Role of BMP-4 and Its Signaling Pathways in Cultured Human Melanocytes

Park, Hee-Young

Hindawi Publishing Corporation

http://hdl.handle.net/2144/3403

Boston University
Role of BMP-4 and Its Signaling Pathways in Cultured Human Melanocytes

Hee-Young Park, Christina Wu, Mina Yaar, Christina M. Stachur, Marita Kosmadaki, and Barbara A. Gilchrest

Department of Dermatology, School of Medicine, Boston University, Boston, MA 02118, USA

Correspondence should be addressed to Hee-Young Park, hypark@bu.edu

Received 22 July 2009; Accepted 1 October 2009

Bone Morphogenetic Protein (BMP-4) was shown to down-regulate melanogenesis, in part, by decreasing the level of tyrosinase [Yaar et al. (2006) JBC:281]. Results presented here show that BMP-4 down-regulated the protein levels of TRP-1, PKC-β, and MC1-R. When paired cultures of human melanocytes were treated with vehicle or BMP-4 (25 ng/ml), MAPK/ERK were phosphorylated within one hour of BMP-4 treatment. Then the activated MAPK/ERK caused an acute phosphorylation of MITF, followed by proteosome-mediated degradation of MITF, the key transcription factor for melanogenic proteins [Wu et al. (2000) Gene & Development:14]. However, prolonged exposure of melanocytes to BMP-4 (up to 48 hours) caused a decrease in the level of MITF-M transcript. In addition, BMP-4 decreased the intracellular level of cAMP, the key regulator of MITF expression. These results demonstrate that BMP-4 activates MAPK/ERK signaling pathway to transiently activate MITF; however, chronic treatment of BMP-4 to melanocytes causes a down-regulation of the expression of MITF, possibly in a cAMP-dependent pathway.

Copyright © 2009 Hee-Young Park et al. This is an open access article distributed under the Creative Commons Attribution License, which permits unrestricted use, distribution, and reproduction in any medium, provided the original work is properly cited.

1. Introduction

Bone morphogenetic proteins (BMPs) are disulfide-linked dimeric proteins, produced as large precursor proteins (reviewed in [1]). BMPs can be further classified into several subgroups by their amino acid similarities. BMP-2 and BMP-4 have 83% amino acid sequence identity and are the best-studied members in the BMP family (reviewed in [1]). BMP-2 and -4 elicit their biological actions by first binding to their specific cell surface receptors, termed type II receptor (BMPR-II) and type I receptor (BMPR-I) [2–6]. Two different types of BMPR-I, termed BMP type IA (BMPR-IA) and type IB (BMPR-IB), have been identified [7–9]. Binding of BMP-2 and BMP-4 to BMPR-II is facilitated by the type I receptors [4, 6]; and upon binding with BMP-2 or -4, BMPR-II subsequently activates type I receptors [10]. Activated BMPR-I phosphorylates Smad-1, -5, or -8 [11], then recruits Smad-4, and the entire complex translocates to the nucleus [12–14] and interacts with specific DNA consensus sequences to either activate or suppress transcription of target genes. Activated BMPR-I can also phosphorylate (activate) extracellular-signal regulated kinase (ERK) via mitogen activated protein kinase (MAPK) and this pathway can also cross-talk with Smads-dependent pathway [15, 16].

Recently, it was reported that BMP-4, a member of the TGF-β superfamily (reviewed in [17]), negatively modulates melanogenesis in part by reducing the level of tyrosinase expression in cultured human melanocytes [18]. Treatment of quail neural crest cultures with BMP-4 caused a dramatic decrease in melanogenesis [19]. Moreover, in a transgenic mouse model which overexpresses noggin, the known physiological inhibitor of BMP-4 [20], the level of microphthalmia-associated transcription factors (MITF), and binding of α-melanocyte stimulating hormone (α-MSH) to melanocortin receptor-1 (MC1-R) were increased, resulting in darkening of the coat color [21].

The synthesis and dispersion of melanin within the epidermis is largely responsible for skin color as well as for protection from sun-induced injuries. Melanogenesis is initiated when the amino acid tyrosine is oxidized to dopa and dopa quinine by the enzyme tyrosinase, the key
and rate-limiting enzyme in melanogenesis (reviewed in [22]). Tyrosinase exists as an inactive form and is activated when protein kinase c-β (PKC-β) phosphorylates the serine residues in the cytoplasmic domain of this enzyme [23]. Activation of tyrosinase by PKC-β was shown to be required for melanogenesis both in vivo [24] and in vitro [23, 25–27], while conversely increased activity and/or expression of PKC-β leads to increased pigmentation both in vitro [28] and in vivo [29].

Human melanogenesis is heavily influenced by both paracrine and autocrine factors such as α-MSH [30], endothelin-1 [31], interleukin-1 [32], and TNF-α [32]. Among these factors, α-MSH and endothelin-1 induce pigmentation in part through increased expression and activity of tyrosinase [30, 31, 33]. The expression of both α-MSH and endothelin-1 was shown to be increased in epidermal keratinocytes and induced when keratinocytes were exposed to ultraviolet irradiation (UV) [31, 34]. Endothelin-1 was shown to induce melanogenesis by activating multiple signaling pathways, including the PKC-dependent pathway [35]. On pig cell melanocytes, α-MSH first binds to its cell surface receptor MC1-R [36, 37], a heptahelical transmembrane protein coupled to the G-protein [38], and thereby increases the intracellular level of cAMP [33]. One way in which cAMP increases pigmentation is by increasing the expression of MITF [39], shown to be a key transcription factor for melanogenic protein, such as tyrosinase and tyrosinase related protein-1 (TRP-1) and PKC-β [40–42].

MITF is a basic-helix-loop-helix (bHLH) and leucine zipper transcription factor that binds to conserved consensus elements in gene promoters, specifically the M-(AGTCATGTGCT) and E-(CATGTG) boxes [43], and regulates the transcription of tyrosinase, TRP-1, TRP-2 [41], PKC-β [40], and MC1R [44]. MITF can bind as a homodimer or a heterodimer with another related family members (reviewed in [45]). MITF comprises a family of at least nine isoforms [46, 47], of which the MITF-M isoform controls tyrosinase and PKC-β transcription in melanocytes [40, 48, 49]. MITF activity and stability are regulated by its phosphorylation state through MAPK/ERK-dependent pathway [50, 51]. Upon phosphorylation, MITF binds to either M-box or E-box consensus sequences to activate transcription [51, 52], but phosphorylation then decreases MITF stability and enhances its degradation in proteosomes [50, 53]. Thus, prolonged activation of MAPK/ERK pathway would result in down-regulation of MITF protein level. More recently, MITF was shown to be a key transcription factor for Rab27a [54], a protein important for melanosome transport, and Pmel17 [48], a protein required for melanosome matrix formation. Therefore, MITF plays a central role in melanin synthesis, as well as melanosome biogenesis and transport.

While factors that increase melanogenesis are well understood, physiological agents that suppress melanogenesis are poorly understood. Recently, it has been shown that interleukin-1α and -6, as well as TNF-α decrease melanogenesis in human melanocytes [32]. TGF-β was also shown to decrease pigmentation [55].

In this paper, we further delineate the mechanism by which the BMP-4-dependent signaling pathway down-regulates melanogenesis. Specifically, the intracellular signaling pathways that mediate the effects of BMP-4 and its target genes/proteins are explored.

2. Materials and Methods

2.1. Materials. Recombinant human BMP-4 was obtained from R&D Systems, Inc., Minneapolis, MN. Dulbecco’s modified essential medium (DMEM), L15, nonessential amino acids, glutamine, Medium 199, and trypsin were purchased from Gibco/BRL, Gaithersburg, MD. Recombinant basic fibroblast growth factor was purchased from Amgen Inc., Thousand Oaks, CA, bovine pituitary extract (BPE) from Clonetics Corp., San Diego, CA, and hydrocortisone from Calbiochem, San Diego, CA. Insulin, transferrin, and dbcAMP were from Sigma Chemical Co., St. Louis, MO. Nylon membranes and Tag-Ready-to-Go PCR Beads were from Amersham Biosciences, Piscataway, NJ. Bovine calf serum and fetal calf serum were from HyClone Laboratories, Inc., Logan, UT. Polyvinyl difluoride (PVDF) membranes were from BioRad Laboratories, Carlsbad, CA. FuGene6 was purchased from Roche Diagnostic Corp., Nutley, NJ. Lipo-fectamine Plus reagent was purchased from Invitrogen, Hercules, CA, and promoterless renilla luciferase construct was purchased from Promega, Madison, WI.

2.2. Antibodies. Monoclonal antibody specific for PKC-β (dilution: 1:50) was purchased from BD Biosciences, San Jose, CA. Polyclonal antibodies against phosphorylated smad 1/5/8 (dilution: 1:10), TRP-1 (dilution: 1:1000), and MC1R (dilution: 1:100) were purchased from Santa Cruz Biotechnology, Santa Cruz, CA. Monoclonal antibody for tyrosinase (dilution: 1:1000) was purchased from Vector Laboratories, Inc., Burlingame, CA. Monoclonal antibody for MITF (dilution: 1:10) was a generous gift from Dr. David Fisher at the Department of Dermatology, Massachusetts General Hospital, Boston, MA.

2.3. Cells and Media. Neonatal foreskins were used to obtain human melanocytes as previously described [23]. In brief, the epidermis was separated from the dermis after overnight incubation in 0.25% trypsin at 4°C. Primary cultures were then established in Medium 199 containing 1.8 mM Ca++ and supplemented with 10−9 M triiodothyronine, 10 μg/mL insulin, 1 × 10−6 M hydrocortisone, BPE (35 μg/mL), 80 U/mL dbcAMP, 10 ng/mL basic fibroblast growth factor, and 5–10% fetal bovine serum. All post primary cultures were maintained in low calcium (0.03 mM). Cells at first or second passage were used for all experiments. LH melanoma cells were maintained in DMEM containing 10% of calf serum.

2.4. Immunoblot Analysis. Immunoblot analysis was performed as previously described [23]. Cells were harvested in RIPA buffer (0.25 M Tris (pH 7.5), 0.5 M NaCl, 2.5% SDS, and 0.1% Triton-X-100) containing protease inhibitors
(100 μM Okadaic acid, 100 μM PMSE, 100 μg/mL Apro- 

tinin, 2 μg/mL Leupeptin, and 100 μM Sodium Ortho-

Vanadate). Protein samples were subjected to 7.5% SDS-

PAGE and transferred to a PVDF membrane electrophoret-

ically. The membrane was preincubated in 100% Blotto 

(5 g nonfat dry milk in 100 mL phosphate buffered saline 

(PBS)) for 1–3 hours at room temperature with shaking, 

followed by an overnight incubation with antiserum (0.5– 

1 μg/mL in 10% Blotto) at 4°C [18, 26, 50]. At the end of the 

incubation, the membrane was washed extensively with PBS 

containing 0.5% Tween-20, and processed using the ECL kit. 

The membrane was then exposed to Kodak X-OMAT film.

2.5. Vectors. PKC-β promoter-CAT reporter construct and 

control expression vectors were obtained from Dr. Denise 

Cooper at the University of South Florida [56].

2.6. Transient Transfection. In order to introduce PKC-β 

promoter-CAT reporter construct into LH melanoma cells, 

4 μg of DNA were pretreated with Lipofectamine Plus reagent 

and as an internal control for the transfection efficiency, 

0.4 μg of promoterless renilla Luciferase was cotransfected. 

24 hours after transfections, cells were treated with vehicle or 

BMP-4 (25 ng/mL). Cells were then harvested from 48 to 72 

hours after BMP-4 treatment. CAT and luciferase assays were 

performed according to the manufacturer’s protocol.

2.7. Semi Quantitative RT-PCR. To determine the effect of 

BMP-4 on the level of PKC-β transcripts, semiquantita-

tive RT-PCR was first performed. Specifically, cDNA was 

made from total RNA isolated from melanocytes using 

oligo pd (N)₆ (Pharmacia Fine Chemicals, Piscataway, NJ). 

Primers specific for PKC-β that recognize both βI and 

βII were purchased from Oxford Biomedical Research, 

Inc. For the internal control, primers against GAPDH 

(forward: 5’ GTCATCATCTCTGGCCCTTC3’ and backward: 

3’ CCGCACTACCGGACCCCCGT5’) were used. 0.1 μg of 

cDNA was amplified with 15 pmol of each forward and 

backward primers. Initially, 37 cycles of amplification were 

performed on cDNA as well as on RNA from each sample 

to assure no contamination of RNA samples with genomic 

dNA. All PCR reactions were amplified for 29 cycles which 

fell within the exponential phase of the amplification for 

optimal detection of PKC-β and BMP receptors and allow-

ing quantitative analysis of mRNA transcript modulation. 

Denaturation was performed at 94°C for 30 seconds, primer 

annealing at 58°C for 1 minute, and DNA polymerization 

at 72°C for 1 minute in a thermal cycler (MJ Research 

Inc., Waltham, MA). PCR products were separated over 

a 1% agarose gel in 1X TAE and stained with ethidium 

bromide. The levels of PCR products were subjected to the 

densitometric analysis. PKC-β transcript levels were then 

normalized against the level of GAPDH transcript.

2.8. Quantitative Real-Time PCR (qRT-PCR). The influence 

of BMP-4 on the level of MITF-M in vehicle vs. BMP- 

4 treated melanocytes was analyzed using MyiQ™ 

optical module Real-Time PCR system (Biorad) and 

iQ SYBR Green Supermix containing the MITF-M 

upstream primer: 5′ CACGGGTCTCTGCTTCC3′ and 

downstream primer: 5′ GGTGGTGTGGAAGGTGATGG3′. A second set of MITF-M primers was designed 

and used to confirm the results: upstream primer: 

5′ ATAGAAAGTAGGGAGGAGTGG5′ and downstream 

primer: 5′ TTATTTGCTAAAGTGTTGAAAGG3′. The PCR 

reaction was performed in a 50 uL volume containing 2–4 ng 
cDNA. The final primer and probe concentrations were 

optimized for each primer/probe combination. Two-step 

PCR cycling was carried as follows: 95°C 5 minutes × 1 

cycle, 95°C 0.5 minute, 55°C 0.5 minute × 40 cycles. As 

all internal control, GAPDH was amplified in parallel for 

normalization.

2.9. Densitometric Analysis. Densitometric analysis was per-

formed by scanning the autoradiograms of western and 

northern blots and PCR products into a computer (PC 

Dell™). Intensity of each scanned band was determined 

using BIO-RAD Gel Doc 1000/2000 imaging densitometer 

after background subtraction.

3. Results

3.1. BMP-4 Decreased the Protein Levels of TRP-1, PKC-β, and 

MC1-R. BMP-4 was shown to down-regulate melanogene-

sis, the key differentiated function of melanocytes, in part 

by decreasing the expression of tyrosinase [18]. To examine 

whether BMP-4 affects the expression of other melanogenic 

proteins, paired cultures of subconfluent primary human 

melanocytes were treated with vehicle (4 mM HCl/0.01% 

BSA) or BMP-4 at 25 ng/mL; the optimal concentration 

to treat melanocytes as previously reported [18], for 24 

and 48 hours and the protein levels of TRP-1, PKC-β, and 

MC1-R were assessed, using specific antibodies against these 

proteins. Results showed that BMP-4 decreased the protein 

level of PKC-β by 49.9 ± 29.7% (P<.04) at 24 hours and 47.7 

± 18.0% (P<.04) at 48 hours (Figure 1(a)), a magnitude of 

reduction known to reduce tyrosinase phosphorylation and 

activity in human melanocytes [23, 27]. The protein level of 

TRP-1 was decreased by 45% ± 5.0% (P<.05) and 36 ± 

6.5% (P<.045) at 24 and 48 hr time points, respectively 

(Figure 1(b)). The protein level of MC1-R was decreased by 

50% ± 2.0% (P<.03) and 55% ± 2.5% (P<.05) at 24 

and 48 hours, respectively (Figure 1(c)). In summary, BMP- 

4 reduced the protein levels of PKC-β, TRP-1, and MC1-R. 

Earlier time points did not show significant changes in the 

levels of these proteins in response to BMP-4 treatment.

3.2. BMP-4 Activated Both Smads-1/5/8 and MAPK/ERK 

Signaling Pathways. It is well documented that BMP-4 

regulates transcription of target genes through Smads 1/5/8-

mediated Smad proteins activation [11–14] and/or MAPK/ERK dependent pathways [15, 16]. To examine whether one or both of these pathways are 

involved in mediating effects of BMP-4 on the expression of melanogenic proteins in human melanocytes, phospho-

rylation (activation) of Smads 1/5/8 involved in mediating e

response to BMP-4 treatment.
with vehicle or BMP-4 (25 ng/mL) for 2 and 4 hours as previously done [18]. Then cells were harvested for immunoblot analysis using a polyclonal antibody that reacts against phosphorylated Smads-1/5/8. Results show that Smads-1/5/8 are phosphorylated by BMP-4 within 2 hours of treatment (Figure 2(a)).

To investigate whether BMP-4 can also activate MAPK/ERK-dependent pathway in human melanocytes, paired cultures of melanocytes were treated with vehicle or BMP-4 (25 ng/mL) for 15 and 30 minutes. Tumor promoter phorbol dibutyrate (PDBu) was used as a positive control as shown before [50]. As shown in Figure 2(b), ERK1/2 was phosphorylated within 30 minutes of BMP-4 treatment. As expected, PDBu phosphorylated ERK1/2 within 15 minutes of treatment. These results suggest that both Smads-1/5/8 and MAPK/ERK-pathways are activated by BMP-4 and may be involved in mediating BMP-4 responses in cultured human melanocytes.

3.3. Level of Phosphorylated MITF is Transiently Increased by BMP-4. It is well known that activated MAPK/ERK pathway can phosphorylate MITF, which results in a transient activation and then degradation of MITF [50, 51]. MITF is the key transcription factor that regulates the expression of TRP-1, PKC-β, and MC1-R. Therefore, it is possible that the effect of BMP-4 on the expression of these melanogenic proteins is mediated in part or totally through MITF. To test whether BMP-4 treatment causes phosphorylation of MITF, paired cultures of melanocytes were treated with BMP-4 (25 ng/mL) for 0, 0.5, 1, 2, and 5 hours. Results show a high basal level of phosphorylated MITF (0 hr). Then the level of phosphorylated MITF increased by a 2–4 fold (\( P < .03 \)) at 1 hour, and there was a subsequent reduction at 5 hours time point (Figure 3(a)). To confirm that phosphorylation of MITF by BMP-4 treatment is mediated through MAPK/ERK-dependent pathway, paired cultures of melanocytes were treated with MAKP inhibitor PD98056 (10 uM) or vehicle
BMP-4 (hrs)  
0 0.5 1 2 5

MITF- P
MITF →
Actin
(a)

MITF- P
MITF →
Actin
(b)

MITF- P
MITF →
Actin
(c)

Figure 3: Effects of BMP-4 on the level of phosphorylated level of MITF. (a) Paired cultures of melanocytes were treated with BMP-4 (25 ng/mL) for 0, 0.5, 1, 2, and 5 hours. At each time point, cells were harvested and immunoblot analysis using monoclonal antibody against MITF was performed (provided by Dr. David Fisher from the Dana Farber Cancer Institute). This antibody has been shown to recognize both phosphorylated and nonphosphorylated MITF. As the loading control, the membrane was immunoblotted for actin. One representative result from three separate experiments is shown. (b) Paired cultures of melanocytes were treated with BMP-4 (25 ng/mL) for 15 and 30 minutes in presence or absence of MEK inhibitor PD98509 at 10 μM. At each time point, cells were harvested and the level of MITF protein was determined. One representative result from three separate experiments is shown. (c) Paired cultures of melanocytes were treated with BMP-4 (25 ng/mL) and cycloheximide (15 μg/mL) for 0, 0.5, 1, 2, and 5 hours in presence or absence of proteosome inhibitor MG-132 at 10 μM. At each time point, cells were harvested and the level of MITF protein was determined using immunoblot analysis. One representative result from three separate experiments is shown.

(DMSO). PD98056 completely blocked the BMP-4 induced MITF phosphorylation (Figure 3(b)), confirming that BMP-4 phosphorylates MITF through MAPK/ERK-dependent pathway.

Since it is well known that phosphorylation of MITF by MAPK/ERK-dependent pathway results in the proteosome-mediated degradation of MITF [50, 51], the decreased level of MITF by BMP-4 treatment at 4 to 5 hours is likely due to degradation of MITF. To further confirm this possibility, paired cultures of melanocytes were treated with BMP-4 (25 ng/mL) for 0, 1, 2, 3, and 5 hours in presence or absence of proteosome inhibitor MG-132 (10 μM). Protein synthesis inhibitor cycloheximide (15 μg/mL) was added to all treatment groups. Results show that BMP-4 first increased the level of phosphorylated MITF (0.5 to 1 hour) as expected and then the level of MITF began to decline at 2 hours and was significantly reduced at 5 hours. Presence of the proteosome inhibitor MG-132 completely blocked the degradation of MITF (Figure 3(c)). These results clearly demonstrate that BMP-4 transiently phosphorylated MITF, thus potentially promote acute melanogenesis, followed by down-regulation.

3.4. Chronic Treatment of BMP-4 Decreased Total MITF. If BMP-4 causes a transient phosphorylation of MITF, followed by proteosome-mediated degradation, then a long term treatment of BMP would ultimately result in a decreased level of total MITF. To test this possibility, paired cultures of
3.5. BMP-4 Reduces the Intracellular Level of cAMP in Melanocytes. Results thus far suggest that BMP-4 transiently activates MITF through MAPK/ERK-dependent pathway. Our results also suggest that BMP-4 transcriptionally regulates the expression of MITF. Recent reports have suggested that cAMP-dependent pathway may mediate BMP-4 induced changes [57]. Because cAMP has been implicated to be the key molecule inducing pigmentation [39, 41], in part through inducing the expression of MITF-M [40], possible effects of BMP-4 on the level of intracellular cAMP in cultured human melanocytes were explored. Paired cultures of subconfluent melanocytes were treated with BMP-4 (25 ng/mL). Then cells were harvested at each time point and the level of cAMP was measured using enzyme-linked immunoblot assay (ELISA). Within 18 hours of BMP-4 treatment, cAMP level began to decrease and by 48 hours, BMP-4 significantly reduced the intracellular level of cAMP (Figure 5). This result suggests that part of BMP-4 effects on the expression of MITF may be mediated through cAMP-dependent pathway.

3.6. Effect of BMP-4 on the Promoter Activity of PKC-β. It is not clear to what extent Smads-1/5/8 pathway is involved in mediating the effects of BMP-4 on the expression of TRP-1, PKC-β, MC1-R, and MITF. Published nucleotide sequences of the promoter regions of tyrosinase, and PKC-β revealed that they both contain consensus sequences for Smad-1 and Smad-5 [58, 59], suggesting that BMP-4 could regulate the expression of these two key melanogenic proteins through Smad-dependent pathway.

To explore the possibility that BMP-4 may regulate PKC-β expression, in part, by suppressing the promoter activity of the PKC-β gene, presumably through Smad-dependent pathway, the level of PKC-β transcripts was first examined using semiquantitative RT PCR. As shown in Figure 6(a), two
transcripts of PKC-β were identified and the levels of both transcripts were significantly decreased by BMP-4 (Figure 6).

To demonstrate whether BMP-4 affects the promoter activity of PKC-β, a PKC-β promoter-CAT reporter construct was transfected into LH human melanoma cells. A promoterless renilla-luciferase reporter construct was cotransfected into each plate to normalize the transfection efficiency among plates, and results were expressed as CAT activity per luciferase activity. Plates were treated with BMP-4 (25 ng/mL) or vehicle alone 24 hours after transfection, and cells were harvested 48 hours after BMP-4 treatment. In eight separate experiments, the promoter activity of PKC-β was reduced by ∼25 ± 5% in BMP-4-treated cells when compared to vehicle-treated cells (Figure 7). A paired Student t-test showed that suppression of the promoter activity of PKC-β by BMP-4 was statistically significant (P < .01). Although the effect of BMP-4 on the promoter activity of PKC-β was statistically significant, the total level of reduction was modest and less than the reduction level seen in both PKC-β mRNA and protein levels. These results suggest that BMP-4 utilizes only partially Smads-dependent pathway to regulate the expression of PKC-β.

4. Discussion

Bone morphogenetic proteins (BMPs) were shown to be important for the development of melanoblasts during embryogenesis (reviewed in [19]). In recent years, it has
been shown that BMPs, BMP 2/4 in particular, may play key roles in the biological function of differentiated epidermal melanocytes as a paracrine and/or autocrine factor by influencing the activities of melanocytes. Both epidermal keratinocytes and melanocytes express BMP-4 and its receptors; BMPR-IA, BMPR-IB, and BMPR-II, and both BMP-4 and all three BMP-4 receptors are down-regulated in these cells when exposed to UV [18]. Moreover, BMP-4 and its signaling pathway have recently been suggested to be a novel modulator of human pigmentation [18].

BMP-4 is able to exert its biological action through Smads- and/or MAPK/ERK signaling pathways [15, 16]. Our results showed that BMP-4 treatment caused activation of both signaling pathways. However, results strongly suggest that BMP-4 preferentially utilizes MAPK/ERK-pathway to regulate synthesis of melanin or melanogenesis once it binds to its cell surface receptors. Experiments presented in this paper demonstrate that one of the major target proteins of BMP-4 activated MAPK/ERK pathway in human melanocytes is MITF. Modulation of activity and/or expression of MITF could have a significant impact on the biology of melanocytes because MITF is the central transcription factor for many of the melanogenic proteins, including tyrosinase [41], TRP-1, [41] PKC-β [40–42], and MC1R [44], as well as antiapoptotic protein bcl2 [63]. When BMP-4 is added to the melanocyte cultures, MITF is transiently phosphorylated by BMP-4 via MAPK/ERK-dependent signaling pathway and this phosphorylation initially activates the transcriptional activity of MITF, as previously shown [50, 51]. Therefore, BMP-4 can acutely promote melanogenesis. However, the phosphorylated MITF results in proteosome-mediated degradation, as previously reported [53]. The eventual decreased level of MITF by BMP-4 would then cause an overall decrease in the levels of melanogenic proteins and thus, cause down-regulation of melanogenesis.

Cell surface receptor c-kit, known to be activated after the interaction with its ligand Stem cell factor (CSF), was one of the first cell surface receptors on melanocytes that was shown to activate MAPK/ERK pathway that causes the transient phosphorylation of MITF and thus, activation of this transcription factor [51]. It was recently shown that c-kit has a soluble form that is released from melanocytes and sequesters CSF and therefore, inhibits melanogenesis [64]. It would be interesting to determined whether BMP-4 and soluble c-kit could work synergistically to down-regulate the level of MITF.

Our results further suggest that effects of BMP-4 on the expression of TRP-1, PKC-β, and MC1-R are mainly mediated indirectly through decreasing the level of MITF. Decreased level of MITF would then lead to a decreased level of transcriptional activities of these genes. In fact, it caused a reduction in the level of the transcripts of PKC-β. However, part of reduction in PKC-β transcripts by BMP-4 is through Smad-dependent pathway, as explained later. Nonetheless, MAPK/ERK-dependant pathway appears to play a larger contribution to the reduced level of PKC-β through MITF.

Our results further suggest that BMP-4 down-regulates the protein level of MITF first by proteosome-mediated degradation. It is then followed by a decrease in the level of MITF-M transcripts, suggesting that BMP-4 can transcriptionally down-regulate the expression of MITF-M. This decrease in the level of MITF-M may be mediated by cAMP-dependent pathway. Results showed that BMP-4 decreased the intracellular level of cAMP. It has been reported that BMP-4 can regulate the intracellular level of cAMP [19, 65, 66]. In melanocytes it is not clear whether BMP-4 down-regulates the intracellular level of cAMP through down-regulating the expression of MC1-R, or by other mechanism. It is unlikely that Smads-dependent pathway is involved in down-regulation of MITF-M transcripts because the promoter region of MITF gene does not contain DNA consensus sequences for Smads binding sites [46].

MITF was also shown to positively regulate the expression of the antiapoptotic protein bcl2 [63]. Therefore, decreased level of MITF by BMP-4 could eventually affect the survival of melanocytes. Yaar et al. have reported that short-term exposure of BMP-4 to melanocytes did not change the proliferation rate and it did not cause apoptosis of melanocytes [18]. However, BMP-4 could potentially increase the susceptibility of melanocytes to damages such as UV and other chemical agents that could reduce survival of melanocytes. Therefore, role of BMP-4 on melanocyte may have implications beyond its role in melanogenesis.

Smads 1, 5, and/or 8 were phosphorylated by BMP-4 within 2 hours of treatment in cultured human melanocytes. Moreover, Yaar et al. have shown that activated (phosphorylated) Smad 1, 5, and/or 8 translocates to nucleus [18]. Among the melanogenic proteins examined, only the promoter regions of PKC-β reveal twelve Smad-1 and one Smad-5 consensus sequences [58], suggesting that BMP-4 could directly regulate the transcriptional activity of the PKC-β gene. Transfection of a PKC-β promoter–CAT reporter construct into LH cells showed modest but significant suppressions of PKC-β promoter–CAT reporter activity. The level of decrease in PKC-β transcript by BMP-4 was greater than the suppression of the PKC-β promoter activity. These results suggest that the effect of BMP-4 on PKC-β expression may involve direct transcriptional regulation. However, clear demonstration of direct interaction between PKC-β promoter and Smads 1, 5, and/or 8 would require additional experiments involving gel-shifts and/or CHIP assays.

In murine melanoma cells, TGF-β decreases pigmentation by decreasing the stability of tyrosinase mRNA and protein [67], as well as decreasing the level and the activity of MITF [55], TGF-β was shown to decrease only the level of phosphorylated MITF [55]. However, results presented in this paper show that the levels of both phosphorylated and nonphosphorylated MITF were decreased by BMP-4 in cultured human melanocytes. Even though BMP-4 belongs to the superfamily of TGF-β, their detailed mechanisms of action may differ with regard to melanogenesis.

The findings of the present in vitro studies are likely of great in vivo relevance, as transgenic mice in which BMP-4 action is antagonized by overexpression of its physiologic inhibitor noggins in hair follicles have a markedly darker coat color [21].

Our results suggest that the BMP-4-dependent pathway is a key physiologic negative modulator of melanogenesis,
suppressing the expression of melanogenic proteins and signaling molecules. BMP-4 may thus provide a novel target for modulation of melanogenesis.

**Abbreviations**

BMP-4: Bone-morphogenetic protein 4  
PKC-β: Protein kinase C-β  
MITF: Microphthalmia-associated transcription factor  
CAT: Chloramphenicol acetyl transferase  
DMEM: Dulbecco’s modified essential medium  
BPE: Bovine pituitary extract  
TRP-1: Tyrosinase related protein-1  
MC1-R: Melanocortin receptor 1  
MAPK: Mitogen-activated protein kinase  
ERK: Extracellular-signal regulated kinase  
UV: Ultraviolet irradiation.

**Acknowledgments**

This work was supported in part by the Amal K. Kurban Career Development Award and National Vitiligo Foundation (H.Y. Park). The authors would like to thank Dr. Denise Cooper at the University of South Florida for providing the PKC-β promoter construct and Dr. David Fisher at the Department of Dermatology, Massachusetts General Hospital, Boston, MA for providing monoclonal antibody against MITF.

**References**


