

# Molecular modulation of osteocalcin and its relevance in diabetes (Review)

JOSÉ RAFAEL VILLAFÁN-BERNAL, SERGIO SÁNCHEZ-ENRÍQUEZ  
and JOSÉ FRANCISCO MUÑOZ-VALLE

Departamento de Biología Molecular y Genómica, Centro Universitario de Ciencias de la Salud,  
Universidad de Guadalajara, Colonia Independencia, CP 44340 Guadalajara, Jalisco, Mexico

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**Abstract.** Osteocalcin (OC) is encoded by the bone  $\gamma$ -carboxyglutamate (Gla) protein (*BGLAP*) gene, and it is released by osteoblasts during osteogenesis. Its expression can be modulated by growth factors, hormones, cytokines and physical stimuli via signal transduction pathways, binding to the *BGLAP* gene promoter or interactions with nuclear transcription factors. It was recently demonstrated that uncarboxylated OC improves glucose tolerance and insulin sensitivity in mice by increasing the expression and secretion of insulin in  $\beta$ -cells and of adiponectin in adipocytes. Humans with type 2 diabetes have significantly lower serum levels of OC than healthy individuals and indeed, serum OC levels have been inversely correlated with fasting plasma glucose, fasting insulin and the homeostasis model assessment of insulin resistance (HOMA-IR) index. Moreover, several drugs have been shown to influence OC expression and its serum concentration. This review summarizes the molecular mechanisms involved in the modulation of OC expression, and discusses the potential relevance of OC in the pathogenesis and treatment of diabetes.

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*Correspondence to:* Dr José Francisco Muñoz-Valle, Sierra Mojada 950, Colonia Independencia, CP 44340 Guadalajara, Jalisco, Mexico

E-mail: biologiamolecular@hotmail.com

**Abbreviations:** OC, osteocalcin; cOC, carboxylated OC; ucOC, uncarboxylated OC; *BGLAP*, bone  $\gamma$ -carboxyglutamate protein; HOMA-IR, homeostasis model assessment of insulin resistance; HbA<sub>1c</sub>, glycated hemoglobin; FPG, fasting plasma glucose; VD, vitamin D; NMP, nuclear matrix proteins; RER, rough endoplasmic reticulum; PTH, parathyroid hormone

**Key words:** osteocalcin, metabolism, bone, insulin, diabetes

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## 1. Introduction

Osteocalcin (OC) is the product of the bone  $\gamma$ -carboxyglutamate (Gla) protein (*BGLAP*) gene, which is primarily expressed in osteoblasts and is regulated in response to physiological or pathological processes (1). Growth factors, hormones, cytokines, load and vibration have all been shown to modulate *BGLAP* gene expression through signal transduction pathways or through the interaction of nuclear transcription factors with the *BGLAP* gene promoter (1-5). OC exists in two main forms, the carboxylated (cOC) and uncarboxylated (ucOC) forms. While cOC, that is considered a chemoattractant and an activator of cells with bone resorption properties (6,7), has a high affinity for hydroxyapatite and mineral ions; ucOC improves glucose tolerance and insulin sensitivity in mice, protecting them against the development of diabetes (8). In humans, OC serum levels have been inversely correlated with glycated hemoglobin (HbA<sub>1c</sub>), fasting plasma glucose (FPG), fasting insulin and the homeostasis-model assessment insulin resistance (HOMA-IR) index; also, they are diminished in type 2 diabetes (9-11). Several drugs used in both clinical and non-clinical settings influence the serum levels of OC (12-15) and hence, might be of potential utility for the treatment of type 2 diabetes.

Here, we present an overview of the organization of the *BGLAP* gene and of the expression of OC; in addition, we describe how OC expression is modulated by various compounds. A synopsis of the biological activities of OC and its activity in humans is presented and finally, we discuss its potential relevance in diabetes pathogenesis and treatment.

## 2. *BGLAP* gene organization

The human *BGLAP* gene is a DNA sequence of ~1700 bp that is located at chromosome 1q25-q31 (GenBank Acc. no. X04143.1), and which shares a high degree of homology

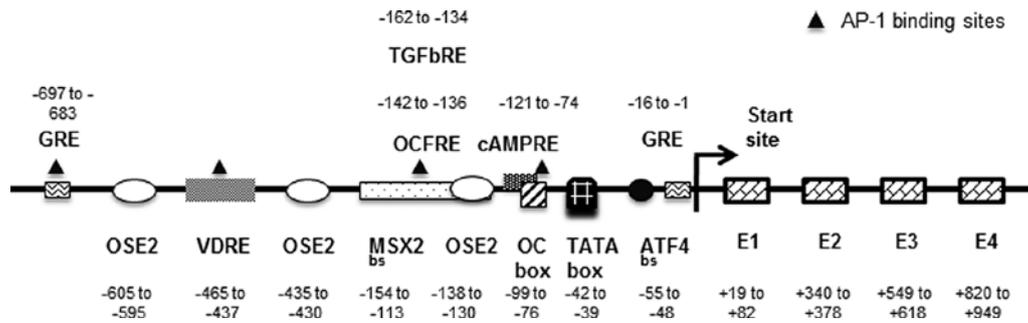


Figure 1. Schematic representation of the organization of the *BGLAP* gene. The *BGLAP* gene exhibits the same basic organizational features of most eukaryotic genes. A TATA box is located in the upstream region (at -42 to -39) of exon 1 and the OC box is a sequence of 24 nucleotides with a CAAT motif located in the proximal promoter region. There are three RUNX2 binding sites, named OSE2, a VDRE enhancer sequence located in the distal promoter region and several DNA sequences that are hormone response elements to glucocorticoids (GRE) and cAMP (cAMPRE). The *BGLAP* gene also has DNA sequences that are response elements for FGF (OCFRE) and TGFb (TGFbRE), four AP-1 binding sites and one DNA sequence for MSX2 binding. E1, Exon 1; E2, Exon 2; E3, Exon 3; E4, Exon 4.

with mammalian counterparts from the mouse, rat, monkey and pig (1,16). The upstream sequences associated with the *BGLAP* gene are approximately 700 bp long from the 5'-end to the start site and they exhibit the same basic organizational features of most eukaryotic genes. These include: i) widely conserved DNA sequences; ii) DNA sequences responsive to hormones, cytokines and growth factors; iii) DNA sequences that bind to tissue specific-transcription factors; and iv) enhancer and silencer DNA sequences (5,17-23). Downstream from the start site, the gene contains four exons and three introns that are approximately 1,000 bp long (Fig. 1).

### 3. OC expression

The *BGLAP* gene encoding for OC is expressed efficiently in osteoblasts and odontoblasts, and more weakly in the ovaries, prostate, testes, skeletal muscle, thyroid and other tissues (24). In this section, we will review the expression of the *BGLAP* gene, including its transcription, post-transcriptional processing, as well as the synthesis and release of OC.

*BGLAP* gene transcription and post-transcriptional processing. Transcription of the *BGLAP* gene is regulated temporarily, depending on the stage of osteogenesis and on the development of osteoblast precursors (25). While the *BGLAP* gene is switched off (inactive) during osteoblast proliferation, it is activated during differentiation (25). During this stage, transcriptional activation involves the basal transcription of the *BGLAP* gene as well as vitamin D (VD)-induced *BGLAP* gene transcription, in which the rate of *BGLAP* gene transcription increases 10- to 20-fold (25,26) (Fig. 2). Indeed, there is evidence that a sequence of events culminates in *BGLAP* gene transcription, including structural gene activation, and the binding of the appropriate transcription factors and DNA polymerase to the gene promoter.

Histone acetylation at the OC locus has been observed during bone-specific OC expression (27) and the interaction of histone deacetylase 3 with RUNX2 is thought to repress the OC promoter (28). Thus, structural activation of the *BGLAP* gene appears to involve histone acetylation by histone acetyltransferases and simultaneous inactivation of histone deacetylases (27,28). Once gene activation occurs, initiation

of basal *BGLAP* gene transcription requires the binding of the specific elements to its cognate DNA sequences, including: i) basal transcription factors TFIID, TFIIA, TFIIB, TFIIF, TFIIE and TFIIH; ii) the osteoblast specific transcription factor RUNX2; iii) the OC box binding protein; and iv) RNA polymerase II (25,29-32).

Basal *BGLAP* gene transcription can be enhanced by activators and co-activators. The most important activator is the VD-vitamin D receptor-retinoid X receptor complex (VD-VDR-RXR), which increases the rate of *BGLAP* transcription by binding to an enhancer sequence located at the distal gene promoter (26,30,33). The binding of VD to VDR induces conformational changes in the receptor, facilitating its interaction with co-activators, such as NCoA-1/SRC-1, NCoA-2/GRIP/TIF2, CBP, p300 and DRIP (31). *BGLAP* gene transcription is repressed by transcription factors such as DLX3, MSX2, DLX5 and MEF, and by the glucocorticoid-glucocorticoid receptor complex. This repression is achieved through interaction with RUNX2 (17,34), binding to OC box (22) and by modulating the expression of other *BGLAP* regulators (35,36), as well as by competition or interference with the binding of other transcription factors to the *BGLAP* gene promoter (37,38). In addition, nuclear matrix proteins (NMP) participate in *BGLAP* gene transcription. Accordingly, NMP-1 and NPM-2 interact transiently with two binding elements for NMP located at the *BGLAP* gene promoter; hence, NMP could be important in concentrating and/or localizing transcription factors that mediate both basal and VD-induced *BGLAP* gene transcription (39,40).

Six OC splice variants exist, such variants retain distinct combination of introns, although a dominant transcript containing all 3 introns has been detected by RT-PCR. These 6 variants encode proteins of 57 to 100 amino acids (aa) that are identical to pre-pro-OC in the N-terminal domain. However, only in bone-related tissue are all 3 OC introns efficiently spliced out (24).

*OC synthesis and release.* Human OC synthesis occurs after the *BGLAP* gene is transcribed and processed to OC mRNA, that is then translated in the rough endoplasmic reticulum (RER) to pre-pro-OC, a protein precursor of 98 aa residues (1,16,41,42). Subsequently, pre-pro-OC undergoes proteolysis

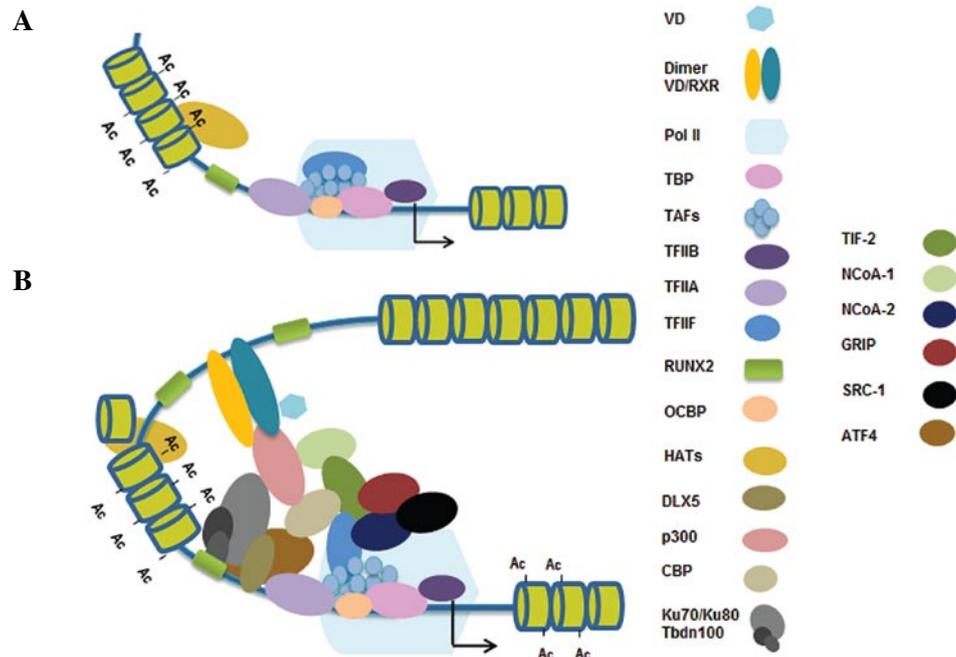


Figure 2. Basal and vitamin D (VD)-induced *BGLAP* gene transcription. (A) The basal *BGLAP* gene transcription that occurs when the TATA box, osteocalcin (OC) box and OSE2 are occupied by appropriate transcription factors. (B) The VD-induced *BGLAP* gene transcription due to the contribution of additional activators and co-activators.

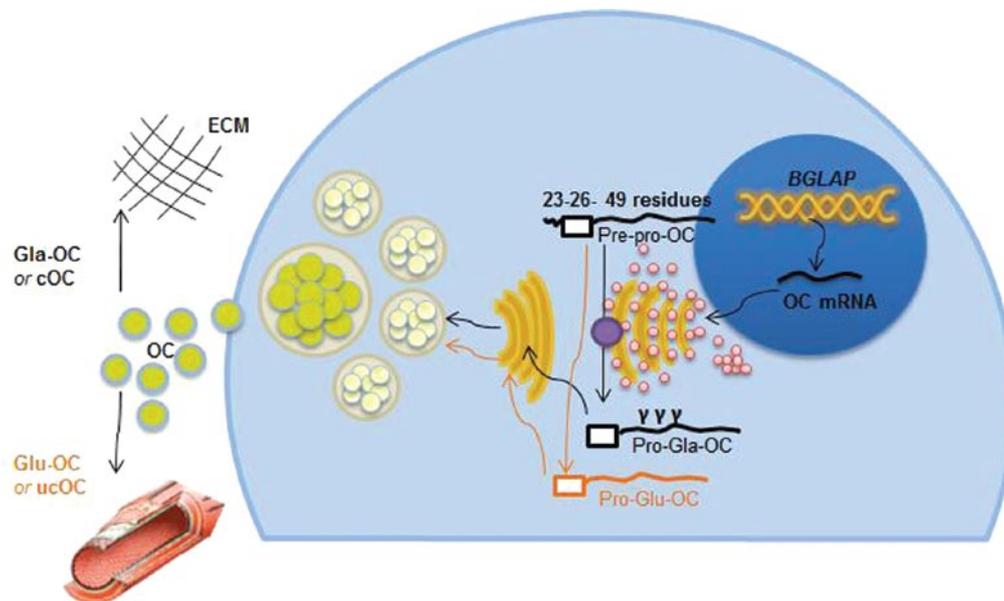


Figure 3. Osteocalcin (OC) synthesis and release. Human OC mRNA is translated in the rough endoplasmic reticulum (RER) as pre-pro-OC. pre-pro-OC is proteolytically cleaved to generate a pre-peptide and pro-OC. pro-OC is then directed to the RER membrane where GGCX carboxylates a proportion of pro-OC, at glutamic acid residues 17, 21 and 24, giving rise to pro-Gla-OC. The remaining pro-OC is not carboxylated and is referred to as pro-Glu-OC. Subsequently, pro-Gla-OC and pro-Glu-OC are cleaved to Gla-OC or cOC, and Glu-OC or ucOC. These two main forms are subsequently released, and a fraction of both reaches the bloodstream where they can be measured. Finally, the cOC is mainly deposited at the bone matrix where it interacts with calcium and hydroxyapatite, whereas ucOC acts on pancreatic and adipose tissues.

to form a pre-peptide (23 aa) and pro-OC (75 aa) (1,16,41,42). The pre-peptide then directs pro-OC to the RER membrane where  $\gamma$ -glutamyl carboxylase (GGCX) carboxylates glutamic acid residues at the 17, 21 and 24 positions, resulting in the formation of Gla-residues in a vitamin-K-dependent process (43,44). However, this process only occurs in a proportion

of newly synthesized pro-OC (1,43). Pro-OC is ultimately subjected to a final proteolytic process that produces a pre-peptide (26 aa) and cOC or ucOC. Both forms are released from osteoblasts in a process which is calcium-dependent and relies on the activity of calcium and potassium channels (Fig. 3) (45,46).

Table I. Modulators of *BGLAP* gene expression and the level of their modulation.

Molecule	Effect	Level of modulation	Experimental model	References
Growth factors				
IGF-1	↑ OC production; ↑ OC mRNA, stabilizes OC mRNA; ↑ VD-induced OC secretion and reversal of the inhibitory effect of dexamethasone	Transcriptional, post-transcriptional	IGF-1 addition to mouse long bone culture and to the human osteosarcoma cell line MG-63	64,66
TGF-β	↓ OC production	Transcriptional	TGF-β addition to mouse long bone culture; rat osteosarcoma cell line ROS 17/2.8; human osteosarcoma cell line MG-63	4,18,66
bFGF	↑ OC mRNA (independent of both VD and retinoic acid activation of the OC promoter)	Transcriptional	bFGF transfection of rat osteosarcoma cell line ROS 17/2.8; bFGF injection to ovariectomized rats <i>in vivo</i>	23,67
BMP-2	↑ OC mRNA	Transcriptional	BMP-2 transfection of human periosteal cells; addition of rBMP-2 to fetal rat calvarial osteoblasts KS483 and to rat primary osteoblastic cells; subcutaneous injection of BMP-2 to mice	68-70
BMP-7	↑ OC mRNA; inhibits osteoblast proliferation and induces differentiation	Transcriptional	BMP-7 addition to the rat osteosarcoma cell line ROS 17/2.8	65
Hormones				
Parathyroid hormone	↑ OC mRNA through an increase in cAMP and PKA which then activates ATF4, with a weaker requirement for the protein kinase C and the MAPK/ERK pathways	Transcriptional	PTH subcutaneous injection to C57BL/6J mice; SaOS-2 osteosarcoma cell line transfected with the OC promoter with addition of PTH to the culture; MC-4 cells or osteoblasts transfected with the OC promoter + PTH addition to the culture	74,75
Calcitonin	↑ OC mRNA <i>in vitro</i> while it has no effect <i>in vivo</i>		Addition of calcitonin to mouse osteoblastic MC3T3-E1 cells; calcitonin administration to rats	76,78
Leptin	↓ OC expression	Unknown	Leptin and leptin receptor deficient mice have increased bone formation and OC expression	79,80
Vitamin D	↑ <i>BGLAP</i> gene transcription <i>in vitro</i> and <i>in vivo</i> , thereby increasing OC mRNA, OC synthesis and release (has an opposite effect in the mouse)	Transcriptional	VD addition to cultures (fetal rat calvarial osteoblasts, ROS 17/2.8) and supplementation to animals	3,33,101
Glucocorticoids	Repression of <i>BGLAP</i> gene expression by direct binding to the OC promoter, thereby decreasing OC mRNA <i>in vivo</i> and <i>in vitro</i>	Transcriptional	Addition of dexamethasone to rat osteosarcoma cells ROS 17/2.8; dexamethasone administration to rats	38,88, 95,101
Estrogens	Repression of <i>BGLAP</i> gene expression	Transcriptional	Administration of estradiol dipropionate to ovariectomized rats and measurement of OC mRNA at the femur	90,91
Progesterone	↑ OC mRNA	Transcriptional	Addition of progesterone to fetal rat calvarial osteoblasts	69,70,92

Table I. Continued.

Molecule	Effect	Level of modulation	Experimental model	References
Thyroid hormone	↑ OC mRNA and protein levels	Transcriptional	Addition of triiodothyronine (T3) to MC3T3-E1 cells; systemic TSH administration	93
Retinoic acid	Enhances the effect of VD on <i>BGLAP</i> gene expression and OC synthesis	Transcriptional	Addition of retinoic acid to rat osteosarcoma cells ROS 17/2.8	89
Cytokines				
IL-1	↓ <i>BGLAP</i> gene expression	Transcriptional	Addition of IL-1 to human osteoblast-like cultures and rat osteoblasts	2
Physical stimuli				
Load	↑ <i>BGLAP</i> gene expression	Transcriptional	Osteoblast differentiation induced by mechanical forces; compressive forces to rat calvarial osteoblasts	47,48
Vibrations	↑ <i>BGLAP</i> gene expression	Unknown	Broad frequency vibration applied to MC3T3-E1 cells	50
Matrix-cell interactions				
Collagen- $\alpha_2\beta_1$ integrin interaction	↑ <i>BGLAP</i> gene expression	Probably post-translational or via an accessory factor	Addition of anti- $\alpha_2$ integrin antibody to MC3T3-E1 cells blocks ascorbic acid-dependent induction of the <i>BGLAP</i> gene promoter	58,59,61
Osteopontin- $\alpha\nu$ integrins	↑ <i>BGLAP</i> gene expression	Unknown	Addition of anti-osteopontin monoclonal antibody inhibited OC expression in MC3T3-E1 cells; addition of the Arg-Gly-Asp peptidomimetic compound or rOPN increased <i>BGLAP</i> gene expression	60
Fibronectin- $\alpha_5\beta_1$ , $\alpha_3\beta_1$ and $\alpha_8\beta_1$ integrins	↑ <i>BGLAP</i> gene expression	Unknown	Addition of anti-integrin $\alpha_5\beta_1$ , $\alpha_3\beta_1$ and $\alpha_8\beta_1$ antibodies to fetal calvarial osteoblasts decreases OC mRNA	62

↑, increase; ↓, decrease. IGF-1, insulin-like growth factor 1; OC, osteocalcin; TGF- $\beta$ , transforming growth factor- $\beta$ ; bFGF, basic fibroblast growth factor; BMP-2, bone morphogenetic protein-2; BMP-7, bone morphogenetic protein 2; PTH, parathyroid hormone; VD, vitamin D.

#### 4. Modulation of OC expression

OC expression is regulated by growth factors, hormones, cytokines and physical stimuli, such as load and vibration (Fig. 4, Table I).

*Physical stimuli modulate OC expression through Wnt/ $\beta$ -catenin and matrix-cell interactions.* Mechanical loading and vibrations have been shown to influence *BGLAP* gene expression. Simulated mechanical loading significantly increases OC expression both *in vivo* and *in vitro* (47,48). Likewise, broad frequency vibration and dynamic stress (to simulate vibration) increase OC gene expression in cell culture (49,50). These changes in OC expression in response to mechanical loading are mediated, at least in part, by

activation of signal transduction pathways such as the Wnt/ $\beta$ -catenin, MAPK and extracellular matrix-integrins-nuclear pathways (20,51,52).

In the canonical Wnt/ $\beta$ -catenin pathway, Wnt (ligand) binding to its receptor causes  $\beta$ -catenin stabilization and its accumulation in the cell cytosol. This  $\beta$ -catenin then translocates to the nucleus where it associates with the Tcf/LEF family of transcription factors, consequently regulating the expression of canonical Wnt target genes (53,54) like *Runx2* (55). Subsequently, RUNX2 (the product of *Runx2*) interacts with three DNA sequences in the *BGLAP* gene promoter to maintain its basal transcription (26,30).

The bone extracellular matrix is composed of many proteins, including, OC, collagen, elastin, fibronectin, osteonectin, osteopontin and bone sialoprotein (56,57). Such proteins

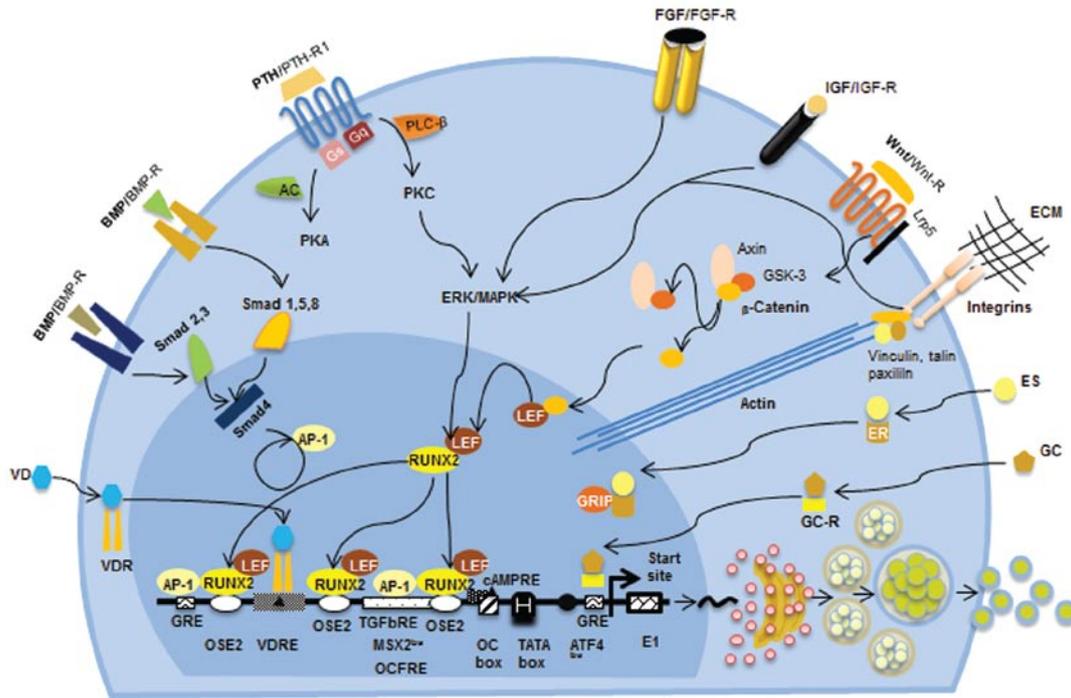


Figure 4. Main signal transduction pathways and protein/DNA interactions involved in modulating *BGLAP* gene expression. Expression of the *BGLAP* gene can be modulated by hormones, cytokines, matrix-cell interactions and growth factors via signal transduction pathways, binding to the *BGLAP* gene promoter or interactions with nuclear transcription factors. The signal transduction pathways involve the PKC/PKA, Smad, ERK/MAPK and the Wnt/ $\beta$ -catenin pathways. VD, vitamin D; VDR, VD receptor; GC, glucocorticoids; GR, glucocorticoid receptor; ES, estrogens; ER, estrogen receptor.

interact with integrins at the cell membrane acting as signals that can modify gene expression, including that of *BGLAP*, through the activation of cytoskeletal and intracellular proteins (actin, talin, vincullin, paxillin and MAPK) (20,58-60). In fact, OC expression can be modified by exposing cells in culture to blocking antibodies, proteins or peptides directed against specific integrin subunits or matrix proteins (59-62).

*Growth factors modulate OC expression through BMP/TGF- $\beta$  signaling.* Growth factors are essential for cell proliferation and organ growth, they are also capable of modulating OC expression (63). The transforming growth factor- $\beta$  (TGF- $\beta$ ) keeps the *BGLAP* gene inactive during the proliferation stage of osteogenesis (18,20). In contrast, the bone morphogenetic protein-7 (BMP-7) and the insulin-like growth factor (IGF-1) induce differentiation and OC expression during the mineralization stage of osteogenesis (64-66). Other growth factors, including the basic fibroblast growth factor (bFGF, also called FGF2), and BMP-2 (67,68) can also modulate OC expression.

Growth factors mediate their effects on OC expression via the BMP/TGF- $\beta$  signaling pathway. Briefly, the binding of BMP to its membrane receptor in osteoblasts triggers a signaling cascade that activates Smad 1, 5 and 8 by phosphorylation. These members of the family form a complex with Smad 4 (also called C-Smad), which is subsequently translocated to the nucleus where it interacts with other transcription factors such as RUNX2 to modulate the activity of target genes including *BGLAP* (69-72). Conversely, TGF- $\beta$  binding to its membrane receptor results in activation of Smad 2 and Smad 3, which form a stable complex with Smad 4. This also translocates to the nucleus and interacts

with transcription factors such as AP-1, VDR and p300/CBP, thereby positively or negatively modulating *BGLAP* gene transcription (20,31,73).

*Peptide hormones modulate OC expression.* Peptide hormones, including the parathyroid hormone (PTH), calcitonin and leptin, are involved in the modulation of *BGLAP* gene expression (74-78). PTH binds to its specific cell membrane receptor and activates several signal transduction pathways, including those mediated by protein kinase A (PKA), protein kinase C (PKC) and MAPK/ERK, resulting in increased *BGLAP* gene expression (see below). Calcitonin stimulates OC expression in a mouse osteoblastic cell line, whereas no such effect is observed in the appendicular and axial skeleton of the rat *in vivo* (76,78). Leptin modifies osteoblast function and down-regulates OC expression through a hypothalamic relay that is mediated by the sympathetic nervous system (79,80).

*PKA/PKC signaling modulates OC expression.* As indicated, PKA and PKC signaling is activated by PTH and binding of PTH to its G-protein-coupled receptor activates adenylyl cyclase and phospholipase C, thereby increasing the intracellular levels of cAMP and/or  $Ca^{2+}$ . Increased levels of these second messengers activate PKA and PKC, which in turn modulate OC expression (74,81,82).

*The MAPK-ERK signaling pathway is a common point of modulation in OC expression.* The extracellular signal regulated kinase (ERK)-mitogen-activated protein kinase (MAPK) pathway represents an important link between the cell surface and the nucleus, and it influences proliferation and

differentiation (83,84). MAPK signaling up-regulates *BGLAP* gene expression through RUNX2 activation, increasing OC mRNA levels (85). This signaling pathway is activated by cell-matrix interactions, cytokines, osteogenic growth factors, such as BMPs and bFGF, mechanical loading and hormones such as PTH (51). In each case, the common event in this cascade is the activation of MAPK, which in turn phosphorylates transcription factors and other proteins, such as RUNX2, ultimately modulating gene expression and other cell functions (86).

*Lipid hormones and thyroid hormones modulate OC expression by interacting with the BGLAP gene promoter and transcription factors.* Lipid-derived hormones are also implicated in the modulation of *BGLAP* gene expression. These include VD, glucocorticoids, estrogens, progesterone, triiodothyronine and retinoic acid. Lipid-derived hormones enter the cell to form complexes with specific cytoplasmic or nuclear receptors. These complexes then bind to specific DNA sequences of the *BGLAP* gene promoter or they interact with transcription factors (repressors or activators), which in turn modulate the rate of transcription (3,33,37,69,87-97).

*Drugs that up-regulate OC expression or serum levels.* Several drugs are known to increase OC expression or serum levels. Glipizide and glybenclamide (blockers of ATP-dependent potassium channels) increase VD-induced OC secretion in cell cultures (45). This suggests that OC is released in osteoblasts by a mechanism similar to that which regulates insulin secretion in pancreatic  $\beta$ -cells (46). The bisphosphonates olpadronate (OPD) and IG-9402, increase OC release through the opening of voltage-dependent  $\text{Ca}^{2+}$  channels in the ROS 17/2.8 cell line, an effect abolished by  $\text{Ca}^{2+}$  channel blockers such as nifedipine and verapamil (15). The active form of VD,  $1\alpha$ -25(OH) $_2$ D $_3$  (VD3), increases OC expression and release in osteoblasts. In fact, two VD metabolites, [25-OH-16,23E-diene-D $_3$ (R) and  $1\alpha$ ,25-OH(2)-16,23E-diene-D $_3$ (A)], are 10 times more potent than VD3 in increasing OC mRNA expression and protein secretion (14). In addition TAK-788, which promotes osteoblast differentiation, increases OC release in rat bone marrow stromal cells (98). Genistein (a natural isoflavone phytoestrogen) significantly increases OC serum concentrations within 6-12 months of its administration to early postmenopausal women (99).

*Drugs that down-regulate OC expression or serum levels.* Other drugs demonstrably decrease OC expression or serum levels. The administration of glucocorticoids to humans decreases OC serum concentrations by decreasing OC expression and release (12,37,100,101). Hydrochlorothiazide, a thiazide diuretic, decreases the serum concentration of OC by inhibition of *BGLAP* gene transcription via the c-fos transcription factor (13). In addition, administration of low-doses of the antiepileptic drug levetiracetam to female rats over 90 days significantly reduces OC serum concentrations (102).

## 5. Biological activities of OC

Since its discovery in chicken bone (103), OC has attracted considerable attention, leading to the elucidation of a number of biological roles for this molecule.

*Roles of cOC in bone.* Human cOC is a protein of 49 aa with carboxyglutamic acid (Gla) residues at positions 17, 21 and 24 (43). It is produced during the mineralization stage of osteogenesis and it possesses high affinity for hydroxyapatite and mineral ions like calcium (103). However, loss and gain-of-function experiments have failed to identify a role for OC in extracellular matrix mineralization *in vivo* (104,105). cOC is considered to be an activator and a chemoattractant molecule in cells with bone resorption properties (6,7). Interestingly, OC-deficient mice exhibit increased bone formation (104), suggesting a central role for cOC in the regulation of bone remodeling.

*Endocrine actions of ucOC.* The first data on the endocrine action of OC were reported in 2007. In their study, Lee *et al* demonstrated, both in cell culture and in mice, that ucOC in the pancreas increased  $\beta$ -cell proliferation, insulin expression and release, resulting in improved glucose tolerance (8). In addition, ucOC increased adiponectin expression and secretion in adipose tissue, which in turn enhanced insulin sensitivity (8). Furthermore, OC (-/-) mice show an increase in fat mass, adipocyte number and serum triglyceride levels, whereas *Esp* (-/-) mice, which lack a protein responsible for down-regulation of OC expression and show high OC serum levels were protected against diet-induced obesity and diabetes (8). In 2008, the same group demonstrated that continuous infusion of recombinant OC (rOC) in wild-type mice i) improved glucose tolerance (at 0.03 and 0.3 ng/h) and insulin sensitivity (at 30 ng/h) and ii) increased insulin secretion (at 0.03 and 0.3 ng/h). In addition, continuous infusion of rOC to mice fed a high-fat diet protected from weight gain and glucose intolerance and maintained triglyceride serum levels within the normal range (106).

## 6. Evidence for the association of OC with markers of glycemic status in humans

Well before the findings of Lee *et al* (8), lower serum concentrations of total OC were described in children with type 1 diabetes (107). Several studies showed lower total OC serum concentrations in patients with type 2 diabetes when compared to healthy subjects (9,108-112). Actually, it was demonstrated that improved glycemic control in poorly controlled diabetic patients increased total OC serum concentrations and decreased HbA $_{1c}$  (108,113,114). Furthermore, the total OC levels were inversely correlated with HbA $_{1c}$  (115).

Up to this point, clinical studies had focused on the effects of diabetes on bone. However, the discovery of endocrine actions of ucOC in mice has focused research efforts on the possible role of OC in glycemic control in humans, prompting cross-sectional studies to identify the association of OC with type 2 diabetes and markers of the glycemic status. A recent study in Korean diabetic postmenopausal women found an inverse correlation between total OC and FPG, HbA $_{1c}$ , fasting insulin and insulin resistance. Moreover, participants with total OC in the highest quartile had significantly lower FPG and HbA $_{1c}$  (10). Furthermore, a study of Japanese men and postmenopausal women with type 2 diabetes also demonstrated an inverse correlation between total OC concentrations, FPG and HbA $_{1c}$ , and a positive correlation between

total OC and total adiponectin, a hormone that enhances insulin sensitivity (11).

In healthy subjects, the association of OC with indicators of glycemic status and insulin resistance has also been studied. Among middle-aged males, elevated serum cOC and ucOC levels are associated with improved glucose tolerance and insulin sensitivity, as well as enhanced  $\beta$ -cell function (116). In healthy older men and women, elevated cOC concentrations are associated with lower insulin resistance (117). In men and women aged 65 and older (only 5% with diabetes), total OC concentrations were inversely correlated with FPG, fasting insulin, HOMA-IR, high-sensitivity C-reactive protein, IL-6, body mass index and body fat (118). In addition, participants with total OC concentrations within the highest tertile had lower FPG concentrations than those in the lowest tertile (118). Furthermore, higher total OC concentration at onset was associated with a significant attenuation in the observed increase in FPG after a 3-year follow-up (118).

## 7. Discussion

OC has become a subject of increased interest in recent years based on its role in glycemic control in mice and the clinical evidence supporting a similar role in humans. However, many questions must be answered before its complete role in human metabolic control is fully understood. In addition, it remains to be determined if drugs that increase OC expression, or its release or accumulation in serum, could modulate glycemic control. Conversely, it must be determined whether those drugs that inhibit OC activity should be avoided in patients with type 2 diabetes.

It is well known that glucocorticoids increase glucose serum levels by enhancing gluconeogenesis and that they should thus be avoided in healthy and diabetic subjects (119). However, their negative effects on OC expression and serum concentrations may provide further reason for caution. Thiazide diuretics, such as hydrochlorothiazide can impair glucose metabolism and exacerbate new-onset diabetes (120). Their administration to both diabetic and non-diabetic subjects has negative effects on pancreatic  $\beta$ -cell and glucose-induced insulin secretion, as well as on OC expression and serum levels (13,120). Thus it is possible that the impairment of glucose metabolism by thiazide diuretics may in part be due to impaired OC expression. Levetiracetam has not been associated with impaired glucose metabolism or diabetes; however, given that it significantly reduces OC serum concentrations in rats (102), its metabolic effect in humans should be determined. Glybenclamide and glipizide are effective in treating type 2 diabetes, and it is possible that their positive effects on glycemic control are in part due to an increase in OC serum concentrations, which could add therapeutic value to these drugs.

Given that bisphosphonates increase OC release *in vitro* (15), it may be interesting to test whether they have positive metabolic effects in postmenopausal type 2 diabetic women, in addition to their beneficial effects on bone mineral density. Conversely, it will be important to discount any deleterious effect of  $\text{Ca}^{2+}$  channel blockers on OC serum levels *in vivo* given their inhibition of OC release (15). Moreover, in light of the effects of VD and its metabolites, as well as TAK-788 and genistein, on OC expression and serum concentrations, it

would be of interest to quantify their effects on glucose serum levels (14,98,99). Some drugs targeting insulin signaling pathways are currently under development to treat type 2 diabetes (121-124). Thus, it may be valuable to test whether compounds targeting signaling pathways modulating OC expression and release could have a positive effect on glycemic control or the management of type 2 diabetes.

To date, neither the OC receptor nor the OC signaling pathways have been identified. The elucidation of these central components of OC activity on beta cells and adipocytes is vital, not only to fully understand the biological roles of OC but also to define additional therapeutic approaches to treat type 2 diabetes.

Dipeptidyl peptidase-IV (DPP-IV) is an enzyme that regulates the half-life of the incretin, glucagon-like peptide-1 (GLP-1). Inhibitors of DPP-IV are used to treat type 2 diabetes (121,125). Given that OC, like GLP-1, increases  $\beta$ -cell proliferation and insulin synthesis and release, increasing the half-life of OC through the inhibition of modulatory proteases may potentially improve glycemic control in type 2 diabetes.

We previously found that patients with type 2 diabetes have a lower ucOC/cOC index than healthy subjects and that this value was negatively correlated with FPG and HOMA-IR levels. Subjects with an ucOC/cOC index within the highest quartile have lower FPG and HOMA-IR values than those in the lowest quartile (126). It remains to be determined whether any changes in  $\gamma$ -glutamyl carboxylase activity occurs in type 2 diabetes, or whether other factors might mediate the observed alterations in ucOC/cOC.

Lower OC levels have been widely reported in patients with type 2 diabetes (9,108-112). However, it remains to be determined if low ucOC concentrations in type 2 diabetes represent a cause and/or a consequence of hyperglycemia. A deleterious effect of hyperglycemia on the bone has been suggested (113,127,128), while a protective effect has been ascribed to high concentrations of OC in humans (118). As such, a causal effect of OC in type 2 diabetes cannot be discounted. In addition, it remains unclear which forms of OC, ucOC, cOC or both, participate in the glycemic control in humans.

In conclusion, there is ample evidence supporting the participation of OC in the glycemic control and energy metabolism. Positive modulation of OC serum concentrations could be valuable for the prevention, delay and treatment of diabetes, while, drugs that have deleterious effects on OC serum levels and release should be used with caution.

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