

## Effect of zinc on gene expression in osteoblastic MC3T3-E1 cells: enhancement of Runx2, OPG, and regucalcin mRNA expressions

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**Abstract** The effect of zinc sulfate on the mRNA expressions in Runx2, osteocalcin,  $\alpha 1(I)$  collagen, insulin-like growth factor-I (IGF-I), transforming growth factor- $\beta 1$  (TGF- $\beta 1$ ), osteoprotegerin (OPG), regucalcin, zinc transporter 1 (ZIP1), or glyceraldehyde-3-phosphate dehydrogenase (G3PDH) in osteoblastic MC3T3-E1 cells in vitro was investigated. Cells with subconfluency were cultured for 48 h in a medium containing either vehicle or zinc sulfate ( $10^{-6}$ – $10^{-4}$  M) without fetal bovine serum. Culture with zinc sulfate ( $10^{-5}$  M) caused a significant increase in Runx2, OPG, or regucalcin mRNA expressions in the cells, while it did not have a significant effect on osteocalcin,  $\alpha 1(I)$  collagen, IGF-I, TGF- $\beta 1$ , ZIP1, or G3PDH mRNA expressions. The effect of zinc sulfate ( $10^{-4}$  M) in increasing Runx2 mRNA expression was seen at 24–72 h after culture. A significant increase in OPG mRNA expression was observed at 24 or 48 h after culture. Regucalcin mRNA expression was significantly increased at 48 or 72 h after culture with zinc sulfate ( $10^{-4}$  M). The stimulatory effects of zinc sulfate on Runx2, OPG, or regucalcin mRNAs were significantly prevented in the presence of cycloheximide ( $10^{-7}$  M), an inhibitor of protein synthesis, or 5,6-dichloro-1- $\beta$ -D-ribofuranosyl-benzimidazole ( $10^{-6}$  M), an inhibitor of transcription activity. Culture with  $\beta$ -alanyl-L-histidinato zinc ( $10^{-5}$  M) caused a significant increase in Runx2 or regucalcin mRNA expressions, while zinc acexamate

( $10^{-5}$  M) did not have a significant effect on Runx2, OPG, ZIP1, or regucalcin mRNA expressions. This study demonstrates that zinc sulfate has a role in the enhancement of Runx2, OPG, or regucalcin mRNA expression in osteoblastic cells in vitro, suggesting its role in the regulation of gene expression in the cells.

**Keywords** Zinc · Runx2 · OPG · Regucalcin · Osteoblast · Bone formation

### Introduction

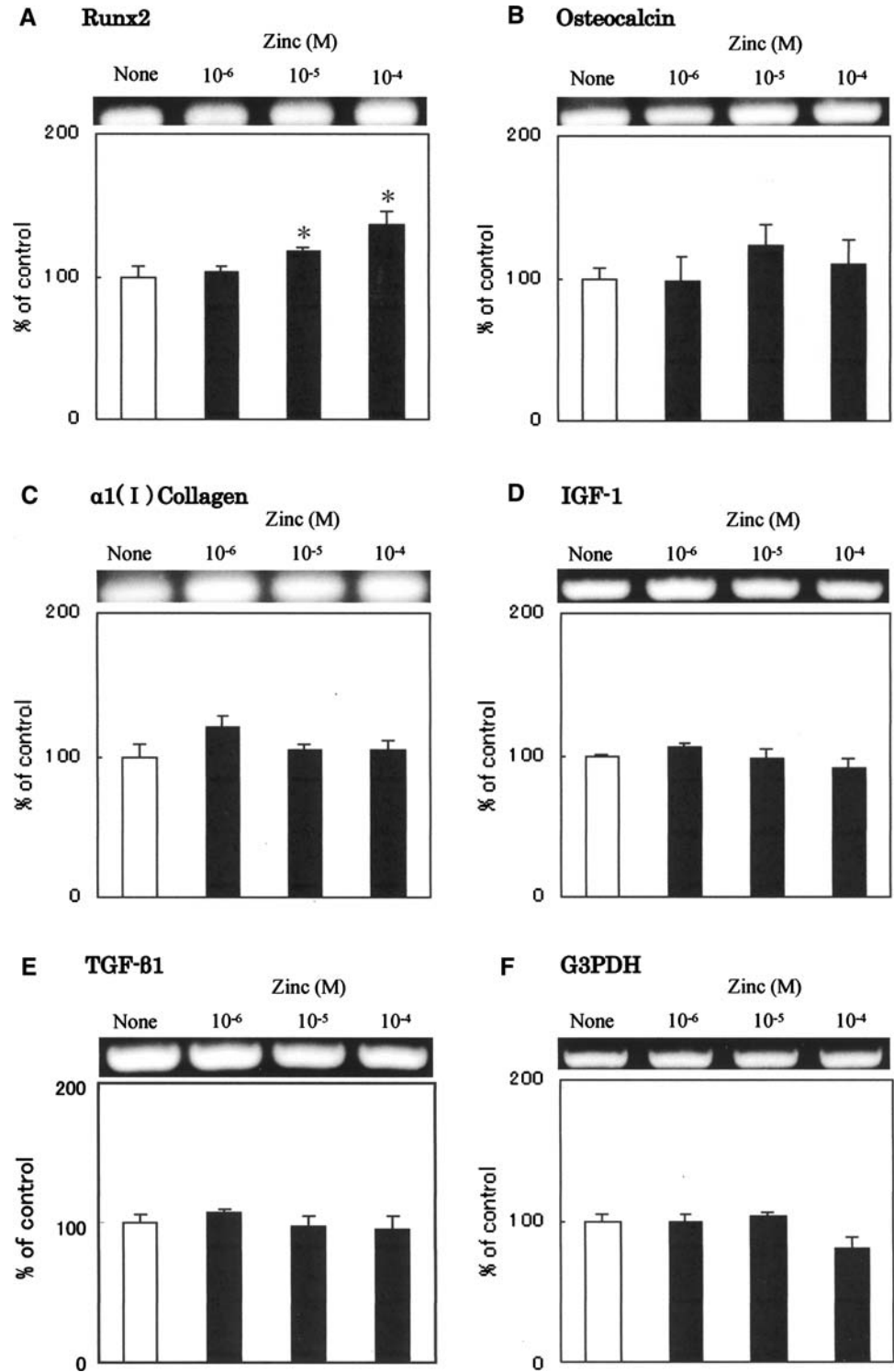
Zinc is known to be an essential trace element for the growth of human and animals [1, 2]. Zinc deficiency results in a retardation of bone growth [3, 4], indicating that the element is required for the growth, development, and maintenance of healthy bone. The pathophysiologic role of zinc in osteopenia and osteoporosis has also been shown. Bone zinc content is reduced with increasing age [5] and skeletal unloading in rats [6]. Osteoporosis patients have been shown to have lower levels of skeletal zinc than normal individuals [7]. Women with osteoporosis excrete a great amount of zinc in urine [8]. Zinc supplementation has been shown to have a preventive and therapeutic effect on bone loss [9, 10]. Zinc has been shown to have a role as a nutritional and pharmacologic tool in the prevention of osteoporosis with increasing age [11, 12].

Zinc can stimulate osteoblastic bone formation [13–15] and inhibit osteoclastic bone resorption [16–18]. Zinc has been shown to have stimulatory effects on alkaline phosphatase activity [19], protein tyrosine phosphatase activity [20], protein [21], and deoxyribonucleic acid (DNA) syntheses [22] in osteoblastic MC3T3-E1 cells, indicating that the metal has anabolic effects on cellular function in vitro.

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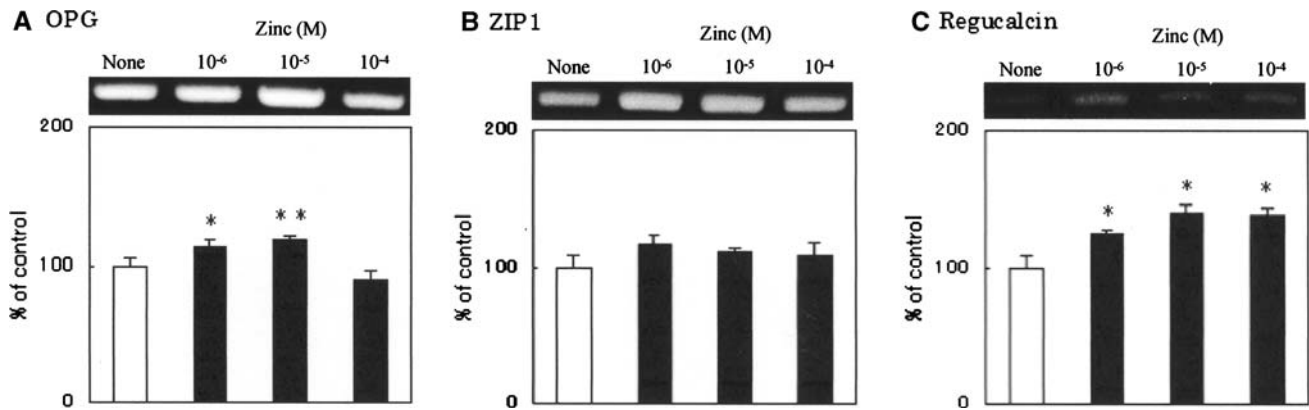
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**Fig. 1** Effect of zinc sulfate on the expression of Runx2, osteocalcin,  $\alpha 1(I)$  collagen, IGF-1, TGF- $\beta 1$ , or G3PDH mRNAs in osteoblastic MC3T3-E1 cells. Osteoblastic cells were cultured for 72 h in a medium containing 10% FBS. Cells with subconfluency were changed to a medium without FBS in the presence or absence of zinc sulfate ( $10^{-6}$ – $10^{-4}$  M). After medium change, cells were cultured for 48 h. Total RNAs (2 or 4  $\mu$ g) extracted from the cells were analyzed using RT-PCR with specific primers. The figure shows one of four experiments with separate culture. The densitometric data for each mRNA level in the cells cultured for 48 h in the presence of zinc sulfate were indicated as % of control (mean  $\pm$  SEM of four experiments). \*  $P < 0.025$  compared with the control (none) value (Student's  $t$ -test)



Zinc has also been shown to inhibit the formation of osteoclastic cells from bone marrow cells [17], and the metal acts on the later stage of differentiation from bone marrow cells to osteoclasts [23]. Zinc can inhibit osteoclastogenesis induced by receptor activator of NF- $\kappa$ B

ligand (RANKL), which plays a pivotal role in differentiation from pre-osteoclast to osteoclasts in bone marrow culture systems [24, 25]. Thus zinc may play a role in the preservation of bone mass due to stimulating bone formation and inhibiting bone resorption [11].



**Fig. 2** Effect of zinc sulfate on the expression of OPG, ZIP1, or regucalcin mRNAs in osteoblastic MC3T3-E1 cells. Osteoblastic cells were cultured for 72 h in a medium containing 10% FBS. Cells with subconfluency were changed to a medium without FBS in the presence or absence of zinc sulfate ( $10^{-6}$ – $10^{-4}$  M). After medium change, cells were cultured for 48 h. Total RNAs (2 or 4  $\mu$ g)

extracted from the cells were analyzed using RT-PCR with specific primers. The figure shows one of four experiments with separate culture. The densitometric data for each mRNA level in the cells cultured for 48 h in the presence of zinc sulfate were indicated as % of control (mean  $\pm$  SEM of four experiments). \*  $P < 0.025$  or \*\*  $P < 0.01$  compared with the control (none) value (Student's *t*-test)

The cellular mechanism of zinc action in osteoblastic cells has not been fully clarified. Zinc has been demonstrated to stimulate proliferation and differentiation in osteoblastic cells [26, 27]. Zinc has stimulatory effects on protein synthesis in osteoblastic cells due to activating aminoacyl-tRNA synthetase, which is a rate-limiting enzyme at transcriptional process [21, 28], and the metal can increase the production of insulin-like growth factor-I (IGF-I), transforming growth factor- $\beta$ 1 (TGF- $\beta$ 1), or osteocalcin in osteoblastic cells, and other protein components [14].

The effect of zinc on gene expression in osteoblastic cells has not been fully determined, however. This study was undertaken to determine whether zinc has a regulatory effect on mRNA expression of various proteins that are expressed in osteoblastic cells in vitro. We paid attention to the gene expressions of Runx2, a stimulator factor in differentiation of osteoblastic cells; osteocalcin and  $\alpha$ 1(I) collagen, a matrix protein in bone; osteoprotegerin (OPG), a suppressor of osteoclastogenesis; IGF-I and TGF- $\beta$ 1, growth factors of osteoblastic cells; ZIP1, a zinc transporter; or regucalcin, a stimulator of osteoclastogenesis and suppressor of osteoblastic bone mineralization in the cells.

We found that culture with zinc has stimulatory effects on the mRNA expressions of Runx2, OPG, or regucalcin expressions in osteoblastic MC3T3-E1 cells in vitro. Zinc may have a role in the regulation of gene expression in osteoblastic cells.

## Materials and methods

### Chemicals

$\alpha$ -Minimal essential medium ( $\alpha$ -MEM) and penicillin-streptomycin (5,000 U/ml penicillin; 5,000  $\mu$ g/ml streptomycin)

were obtained from Gibco Laboratories. Fetal bovine serum (FBS) was obtained from Bioproducts, Inc. 5,6-Dichloro-1- $\beta$ -D-ribofuranosyl-benzimidazole (DRB), cycloheximide were purchased from Sigma Chemicals (St. Louis, MO).  $\beta$ -Alanine-L-histidinato zinc (AHZ) was supplied from Zeria Pharmaceutical Co., Ltd. (Tokyo, Japan). Zinc acexamate (ZA) was obtained from Nipro Corporation (Kusatsu, Japan). Zinc sulfate and other chemicals were of reagent grade and were obtained Wako Pure Chemical Industries (Osaka, Japan). All water used were glass distilled.

### Cell culture

Osteoblastic MC3T3-E1 cells were cultured at 37°C in a CO<sub>2</sub> incubator in plastic dishes containing  $\alpha$ -MEM supplemented with 10% FBS. They were subcultured every 3 days using 0.2% trypsin plus 0.02% EDTA in Ca<sup>2+</sup>/Mg<sup>2+</sup>-free phosphate-buffered saline (PBS). For experiments, about  $1.0 \times 10^5$  cells per dish were cultured for 72 h to obtain subconfluent monolayers in 35-mm plastic containing 2-ml  $\alpha$ -MEM with 10% FBS. After the cells were rinsed with PBS, the medium was exchanged for medium without FBS containing either vehicle or zinc sulfate ( $10^{-6}$ – $10^{-4}$  M), and the cells were cultured further for 24–72 h. Cell viability was estimated by staining with trypan blue. Experiments were repeated using a separate batch of cells with four dishes to ensure reproducibility of results.

### Cell counting

After trypsinization of each of the culture dishes using 0.2% trypsin plus 0.02% EDTA in Ca<sup>2+</sup>/Mg<sup>2+</sup>-free PBS for 2 min at 37°C, cells were collected and centrifuged in a

PBS solution at 100g for 5 min. The cells were resuspended on PBS solution and stained with eosin. Cell numbers were counted under a microscope using a Hemacytometer plate. For each dish, we took the average of two countings.

#### Preparation of RNA

Total RNAs were prepared as described previously [29]. Osteoblastic MC3T3-E1 cells with subconfluency were cultured for 24, 48, or 72 h in a medium containing either vehicle or zinc sulfate ( $10^{-6}$ – $10^{-4}$  M). After culture, cells were washed three times ice-cold PBS, and then cells were homogenized in buffer solution containing 4 M guanidinium thiocyanate, 24 mM sodium citrate (pH 7.0), 0.5% sarcosyl, and isoamyl alcohol, and the phases were separated by centrifugation at 10,000g for 20 min at 4°C. RNA located in the aqueous phase was precipitated with isopropanol at  $-20^{\circ}\text{C}$ . RNA precipitates were pelleted by centrifugation, and the pellets were redissolved in diethylpyrocarbonate-treated water.

#### RT-PCR analysis

Primers for amplification of mouse Runx2 cDNA were: 5'-GTATGAGAGTAGGTGTCCCG-3' (sense strand, positions 992–1,011 of cDNA sequence) and 5'-ACATCCCATCCACTC-3' (antisense strand, positions 1,156–1,175) [30]. The pair of oligonucleotide primers was designed to amplify a 1,834-bp sequence from the mRNA of mouse Runx2.

Primers for amplification of mouse osteocalcin cDNA were: 5'-GGGGAAGGGACAACACATGA-3' (sense strand, positions 188–207 of cDNA sequence) and 5'-TCCTGGACATGGGGATTGA-3' (antisense strand, positions 580–599) [31]. The pair of oligonucleotide primers was designed to amplify a 412-bp sequence from the mRNA of osteocalcin.

Primers for amplification of mouse  $\alpha 1(\text{I})$  collagen cDNA were: 5'-TTCTCCTGGTAAAGATGGTGC-3' (sense strand, positions 2,232–2,252 of cDNA sequence) and 5'-GGACCAGCATCACCTTTAACA-3' (antisense strand, positions 2,466–2,486) [32]. The pair of oligonucleotide primers was designed to amplify a 254-bp sequence from the mRNA of mouse  $\alpha 1(\text{I})$  collagen.

Primers for amplification of mouse IGF-I cDNA were: 5'-GCAAGCTTCAGCCACCTTAC-3' (sense strand, positions 955–974 of cDNA sequence) and 5'-GGGTCGTTTACACAGCAGGT-3' (antisense strand, positions 1,466–1,485) [33]. The pair of oligonucleotide primers was designed to amplify a 531-bp sequence from the mRNA of mouse IGF-I.

Primers for amplification of mouse TGF- $\beta 1$  cDNA were: 5'-CTCTCCACCTGCAAGACCAT-3' (sense strand, positions 633–652 of cDNA sequence) and 5'-CTGCCGTACAACCTCCAGTGA-3' (antisense strand, positions 1,312–1,331) [34]. The pair of oligonucleotide primers was designed to amplify a 699-bp sequence from the mRNA of mouse TGF- $\beta 1$ .

Primers for amplification of decoy receptor OPG were: 5'-CGTTACCTGGAGAT-3' (sense strand, positions 421–435 of cDNA sequence) and 5'-GTTCTACCAAGATT-3' (antisense strand, positions 721–725) [35]. The pair of oligonucleotide primers was designed to amplify a 305-bp sequence from the mRNA of mouse OPG.

Primers for amplification of ZIP1 cDNA were: 5'-GACGTGGTCAGGGACATTAG-3' (sense strand, positions 1,270–1,289 of cDNA sequence) and 5'-AAAGGTGAGGACAGGAGAGG-3' (antisense strand, positions 1,582–1,600) [36]. The pair of oligonucleotide primers was designed to amplify a 331-bp sequence from the mRNA of mouse ZIP1.

Primers for amplification of regucalcin cDNA were: 5'-AGATGAACAAATCCCAGAT-3' (sense strand, positions 618–636 of cDNA sequence) and 5'-TCACCCTGCATAGGATAT-3' (antisense strand, positions 906–924) [37]. The pair of oligonucleotide primers was designed to amplify a 307-bp sequence from the mRNA of mouse regucalcin.

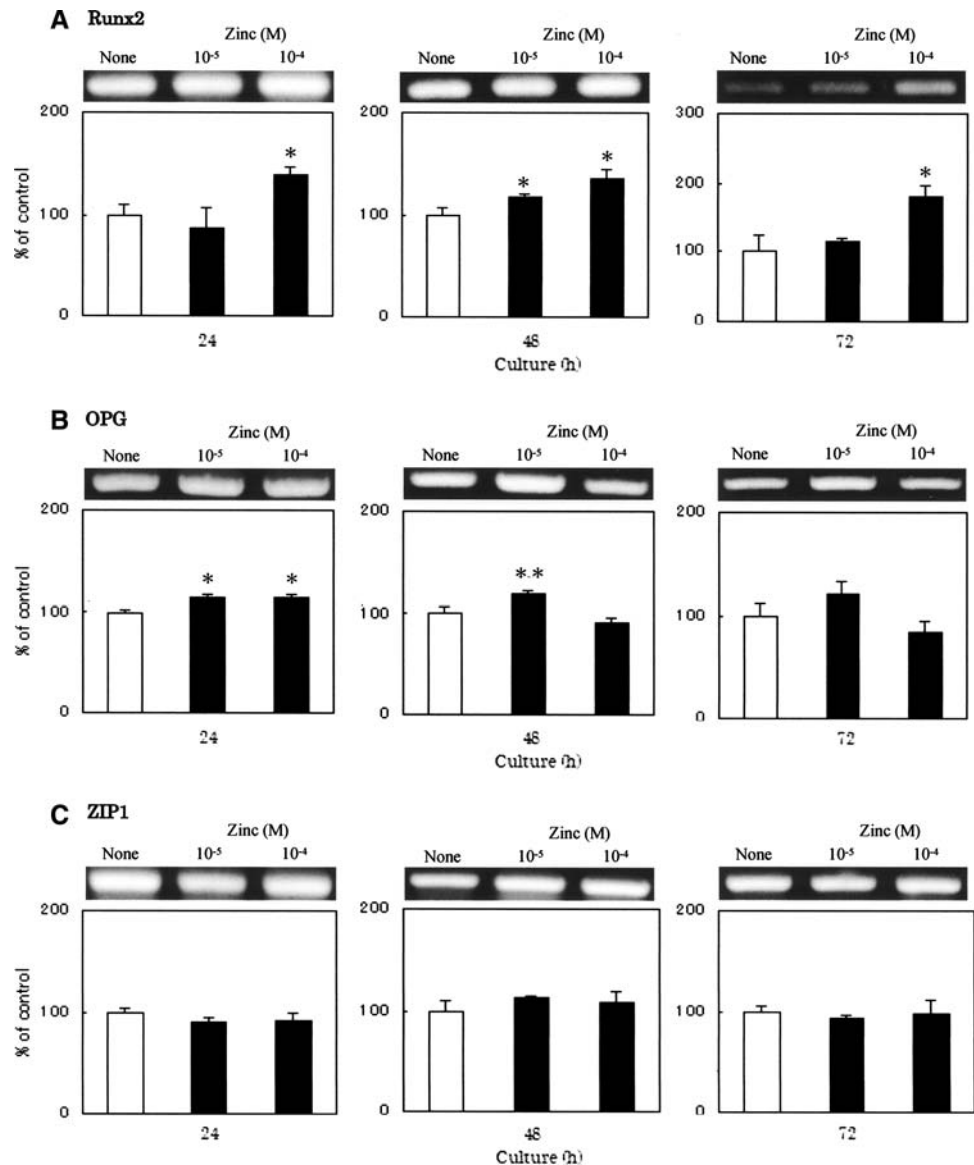
Glyceraldehyde-3-phosphate dehydrogenase (G3PDH) was used as an internal control to evaluate total RNA input. Primers for amplification of G3PDH cDNA were 5'-GATTTGGCCGTATCGGACGC-3' (sense strand) and 5'-CTCCTTGGAGGCCATGTAGG-3' (antisense strand). The pair of oligonucleotide primers was designed to amplify a 977-bp sequence from the mRNA of rat G3PDH.

RT-PCR was performed using reaction mixture (20  $\mu\text{l}$ ) containing 2 or 4  $\mu\text{g}$  of total RNAs, supplied RT-PCR buffer, Titan<sup>TM</sup> enzyme mix (AMV and Expand<sup>TM</sup> High Fidelity), 0.2 mM dNTP, 5 mM dithiothreitol, 5 U RNase inhibitor, and 0.3  $\mu\text{M}$  primers. Samples were incubated at 50°C for 30 min, and then amplified for 30 cycles under the following conditions: denaturation for 30 s at 94°C, annealing for 30 s at 56°C, and extension for 60 s at 68°C. The conditions for all genes used in this experiment of the RT-PCR reaction showed the linearity. The amplified products were separated by electrophoresis on a 1.5% agarose gel and visualized by ethidium bromide staining. Image density was quantified with a FluoroImager SI (Amersham Pharmacia Biotech).

#### Statistical analysis

Data are expressed as the mean  $\pm$  SEM. Statistical differences were analyzed using Student's *t*-test. *P* values less than 0.05 were considered to indicate statistically

**Fig. 3** Effect of zinc sulfate on the expression of Runx2, OPG, or ZIP1 mRNAs in osteoblastic MC3T3-E1 cells. Osteoblastic cells were cultured for 72 h in a medium containing 10% FBS. Cells with subconfluency were changed to a medium without FBS in the presence or absence of zinc sulfate ( $10^{-5}$  or  $10^{-4}$  M). After medium change, cells were cultured for 24, 48, or 72 h. Total RNAs (2  $\mu$ g) extracted from the cells were analyzed using RT-PCR with specific primers. The figure shows one of four experiments with separate culture. The densitometric data for each mRNA level in the cells cultured for 48 h in the presence of zinc sulfate were indicated as % of control (mean  $\pm$  SEM of four experiments). \*  $P < 0.025$  or \*\*  $P < 0.01$  compared with the control (none) value (Student's *t*-test)



significant differences. Also, we used an ANOVA multiple comparison test to compare the treatment groups.

## Results

Effect of zinc sulfate on gene expression in osteoblastic cells

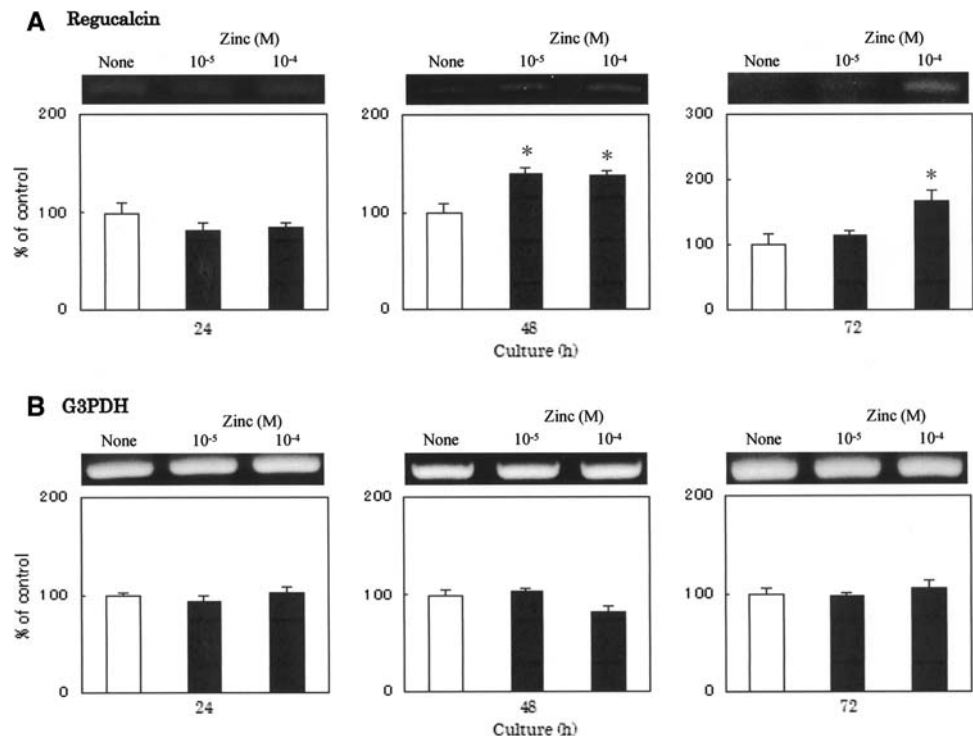
Osteoblastic MC3T3 cells with subconfluency were cultured for 48 h in a medium containing either vehicle or zinc sulfate ( $10^{-6}$ – $10^{-4}$  M), and the expression of various genes were examined (Fig. 1). Culture with zinc sulfate caused a significant increase in the mRNA expression of Runx2 (Fig. 1a), OPG (Fig. 2a), or regucalcin (Fig. 2c). Runx2 mRNA expression was significantly increased 25 or

45% after culture with  $10^{-5}$  or  $10^{-4}$  M zinc sulfate, respectively. OPG mRNA expression was significantly increased 25 or 30% after culture with  $10^{-6}$  or  $10^{-5}$  M zinc sulfate, respectively. Culture with zinc sulfate of  $10^{-6}$ – $10^{-4}$  M caused a significant increase in about 30 to 50% of regucalcin mRNA expression. Meanwhile, culture with zinc sulfate ( $10^{-6}$ – $10^{-4}$  M) did not cause a significant increase in mRNA expression of osteocalcin (Fig. 1b),  $\alpha 1(I)$  collagen (Fig. 1c), IGF-I (Fig. 1d), TGF- $\beta 1$  (Fig. 1e), G3PDH (Fig. 1f), or ZIP1 (Fig. 2b).

The time course of zinc sulfate effect on the mRNA expression of Runx2, OPG, ZIP1, regucalcin, or G3PDH in osteoblastic MC3T3-E1 cells with subconfluency was examined (Figs. 3, 4). Runx2 mRNA expression was significantly increased at 24, 48, or 72 h after culture with zinc sulfate ( $10^{-4}$  M) (Fig. 3a). OPG mRNA expression



**Fig. 4** Effect of zinc sulfate on the expression of regucalcin or G3PDH mRNAs in osteoblastic MC3T3-E1 cells. Osteoblastic cells were cultured for 72 h in a medium containing 10% FBS. Cells with subconfluency were changed to a medium without FBS in the presence or absence of zinc sulfate ( $10^{-5}$  or  $10^{-4}$  M). After medium change, cells were cultured for 24, 48, or 72 h. Total RNAs (2  $\mu$ g) extracted from the cells were analyzed using RT-PCR with specific primers. The figure shows one of four experiments with separate culture. The densitometric data for each mRNA level in the cells cultured for 48 h in the presence of zinc sulfate were indicated as % of control (mean  $\pm$  SEM of four experiments). \*  $P < 0.025$  compared with the control (none) value (Student's *t*-test)



was significantly increased at 24 or 48 h after culture with  $10^{-5}$  M zinc sulfate (Fig. 3b). A significant increase in regucalcin mRNA expression was seen at 48 or 72 h after culture with  $10^{-4}$  M zinc sulfate (Fig. 4a). ZIP1 (Fig. 3c) or G3PDH (Fig. 4b) mRNA expressions were not significantly changed after culture with zinc sulfate ( $10^{-5}$  or  $10^{-4}$  M) for 24, 48, or 72 h.

Effect of cycloheximide or DRB on zinc sulfate-increased gene expression in osteoblastic cells

The effect of cycloheximide, an inhibitor of protein synthesis, or DRB, an inhibitor of transcription activity, on the zinc sulfate-increased mRNA expression of Runx2, OPG, or regucalcin was examined in osteoblastic cells with subconfluency cultured with zinc sulfate ( $10^{-5}$  M) for 48 h. The effect of zinc sulfate ( $10^{-5}$  M) in increasing Runx2 (Fig. 5a), OPG (Fig. 5b), or regucalcin (Fig. 6a) mRNA expressions was significantly inhibited in the presence of cycloheximide ( $10^{-7}$  M) or DRB ( $10^{-6}$  M).

Effect of various zinc compounds on gene expression in osteoblastic cells

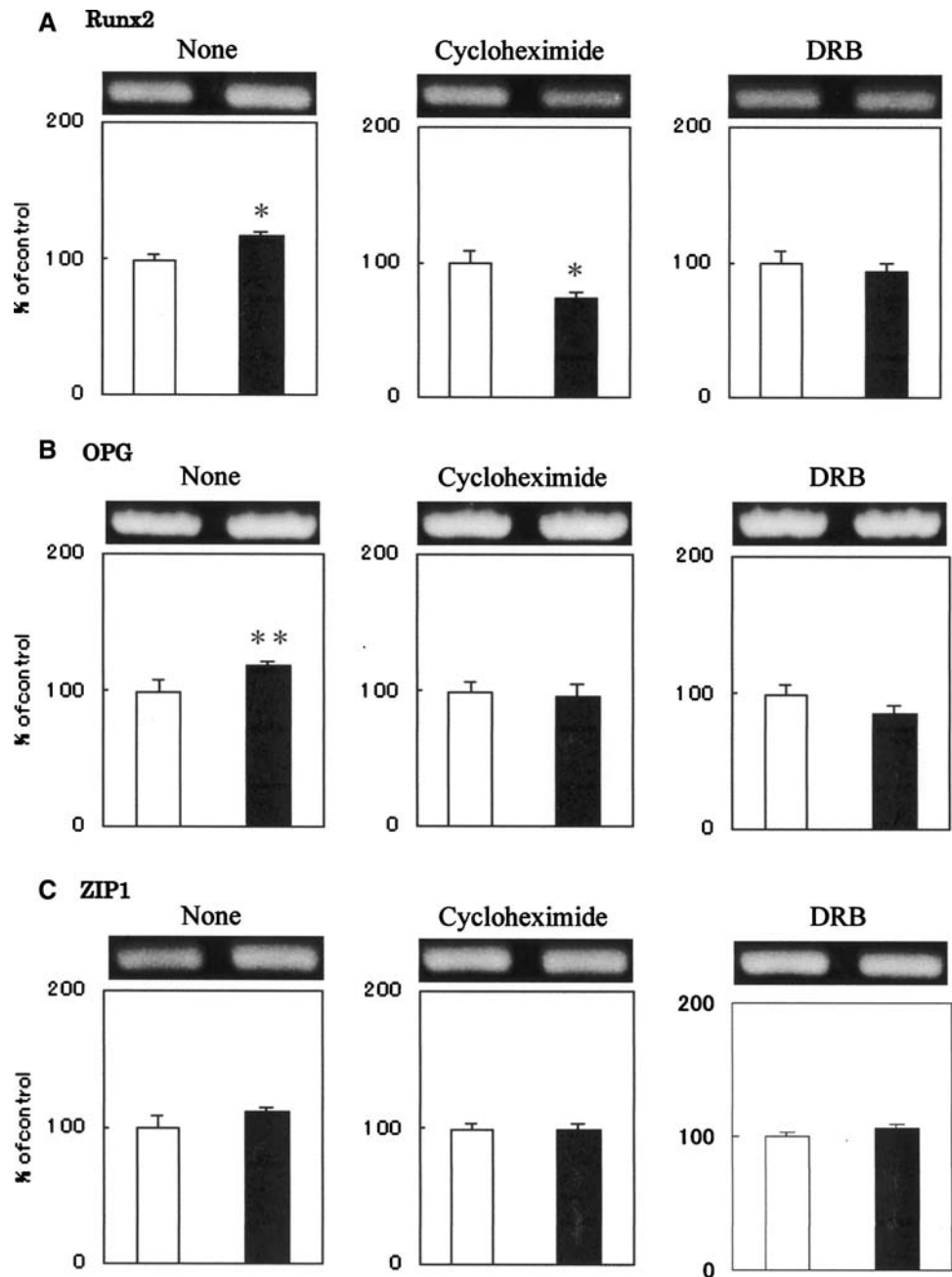
The effect of zinc sulfate, AHZ, or ZA on the mRNA expression of Runx2, OPG, ZIP1, regucalcin, or G3PDH in osteoblastic MC3T3-E1 cells with subconfluency cultured with each zinc compound ( $10^{-5}$  M) for 48 h was compared

(Fig. 7). Runx2 mRNA expression was significantly increased after culture with zinc sulfate or AHZ. OPG mRNA expression was significantly increased with zinc sulfate. Regucalcin mRNA expression was significantly increased with zinc sulfate or AHZ. Culture with zinc sulfate, AHZ, or ZA did not cause a significant change in ZIP1 or G3PDH mRNA expressions. The effect of zinc sulfate on the mRNA expression of Runx2, OPG, ZIP1, or regucalcin in osteoblastic MC3T3-E1 cells was potent in among zinc compounds used.

## Discussion

Zinc has been shown to have anabolic effects on osteoblastic MC3T3-E1 cells [14, 19–22, 26, 27]. Zinc can stimulate protein [14, 21] and DNA [22] syntheses in osteoblastic cells. The effect of zinc on gene expression in osteoblastic cells has not been fully clarified, however. This study was undertaken to determine the effect of zinc on the mRNA expressions of Runx2, osteocalcin,  $\alpha 1(I)$  collagen, OPG, IGF-I, TGF- $\beta 1$ , ZIP1, regucalcin, or G3PDH using RT-PCR. We found that culture with zinc sulfate caused a significant increase in Runx2, OPG, or regucalcin mRNA expressions in the cells, although it did not cause a significant increase in the mRNA expressions of osteocalcin,  $\alpha 1(I)$  collagen, IGF-I, TGF- $\beta 1$ , or ZIP1.

**Fig. 5** Effect of cycloheximide or DRB on the zinc sulfate-induced increase in Runx2 or OPG mRNA expressions in osteoblastic MC3T3-E1 cells. Osteoblastic cells were cultured for 72 h in a medium containing 10% FBS. Cells with subconfluency were changed to a medium without FBS in the presence or absence of zinc sulfate ( $10^{-5}$  M) with or without cycloheximide ( $10^{-7}$  M) or DRB ( $10^{-6}$  M). After medium change, cells were cultured for 48 h. Total RNAs (2  $\mu$ g) extracted from the cells were analyzed using RT-PCR with specific primers. The figure shows one of four experiments with separate culture. The densitometric data for each mRNA level in the cells cultured for 48 h in the presence of zinc sulfate were indicated as % of control (mean  $\pm$  SEM of four experiments). \*  $P < 0.025$  or \*\*  $P < 0.01$  compared with the control value without zinc addition (Student's *t*-test or ANOVA). White bars, without zinc; black bars, with zinc



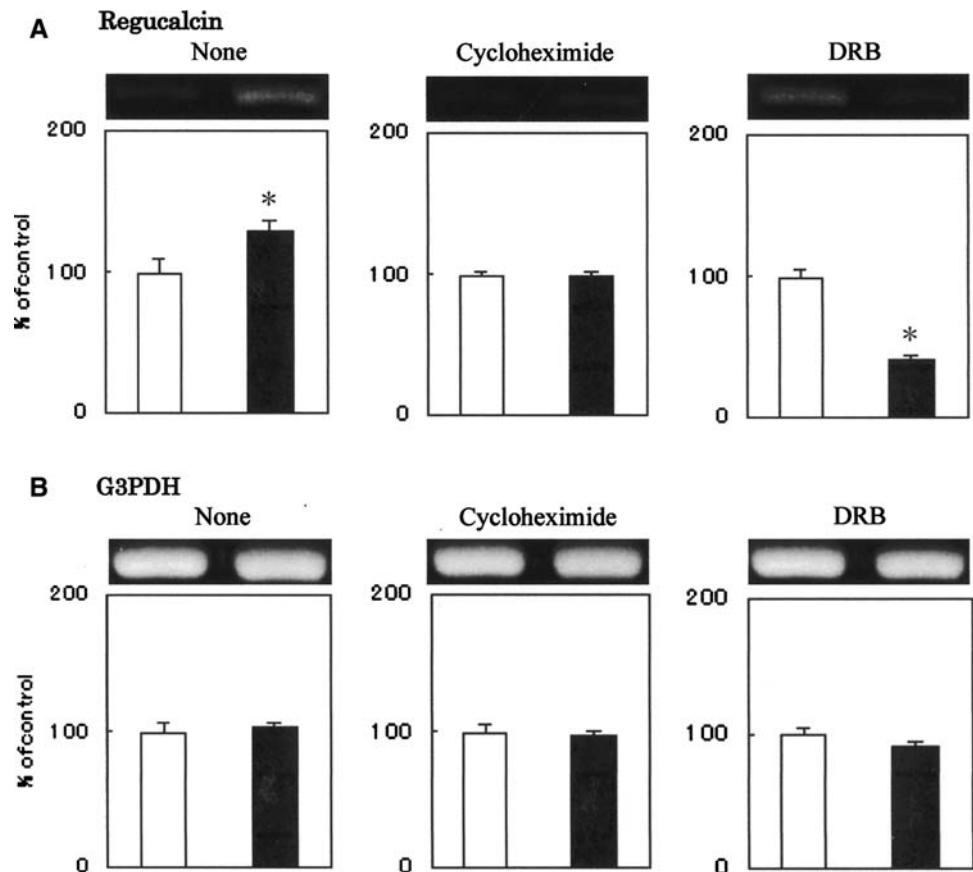
Runx2 is a member of the runt domain family of transcription factors, and it is involved in bone development and osteoblastic cell differentiation [38]. OPG involves in the inhibition of RANKL action that stimulates osteoclastogenesis [39]. Zinc has a role in the enhancement of these gene expressions in osteoblastic cells.

Zinc can stimulate differentiation in osteoblastic cells [27], and the metal has an inhibitory effect on RANKL-induced osteoclastogenesis in mouse marrow culture [25]. These effects of zinc may be partly mediated through the production of Runx2 or OPG in osteoblastic cells.

Regucalcin is a regulatory protein in intracellular signaling [40]. Regucalcin has been shown to suppress osteoblastic mineralization [41] and to stimulate osteoclastogenesis [42], thereby decreasing bone mass in regucalcin transgenic rats [43]. The physiologic significance which zinc stimulates regucalcin mRNA expression is unknown. However, it is possible that zinc has a role in the regulation of osteoblastic mineralization and osteoclastogenesis which is mediated through regucalcin expression in osteoblastic cells.

Culture with zinc stimulates protein production of IGF-I, TGF- $\beta$ 1, or osteocalcin in osteoblastic MC3T3-E1 cells

**Fig. 6** Effect of cycloheximide or DRB on the zinc sulfate-induced increase in regucalcin or G3PDH mRNA expressions in osteoblastic MC3T3-E1 cells. Osteoblastic cells were cultured as described in the legend of Fig. 5. Total RNAs (2  $\mu$ g) extracted from the cells were analyzed using RT-PCR with specific primers. The figure shows one of four experiments with separate culture. The densitometric data for each mRNA level in the cells cultured for 48 h in the presence of zinc sulfate were indicated as % of control (mean  $\pm$  SEM of four experiments). \*  $P < 0.025$  compared with the control value without zinc addition (Student's *t*-test or ANOVA). White bars, without zinc; black bars, with zinc



using ELISA assay [14]. However, zinc did not enhance mRNA expressions of IGF-I, TGF- $\beta$ 1, or osteocalcin in osteoblastic MC3T3-E1 cells using RT-PCR analysis. Zinc has been shown to activate aminoacyl-tRNA synthetase, which is a rate-limiting enzyme in translational process of protein synthesis, in osteoblastic cells, and the metal could induce protein synthesis [21, 26]. Presumably, the effect of zinc in increasing IGF-I, TGF- $\beta$ 1, or osteocalcin in osteoblastic cells is mainly resulted from the stimulation of protein synthesis that acts on translational process. However, it cannot exclude the possibility that zinc has a partially stimulatory effect on transcriptional process in osteoblastic cells.

Culture with zinc sulfate, AHZ, or ZA, which are zinc compounds, increases bone components in rat femoral tissues in vitro [13, 44, 45]. Among these compounds, culture with zinc sulfate had a potent effect on the mRNA expressions of Runx2, OPG, or regucalcin in osteoblastic cells. However, AHZ has been shown to have potent effects on protein [14, 21] and DNA syntheses [22] in osteoblastic cells as compared with zinc sulfate. The effect of zinc in stimulating gene expression of transcriptional process may differ with chemical form of zinc compounds in osteoblastic cells. It is speculated that zinc ion in zinc

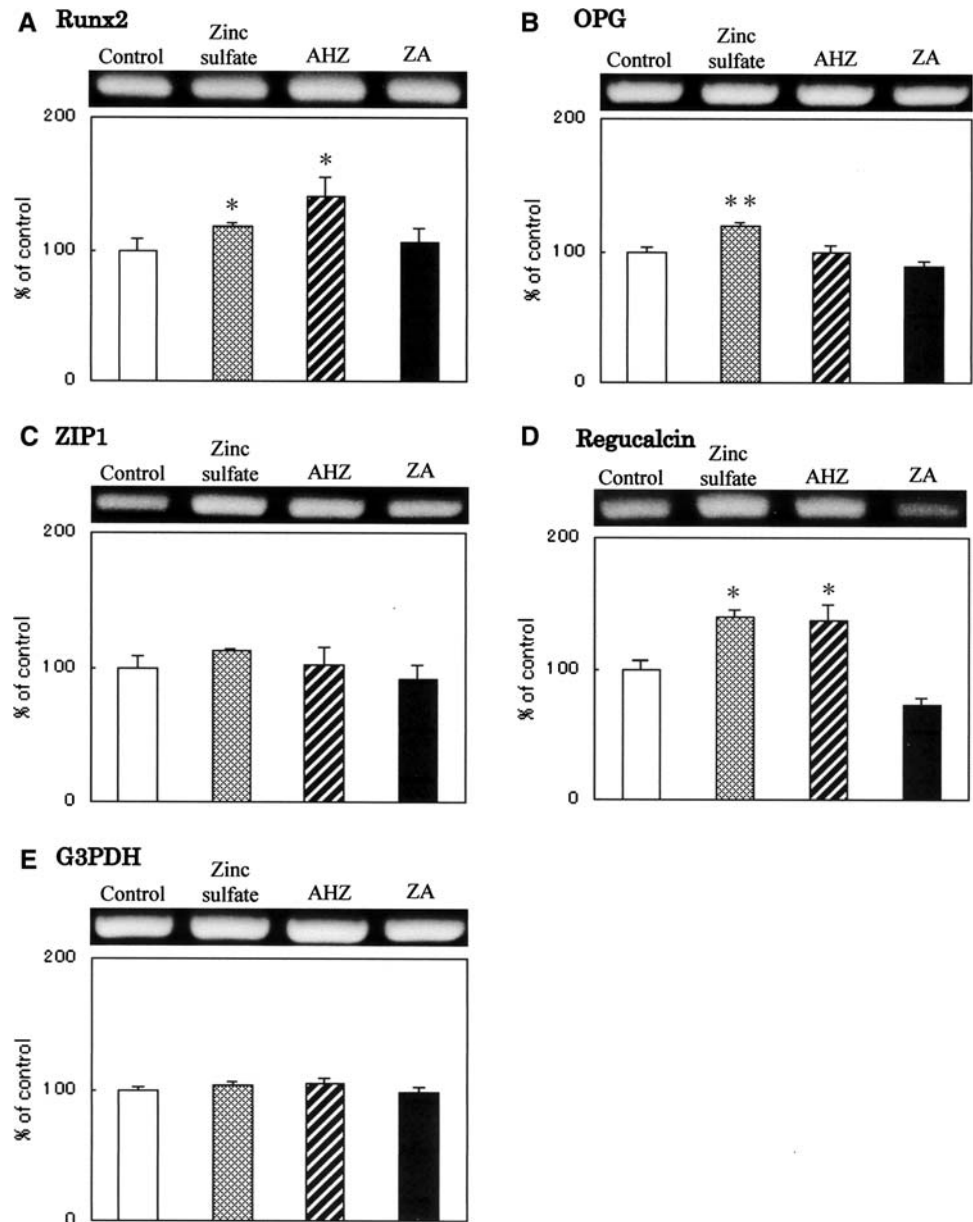
sulfate is potentially translocated to the nucleus in osteoblastic cells and that the metal ion can bind to transcriptional proteins and/or DNA in the cells as compared with that of AHZ or ZA.

The effect of zinc sulfate in increasing the expression of Runx2, OPG, or regucalcin mRNAs in osteoblastic cells was completely prevented in the presence of cycloheximide, an inhibitor of protein synthesis, or DRB, an inhibitor of transcription activity. This result suggests that the zinc-increased gene expression was resulted from newly synthesized protein components in osteoblastic cells. Zinc is known to stimulate protein synthesis in osteoblastic cells in vitro [14, 21, 27]. It is speculated that zinc ion stimulates protein synthesis including transcription factors (proteins), that the synthesized proteins enhance gene expression in osteoblastic cells, and that zinc directly acts on gene expression due to activating transcription factors in the cells.

In conclusion, it has been demonstrated that culture with zinc sulfate has stimulatory effects on the mRNA expressions of Runx2, OPG, or regucalcin in osteoblastic MC3T3-E1 cells in vitro. Zinc may have a role in the regulation of gene expression in osteoblastic cells.



**Fig. 7** Effect of various zinc compounds on the expression of Runx2, OPG, ZIP1, regucalcin, or G3PDH mRNAs in osteoblastic MC3T3-E1 cells. Osteoblastic cells with subconfluency were cultured for 48 h in a medium containing either vehicle, zinc sulfate ( $10^{-5}$  M),  $\beta$ -alanyl-L-histidinato zinc (AHZ;  $10^{-5}$  M), or zinc acexamate (ZA;  $10^{-4}$  M) without FBS. Total RNAs (2  $\mu$ g) extracted from the cells were analyzed using RT-PCR with specific primers. The figure shows one of four experiments with separate culture. The densitometric data for each mRNA level in the cells cultured for 48 h in the presence of zinc compound were indicated as % of control (mean  $\pm$  SEM of four experiments). \*  $P < 0.025$  compared with the control (none) value (Student's *t*-test)



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