Table 4.2: Changes in expression of genes of interest during osteoblast differentiation.....	119
Table 4.3: Fold changes in gene expression between non-pagetic and pagetic samples in the 12 samples that were analysed on the microarrays determined using microarray analysis and real time PCR .....	121

# LIST OF FIGURES

Figure 1.1: Macrostructure of a human femur .....	2
Figure 1.2: Formation and growth of long bones by endochondral ossification.....	3
Figure 1.3: Bone remodelling in a BMU.....	5
Figure 1.4: Osteocyte cells and networks.....	9
Figure 1.5: The mesenchymal cell lineage.....	10
Figure 1.6: Osteoblastic cell differentiation.....	12
Figure 1.7: Wnt/β-catenin signalling pathway .....	13
Figure 1.8: Transmission electron microscopy image of a primary rat osteoclast on bone....	16
Figure 1.9: Schematic showing the roles of RANKL and OPG in osteoclastogenesis .....	18
Figure 1.10: The mechanism of osteoclastic bone resorption.....	19
Figure 1.11: Sketch of a patient with Paget's disease .....	27
Figure 1.12: Variable prevalence of Paget's disease in the UK.....	28
Figure 1.13: Imaging of Paget's disease .....	31
Figure 1.14: Bisphosphonate structure .....	32
Figure 1.15: Radiographic progression of Paget's disease .....	34
Figure 1.16: Scanning electron micrographs of normal and pagetic trabecular bone.....	35
Figure 1.17: Structure and function of SQSTM1/p62.....	39
Figure 1.18: Transmission electron microscopy images of typical inclusions in nuclei of pagetic osteoclasts .....	42
Figure 1.19: Photograph of a patient suffering from familial expansile osteolysis .....	43
Figure 2.1: Human osteoblast outgrowth cultures from trabecular bone explants.....	50
Figure 3.1: Alkaline phosphatase staining in osteoblast cultures.....	70
Figure 3.2: Pagetic bone marrow culture stained for alkaline phosphatase and TRAP .....	70
Figure 3.3: Gel and electropherogram images of selected RNA samples examined using the bioanalyser .....	72
Figure 3.4: Box plots showing distribution of gene expression in the twelve microarrays ....	74
Figure 3.5: Dendrograms for clustering experiments using all expressed probesets .....	75
Figure 3.6: Dendrograms for clustering experiments using the 269 most changed probesets.	76
Figure 3.7: Heat map showing expression levels of genes with the most significant changes or highest fold changes between pagetic and non-pagetic gene expression overall.....	79
Figure 3.8: Heat map showing most highly changed genes in the paired samples .....	81

Figure 4.1: Relative expression levels of <i>RANKL</i> , OPG and <i>RANK</i> in non-pagetic and pagetic cells.....	90
Figure 4.2: Relative expression levels of <i>IL-6</i> , <i>IL-1<math>\beta</math></i> and <i>IL-11</i> mRNA in non-pagetic and pagetic cells .....	91
Figure 4.3: Relative expression of <i>BMP2</i> and <i>IGF1</i> mRNA in non-pagetic and pagetic cells	91
Figure 4.4: Relative expression of osteoblast matrix protein mRNA in non-pagetic and pagetic cells.....	93
Figure 4.5: Relative expression of osteoblast transcription factor mRNA in non-pagetic and pagetic cells .....	94
Figure 4.6: Expression of genes involved in the Wnt signalling pathway in non-pagetic and pagetic cells .....	96
Figure 4.7: Relative expression of <i>CCL2</i> , <i>FGFR2</i> , <i>PTGDS</i> and <i>MMP13</i> mRNA in non-pagetic and pagetic cells .....	98
Figure 4.8: Relative mRNA expression of genes with unknown functions in bone biology in non-pagetic and pagetic cells .....	99
Figure 4.9: Summary of changes in gene expression in pagetic cells .....	101
Figure 4.10: Differences in gene expression between cell types and relative expression levels of different genes.....	102
Figure 4.11: Correlation of expression of different pairs of genes in osteoblasts and bone marrow .....	103
Figure 4.12: Correlation of expression of different pairs of genes in osteoblasts only.....	104
Figure 4.13: Mineralisation in MC3T3-E1 subclone 4 cells over time.....	107
Figure 4.14: Mineralisation in human osteoblasts .....	108
Figure 4.15: Expression of osteoblast marker genes in differentiating MC3T3-E1 and human osteoblast cells.....	110
Figure 4.16: Expression of matrix gla protein, tenascin C and osteomodulin in differentiating MC3T3-E1 and human osteoblast cells .....	111
Figure 4.17: Expression of local regulators of bone metabolism <i>OPG</i> , <i>IL-6</i> and <i>BMP2</i> in differentiating MC3T3-E1 and human osteoblast cells.....	112
Figure 4.18: Expression of osteoblast transcription factors <i>RUNX2</i> and <i>DLX5</i> in differentiating MC3T3-E1 and human osteoblast cells.....	113
Figure 4.19: Expression of genes involved in Wnt signalling during human osteoblast differentiation .....	114

Figure 4.20: Expression of other genes involved in bone metabolism <i>CCL2</i> , <i>FGFR2</i> , <i>PTGDS</i> and <i>MMP13</i> in differentiating MC3T3-E1 and human osteoblast cells.....	115
Figure 4.21: Expression of <i>GATA6</i> , <i>IFI27</i> , <i>RGS4</i> and <i>SGK1</i> in differentiating MC3T3-E1 and human osteoblast cells.....	117
Figure 4.22: Expression of <i>KRT18</i> (a) and <i>MAFB</i> (b) in differentiating human osteoblast cells .....	118
Figure 4.23: Effects of genes with altered transcript abundance in pagetic osteoblasts on bone cell differentiation .....	120
Figure 5.1: Hierarchical structure of an intermediate filament .....	133
Figure 5.2: Effect of Dkk1 on osteoblast proliferation and osteoclastogenesis .....	134
Figure 5.3: Structure and gene transduction pathway of an adenovirus vector .....	135
Figure 5.4: Primary human osteoblasts transduced with adenoviral vectors with and without GeneJammer transfection reagent .....	136
Figure 5.5: Transduction of various osteoblastic cell types with keratin 18 adenovirus .....	137
Figure 5.6: Western blot showing keratin 18 expression in transduced cells .....	138
Figure 5.7: Human osteoblast proliferation in cells transduced with keratin 18.....	139
Figure 5.8: SaOS2 proliferation in cells transduced with keratin 18 .....	140
Figure 5.9: Relative expression of <i>BMP6</i> , <i>DKK1</i> and <i>M-CSF</i> in human osteoblasts transduced with keratin 18.....	142
Figure 5.10: Relative expression of <i>CCL2</i> , <i>IL-1<math>\beta</math></i> and <i>IL-6</i> in human osteoblasts transduced with keratin 18.....	143
Figure 5.11: Relative expression of <i>FGF2</i> , <i>IL-6</i> and osteopontin in SaOS2 cells transduced with keratin 18.....	145
Figure 5.12: Unusual cellular structures in keratin 18-transduced cells .....	146
Figure 5.13: GFP-positive cells after four weeks growth on collagen scaffolds .....	147
Figure 5.14: Live/dead staining of cells cultured on collagen scaffolds for four weeks.....	148
Figure 5.15: Sections of collagen scaffolds stained for alkaline phosphatase, mineralisation, and cell number after four weeks in culture .....	149
Figure 6.1: Parts of the <i>SQSTM1</i> cDNA sequence (genbank accession number NM_003900) showing the introduction of restriction sites for cloning.....	156
Figure 6.2: Gel showing the plasmid after insertion of the <i>SQSTM1</i> sequence.....	156
Figure 6.3: Expression of <i>SQSTM1</i> in osteoblastic cells .....	158
Figure 6.4: Functional SQSTM1 is over-expressed by the plasmids .....	159
Figure 6.5: Proliferation in SaOS2 cells transfected with wild-type or mutant SQSTM1.....	160

Figure 6.6: NF $\kappa$ B activity in cells transfected with wild-type and mutant SQSTM1 .....	161
Figure 6.7: Effects of TNF $\alpha$ on osteoblast-like cells .....	162
Figure 6.8: NF $\kappa$ B activity in transfected osteoblast-like cells treated with TNF $\alpha$ .....	163
Figure 6.9: Phospho-JNK levels detected using Luminex in transfected SaOS2 cells .....	165
Figure 6.10: Gene expression in SaOS2 cells transfected with SQSTM1 and mutants.....	167
Figure 7.1: Relative expression of osteoclast precursor marker genes in bone marrow cell cultures .....	173
Figure 8.1: Electropherograms showing the heterozygous 1215C/T mutation.....	178
Figure 8.2: Allelic discrimination standard curve amplification plots.....	179
Figure 8.3: Allelic discrimination standard curve endpoint results .....	180
Figure 8.4: Allelic discrimination results in patient cDNA samples.....	181
Figure 9.1: Schema showing possible effects and interactions of changes in interleukin and DKK1 expression identified in this study .....	186

# ABBREVIATIONS

1,25(OH) <sub>2</sub> D <sub>3</sub>	1,25 dihydroxyvitamin D <sub>3</sub>
A2P	L-ascorbic acid 2-phosphate
AAA	ATPases associated with a variety of activities
ALP	alkaline phosphatase
aPKC	atypical protein kinase C
ATF	activating transcription factor
αMEM	minimum essential medium, alpha
BM	bone marrow
BMD	bone mineral density
BMP	bone morphogenetic protein
BMU	basic multicellular unit
bp	base pair
BSA	bovine serum albumin
BSP	bone sialoprotein
C/EBP	CAAT/enhancer-binding protein
cAMP	cyclic adenosine monophosphate
CAR	Coxsackie-adenovirus receptor
Cath K	cathepsin K
CBP	CREB binding protein
CCL2	monocyte chemotactic protein 1
CFU-GM	granulocyte macrophage colony forming unit
CHI3L1	chitinase 3-like 1
Col-I/II/X	type I/II/X collagen
CREB	cAMP response element binding
CTx	C-telopeptide of collagen crosslinks
DCt	delta Ct
DKK1	dickkopf 1
DMEM	Dulbecco's modified Eagle's medium
DMP1	dentin matrix protein 1
Dpd	deoxypyridolines
Dsh	dishevelled
ER	oestrogen receptor

ERK	extracellular signal-regulated kinase
FBS	foetal bovine serum
FEO	familial expansile osteolysis
FGF	fibroblast growth factor
FGFR	fibroblast growth factor receptor
FPPS	farnesyl pyrophosphate synthase
Fra	fos-related antigen
GATA6	GATA binding protein 6
GCOS	GeneChip operating software
GFP	green fluorescent protein
gla	$\gamma$ -carboxylic acid
GM-CSF	granulocyte macrophage colony stimulating factor
GO	gene ontology
GSK	glycogen synthase kinase
HOB	human osteoblasts
hr	hour
IBMPFD	inclusion body myopathy, Paget's disease and frontotemporal dementia
IGF	insulin-like growth factor
Ihh	Indian hedgehog
IKK	I $\kappa$ B kinase
IL	interleukin
IL-1R	Interleukin 1 receptor
I-Smad	inhibitory Smad
I $\kappa$ B	inhibitor of NF $\kappa$ B
JAK	Janus kinase
JNK	jun N-terminal kinase
KEGG	Kyoto encyclopaedia for genes and genomes
LRP	low-density lipoprotein receptor-related protein
M gene	matrix gene
MAPK	mitogen activated protein kinase
M-CSF	macrophage colony stimulating factor
MEF2	myocyte-enhancer factor 2
MEM	minimum essential medium
MEPE	matrix extracellular phosphoglycoprotein

MGB	minor groove binder
MGP	matrix gla protein
min	minute
MMP	matrix metalloproteinase
MOI	multiplicity of infection
MRFs	myogenic regulatory factors
MV	measles virus
N gene	nucleocapsid gene
NCoR	nuclear receptor corepressor
NFAT	nuclear factor of activated T cells
NF $\kappa$ B	nuclear factor kappa B
NGFR	nerve growth factor receptor
NIBSC	National Institute of Biological Standards and Control
NTC	no template control
NTx	N-telopeptide of collagen crosslinks
OB	osteoblast
OC	osteocalcin
ONPG	o-nitrophenyl- $\beta$ -D-galactopyranoside
OPG	osteoprotegerin
Osx	osterix
P1CP	carboxy-terminal propeptide of type I collagen
P1NP	amino-terminal propeptide of type I collagen
p62	sequestosome 1
PB1	phox and bem1p-1
PBMC	peripheral blood mononuclear cells
PBS	phosphate buffered saline
PCOT2	principal coordinates and Hotelling's T squared
PCR	polymerase chain reaction
PDGF	platelet-derived growth factor
pfu	plaque-forming units
PGD <sub>2</sub>	prostaglandin D <sub>2</sub>
PGE <sub>2</sub>	prostaglandin E <sub>2</sub>
PI3K	phosphatidylinositol-3 kinase
PPAR $\gamma$	peroxisome proliferator-activated receptor $\gamma$

PTGDS	prostaglandin D2 synthase, brain isoform
PTH	parathyroid hormone
PTHrP	parathyroid hormone related protein
Pyd	pyridinolines
RANK	receptor activator of nuclear factor κB
RANKL	RANK ligand
RGS	regulator of G-protein signalling
RIN	RNA integrity number
RIP	receptor interacting protein
RMA	robust multichip analysis
R-Smad	receptor-activated Smad
RT-PCR	reverse transcription polymerase chain reaction
s	second
SAFE	significance analysis of function and expression
SERM	selective oestrogen receptor modulators
sFRP	secreted frizzled-related protein
SGK1	serum/glucocorticoid regulated kinase 1
SIBLING	small integrin-binding ligand, N-glycosylated proteins
SMRT	silencing mediator of retinoid and thyroid receptors
SOST	sclerostin
SQSTM1	sequestosome 1
SSPE	subacute sclerosing panencephalitis
STAT	signal transducers and activators of transcription
TBS	tris-buffered saline
TBS-T	tris-buffered saline with Tween20
TFGβ	transforming growth factor β
TNF	tumour necrosis factor
TNFR1	TNFα receptor
TRADD	TNFR1-associated via death domain
TRAF6	TNF receptor-associated factor 6
TRAP	tartrate resistant acid phosphatase
UBA	ubiquitin associated
VCP	valosin-containing protein
VDR	vitamin D receptor

WIF	Wnt inhibitory factor
WISP	WNT1 inducible signalling pathway protein

Selected amino acid abbreviations

D	aspartic acid
G	glycine
K	lysine
L	leucine
P	proline
R	arginine
X	truncation mutation/stop codon