

1,25 Dihydroxyvitamin-D₃ Attenuates the Confluence-Dependent Differences in the Osteoblast Characteristic Proteins Alkaline Phosphatase, Procollagen I Peptide, and Osteocalcin

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Abstract. In the present study a cell culture model of primary human osteoblasts based on degrees of confluence was investigated by measuring basal and 1,25(OH)₂D₃-stimulated levels of the osteoblast characteristic proteins alkaline phosphatase (AP), procollagen I-peptide (PICP), and osteocalcin (OC), as well as the corresponding gene expression. Primary osteoblast-like cell cultures from seven donors were treated in the second passage with 1,25(OH)₂D₃ (5×10^{-8} M for 48 hours) and investigated at four stages of confluence (stage I 50%, stage II 75%, stage III 100%, and stage IV 7 days postconfluence). In untreated cultures passing through the different stages of confluence, we saw a 1.8-fold increase of AP activity, a 2.3-fold increase of OC secretion, but a decrease of PICP levels to 0.36-fold. Gene expression showed only minor variation between the different confluence stages. 1,25(OH)₂D₃ did not significantly affect PICP production. Alkaline phosphatase protein was stimulated during proliferation until confluence, with no effect thereafter. Surprisingly, OC secretion and mRNA expression were stimulated in all four stages to the same absolute level independent of basal values. We conclude that our results correspond to other studies showing differentiation-stage dependent changes of basal levels of osteoblast-specific proteins. However, 1,25(OH)₂D₃ stimulation decreased the confluence-dependent difference for AP and abolished it for osteocalcin, thus leading to a more differentiated phenotype of the osteoblast. Therefore, 1,25(OH)₂D₃ stimulation might improve the reproducibility of results obtained at different confluence stages from cultures of clinical samples.

Key words: Human osteoblast — Differentiation — 1,25(OH)₂D₃ — Osteocalcin — Procollagen I — Alkaline phosphatase.

Osteogenic differentiation has been extensively studied *in vitro* using rat calvarial cells cultured in conditions that facilitate the formation of a mineralized bone matrix [1, 2]. The progressive differentiation of osteoblasts in culture is associated with the expression of alkaline phosphatase, an early marker of the osteoblast phenotype. The subsequent deposition of a collagenous extracellular matrix is accom-

panied by a decrease of procollagen I mRNA [3] and an increase of osteocalcin expression, a late marker of differentiated osteoblasts [4]. It is not yet clear if human osteoblasts behave in a similar manner *in vivo* and *in vitro*, but initial studies in developing human bone appear to indicate an analogous process [5, 6]. Because differences in hormonal regulation between species have been described [7], characterization and evaluation of human osteoblast systems seem to be essential for the study of human bone biology.

Studies on the differentiation of human bone have been greatly facilitated by the recent development of culture methods for human bone cells [8–14]. Normal adult human cells of the osteoblast lineage derived from various bone compartments are routinely established in short-term cultures. However, over the last years it emerged that all these cultures are a mixture of cells, composed of different subpopulations of bone cells, if not also contaminated with other cell types [10, 15]. The subpopulations seem to be primary osteoblasts in varying differentiation stages, modulated by hormones, depending on the phenotype as well as cell density in culture [16–20] probably accounting for the high variation in results [15, 21–23]. However, the differences in growth and differentiation of these cells in culture is thought to give the most realistic insight into the physiology and possibly pathophysiology of bone differentiation. Therefore, methods were developed to compare early and late stages of differentiation of primary osteoblasts in culture. The effect of hormones was studied on different stages of developing osteoblasts, on different clones of the same tumor cell line [16], on cells treated with dexamethason [24], on osteoblasts derived from spongiosa and marrow [25], and on cells plated at low and high densities [18, 19, 26]. In models of different cell density the influence of cell to cell contacts onto the differentiation was investigated. However, the possible effect of different culture time on a defined stage of confluence was not considered. Lately, we tested the influence of degree of confluence on the growth and differentiation of primary human osteoblasts *in vitro* [27]. In that study, we were able to show an increasing differentiation characterized by the increasing number of alkaline phosphatase-positive cells with decreasing proliferation of cultured cells in four stages of confluence. According to results in the chicken osteogenic system [3], collagen secretion was maximum at early proliferation, decreasing with later confluence stages.

1,25(OH)₂D₃ is a hormone known to differentially in-

fluence primary human osteoblast in culture [19] and also to promote differentiation [28–30]. In this study we used the confluence model to investigate the influence of 1,25(OH)₂D₃ on proteins and gene expression of alkaline phosphatase (AP), osteocalcin (OC), and procollagen I (PICP) in primary human osteoblasts.

Materials and Methods

All culture media and fetal calf serum (FCS) were purchased from Biochrom (Berlin, Germany). Cell culture disposables were purchased from Nunc (Wiesbaden-Biebrich, Germany) or Greiner (Solingen, Germany), and medium supplements (antibiotics, glutamine) from GIBCO/BRL (Eggenstein, Germany). Standard laboratory reagents were purchased from Sigma (Deisenhofen, Germany) if not noted otherwise.

Cell Culture

Primary human bone cell cultures were established from bone specimens as previously described [11, 12]. Trabecular bone sections, obtained during hip or knee replacement surgery, were treated identically. The explants were cleaned of adherent tissue and periosteum, cut in 1–3 mm³ pieces, thoroughly rinsed, and maintained in Dulbecco's Modified Eagle Medium (DMEM) containing 10% FCS, glutamine (58.5 µg/ml), penicillin (100 U/ml), and streptomycin (100 µg/ml). Cultures were initiated within 24 hours of bone specimen sampling.

The grown cells were released from the surface of the culture dish with 0.25% trypsin after 14–21 days. Cells were divided in two 75 cm² tissue culture flasks and are referred to as 1st passage cells. After growing near confluence, the cells were released from the culture dish as described above, counted, plated in 75 cm² flasks at a density of 2 × 10⁵ cells/dish, and evenly distributed. These cells are referred to as 2nd passage cells. They were analyzed in different growth phases, defined by the extent of cell confluence. We selected cells at a confluence of 50% and 75% as being in the stage of fast proliferation; cells at 100% and 7 days 100% confluence were used to describe the period of diminished proliferation and increasing differentiation. To assess cell density, daily control of all culture dishes was necessary.

Cultures were controlled every day to establish the stage of confluence. At each time point the cells were washed twice with incubation medium (DMEM with 1% BSA) (Sigma), MgSO₄ 0.02% and penicillin (100 U/ml), and streptomycin (100 µg/ml) to remove the FCS followed by incubation with 5 × 10⁻⁸ M 1,25(OH)₂D₃ or solvent (ethanol <0.01%) for 48 hours at 37°C. After harvesting, cell numbers were determined for each time point by hemocytometer counting. The remaining cells were frozen at -20°C for determination of total protein and AP activity. All cells were regularly tested for mycoplasma contamination by PCR (Mycoplasma PCR Primer Set, Stratagene, Heidelberg, Germany).

Biosynthesis of Proteins

For the determination of total protein and AP, the medium was decanted and the adherent cell monolayer was washed with phosphate-buffered saline (PBS). Cells were lysed in distilled water by repeated freeze-thawing and sonication (Ultrasonics W 185F, 40–60 W, 3 × 30 seconds). The lysate was clarified by centrifugation (10 minutes at 10,000 × g) and the soluble protein fraction was quantified using the BioRad protein assay with an albumin standard (BioRad, München, Germany). AP activity was assayed in cell lysates by determining the release of p-nitrophenol from p-nitrophenyl phosphate at 37°C and pH of 10.5. The substrate solution contained 8 mM p-nitrophenyl phosphate in 0.5 M AMP-buffer (2-amino-2-methyl-1-propanol), pH 10.5, supplemented with 0.2 mM MgCl₂. After 10 minutes at 37°C the reaction (volume 110 µl) was stopped with 50 µl 1 N sodium hydroxide. AP

determinations were done in duplicate and are reported as units per mg protein per minute incubation time (modified after [31]). Secreted osteocalcin (OC) and the carboxyterminal peptide of procollagen I (PICP) were measured in culture supernatants after the incubation with 5 × 10⁻⁸ M 1,25(OH)₂D₃ or solvent for 48 hours at 37°C. Osteocalcin was determined in duplicate with an immunoradiometric assay (Nichols, Bad Nauheim, Germany); the detection range of the kit was 0.9–267 ng/ml. Values were corrected for interfering substances present in the culture medium. Collagen type I content was calculated from the c-terminal propeptide concentration [32, 33] measured in duplicate samples by ELISA (Metra Biosystems, Palo Alto, CA, USA). Values were corrected for the presence of background levels of PICP in fresh culture medium. Results are reported as the mean of two supernatants for each time point.

RNA Isolation and Northern Blot Analysis

Total RNA was isolated from cells by the Caesium-chloride method [34]. For Northern blot analysis, 10 µg per lane of total cellular RNA was separated electrophoretically in a 1% agarose gel (2% formaldehyde, 0.3 µg/ml ethidium bromide), blotted to nylon membranes (Zeta Probe, BioRad, München, FRG) by capillary diffusion, and immobilized by UV-crosslinking (Stratagene, Heidelberg). ³²P-labeled cDNA probes were synthesized by random priming (Stratagene, Heidelberg). Prehybridization and hybridization were each performed in 50% formamide, 4 × SSPE, 5 × Dehnhardt's, 1% SDS, and 500 µg/ml salmon sperm DNA at 42°C for 24 hours. For hybridization, approximately 10⁶ cpm/ml of ³²P-labeled probe was added. Following hybridization, blots were washed for 2 × 30 minutes at room temperature in 2 × SSC/0.1% SDS and 5–30 minutes at 55°C (rat cDNA probes) or 63°C (human cDNA probes) in 0.1 × SSC/0.1% SDS. Autoradiographs were quantified by laser densitometry (Biometra, Göttingen, FRG) and normalized to the ethidiumbromide staining of 18S and 28S ribosomal RNA, using a negative film for densitometric analysis. Values are expressed in relation to the maximal absorbency of each transcript. For each mRNA the maximal absorbency of stimulated and unstimulated values was set to 100.

Values represent the mean of five (OC), seven (PICP), or three (AP) independent experiments. For analysis of GAPDH mRNA expression, seven experiments and two sets of blots were analyzed. Two sets of Northern blots were rehybridized without stripping for each experiment.

cDNA Probes

Procollagen type I expression was measured using the 372 bp fragment of the rat proalpha(I) collagen cDNA clone pHCAL1U [35]. This fragment binds to α1(I) and α2(I) procollagen. The glycerol-aldehydephosphatase-dehydrogenase (GAPDH) human cDNA probe was a 1.2 kb insert from clone pHcGAP [36] and the AP human cDNA probe was a 2.5 kb insert of pAT153 [37]. Osteocalcin expression was detected using the human 1.2 kb Sac I fragment of SP 65 [38].

Statistical Methods and Mathematics

Statistical testing was performed with Stat Works statistical package. Student's paired *t*-test, Wilcoxon, and Whitney-U-Test were used where appropriate and as noted. All values are expressed as mean ± SEM. (See Figs. 3 and 4 for significant differences.)

Results

Cell Culture

The bone used to initiate explant cultures was taken from

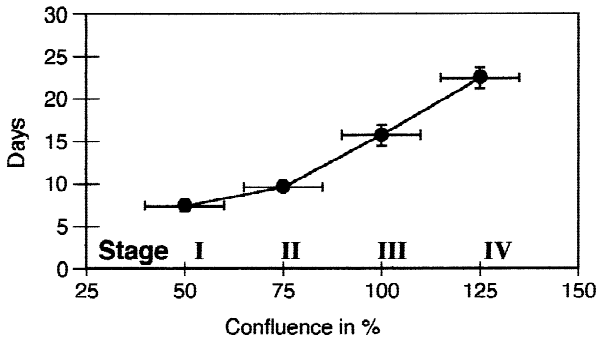


Fig. 1. Stages of confluence in relation to time in culture. For reasons of presentation, stage IV is designated as 125% confluence. SEM of days (vertical) and confluence (horizontal) are depicted. Values are mean \pm SEM ($n = 7$).

seven male patients who underwent an arthroplasty (3 hip, 3 knee, 1 talus). The mean age of the patients was 62 ± 4.36 years, range 46–80, median 60 years. The cultures grew predominantly homogeneous and reached the stages of confluence within 7.43 ± 0.63 days (50%, stage I), 9.57 ± 0.49 days (75%, stage II), 15.71 ± 1.27 days (100%, stage III), and 22.43 ± 1.23 days (seven days 100%, stage IV), respectively, showing decreasing proliferation, as demonstrated in Figure 1. The morphological appearance of the four confluence phases is shown in Figure 2. During proliferation, the cells showed a long and narrow fibroblast-like appearance; with further confluence, cells became flattened and wide-spread. At 100% confluence, cells changed again to a long and extended shape before they started to grow in multilayer formation.

Protein Synthesis in the Confluence Model

Cultures showed higher proliferation while passing through stages I and II and growth slowed when reaching stages III and IV (Fig. 1). The secretion of PICP in the supernatant decreased from early proliferation (50%, stage I) to late confluence (stage IV) to 78% of the starting level (factor 0.36, $P < 0.05$, Wilcoxon) (Fig. 3). Alkaline phosphatase (AP) increased from basal level at early proliferation to 1.8-fold values at stage IV (Wilcoxon, $P < 0.05$). In five of the cultures the unstimulated cells were already able to synthesize osteocalcin (OC) during proliferation, with a 2.3-fold increase towards confluence and thereafter ($1.27 \text{ ng} \pm 0.4 \text{ ng/mg}$ total protein stage I versus 2.91 ± 0.84 stage IV). Due to high culture-dependent variance results did not reach significance levels.

Gene Expression in the Confluence Model

To plot the mRNA content in relation to a housekeeping gene we examined GAPDH gene expression in addition to the specific genes. Analyzing seven experiments with a double set of northern blots we found a decrease in GAPDH gene expression from early proliferation to late confluence in unstimulated cultures ($P < 0.05$) (Figs. 4 and 5). Therefore, we adjusted the specific gene expression to the amount of loaded mRNA depicted by the ethidium bromide staining (Figs. 4 and 5). Procollagen I mRNA increased (1.3-fold) significantly in unstimulated cultures from early to late pro-

liferation (stages I and II, $P < 0.05$). There was no further change in stages III and IV (Figs. 4 and 5). Unstimulated OC gene expression was detectable only in two of five cultures at confluence (stage III) with a further 2.2-fold increase at stage IV. During proliferation (stages I and II) no basal osteocalcin expression was detectable (stages I and II, Figs. 4 and 5). Parallel to procollagen I, AP showed an increase to 1.5 levels of basal gene expression from stages I to II and stayed constant thereafter (Figs. 4 and 5).

Influence of 1,25(OH)₂D₃ on Specific Proteins

Alkaline phosphatase/mg protein was increased by 1,25(OH)₂D₃, the effect being more pronounced in early proliferation; the stimulation was significant only at confluence ($P < 0.05$, Wilcoxon).

Osteocalcin greatly increased after stimulation with 1,25(OH)₂D₃ independently of the stage of confluence or the increasing basal osteocalcin synthesis, and the maximal level was similar in all stages ($P < 0.05$) (Figs. 3–5).

There were only minor effects on the PICP content in the conditioned medium. PICP/mg protein was inhibited to 0.74-fold by 1,25(OH)₂D₃ during early proliferation (50% confluence, $P = 0.058$) with no effect thereafter.

1,25(OH)₂D₃ Influence on Gene Expression

Analyzing seven experiments with a double set of northern blots we found a highly significant inhibition of GAPDH expression by 1,25(OH)₂D₃ in all stages (to 70% of control). Furthermore, expression decreased from proliferation to late confluence in stimulated as in unstimulated cultures (1.2-fold). Procollagen I gene expression increased 1.2-fold in stimulated cultures from early to late proliferation (stages I to II, $P < 0.05$) with no further change thereafter. 1,25(OH)₂D₃ slightly stimulated procollagen I gene expression in all stages, the stimulatory effect being significant only at early proliferation and at confluence (1.1-fold, stage III, Wilcoxon, $P < 0.05$) (Figs. 4 and 5). Osteocalcin mRNA was already stimulated by 1,25(OH)₂D₃ in stages I and II, this increase being 30-fold of the basal level at confluence and 12-fold thereafter (stage IV, $P < 0.01$) (Figs. 4 and 5). The gene expression of AP was detectable in only three cultures. We saw a stimulation of AP gene expression by 1,25(OH)₂D₃ in all stages which was not significant due to wide scatter and the small number of cases.

Influence of 1,25(OH)₂D₃ on Variation of Results

Overall variation did not show a difference between unstimulated and stimulated cultures, the coefficient of variation for GAPDH gene expression being 6–29% depending on the stage of confluence. The respective values for stimulated OC expression were 27–44% (analysis for unstimulated cultures not done) and 35–58% for procollagen I expression. Variation of basal cell culture analysis was 47–60% for cell number, 32–120% for total protein, 39–106% for AP, 37–121% for osteocalcin, and 38–149% for PICP values.

Discussion

In this paper we can show decreasing proliferation and in-

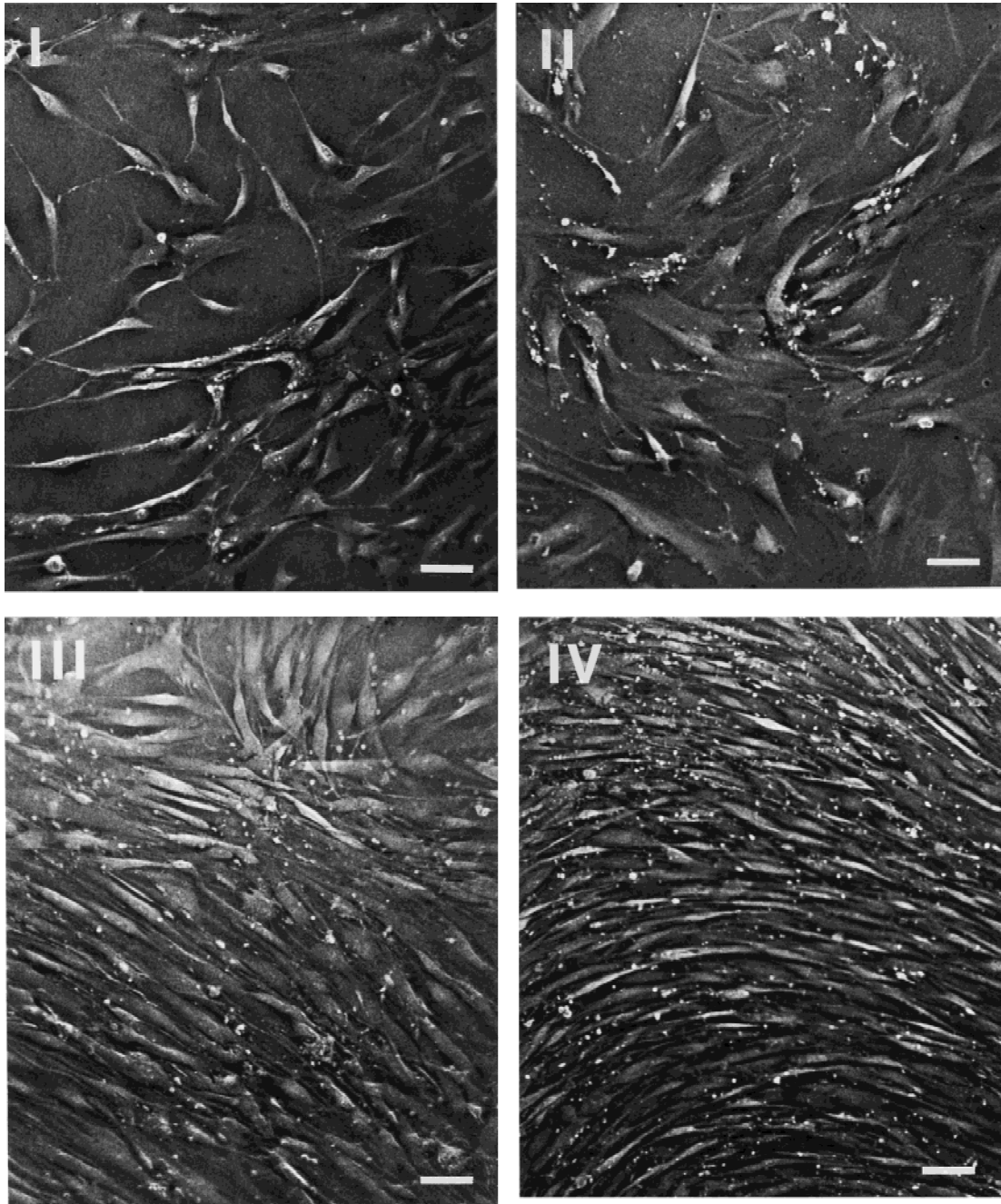


Fig. 2. Representative light field microscopy of primary human osteoblast-like cells depicting the four different stages of confluence (single culture at different time points). Stage I depicts 50% confluence, stage II 75%, stage III 100%, and stage IV 7 days 100% confluence. Bar 50 μ m.

creasing differentiation in pHOB, characterized by different stages of confluence. The development of a more mature phenotype was shown by an increase in AP activity, decreasing collagen I secretion when cultures became confluent, and increasing OC secretion with further confluence. Finally, OC mRNA expression was detected at a very low level beginning at confluence. So far, these results are in accordance with data from other species and therefore the regulation of synthesis of characteristic proteins seemed to be partly confluence dependent. Concerning the analysis of AP and procollagen I gene expression we did not find any

confluence-dependent differences and we conclude that mRNA expression of these proteins does not contribute systematically to the characterization of differentiation stages in pHOB.

Furthermore, we investigated whether the observed stage-dependent differences in protein expression and secretion could be influenced by 1,25(OH)₂D₃, a hormone that is known to regulate the differentiation of osteoblasts. We found that confluence-dependent differences were attenuated by 1,25(OH)₂D₃, because the hormone caused the proliferating osteoblasts to behave like more differen-

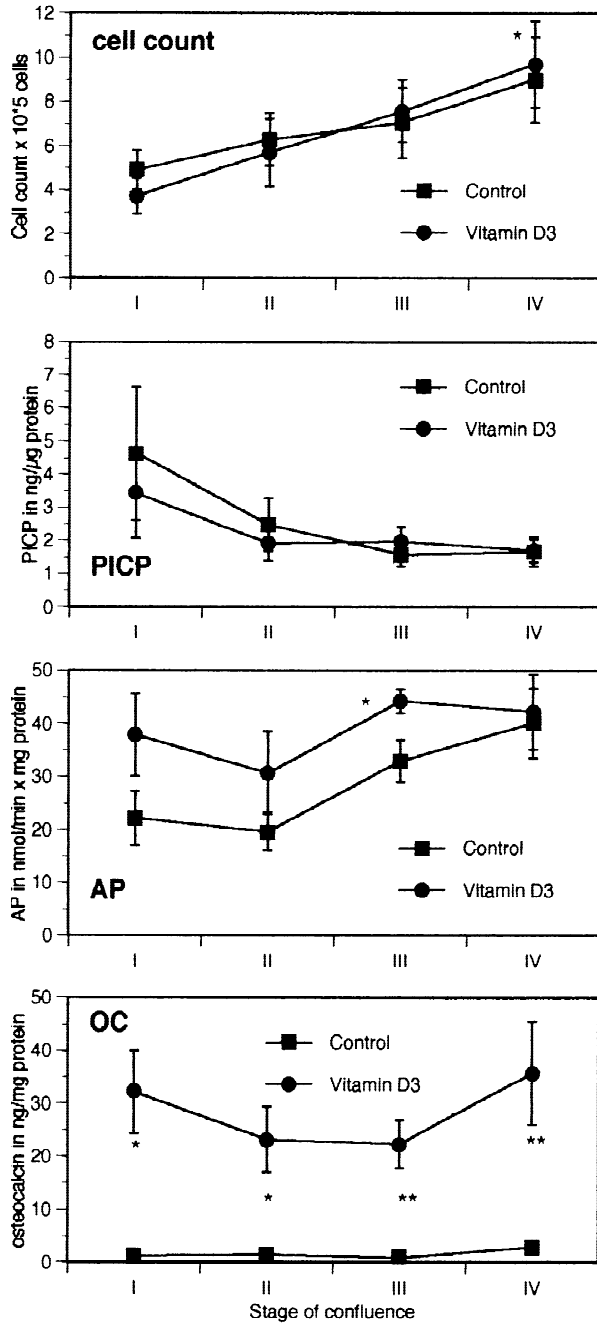


Fig. 3. Growth and specific proteins in four different stages of confluence under the influence of 5×10^{-8} M $1,25(\text{OH})_2\text{D}_3$ in comparison to control values. Stage I means 50% confluence, stage II 75%, stage III 100%, and stage IV 7 days 100% confluence. Procollagen I secretion measured by production of procollagen-I-peptide (PICP) in the supernatant. Alkaline phosphatase (AP), osteocalcin (OC). Significant values: * $P < 0.05$, ** $P < 0.01$, *** $P < 0.001$.

tiated cells. In other words, the differentiating influence of $1,25(\text{OH})_2\text{D}_3$ was strongest during proliferation whereas only minor effects were seen after cells became more confluent and differentiated. In the following we discuss the different aspects in more detail.

A stimulation of AP by $1,25(\text{OH})_2\text{D}_3$ is seen in rat [16, 39, 40], in human osteosarcoma cell lines [41], and in nor-

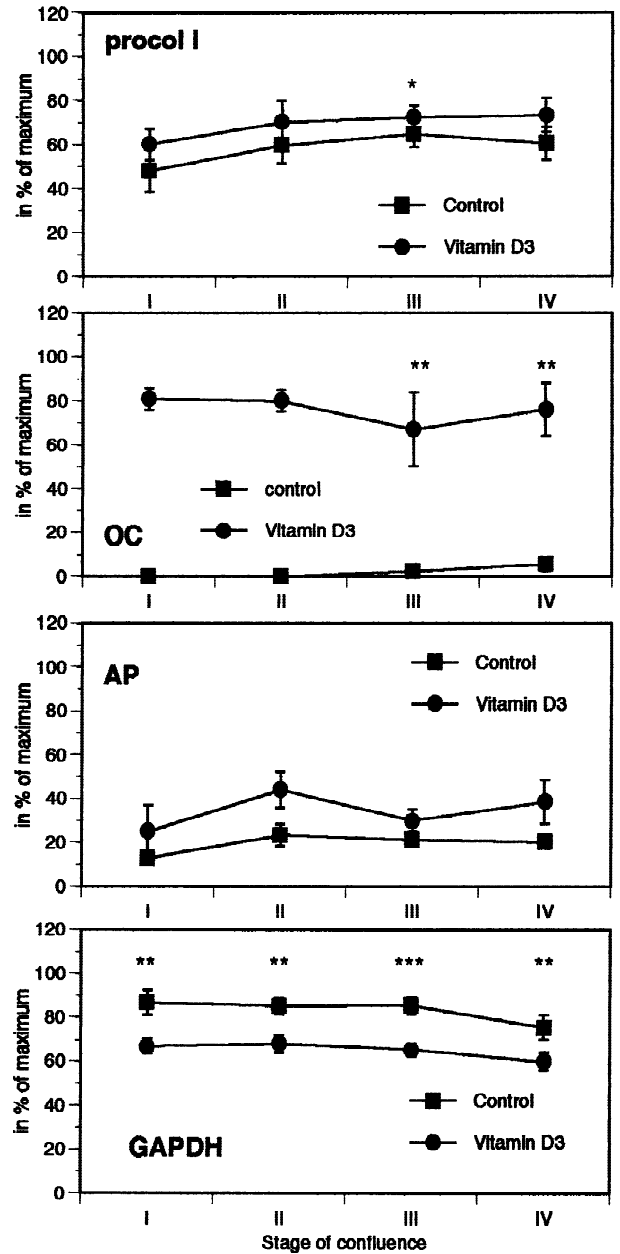


Fig. 4. Gene expression of osteoblast specific genes and GAPDH in four different stages of confluence under the influence of 5×10^{-8} M $1,25(\text{OH})_2\text{D}_3$ compared to control values. Stage I means 50% confluence, stage II 75%, stage III 100%, and stage IV 7 days 100% confluence. Cellular RNA from seven independent experiments was isolated and assayed by northern blot analysis. Expression was quantified by laser densitometry and results are plotted against the maximum expression of each transcript. The panel shows the mean values \pm SEM for each specific transcript. $\alpha 1$ and $\alpha 2$ (I) procollagen (procol I, $n = 7$), alkaline phosphatase (AP, $n = 4$), osteocalcin (OC, $n = 7$), and glycerol-aldehyde phosphate dehydrogenase (GAPDH, $n = 14$). Significant values: * $P < 0.05$, ** $P < 0.01$, *** $P < 0.001$.

mal human bone cells [26]. Most of the data concerning the differentiation-dependent AP regulation by $1,25(\text{OH})_2\text{D}_3$ have been investigated in rat osteoblast or rat osteosarcoma cells. Early studies in ROS 17/2.8 cells and subclones [16, 39] showed a biphasic regulation for AP by $1,25(\text{OH})_2\text{D}_3$,

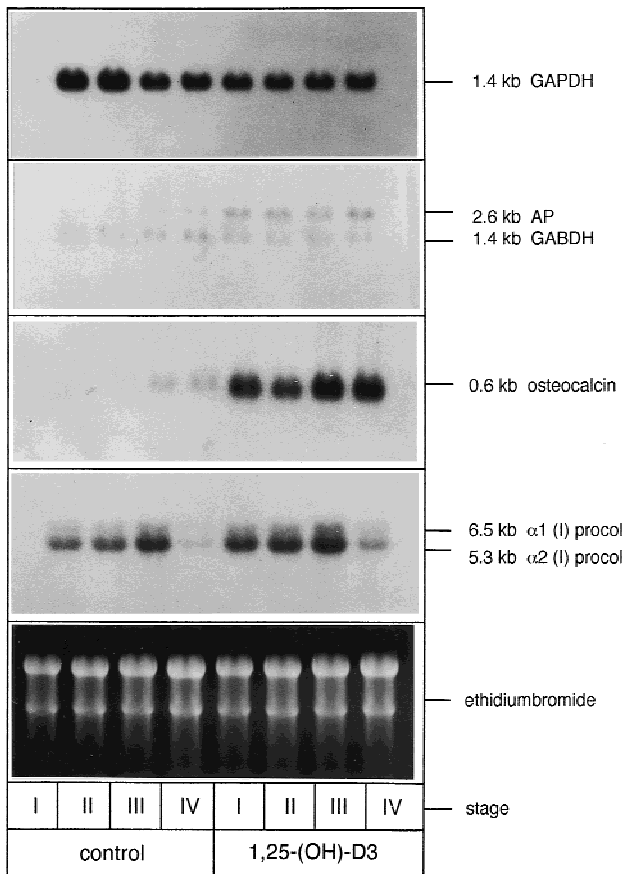


Fig. 5. Steady state gene expression during four different stages of confluence under the influence of 5×10^{-8} M 1,25(OH)₂D₃ compared with control values. Stage I means 50% confluence, stage II 75%, stage III 100%, and stage IV 7 days 100% confluence. Cellular RNA from seven independent experiments was isolated and assayed by northern blot analysis. Representative northern blots are depicted to illustrate alterations in mRNA expression. The ethidium bromide staining, which shows equal gel loading, was used as reference. Glycerol-aldehyde phosphate dehydrogenase (GAPDH), alkaline phosphatase (AP), osteocalcin (OC), and procollagen I (procol I) are shown. Due to repeated hybridization, AP has to be shown on a blot together with GAPDH.

dependent on the basal level of enzyme activity. With low basal AP levels, a stimulation of enzyme activity and inhibition of proliferation were seen, an effect that was inverted with high basal AP-enzyme activity. In our confluence model, 1,25(OH)₂D₃ stimulated AP synthesis from early proliferation to confluence; after confluence, no effect of 1,25(OH)₂D₃ was seen. The AP mRNA expression in human osteoblasts is known to be increased after 1,25(OH)₂D₃ stimulation in pHOB [18]. However, we do not have enough good quality data to support these results although we saw an increase in one case. These data on AP protein synthesis support the finding that 1,25(OH)₂D₃ can positively and negatively regulate expression of osteoblast phenotypic markers as a function of basal levels of expression, which is a reflection of the differentiated state of the osteoblast.

Osteocalcin protein and mRNA synthesis are highly stimulated by 1,25(OH)₂D₃, as shown in different osteoblast-like cell models [22, 26, 42, 43]. However, in contrast

to other studies our data show that the increase was not dependent on confluence, e.g., there were no differences in the stimulated levels between stages [44]. This was also evident for the mRNA expression which was not dependent on basal expression of OC mRNA or the stage of confluence in our model, in contrast to published results in primary rat osteoblasts [44]. Gene expression and protein secretion are similarly stimulated by 1,25(OH)₂D₃, supporting the hypothesis that 1,25(OH)₂D₃ controls OC protein synthesis by direct transcriptional mechanisms [45].

Procollagen secretion was measured by a PICP-specific antibody, which binds the intact procollagen I molecule and the free c-terminal propeptide which is split off during collagen I formation. Therefore, we only have information on the release of procollagen from the osteoblasts and the fraction that is transformed to collagen but not on the accumulation of collagen matrix. Gerstenfeld et al. [3] demonstrated in chicken embryo osteoblasts that synthesis of new collagen protein increased during proliferation but decreased thereafter, whereas collagen in the matrix further increased [3]. Our results, showing a decrease in PICP secretion with further confluence, would therefore be in agreement with the decrease of newly synthesized collagen seen in the chicken system. In contrast, the increasing procollagen I mRNA expression until confluence is surprising. Indeed, we have already found a similar expression sequence under basal conditions in cultures from 10 other donors of different ages where the decrease of PICP and increase from procollagen I mRNA was significant from stages I to III [27]. We assumed that the lack of ascorbate, which is a prerequisite for the full expression of collagen protein, was responsible for the expression sequence of procollagen mRNA and protein secretion. This is supported by the finding that in pHOB, ascorbate stimulates PICP secretion up to 5.8-fold values after confluence, probably by stimulating hydroxylation and secretion of accumulated procollagen I peptides [46, 47]. In rat and chicken calvarial cells the total collagen accumulation was increased as a function of ascorbic acid concentration, with no collagen in the matrix without the addition of ascorbate [3, 4]. Alternatively, the cultures might not have been analyzed long enough after confluence to see the decrease in procollagen I mRNA.

There was only a minor inhibitory effect of 1,25(OH)₂D₃ on procollagen I-peptide secretion in the early proliferative phase but there was no effect thereafter. We found a small but confluence-independent stimulation of the procollagen mRNA message by this hormone. This supports findings in the human osteosarcoma cell line MG 63 that the effect of 1,25(OH)₂D₃ on collagen production is only partly regulated by mRNA, but mainly at a post transcriptional level [28]. In rat calvarial cells, collagen I mRNA was inhibited by 1,25(OH)₂D₃ at its peak level during proliferation and stimulated at its lowest basal level during the mineralization period, demonstrating that the effect of 1,25(OH)₂D₃ is dependent on the proliferative and differentiated phenotype of the osteoblast [44]. In pHOB cells only marginal influence [48] or an increase independent of the seeded cell density on procollagen I synthesis after 1,25(OH)₂D₃ stimulation [26] was seen. In addition, procollagen I mRNA was stimulated at high cell density [18]. In human bone cells cultivated from periosteum, 10^{-8} M 1,25(OH)₂D₃ stimulated collagen I synthesis and mRNA expression to the same extent after confluence [49]. In immortalized human bone marrow cells, 1,25(OH)₂D₃ stimulated collagen mRNA and protein secretion into the same direction as proliferating and differenti-

ating cells [50]. Regarding the differences in results between the rat and the human system we suggest that differences in the expression sequence of collagen I might be due to species differences.

In conclusion, when regarding the basal levels of the osteoblast specific proteins in correlation to the stage of confluence we show an increase of AP and OC and a decrease of procollagen I secretion, representing the development from proliferation to differentiation. These data support the current view on osteoblast differentiation in culture. Stimulation with 1,25(OH)₂D₃ decreased the difference between confluence stages by causing the proliferating cultures to behave like more differentiated cells, whereas there was little effect except on osteocalcin after confluence. In other words, the functional status of differently differentiated cell populations became more alike under the influence of 1,25(OH)₂D₃. Therefore 1,25(OH)₂D₃ might improve the reproducibility of results obtained from clinical samples.

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