Glyceollins, a novel class of soybean phytoalexins, inhibit SCF-induced melanogenesis through attenuation of SCF/c-kit downstream signaling pathways

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The anti-melanogenesis effect of glyceollins was examined by melanin synthesis, tyrosinase activity assay in zebrafish embryos and in B16F10 melanoma cells. When developing zebrafish embryos were treated with glyceollins, pigmentation of the embryos, melanin synthesis and tyrosinase activity were all decreased compared with control zebrafish embryos. In situ expression of a pigment cell-specific gene, Sox10, was dramatically decreased by glyceollin treatment in the neural tubes of the trunk region of the embryos. Stem cell factor (SCF)/c-kit signaling pathways as well as expression of microphthalmia-associated transcription factor (MITF) were determined by western blot analysis. Glyceollins inhibited melanin synthesis, as well as the expression and activity of tyrosinase induced by SCF, in a dose-dependent manner in B16F10 melanoma cells. Pretreatment of B16F10 cells with glyceollins dose-dependently inhibited SCF-induced c-kit and Akt phosphorylation. Glyceollins significantly impaired the expression and activity of MITF. An additional inhibitory function of glyceollins was to effectively downregulate intracellular cyclic AMP levels stimulated by SCF in B16F10 cells. Glyceollins have a depigmentation/whitening activity in vitro and in vivo, and that this effect may be due to the inhibition of SCF-induced c-kit and tyrosinase activity through the blockade of downstream signaling pathway.

INTRODUCTION

Melanin is a dark pigment that has an essential role in protection against ultraviolet radiation, and its production is restricted to ~5% of skin cells that have a common embryological origin and a unique cell type: melanocytes. Melanocytes differentiate from neural crest cells in the neural tubes during the early embryonic stage. Many cytokines that influence survival, proliferation, differentiation and function have been identified in melanocytes. Besides their gross phenotypic reaction, the response of melanocytes to cytokines is in general poorly understood. However, the best studied paracrine factor with actions on melanocytes is probably stem cell factor (SCF).

SCF and a kit receptor, c-kit (also known as SCF receptor, or SCFR), are encoded by the steel (Sl) locus and the dominant white spotting (W) locus, respectively. SCF/c-kit signaling is known to be involved in melanocyte development during embryogenesis by the analysis of Sl and W knockout mice. Soluble form of kit, s-kit, diminishes SCF-induced melanogenesis in human melanocytes. Several lines of evidence demonstrate the role of SCF/c-kit signaling in the regulation of epidermal melanogenesis under homeostatic, stimulatory or pathogenic conditions, including ultraviolet B exposure and pigmentation disorders. In the epidermis, keratinocytes produce increased amounts of SCF in response to several stimuli including aging. Binding of SCF to the extracellular domain of c-kit, a tyrosine kinase receptor, induces dimerization of the receptor. Receptor dimerization is followed by autophosphorylation, with the subsequent activation of a downstream signaling cascade. This cascade involves the stimulation of phosphatidylinositol 3’-kinase and Ras-mitogen-activated protein kinase (Ras-MAPK) through
the Shc and Grb2 adaptor proteins, as well as the guanine nucleotide exchange factor, SOS. The end point of this cascade is the activation of the tyrosinase enzyme. Tyrosinase is in turn required for melanocyte survival and melanin biosynthesis.6–10

The MAPKs, including extracellular responsive kinase (ERK) and p38 MAPK signaling cascade have been suggested as the signaling pathways modulating melanogenesis.11–13 Activation of p38 MAPK positively modulated melanin synthesis12,14 by activating cyclic AMP (cAMP) response element-binding protein (CREB), which in turn activates microphthalmia-associated transcription factor (MITF) expression), a critical melanocyte differentiation and survival transcription factor.15 The c-kit receptor also phosphorylates itself through the MAPK pathway.16,17 This suggests that protein kinase A signaling is also involved in SCF-induced melanogenesis. Protein kinase A is activated by the elevated cellular cAMP, which leads to the activation of MITF through the activation of CREB, resulting in the expression of tyrosinase, tyrosinase-related protein 1 (TRP-1) and TRP-2 genes.18,19 As noted above, it has been shown that SCF induces tyrosinase,20 which catalyzes the first two steps of the biosynthesis of eumelanin or pheomelanin, that is, the hydroxylation of tyrosine to dihydroxyphenylalanine (DOPA) and then the oxidation of DOPA to DOPA-quinone.21 For this reason, we studied the role of putative inhibitors of SCF/c-kit signaling in cultured melanoma cells, which provide an in vitro model for melanocytes, and in a zebrafish animal model, as detailed below.

This study used zebrafish as an experimental animal model in a phenotype-based screening for pigmentary inhibitors for the following reasons. The zebrafish system has several advantages, such as numerous quantities of embryos relative to other vertebrates, inducible spawning by light, convenience in observing melanin development, a rapid pigmentation process and high permeability to small molecules. In addition, it possesses epidermal melanocyte equivalents that have similar structural and functional characteristics to those of mammals.22 The characteristic external pigment pattern of the zebrafish is generated by an array of three types of pigment cells, all of which are derived from the neural crest. These include melanophores (melanin-containing melanocytes), xanthophores (containing yellow pigment) and iridophores (containing reflecting platelets).23 The combination of xanthophores and iridophores leads to the yellow–silver interstripes of the zebrafish, while the melanophores contribute to the longitudinal dark stripes of the epidermis.24,25

Glyceollins are a group of phytoalexins that are produced by the soybean plant (Glycine max) under the conditions of stress26,27 and abundant in fermented traditional Korean soybean product, such as Chungkookjang, Meju,28 which have been used for soothing skin burn or damage traditionally. Glyceollins (a mixture of glyceollin I, II and III) exert antimicrobial activity against several plant pathogens.29,30 Moreover, glyceollins function as antiestrogenic agents and may be used for the prevention or treatment of prostate, breast and ovarian carcinomas.31,32 The biological functions of glyceollins include anticontractile activity in vascular smooth muscle through the RhoA/Rho kinase signaling pathway,33 increased insulin sensitivity34 and antimelanin synthesis activity through the inhibition of cAMP in vitro.35 However, the biological activity of glyceollins and their underlying mechanisms of action in regard to SCF-induced pigment formation in vitro and in vivo are largely unknown.

In the present study, we isolated glyceollins from elicited soybeans and evaluated the inhibitory activity of glyceollins against SCF-induced tyrosinase activity, MITF expression and cAMP production in B16F10 melanoma cells. Glyceollins effectively suppressed SCF-induced signaling pathways in B16F10 cells. We further examined whether glyceollins could inhibit the skin pigmentation in the zebrafish system through the inhibition of tyrosinase activity, Sox10 expression, a neural crest marker and a key transcription factor that induces MITF gene expression during the differentiation of melanocytes from precursor cells,36 was clearly diminished in zebrafish trunk neural tubes by glyceollin treatment. Thus, the results suggest that glyceollins have a strong depigmentation effect in vitro and in vivo and function by inhibiting the SCF-mediated pathway. As such, they could be potential therapeutic agents for the treatment of post-inflammatory hyperpigmentation or skin-whitening agents for cosmetic use.

MATERIALS AND METHODS

Cell culture and reagents

B16F10 melanoma cells were cultured on tissue culture plates in Dulbecco modified Eagle’s medium (Hyclone, Logan, UT, USA) supplemented with 10% fetal bovine serum (Hyclone, Logan, UT, USA) and 1X antibiotics (100 U ml⁻¹ penicillin, 100 µg ml⁻¹ streptomycin, all from Invitrogen, Carlsbad, CA, USA) at 37°C in a humidified atmosphere incubator containing 5% CO₂. Stock solutions of SCF (Peprotech, Rocky Hills, NJ, USA) and glyceollins were prepared at 1, 5 and 10 µg ml⁻1 in Dulbecco modified Eagle’s medium. Antibodies against c-kit, phosphorylated c-kit, ERK and phosphorylated-ERK were purchased from Cell Signaling Technology (Beverly, MA, USA), and antibodies against p38, phosphorylated-p38 and tyrosinase were purchased from Santa Cruz (Santa Cruz, CA, USA).

Isolation of glyceollins

Glyceollins were isolated using a procedure developed by Boué et al. Briefly, soybean seeds were scarified and inoculated with Aspergillus sojae. After 3 days, the glyceollins were extracted from the seeds with 80% ethanol and then isolated using high-performance liquid chromatography (PerkinElmer series 200, PerkinElmer, Waltham, MA, USA). The chemical structures for glyceollin I, II and III are shown in Figure 1.

Zebrafish growth conditions and chemical treatment

Standard AB strain zebrafish (Danio rerio) and hemizygous transgenic Tg(fkl-EGFP) embryos were maintained at 28°C in an oxygenated fish tank on a 14/10 h (light/dark) cycle as previously described. Embryos were collected from natural matings, dechorionated with pronase at 24, 36 and 48 h post fertilization (h.p.f.) developmental stages, and maintained. Bud stage zebrafish embryos were treated with glyceollins at 1, 5 and 10 µg ml⁻1 that were added to the tank water for 40 h. The developmental phenotypes of the embryos were observed with a Zeiss Imager Z1, Zeiss Axioskop (Zeiss,
After quantification, total protein (250 m to 48 h.p.f. and sonicated in Pro-Prep protein extraction solution. Total proteins were transferred into 96-well plates, and 2 mg ml\(^{-1}\) of lysates were clarified by centrifugation. After quantification, 30 m containing 10% dimethylsulfoxide at 80 for 1 h. The absorbance was measured at 450 nm using a spectrophotometer.

Tyrosinase activity
Glyceollin-treated B16F10 melanoma cells were sonicated in a standard protein extraction solution (20 mm Tris (pH 7.5), 150 mm NaCl, 1 mm EDTA, 1% Triton X-100 and 1X protease inhibitor cocktail) supplemented with 1 mm phenylmethylsulfonyl fluoride, and the lysates were centrifuged. After centrifugation, 30 m total proteins were transferred into 96-well plates, and 2 mg ml\(^{-1}\) L-DOPA in phosphate-buffered saline. Tyrosinase activity was expressed as a percentage of the control.

Measurement of cAMP concentration
The cAMP concentration was measured using a cAMP immunoassay kit (Assay Designs, Ann Arbor, CA, USA). Briefly, B16F10 melanoma cells (3 \(\times\) 10\(^5\)) were lysed in 0.1 M HCl to inhibit the phosphodiesterase activity. The supernatants were then collected, neutralized and diluted. After neutralization and dilution, a fixed amount of cAMP conjugate was added to compete with cAMP in the cell lysate for sites on a rabbit polyclonal cAMP antibody immobilized on a 96-well plate. Cells were washed, and a substrate solution was added to determine the activity of the bound enzyme. The absorbance was read at 405 nm. The intensity of the color was inversely proportional to the concentration of cAMP in the cell lysate.

Western blot analysis
Glyceollin-treated cells were incubated in the presence of SCF (50 ng ml\(^{-1}\), Peprotech, Rocky Hill, NJ, USA) for 4 min and lysed. Total cell lysates were analyzed by sodium dodecyl sulfate polyacrylamide gel electrophoresis followed by western blotting with the antibodies against the phosphorylated or unphosphorylated form of p42/44-ERK, AKT and p38 MAPK. Westerns were incubated with horseradish peroxidase-conjugated rabbit or mouse immunoglo- bulins and then developed by the West Pico Chemiluminescent Substrate (Pierce, Woburn, MA, USA) using standard protocols.

Whole-mount in situ hybridization in zebrafish
Whole-mount in situ hybridization was performed as previously described. Briefly, dechorionated embryos at specific stages were fixed with 4% paraformaldehyde in phosphate-buffered saline. After hybridization, the embryos were incubated with antioxygenin-Dig-AP F (ab\(^{+}\)) (Boehringer, Ingelheim, Germany) and stained with a staining buffer (pH 9.5, 0.1 m Tris–HCl, 0.1 m NaCl, 0.05 m MgCl\(_2\), 0.1% Tween-20) containing BCIP and NBT. Stained embryos were mounted in glycerol and observed under a microscope with a Nomarski differential interference contrast lens (Axioskop, Zeiss).

Statistical analysis
One-way analysis of variance was performed to assess the significance of differences among the experimental groups. The level of significance was set at \(P < 0.05\) versus control, \(P < 0.05\) versus SCF. Results are represented as the means \(\pm\) s.e.m.

RESULTS
Glyceollins inhibit pigment production in zebrafish
To investigate the effect of glyceollins on specific animal pigmentation in the development of melanocyte, we utilized zebrafish embryos. In a preliminary experiment, we tested zebrafish embryos for their feasibility as a model system for melanin synthesis. As a positive control, 1-phenyl-2-thiourea (PTU, 0.3 mg ml\(^{-1}\)), a typical tyrosinase inhibitor, was used. About 200 embryos at the bud stage were treated with each concentration of glyceollins (1, 5 and 10 \(\mu\)m) from bud stage to 48 h.p.f. and sonicated in Pro-Prep protein extraction solution. After centrifugation, total protein (250 \(\mu\)g) was added to 2 mg ml\(^{-1}\) L-DOPA in a phosphate solution and incubated. Absorbance was then measured at 450 nm using a spectrophotometer.

To determine whether the effect of depigmentation was reversible, the embryos were washed intensively at 48 h.p.f. after treatment with PTU and glyceollins (10 \(\mu\)m). Embryos were further grown so as to observe body pigmentation at 72 h.p.f. As compared with the no-treatment control group, glyceollin and PTU treatment resulted in a marked inhibition of body pigmentation (Figure 2C), although the effect of PTU was more pronounced, because the concentration of PTU was much higher than that of glyceollins. Removal of PTU and glyceollins allowed the transparent zebrafish to recover some, but not all, of their pigmentation. Therefore, the depigmentation effects of glyceollins were maintained at least for some period of time (24–36 h).
Glyceollins affect tyrosinase activity and melanin synthesis in zebrafish

To investigate the molecular targets of glyceollins that inhibit pigmentation in the zebrafish, we determined total melanin content and tyrosinase activity. There were considerable decreases in total melanin content, as well as tyrosinase activity and expression, after treatment with glyceollins (Figures 3A–C). As a positive control, we employed PTU and showed that it remarkably decreased melanin content, tyrosinase activity and expression. The inhibitory effects of glyceollins at 1, 5 and 10 μM on total melanin content and tyrosinase activity were comparable with those of PTU (Figures 3A and B). Their ability at 1 and 5 μM to decrease tyrosine expression was less than that of PTU, but glyceollins at a concentration of 10 μM showed comparable activity to PTU. These results suggest that glyceollins impair melanogenesis through the inhibition of tyrosinase activity in zebrafish.

Glyceollins inhibit Sox10 expression in zebrafish embryos

In both zebrafish and mice, one transcription factor, Sox10, controls the expression of another, MITF, which in turn regulates a set of genes critical for pigment cell development and pigmentation. Mutations in either Sox10 or MITF impair pigment cell development. To investigate whether glyceollins affect the differentiation of neural crest cells into melanocytes, we performed whole-mount in situ hybridization with a Sox10 antisense probe in zebrafish embryos. At 24 h.p.f., Sox10 was expressed in melanocyte-sprouting cells in the trunk region (Figure 3D, a–a'). However, both PTU (Figure 3D, b–b') and the glyceollins (Figure 3D, c–c') attenuated and dispersed the expression of Sox10 at the neural tubes in the trunk region (Figure 3, see the arrows).

Glyceollins inhibit SCF-induced melanogenesis

To investigate the underlying molecular mechanism to inhibit melanogenesis, we used SCF for induction of melanogenesis and treated B16F10 melanoma cells with various concentrations of SCF (5, 10, 50 and 100 ng ml⁻¹) for 48 h. SCF increased melanin contents in a dose-dependent manner. The concentration of 50 ng ml⁻¹ SCF induced fairly high melanin content (Figure 4a) and, thus, we used this concentration for further experiments. Next, we investigated whether
glyceollins can inhibit SCF-induced melanin synthesis. Both extracellular, secreted melanin contents and intracellular melanin contents were significantly reduced by the treatment with glyceollins (Figures 4b and c). Furthermore, the colors of the melanoma cell pellet treated with glyceollins become lighter in a dose-dependent manner compared with SCF-treated positive control cells (Figure 4d). These results indicate that glyceollins inhibited SCF-induced melanin biosynthesis in B16F10 melanoma cells.

Glyceollins inhibit SCF-induced signaling pathways in B16F10 cells

Treatment of B16F10 melanoma cells with SCF resulted in the phosphorylation of c-kit. Phosphorylation of c-kit was not observed in the absence of SCF, and pretreatment with glyceollins (1, 5 and 10 μM) significantly inhibited SCF-induced c-kit phosphorylation in a dose-dependent manner (Figure 5a). It is well known that the events downstream of c-kit phosphorylation are related to p44/42 ERK MAPK, and the c-kit/p44/42 ERK MAPK pathway is a common signaling pathway in melanocytes,7,16 endothelial cells42 and hematopoietic cells.43 SCF treatment in melanoma cells induced p44/42 ERK phosphorylation, which was inhibited by glyceollins in a dose-dependent manner (Figure 5a). Glyceollins also suppressed SCF-induced AKT and p38-MAPK phosphorylation (Figure 5a). These results suggest that glyceollins successfully suppressed SCF-induced c-kit activation and downstream kinase signaling pathways involved in the function of melanocytes and melanin synthesis. In addition, SCF increased MITF, a melanocyte differentiation factor,16 and tyrosinase expression in B16F10 cells. On the other hand, the glyceollins downregulated the SCF-induced expression levels of MITF and tyrosinase (Figures 5b and c). It has been reported that SCF stimulates tyrosinase activation and melanogenesis, which are mediated by the activation of adenylyl cyclase in B16F10 melanoma cells.44 Glyceollins significantly suppressed
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SCF-induced tyrosine’s activity (Figure 5d). To investigate whether glyceollins could modulate adenylate cyclase activity, we next measured the intracellular cAMP level stimulated by SCF. Glyceollins did indeed significantly inhibit cAMP production induced by SCF (Figure 5e). Taken together, these data indicate that the degeneration of SCF receptor expression in melanocytes may result from the inhibition of tyrosinase and adenylate cyclase activity involved in the signaling pathway induced by SCF/c-kit activation. 

Taken together, these results clearly indicate that glyceollins can inhibit melanin biosynthesis induced by SCF. This function is likely to occur via the suppression of c-kit-mediated signaling pathways, including adenylate cyclase and Sox-10 pathways.

DISCUSSION
Glyceollins are novel phytoalexins derived from soybeans that exist as three isomers. This group of compounds is generated as a plant defense mechanism by an interaction between the soybean plant and many microorganisms. Accordingly, glyceollins exhibit a wide range of toxicities against nematodes, fungi, insects and bacteria for soybean self-protection.30 It has also recently been recognized that glyceollins can suppress human breast and ovarian carcinoma tumorigenesis and may modulate potential estrogenic properties in the breast through antiestrogenic effects.32 Glyceollins were additionally identified in our earlier study as effective inhibitors of tyrosinase, which was induced in vitro by α-melanocyte-stimulating hormone.35 SCF is highly associated with melanogenesis by ultraviolet-irradiated inflammation in keratinocytes4,45 and photoaged pigmentation in fibroblasts.46 Therefore, we tried to investigate whether glyceollins inhibit SCF-induced melanogenesis. Here, we provide concrete evidence that glyceollins have a potent inhibitory effect against SCF-stimulated melanogenesis, and we demonstrated their molecular mechanisms of action in B16F10 melanoma cells, as well as in an in vivo zebrafish animal model. We did not show that SCF/c-kit signaling pathway is involved in glyceollins-induced inhibition of zebrafish pigmentation, but administration of anti-c-kit antibody dose-dependently suppressed hair pigmentation in association with decreased melanocyte proliferation and differentiation in mice,47 suggesting SCF/c-kit signaling pathway is necessary for the hair or skin pigmentation in mice. Our current study revealed that glyceollins have an inhibitory effect on melanin content in vitro and in vivo. We confirmed that the intracellular and secreted melanin contents and tyrosinase activity were significantly reduced by glyceollins in melanoma cells. This effect of glyceollins was associated with a significant downregulation of the expression of melanocyte-specific proteins, that is, tyrosinase as well as MITF, compared with the positive control, SCF-treated B16F10 melanoma cells. Because SCF/c-kit is known to enhance the expression of tyrosinase, we hypothesized that the downregulation of SCF/c-kit signaling pathways induced by glyceollins might suppress the expression of tyrosinase. It has been reported that SCF causes tyrosinase activation and melanogenesis through cAMP generation by adenylate cyclase in melanoma cells.48 Therefore, inhibition of SCF-stimulated cAMP production in melanoma cells is a key molecular event in the inhibition of melanogenesis.

cAMP generated by the activation of adenyl cyclase induces CREB to activate MITF gene expression in collaboration with Sox10.49 Therefore, the attenuated expression of MITF caused by glyceollins in this study can likely be

Figure 4 Effect of glyceollins on melanin synthesis and tyrosinase activity in melanoma cells. (a) B16F10 cells were treated with various concentrations of stem cell factor (SCF). Melanin contents were then determined using a spectrophotometer. (b and c) The melanoma cells were incubated with glyceollins (1, 5 and 10 μM) in the presence of SCF (50 ng/ml) for 48h, and the extracellular and intracellular melanin contents were detected and quantified using a spectrophotometer. The results are shown as a percentage of control. All experiments were repeated three times. (d) The image shows the pellet color of glyceollin-treated melanoma cells cultured in vitro. *P<0.05 versus control (negative control); #P<0.05 versus SCF (positive control). PTU, 1-phenyl-2-thiourea.
attributed to the reduced levels of cAMP and Sox10. MITF expression induced by α-melanocyte-stimulating hormone is also inhibited by glyceollins.50 As such, glyceollins might also inhibit α-melanocyte-stimulating hormone-induced adenylate cyclase activity. In addition, MAPKs such as ERK1/2 and p38 MAPK are also involved in tyrosinase expression in B16F10 cells.51 Hence, the inhibition of these MAPKs is an important mechanism by which glyceollins inhibit tyrosinase expression and activity induced by SCF. Because involvement of MAPKs in melanocyte proliferation or differentiation is evidenced only by cell culture system in previous reports and in our study, further investigation whether activity of these MAPKs is inhibited in glyceollins-treated zebrafish embryos should be performed.

Figure 5 Inhibition of stem cell factor (SCF)-induced c-kit phosphorylation and downstream signaling pathways by glyceollins in B16F10 melanoma cells. (a) Cells were pretreated with glyceollins (1, 5 and 10μM) for 48 h, and c-kit phosphorylation was stimulated with recombinant human SCF (rhSCF) (50 ng ml−1) for 4 min. Western blot analysis was performed. (b, c) Melanoma cells were treated with or without glyceollins for 48 h. The microphthalmia-associated transcription factor (MITF) (b) and tyrosinase (c) and β-actin protein level was determined by western blot analysis. (d) Tyrosinase activity was determined from the melanoma cells incubated with 2 mg ml−1 L-DOPA and then quantified using a spectrometer. (e) The concentration of cAMP was measured using a cAMP assay kit. *P<0.05 versus control; #P<0.05 versus SCF (positive control). cAMP, cyclic AMP; DOPA, dihydroxyphenylalanine; ERK, extracellular responsive kinase.
MITF itself is not sufficient to promote increased expression of its target genes. Phosphorylation of MITF serine residue 409 by p90 ribosomal S6 kinase (p90RSK) and serine residue 73 by MAPK1 activated by SCF/c-kit signaling might contribute synergistically to its activation by promoting an interaction with the p300 coactivator.\textsuperscript{16,17} If so, coactivators of the MAPK cascades will thus confer on MITF the ability to contribute significantly to the expression of genes critical for the phenotype and function of the mature melanocyte, such as tyrosinase and TRP1.\textsuperscript{52}

Zebrafish pigmentation and the development of melanocytes derived from neural crest cells were also suppressed by glyceollins, probably through the same mechanism as in B16F10 melanoma cells. It stands to reason that SCF might therefore be the primary stimulator of melanogenesis during embryonic development. For example, Sox10 is expressed in migrating neural crest cells in the trunk neural tubes, which differentiate into sensory neurons in the trunk.\textsuperscript{53} The diminished expression of Sox10 in glyceollin-treated embryos might lead to an ectopic migration path and defective differentiation of neural crest cells. More in detail study using primary neural crest cell culture from embryos homozygous for a Kit null allele, it is shown that the onset of MITF in melanoblast does not require Kit. And treatment of choleratoxin, CAMP activator mimics part of the Kit signaling pathway increases MITF and tyrosine expression in Kit null cells.\textsuperscript{54} This study suggests that MITF is not sufficient for tyrosinase expression in melanoblast and that Kit signaling regulates gene expression during the melanocyte development \textit{in vivo}. From our results, normal Sox10-induced melanocyte differentiation from neural crest cells likely occurs in collaboration with SCF through increased MITF transcription in the zebrafish. Here we suggest that SCF/c-kit activates three different downstream pathways, adenyl cyclase, p38 MAPK and MEK/ERK MAPK pathways. Activation of adenyl cyclase induces CAMP and protein kinase A to activate CREB, and p38 and ERK MAPK pathways activate MITF, MITF and CREB collaborate to activate tyrosinase and TRP1/2.

CONFLICT OF INTEREST

The authors declare no conflict of interest.

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