Effect of recombinant vascular endothelial growth factor and translationally controlled tumor protein on 2-hydroxyethyl methacrylate-treated pulp cells

CHUNYANUT WONGKHUM¹, WILAIWAN CHOTIGEAT¹ and UREPORN KEDJARUNE-LEGGAT²

Departments of ¹Molecular Biotechnology and Bioinformatics, Faculty of Science and ²Oral Biology and Occlusion, Faculty of Dentistry, Prince of Songkla University, Hat Yai, Songkhla 90112, Thailand

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Abstract. Vascular endothelial growth factor (VEGF)-A is a potential signaling protein that may promote angiogenesis. VEGF also helps cells survive in stressful or hazardous conditions. The present study aimed to compare the effect of VEGF with translationally controlled tumor protein (TCTP), an anti-apoptotic protein in human dental pulp cells (HDPCs), following exposure to 2-hydroxyethyl methacrylate (HEMA), which is a major residual monomer from resin restorative dental materials. Cell viability, alkaline phosphatase (ALP) activity, mineralization and gene expressions for odontogenic and osteogenic differentiation markers of HDPCs were investigated, following exposure to HEMA and in combination with TCTP and VEGF. The results revealed that TCTP at 1 ng/ml and VEGF at 10 ng/ml significantly promoted the proliferation of HDPCs ($P < 0.05$). TCTP (1 ng/ml) and VEGF (10 ng/ml) maintained the cell viability of 4 mM HEMA-treated cells at the same percentage as the control. However, cells treated with HEMA+TCTP+VEGF had a lower cell viability than the groups treated with HEMA and TCTP or VEGF alone. TCTP and VEGF promoted cell proliferation, ALP activity and mineralization, and upregulated of DSPP, DMP-1, BMP-2, and ALP mRNA expression compared with the control. Furthermore, the HEMA+TCTP and HEMA+VEGF groups had significantly higher percentages of calcium deposition than HEMA-treated cells ($P < 0.001$). HEMA was cytotoxic to HDPCs, reduced ALP activity and caused the significant downregulation of odontogenic and osteogenic gene expressions ($P < 0.05$). It was concluded that VEGF and TCTP

promoted pulp cell growth and the survival of HEMA-treated cells without synergistic effects. TCTP was required in lower concentrations for these effects. VEGF and TCTP enhanced cell differentiation and mineralization.

Introduction

Growth factor typically acts as signaling molecules between cells and is an important component required for tissue engineering. There have been many attempts at developing dental biomaterial or scaffold supplemented with a single or combination of active molecules to induce dentine or pulp regeneration. One growth factor that was widely selected for this purpose is vascular endothelial growth factor (VEGF).

VEGF-A is one of the four subtypes of VEGF (-A, -B, -C, and -D) that are expressed in dental pulp with autocrine and paracrine effects in blood vessels and immune cells (1). It is a potential signaling protein that can promote angiogenesis and activate dentin regeneration (2). VEGF has been used in growth factor delivery-based tissue engineering (3), including pulp tissue regeneration for regenerative endodontics (4). Moreover, VEGF has been shown to induce proliferation and differentiation of human pulp cells into odontoblasts (5). Another interesting property of VEGF is helping cells survive from stress or hazardous conditions. There have been some studies reporting that VEGF played a role in survival of serum-starved endothelial cells (6,7), as well as survival of these cells in hypoxic conditions (8). A previous study has also revealed that VEGF expression was upregulated in mouse odontoblast-like cells (MDPC-23) exposed to 2-hydroxyethyl methacrylate (HEMA) (9).

HEMA is a major monomer released from incomplete polymerization of resin-based dental restorative materials. It is cytotoxic to cells and can damage DNA (10), leading to proliferation impairment (11) and delayed cell differentiation (12).

Translationally controlled tumor protein (TCTP) is a conserved protein found in many eukaryotic cells ranging from plant to animal kingdoms (13). TCTP has many functions, including proliferation, maturation, and antiapoptosis (14). TCTP from *Penaeus merguensis* (Pmer-TCTP) has been shown to have an antiapoptotic property against HEMA-treated dental pulp cells (15). A new formula of resin

Correspondence to: Dr Urepor Kedjarune-Leggat, Department of Oral Biology and Occlusion, Faculty of Dentistry, Prince of Songkla University, 15 Karnjanavanich Road, Hat Yai, Songkhla 90112, Thailand

E-mail: ureporn.l@psu.ac.th

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modified glass ionomer cement supplemented with TCTP can reduce cell death from residual HEMA release (16). The aim of this study was to compare cell proliferation, anticytotoxicity, differentiation and mineralization of VEGF-A with TCTP on HEMA-treated human dental pulp cells (HDPCs). This will assist the further development of a restorative material that may have less toxicity and be able to regenerate pulpal dentine complex.

Materials and methods

Expression and purification of recombinant TCTP and recombinant VEGF. Expression of *Pmer-TCTP* gene was performed in the bacteria *Escherichia coli* (*E. coli*) strain BL21. The protein was purified according to methods previously described (14). Briefly, *E. coli* strain BL21 harboring pGEX-*Pmer-TCTP* was inoculated and induced by 1 mM IPTG (isopropyl β -D-thiogalactopyranositol) for stimulating protein expression. After induction for 3 h, the bacterial cells were harvested by centrifugation and the soluble glutathione S-transferase (GST)-TCTP protein was purified by using Glutathione Sepharose 4 Fast Flow (GE Healthcare Bio-Science, Piscataway, NJ, USA) and thrombin was used for splitting of GST-tagged protein. The purified *Pmer-TCTP* protein with molecular mass about 19.2 kDa (168 amino acid residues) was determined by SDS-PAGE and protein concentration was analyzed by a BCA protein assay kit (Pierce; Thermo Fisher Scientific, Inc., Waltham, MA, USA). Recombinant human VEGF was purchased from Gibco® (Gibco; Thermo Fisher Scientific, Inc.). This protein has about 40 kDa (homodimer) with 165 amino acid residues/subunit.

Cell culture. HDPCs were cultured from pulp tissue of sound third molar teeth of adults aged 18-25 years at the Dental Hospital, Faculty of Dentistry, Prince of Songkla University, with consent forms approved by the Research Ethics Committee (Number EC5703-09-P-LR), Faculty of Dentistry, Prince of Songkla University. Primary culture of pulp cells was performed using an enzymatic method. Briefly, the pulp tissue was cut into small pieces before digested in 3 mg/ml of collagenase type I and 4 mg/ml of dispase for 1 h at 37°C. After centrifugation, cells were cultured in α -MEM, supplemented with 10% fetal calf serum (FCS), 100 μ M L-ascorbic acid 2-phosphate, 100 μ M L-glutamate, 100 U/ml penicillin, and 100 μ g/ml streptomycin and incubated at 37°C with 5% CO₂. Pulp cell passages between three and five were used for this study.

Finding effective concentrations of TCTP and VEGF on HDPCs. The effect of TCTP and VEGF at various concentrations on HDPCs was examined by the 3-(4, 5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide (MTT) assay. TCTP and VEGF were diluted with cell culture medium at various concentrations varying from 1 ng/ml to 15 μ g/ml (n=5 in each group). HDPCs at a density of 5x10³ cells/well were cultured in 96-well plates in a humidified atmosphere of 5% CO₂ at 37°C for 24 h and then the culture medium supplemented with TCTP and VEGF were replaced. After 24 and 72 h, the medium was removed, 200 μ l of fresh medium containing 10 mM HEPES (pH 7.4) and 50 μ l MTT

solution (5 mg/ml in PBS) were added to each well and incubated in the dark for 4 h at 37°C. The medium and MTT were then removed and 200 μ l of DMSO and 25 μ l of Sorensen's glycine buffer (0.1 M glycine plus 0.1 M NaCl equilibrated to pH 10.5 with 0.1 M NaOH) were added. The optical density (OD) of formazan production was measured at 570 nm. The OD values adjusted for a blank (medium only) of the experimental groups were divided by the control (cell cultured in normal medium without TCTP and VEGF) and expressed as percentages of the control, which represented the percentages of viable cells.

Cytotoxicity of HEMA. MTT assay was used to evaluate the cytotoxicity of HEMA at two concentrations, which were 4 mM and 6 mM. HDPCs were seeded at 8x10³ cells/well in a 96-well plate at a humidified atmosphere of 5% CO₂ at 37°C. After 24 h, the media was replaced with HEMA mixed in culture medium and left for 24 h. Then the media was refreshed with normal media and incubated for 48 h before MTT assay was performed as described previously (n=5 in each group).

The results from HEMA cytotoxicity testing led to the selection of 4 mM HEMA as a cytotoxic reagent. In addition, the result from effective concentrations of TCTP and VEGF suggested the use of TCTP at 1 and 10 ng/ml of VEGF for further investigation.

The recovery effect of TCTP and VEGF on HEMA-treated HDPCs. There were five groups of HDPCs according to different medium conditions: HEMA, HEMA+TCTP, HEMA+VEGF, HEMA+TCTP+VEGF, and control group (culture medium only). HDPCs were seeded at 8x10³ cells/well on a 96-well plate at a humidified atmosphere of 5% CO₂ at 37°C and fed with normal medium. After 24 h, the culture medium was replaced with media supplemented with substances as described above for 24 h. After that, the fresh media was replaced and incubated for 48 h. The viable cells were determined by the MTT assay (n=5 in each group).

Alkaline phosphatase (ALP) activity assay. ALP activity assay was used to determine odontoblast-like differentiation. The experiment was divided to six groups according to substances added to the medium: HEMA, HEMA+TCTP, HEMA+VEGF, TCTP, VEGF and cells cultured in normal medium as a control group. HDPCs were seeded at 1x10⁴ cells/well on a 24-well plate and fed with normal culture medium using 15% FCS at a humidified atmosphere of 5% CO₂ at 37°C (n=3 in each group). The medium was changed every 2 days and the ALP activity of cells was determined after being cultured for 3, 7 and 14 days. The ALP activity was measured by using *p*-nitrophenol phosphate as a substrate. Cells were washed twice with PBS pH 7.4 and lysed in 20 μ l/well of lysis buffer (100 mM MgCl₂·6H₂O, 1 M Tris-HCl pH 7.4, 0.1% Triton-X 100, pH 10) on ice and centrifuged at 14,000 g at 4°C for 10 min. The supernatant was collected for total protein and ALP activity determinations. The total protein was determined by using BCA kit. For ALP activity assay, 5 μ l of the supernatant of each sample was incubated in 50 μ l of ALP substrate solution containing 0.2 M 2-amino-2-methyl-1-propanol (AMP), 4 mM MgCl₂ and 50 μ l of 4 mg/ml *p*-nitrophenol phosphate at 37°C for 30 min. Then, the reaction was stopped with 100 μ l of 0.1 M NaOH. The

Table I. qPCR F and R primer sequences.

Gene	Primer sequence (5'- 3')	GenBank accession no.
DSPP	F: GGGATGTTGGCGATGCA R: CCAGCTACTTGAGGTCCATCTTC	NM_014208.3
DMP1	F: GCAGAGTGATGACCCAGAG R: GCTCGCTTCTGTCTATCTTCC	NM_004407.3
ALP	F: CCACAAGCCCGTGACAGA R: GCGGCAGACTTTGGTTTC	NM_001127501.3
BMP-2	F: GCTTCCGCCTGTTTGTGTTTG R: AGAGACATGTGAGGATTAGCAGGT	NM_007553.2
GAPDH	F: GCACCGTCAAGGCTGAGAAC R: ATGGTGGTGAAGACGCCAGT	NM_002046

F, forward; R, reverse; DSPP, dentin sialophosphoprotein; DMP1, dentin matrix protein 1; ALP, alkaline phosphatase; BMP-2, bone morphogenetic protein-2.

absorbance was measured at 405 nm. ALP activity was calculated as nanomolar of *p*-nitrophenol per μg of total protein and adjusted to the increased percentage compared to the control (medium only).

Alizarin red S (ARS) staining assay. Alizarin red S (ARS) staining assay was used to evaluate calcium deposition. There were six groups according to different substances added in the inductive medium, which were HEMA, HEMA+TCTP, HEMA+VEGF, TCTP, VEGF and the last group was cells cultured in an inductive medium only as a positive control group. HDPCs at 1.5×10^5 cells/well were seeded on a 12-well plate and fed with 500 μl /well of inductive medium, which was composed of 1 M β -glycerophosphate, 10 mg/ml ascorbic acid, 100 U/ml penicillin, 100 mg/ml streptomycin and 100 mg/ml antibiotic-antimycotic, in α -MEM with 20% fetal calf serum ($n=3$ in each group) at 37°C in a humidified atmosphere containing 5% CO_2 . After the first 24 h, the medium was refreshed and then the medium was replaced every 2 days. After 7, 14 and 21 days, the medium was removed and cells were washed with PBS pH 7.4 and fixed in 10% (v/v) formaldehyde (Sigma-Aldrich; Merck KGaA, Darmstadt, Germany) at room temperature for 15 min and washed twice with 1 ml/well of ultrapure water at room temperature for 5 min. Then Alizarin Red solution was added 1 ml/well and incubated on shaker for 30 min. After removal of ARS, the plate was washed four times with ultrapure water, while shaking for 5 min, and then Store plates at -20°C prior to dye extraction. The monolayers were incubated by 200 μl cetylpyridinium chloride (CPC) buffer for 2 h on a shaker. After that, it was centrifuged at 20,000 g for 15 min and 100 μl of the supernatant was then transferred to a 1.5 ml tube. It was diluted (10 μl of sample and 90 μl of CPC) to a 96-well plate and the OD values corrected for a blank (CPC only). The absorbance of the sample was measured in an ELISA reader at 550 nm.

Gene expression. Quantitative real-time reverse transcription polymerase chain reaction (qRT-PCR) was used to investigate the effect of each sample on pulp cells differentiation. Dentin

sialophosphoprotein (DSPP), dentin matrix protein 1 (DMP-1), ALP and bone morphogenetic protein-2 (BMP-2) expressions were examined and gene glyceraldehyde-3-phosphate dehydrogenase (GAPDH) was selected to be an internal control. The primers (5'- and 3'-) were designed and shown in Table I. There were six groups in the experiment: HEMA 4 mM alone and with (1 ng/ml TCTP, 10 ng/ml VEGF) with pulp cells, 1 ng/ml TCTP, 10 ng/ml VEGF with pulp cells and cells cultured in normal medium was set as the control. The experiments were repeated three times. HDPCs at 1.5×10^6 cells were seeded on six-well plate with 2,000 μl /well of normal medium. After 24 h, the mixed-medium with selected supplements was replaced and then changed every 2 days. After 1, 3 and 7 days the medium was removed and cells were washed with PBS pH 7.4. The total RNA of cells was extracted and purified by RNeasy® Plus Micro kits (Qiagen, Inc., Valencia, CA, USA) and converted to cDNA with SuperScript™ III RT (Invitrogen; Thermo Fisher Scientific). The thermal profile for RT-PCR, under denaturing conditions was 65°C for 5 min followed by cDNA synthesis at 50°C for 30 min. The reaction was terminated at 85°C for 5 min. Then the remaining RNA was removed by adding 1 μl of RNase H at 37°C for 20 min and 3 μl of each cDNA product was fractionated by 1.2% agarose gel electrophoresis and visualized by ethidium bromide staining by ultraviolet (UV) transillumination. SYBR Green PCR master mix (Roche Diagnostics GmbH, Mannheim, Germany) was used for qPCR analysis. The qPCR mix consisted of 1 μl of cDNA, 2 μl of forward and reverse primer, 10 μl of SYBR Green and 5 μl of nuclease-free water, and the negative control was distilled water instead of cDNA sample. The reaction was performed manually in triplicates in each sample, at a final volume of 20 μl . The thermal profile for amplification of the investigated gene was followed by 40 cycles of denaturation at 95°C for 15 sec, annealing at 60°C for 30 sec and then elongation at 72°C for 30 sec. After the end of the last cycle, the final quantification was reported as an amount after normalization to reference gene GAPDH. Relative values were analyzed using comparative cycle threshold (CT) method ($\Delta\Delta\text{CT}$ method).

Figure 1. Effect of TCTP and VEGF on HDPCs viability investigated by MTT assay. Cells were treated with various concentrations of TCTP and VEGF at (A) 24 and (B) 72 h. Data were presented as the mean $\pm$ standard deviation ($n=5$). * $P<0.05$. ^a $P<0.05$ vs. TCTP control and ^b $P<0.05$ vs. VEGF control. TCTP, translationally controlled tumor protein; VEGF, vascular endothelial growth factor; HDPCs, human dental pulp cells.

Statistical analysis. The results of MTT were analyzed using the one way analysis of variance (ANOVA) and Tukey's post hoc-test to investigate the differences among the experimental groups. Unpaired t-test was used to compare the difference of means between two groups of TCTP and VEGF. Kruskal-Wallis and Dunnett multiple comparison were used to analyze the results of ALP assay, ARS assay and qRT-PCR. $P<0.05$ was considered as statistically significant.

Results

Effective concentration of TCTP and VEGF on HDPCs. HDPCs were treated with various concentrations from 1 to 15 $\mu\text{g/ml}$ of TCTP and VEGF in order to find the effective concentrations that are not cytotoxic to cells and can also promote cell growth. After culturing for 24 h, most of the various concentrations of TCTP can promote cell growth because the percentage of viable cells was over 100% compared to the control (Fig. 1A), especially at 1 ng/ml of TCTP, which gave the highest percentages of

cell viability ($P<0.05$). Cells cultured in medium supplemented with VEGF did not have a higher percentage of viable cells than the control after culturing for 24 h. However, after 72 h, it was found that at low concentration, especially at 1 ng/ml of TCTP and 10 ng/ml of VEGF, had significantly higher cell viability than the control and compared within groups ($P<0.05$). High concentration of VEGF at 15 $\mu\text{g/ml}$ significantly reduced cell viability (Fig. 1B). The results led to the determination to use 1 ng/ml of TCTP and 10 ng/ml of VEGF in culture medium for further studies.

The recovery effect of TCTP and VEGF on HEMA-treated HDPCs. HEMA-treated pulp cells at concentration of 4 mM and 6 mM demonstrated its high level of cytotoxicity, because the percentages of viable cells were all less than 10% (Fig. 2A). The result from Fig. 2B showed that HDPCs cultured in 4 mM of HEMA with either 1 ng/ml TCTP or 10 ng/ml VEGF and both of combinations TCTP and VEGF had percentages of cell viability that were not statistically different from the

Figure 2. MTT assay of HDPCs. (A) Cytotoxicity test of 4 mM and 6 mM HEMA on HDPCs. (B) The recovery effect of TCTP and VEGF on HEMA-treated HDPCs. (C) The morphological features of HDPCs in each group taken by inverted microscope, scale bar represented 500 μ m. Data were presented as the mean $\pm$ standard deviation (n=5). *P<0.05 vs. control group. HDPCs, human dental pulp cells; HEMA, 2-hydroxyethyl methacrylate. TCTP, translationally controlled tumor protein; VEGF, vascular endothelial growth factor.

control group, while cell viability in HEMA-treated cells was significantly lower than other groups ($P<0.05$) (Fig. 2B). The morphological features of HDPCs were shown in Fig. 2C. The results suggested that TCTP and VEGF had an ability to recover the viability of HEMA-treated pulp cells. It was noted that HEMA-treated cells supplemented with TCTP+VEGF did not increase the percentages of cell viability, when compared to either TCTP or VEGF.

ALP activity. The result of ALP activity was revealed in Fig. 3. HEMA significantly reduced ALP activity compared to the control and TCTP or VEGF groups in all three periods (day 3 $P<0.01$), (day 7 and day 14 $P<0.001$). HEMA+TCTP and HEMA+VEGF groups had slightly higher ALP activity than HEMA within each time period.

Mineralization effect of TCTP and VEGF on HEMA-treated HDPCs. The Alizarin red S (ARS) assay was used to determine calcium deposition and the results have been shown in Fig. 4. After culture for 7 days, the levels of calcium

deposition in TCTP and VEGF groups were higher than the others ($P<0.001$). However, calcium in group HEMA+VEGF was significantly higher than HEMA+TCTP and HEMA alone ($P<0.01$). In addition, after culture for 14 days, cells in groups HEMA+TCTP, HEMA+VEGF, TCTP, and VEGF had increased calcium deposition compared to the control. In particular, the HEMA+TCTP group had the highest calcium deposition ($P<0.001$). At this period, the level of calcium deposition in cells treated with TCTP and VEGF was reduced from 7 days. After culture for 21 days, HDPCs cultured in an inductive medium with added TCTP had increased calcium deposition and became the highest compared to other groups.

It was noted that HDPCs cultured in inductive medium only (control) had increased calcium deposition by time and reached its highest after 21 days. Cells treated with HEMA+TCTP had reached its highest calcium deposition after 14 days and then reduced, but were still the same level as control after 21 days. Furthermore, the groups of HEMA+TCTP and HEMA+VEGF had higher calcium deposition than the HEMA group at all three time periods.

Figure 3. ALP activity assay. Data were presented as the mean $\pm$ standard deviation (n=3). *P<0.05, **P<0.01 and ***P<0.001 vs. the control group within each time period; #P<0.05 and ##P<0.01 vs. the corresponding 3 day group. ALP, alkaline phosphatase.

Figure 4. Cell mineralization measured by ARS assay. Data were presented as the mean $\pm$ standard deviation (n=3). **P<0.01 and ***P<0.001 vs. the control group within each time period; #P<0.01, ##P<0.001 vs. the corresponding 7 day group. ARS, Alizarin red S.

Differentiation-related gene expression of HDPCs by qRT-PCR. The result of qRT-PCR was shown in Fig. 5. The response of DSPP gene was reported as an average of fold expression (Fig. 5A). At day 1, the expression of DSPP gene in cells treated with HEMA was lower than control, while upregulation was found in cells treated with TCTP and VEGF.

At day 3, cells treated with HEMA+TCTP (P<0.001) and HEMA+VEGF (P<0.01) had significantly higher upregulation than other groups. However, downregulation was found in all groups on day 7.

The high level of BMP-2 gene expression was observed in cells treated with HEMA+TCTP on day 1 (P<0.01) and on day

Figure 5. The results of RT-qPCR. (A) The fold change of DSPP expression, (B) BMP-2 expression, (C) ALP expression, and (D) DMP-1 expression. Gene expression was described as fold changes relative to the control group at the same time-point. Data were presented as the mean $\pm$ standard deviation (n=3). *P<0.05, **P<0.01 and ***P<0.001 vs. the control group within each time period; #P<0.05, ##P<0.01 and ###P<0.001 vs. the corresponding 1 day group. DSPP, dentin sialophosphoprotein. BMP-2, bone morphogenetic protein-2; ALP, alkaline phosphatase; DMP-1, dentin matrix protein 1.

3 (P<0.001) and HEMA+VEGF on day 1 and day 3 (P<0.001) (Fig. 5B). In addition, the expression of BMP-2 gene upregulated significantly highest in group HEMA+VEGF on day 1 and day 3 (P<0.001). BMP-2 gene was down regulated in all groups on day 7.

On day 1, the expression of ALP gene, in control group, HEMA+TCTP and TCTP was higher than other groups (Fig. 5C). Moreover, the upregulation of ALP expression in TCTP and VEGF groups increased immediately on day 3, and ALP expression in VEGF groups was significantly higher (P<0.01) compared to others which corresponded with the result of ALP activity in Fig. 3. ALP gene expression down regulated in all groups on day 7.

DMP-1 expression was reduced by time in all groups (Fig. 5D). DMP-1 expression in the control group, HEMA+VEGF and VEGF, were higher than other groups on day 1. The expression of DMP-1 gene upregulated significantly highest in VEGF group on day 3 (P<0.05). HDPCs treated with HEMA+VEGF (P<0.05), TCTP (P<0.05), and VEGF (P<0.01) were significantly higher compared to others on day 7.

Discussion

This study evaluated some biological properties of VEGF-A or VEGF, especially the anticytotoxicity, differentiation and mineralization in pulp cells compared to TCTP. Previous studies have reported that VEGF (3,5) and *Pmer*-TCTP (15) can promote pulp cell proliferation and differentiation, which

were confirmed by the results of this study. The VEGF used in this study is in a homodimer form, which has 165 amino acid residues per monomer. *Pmer*-TCTP is composed of 168 amino acid residues and the molecular mass is not much different from this VEGF monomer. The effective concentration of VEGF that can promote cell growth after 72 h and also used to promote survival of HEMA-treated HDPCs was 10 ng, which was the working concentration that induced stem cells to differentiate (3). The concentration that promoted growth of HDPCs of TCTP was lower (1 ng/ml) than VEGF (10 ng/ml), which may suggest that TCTP has a higher potential than VEGF in promoting HDPCs growth.

Both TCTP and VEGF can survive HEMA-treated pulp cells. But they did not have a synergistic effect, as pulp cells treated with HEMA+TCTP+VEGF did not increase the percentages of viable cells compared to HDPCs exposed to either HEMA+TCTP or HEMA+VEGF. This result may suggest that the mechanism whereby both VEGF and TCTP survive HEMA-treated pulp cells might relate to each other or share the same carrier or pathway. However, further investigation is required.

The mechanism of the added VEGF that can promote HDPCs proliferation and survive cells from HEMA exposure may involve signal transduction through receptor tyrosine kinases, vascular endothelial growth factor receptor 2 (VEGF-R2), and lead to cell proliferation by either activation of the mitogen-activated protein kinases or MAPK cascade via Raf stimulation or PLC (phospholipase C)- γ .

Activation of PI3K (phosphatidylinositol 3-kinase) leads to activation of PKB (protein kinase B) and the cell survival process (7,17). VEGF can also survive cells from hypoxic conditions via the PI3K/Akt/FoxO1 pathway (8). VEGF induces expression of the anti-apoptotic proteins Bcl-2 in vascular endothelial cells (18) and in leukemic cells (19).

TCTP and its anti-apoptotic effect against HEMA-treated pulp cells has previously been reported (15). The mechanism whereby TCTP can survive HEMA-treated pulp cells remains unknown. It was shown that TCTP can inhibit apoptosis by maintaining mitochondria stability and inhibiting Bax function (20), as well as forming heterocomplexes with Bcl-xL (21). However, TCTP also inhibits apoptosis by stabilizing anti-apoptotic Bcl-2 family proteins, MCL1 and by inhibiting p53-dependent apoptosis by down regulating this protein (14).

The results of ALP activity assay and cell mineralization by the ARS assay confirmed the results of other studies, which showed that TCTP (16) and VEGF (2) can enhance differentiation and mineralization of pulp cells. This was also consistent with qRT-PCR results that both TCTP and VEGF had up regulated ALP expression on day 3 compared to the control. TCTP may promote cell mineralization longer than VEGF, as shown in the results of calcium deposition after 21 days. DMP-1 is a phosphoprotein that activates both bone and dentin mineralization by inducing the deposition of mineral particles along the collagen fibrils (22). HDPCs had lower expression of DMP-1 in longer culture time in all groups, but it was noticed that TCTP and VEGF groups still had higher expression than the control after culture for 3 and 7 days, respectively.

This study revealed that HEMA at low concentration (4 mM) was cytotoxic to cells and caused down regulation of DSPP, BMP-2, ALP and DMP-1 expression. TCTP and VEGF up regulated DSPP expression on day 1 and BMP-2 expression on day 1 and day 3, which supports that both proteins can promote pulp cells differentiation, especially in TCTP/VEGF supplemented in HEMA-treated pulp cells. A recent study revealed that VEGF may have synergist effect to BMPs, including on cell survival and mineralization (23). Further investigation may be required concerning the potential connection of the two mechanisms protecting cell death and differentiation, since both VEGF and TCTP had anticytotoxicity and promoted cell differentiation.

In conclusion both TCTP and VEGF promoted pulp cells proliferation and survived HEMA-treated cells, but TCTP was required a lower concentration for these functions. They did not have a synergistic effect; however both of them can enhance cell differentiation and mineralization.

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Availability of data and material

The datasets generated and analyzed during the current study are not currently publicly available due to work in progress on a suitable repository in Thailand, however they are available from the corresponding author on reasonable request.

Authors' contributions

The current study was conceived by UK. CW performed the experiments, analyzed and interpreted the data in conjunction with WC and UK. CW and UK were the major contributors in writing the draft manuscript. All authors read and approved the final manuscript.

Ethics approval and consent to participate

The study was approved by the Human Research Ethics Committee, Faculty of Dentistry, Prince of Songkla University (approval number EC5703-09-P-LR) and by the Dental Hospital, Faculty of Dentistry, Prince of Songkla University. This included the use of approved consent forms for the collection of specimens.

Consent for publication

Not applicable.

Competing interests

The authors declare that they have no competing interests.

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