Wnt3a inhibits proliferation but promotes melanogenesis of melan-a cells

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Abstract. Melanocytes are pigment-producing cells responsible for coloration of skin and hair. Although the importance of Wnt3a in melanocyte development has been well recognized, the role of Wnt3a in mature melanocytes has not been elucidated. This study was conducted to further explore the effects of Wnt3a on melanocyte proliferation and melanogenesis, and to elucidate the possible mechanisms involved. We infected melan-a cells with AdWnt3a to serve as the production source of the Wnt3a protein. MTT assay, 5-bromodeoxyuridine incorporation assay and flow cytometric analysis showed that Wnt3a inhibited the proliferation of melan-a cells and this was associated with decrease of cells in the S phase and increase of cells in the G1 phase. Melanin content and tyrosinase activity assay revealed that Wnt3a significantly promoted melanogenesis of melan-a cells. Furthermore, western blot analysis showed that Wnt3a upregulated the expression of microphthalmia-associated transcription factor and its downstream target genes, Wnt3a significantly promoted melanogenesis of melan-a cells. Furthermore, western blot analysis showed that Wnt3a upregulated the expression of microphthalmia-associated transcription factor and its downstream target genes, Wnt3a significantly promoted melanogenesis of melan-a cells. Collectively, our results suggest that Wnt3a plays an important role in melanocyte homeostasis.

Introduction

Melanocytes play pivotal roles in skin and hair pigmentation by producing melanin (1,2). They originate from neural crest-derived melanoblasts and migrate into the epidermis and hair follicles during embryogenesis (3,4). Melanocytes synthesize melanin in melanosomes and transfer the melanin granules to the adjacent keratinocytes, where melanins are accumulated to generate pigmented skin or hairs. High melanin content protects the skin from harmful ultraviolet rays owing to the ability to absorb UV radiation and quench the UV-induced intracellular free radicals (1,5,6). Defects in or a lack of melanocytes can lead to pigment disorders, such as piebaldism, albinism, vitiligo, and hair graying (7-9).

Wnts are a large family of secreted glycoproteins that act as ligands to activate receptor-mediated signaling pathways that play important roles in cell fate, proliferation, differentiation and migration (10-13). Wnt signaling can be divided into at least two distinct pathways: canonical Wnt/β-catenin signaling and noncanonical signaling. In the best characterized canonical pathway, Wnt ligands bind to seven-pass transmembrane receptors of the Frizzled (Fzd) family and co-receptors, low density lipoprotein-related protein (LRP) 5 and 6, leading to the inhibition of the APC/Axin/CK1/GSK3b destruction complex and stabilization and translocation of β-catenin to the nucleus where it interacts with TCF/Lef family transcription factors to regulate the transcription of target genes (11,12). Contrary to canonical Wnt signaling, noncanonical Wnt signaling is transduced independently of β-catenin. Noncanonical Wnt signaling pathways are diverse and less well characterized, and have been termed the Wnt/calcium, Wnt/JNK pathway and Wnt/planar cell polarity pathway (PCP) (11).

Previous studies revealed that Wnt signaling plays a critical role in melanocyte development, specifically Wnt1 and Wnt3a. Wnt1 and Wnt3a promote the development of neural crest cells into pigment cells (14,15). Neural crest cells depleted of these two proteins become neuronal rather than melanocytes (14). Wnt1 acts on melanoblasts to increase melanocyte numbers, while Wnt3a and β-catenin can specify neural crest cells to become melanocytes (16,17). Mutant mice deficient in Wnt1 and Wnt3a are almost completely devoid of pigment cells (18). Furthermore, Wnt3a acts on melanoblasts to maintain microphthalmia-associated transcription factor (MITF) expression and promote melanoblasts to differentiate to become melanocytes (16). In humans, high levels of DKK1 (an inhibitor of the canonical Wnt signaling pathway) inhibits melanocyte growth, pigmentation and induces a less pigmented skin on the palms (19,20).

Although the importance of Wnt3a in melanocyte development has been well recognized, the role of Wnt3a in mature melanocytes remains undetermined. To address this issue, adenoviral gene delivery of Wnt3a was adopted to investigate the effects of Wnt3a on melan-a melanocyte proliferation and melanogenesis, and to elucidate the possible mechanisms involved.
Materials and methods

Cell culture. An immortal line of melanocytes, melan-a, were a kind gift of Dr D.C. Bennett (21). The cells were cultured in RPMI-1640 medium supplemented with 10% FBS (Gibco, USA), 2 mM L-glutamine, 200 nM 12-0-tetradecanoyl phorbol-13-acetate (Sigma, USA), 100 IU/ml penicillin, 50 µg/ml streptomycin, and grown in a humidified atmosphere containing 10% CO₂ in air at 37°C.

Adenovirus (Ad) amplification and infection. The adenoviruses expressing green fluorescent protein (AdGFP), Wnt3a protein (AdWnt3a, also expressing GFP), and SimMITF (AdSimMITF, a small interfering RNA of MITF mediated by adenovirus, also expressing RFP) were kindly provided by Dr T.C. He (Chicago University). The adenoviruses were propagated and purified as previously described (22). Briefly, the adenoviruses were propagated in HEK293 cells, which were collected upon detection of viral cytopathic effect-successful infections were verified by observable GFP incorporation. Cell pellets were resuspended in PBS and lysed by four freeze-thaw-vortex cycles. After being purified by cesium chloride (Amresco, USA) density gradient centrifugation, adenoviruses were dialyzed into storage buffer, then their titers were determined and diluted with storage buffer to the ultimate titer of 10⁸ plaque-forming unit (PFU)/ml. For infection, melan-a cells were plated onto 6- or 24-well plates at a density of 2x10⁴ cells/cm² in the growth medium for 12 h, the cells were then grown in medium supplemented with adenoviruses for 72 h.

Isolation of total-RNA and RT-PCR. Total cellular RNA was isolated at the indicated time-points using TRIzol reagent (Invitrogen, USA). Single-stranded cDNA was synthesized by using ReverTra Ace reverse transcriptase (Toyobo, Japan) and oligo(dt) primers according to the manufacturer's protocol. Semi-quantitative PCR was performed using primers for Fzd1-10, LRP5, LRP6, MITF, tyrosinase-related protein (TRP)1, TRP2, tyrosinase (Table I lists the primer sequences and amplicon size). PCR reaction was performed by using a touchdown protocol previously described (23). Briefly, touchdown PCR was performed with the following program: 1 cycle at 94°C for 2 min, 12 cycles at 92°C for 20 sec, 68°C for 30 sec, and 70°C for 45 sec with a decrease of one degree per cycle, and 25 cycles at 92°C for 20 sec, 55°C for 30 sec, and 70°C for 45 sec. PCR products were analyzed by gel electrophoresis and stained with ethidium bromide.

TOP/FOP-flash luciferase reporter assay. Melan-a cells were seeded onto 24-well plates overnight and subsequently infected with AdWnt3a or AdGFP. After 12 h, the cells were transfected with TOP/FOP-flash luciferase reporter plasmid. For each well, 3 µg TOP- or FOP-Flash Firefly Luciferase reporter plasmid, which contain several TCF4-binding elements (TOP) or mutant sequences (FOP), respectively, were transfected together with 0.03 µg phRG-TK Renilla Luciferase standard plasmid (Promega, USA). The transfection reagent, Lipofectamine 2000 (Invitrogen), was used according to the standard protocol. Cell lysates were harvested 24 h after transfection and the Dual-Luciferase Reporter Assay System (E1910; Promega) was used to detect luciferase activity according to the manufacturer's protocol. The experiments were performed in triplicate.

Cytometry and MTT assays. To study the effect of Wnt3a on melan-a proliferation, 2x10⁴ melan-a cells were plated onto 24-well plates overnight and grown in culture medium supplemented with AdWnt3a at various doses (1, 2, 3 µl) or 2 µl AdGFP as the vehicle control. After 72 h, the cells were detached with trypsin and counted in a hemocytometer. For the MTT assay, melan-a cells were plated onto 96-well plates and treated with AdWnt3a at various doses (1, 2, 3 µl) or 2 µl AdGFP or 0.5 µl AdGFP for 72 h. Then MTT (Sigma) was added, and the cells were incubated at 37°C for 4 h. The medium was removed and dimethyl sulfoxide (DMSO) was added to dissolve the formazan crystals. The absorbance was then measured at 490 nm with a common ELISA reader.

5-Bromodeoxyuridine incorporation assay. Melan-a cells were plated on glass coverslips and were treated with AdWnt3a or AdGFP. After 72 h, 5-bromodeoxyuridine (BrdU, Sigma; stock of 10 mM in PBS) was added at 10 µM. The cells were then incubated at 37°C for 2 h and fixed in cold acetone for 10 min. After washing three times in PBS, the cells were incubated in 2 M HCl for 45 min to denature the DNA and then neutralized

| Table I. Primer sequences for RT-PCR analysis. |
|-----------|-----------|-----------|
| Primer | Sequence (5’-3’) | GenBank ID |
| LRP5 | F: GGTCACCTGGACTTGCTCAT<br>R: TCCAGCGTTGATGTGAAGC | NM_008513.3 |
| LRP6 | F: ACAGAGCCTGACATCC<br>R: TGGATTTGGCACTTGTTG | NM_008514.4 |
| Fzd1 | F: CAAAGTTACCCGCTGCTG<br>R: GTACACGCGGAGGAGAAA | NM_002145.7 |
| Fzd2 | F: TTAGGCGCCTGAGATG<br>R: CAGGGAGGCGTGAAGC | NM_002051.2 |
| Fzd3 | F: GCTCCAGAAMCTGGCTTTG<br>R: GACACTCCCTGGTCTT | NM_002145.8 |
| Fzd4 | F: AACCTGGCTTACACGAC<br>R: TGGCACATAAACCAGAAA | NM_00855.4 |
| Fzd5 | F: AGGCATCCTGGATTTTCT<br>R: TGAGCGGGCAGAGATT | NM_022721.3 |
| Fzd6 | F: TTCAGGCGCTTGAAGAAC<br>R: CAACCCCCGCTTCTCA | NM_001162494.1 |
| Fzd7 | F: ATCATCTTCTGGCTGGTT<br>R: AAGCAACGGAAAGGAT | NM_00850.7 |
| Fzd8 | F: CTGGTCGCCATCCTGAG<br>R: CCGTGGTGCTGCTCATAG | NM_00858.2 |
| Fzd9 | F: TTATGGTTGCTCTCCTCT<br>R: CACCTCCCTCATGAGACA | NM_010246.1 |
| Fzd10 | F: TCTCTACACCTCATTGGT<br>R: GCTGCCCACTATACAC | NM_175284.3 |

F, forward; R, reverse.
with 0.1 M Na$_2$B$_4$O$_7$ (pH 8.5) for 30 min. The detection of BrdU was performed with a mouse anti-BrdU antibody (1:100; Zhongshan, China) at 4˚C overnight. Then, cells were incubated with goat anti-mouse FITC-conjugated secondary antibody (1:100; Zhongshan) at 37˚C for 1 h. After washing, the cells were stained with DAPI for 10 min at room temperature. Six areas/well were randomly selected and counted with an upright microscope BH2 (Olympus, Japan).

**Flow cytometric analysis.** Melan-a cells were plated onto 6-well plates and treated with 2 µl AdWnt3a or 2 µl AdGFP for 72 h. Afterward, the cells were dissociated with trypsin, washed in PBS, and fixed in 1 ml of 70% methanol at 4˚C for at least 24 h. The fixed cells were washed twice in PBS, treated with 100 µg/ml RNase and 50 µg/ml propidium iodide (PI) in PBS for 30 min at 37˚C, and the cell cycle was detected with a flow cytometer. All proliferation assays were performed in triplicate.

**Tyrosinase enzymatic assay.** Tyrosinase activity assays were performed according to the method previously reported (24). Melan-a cells in 6-well plates were infected with AdWnt3a or AdGFP for 72 h, then trypsinized and counted. Cells (1x10$^5$) were treated with 200 µl of 1% Triton X-100/PBS at -70˚C for 30 min and thawed at 37˚C. Then, the extracts were clarified by centrifugation, 50 µl of the supernatant were transferred into 96-well plates and 10 µl of 2 mg/ml L-DOPA (Sigma) were added. After incubation for 2 h at 37˚C, absorbance was measured at 490 nm. The experiments were performed at least three times.

**Western blot analysis.** Cells were lysed in RIPA lysis buffer (Beyontime, China), determined by the Enhanced BCA Protein Assay kit (Beyontime) and denatured by boiling. Protein of 80 µg per lane was loaded on 10% SDS-PAGE and then transferred onto a PVDF membrane. Membranes were blocked with 5% fat-free milk in Tris-buffered saline-Tween-20 (TBST; 20 mM Tris/HCl, pH 7.6, 137 mM NaCl, 0.1% Tween-20) for 2 h, then membranes were probed with rabbit anti-Wnt3a antibody (1:1,000; Abcam, USA), goat anti-TRP1 antibody (1:1,000), rabbit anti-TRP2 antibody (1:1,000), goat anti-tyrosinase antibody (1:1,000; Santa Cruz Biotechnology, Inc., USA), and mouse anti-MITF antibody (1:500, Millipore, USA) at 4˚C overnight. Blots were then incubated with HRP-conjugated secondary antibody. Peroxidase activity on the membrane was visualized on X-ray film by means of the ECL western blotting detection system.

**Statistical analysis.** Data were presented as means ± SD for the three independent experiments. Statistical differences were evaluated by the t-test, and P<0.05 was considered to be statistically significant.

**Results**

Wnt3a activates Wnt/β-catenin signaling in melan-a cells. To explore the potential role of Wnt3a in mouse mature melanocytes, we used a mouse melanocyte cell line, melan-a (21), as an in vitro cell model. DOPA staining was used to confirm that the cells were indeed melanocytes (Fig. 1A). RT-PCR
analysis showed that the Wnt receptors Fzd1 to Fzd10 and the co-receptors LRP5 and LRP6 were all expressed in melan-a cells, indicating that melan-a cells could receive and transduce Wnt signals (Fig. 1E). We infected melan-a cells with AdWnt3a at a predetermined optimal titer. As observed in Fig. 1B, AdWnt3a mediated efficient gene transfer in melan-a cells. The expression of Wnt3a was confirmed by western blot analysis in AdWnt3a-infected cells but not in AdGFP-infected cells (Fig. 1C).

To evaluate the ability of AdWnt3a to activate Wnt/β-catenin signaling, western blot analysis and the TOP/FOP-flash luciferase reporter assay were performed. As shown in Fig. 1C, AdWnt3a upregulated the expression of β-catenin at the protein level. We infected melan-a cells with AdWnt3a or AdGFP and co-transfected with either the β-catenin responsive TCF4 reporter plasmid (TOP-flash) or with negative control reporter plasmids (FOP-flash). The luciferase activities showed that AdWnt3a-infected cells displayed higher ratios of TOP-flash/FOP-flash compared to AdGFP-infected cells (Fig. 1D). The data demonstrate that AdWnt3a efficiently activates Wnt/β-catenin signaling in melan-a cells.

Wnt3a inhibits the proliferation of melan-a cells. To analyze the effect of Wnt3a on melan-a cell proliferation, we infected melan-a cells with different doses of AdWnt3a or AdGFP as control. After 72 h, the MTT cell proliferation assay and manual cell count both showed that Wnt3a inhibited the proliferation of melan-a cells in a dose-dependent manner compared to GFP (Fig. 2C and D).

BrdU is a synthetic thymidine analog that gets incorporated into a DNA cell when the cell is dividing, so it is commonly used for the detection of proliferating cells. We measured the BrdU incorporation of AdWnt3a-infected melan-a cells and AdGFP-infected cells. The incorporated BrdU was detected by using the anti-BrdU antibody (Fig. 2A). The percentage of BrdU positive cells in AdWnt3a-infected cells was decreased by 14.9% (from 32.7±4.45 to 17.8±1.32%, P<0.05) (Fig. 2B), which implied that Wnt3a inhibited the proliferation of melan-a cells.

We also performed cell cycle analysis and found that AdWnt3a-infected cells exhibited an increased population in the G1 phase and a decreased population in the S phase compared to AdGFP-infected cells (Fig. 3).
Taken together, these results showed that Wnt3a inhibited melan-a cell proliferation and this was associated with decreased population of cells in the S phase and an increase in the G1 phase.

Wnt3a promotes the melanogenesis of melan-a cells. We analyzed the effect of Wnt3a on the melanogenesis of melan-a cells by assessing melanin production and tyrosinase activity. Although almost all the cells contained melanin pigment, AdWnt3a-infected cells showed obvious higher level of melanin accumulation compared to AdGFP-infected cells (Fig. 4A and B).

To analyze whether melanin synthesis is activated via a direct effect on tyrosinase, tyrosinase activity assay was adopted to analyze tyrosinase activity in melan-a cells. As shown in Fig. 4C, a greatly significant increase of tyrosinase activity was detected in AdWnt3a-infected cells compared with AdGFP-infected cells.

To investigate how Wnt3a promotes melanin synthesis, we studied the expression of microphthalmia-associated transcription factor (MITF), the transcriptional master regulator of melanocytes, and its downstream target genes, including tyrosinase, TRP1, and TRP2 (25-27). Western blot analyses showed that Wnt3a significantly increased the expression levels of MITF, tyrosinase, and TRP1 in melan-a cells, but did not affect the expression of TRP2 (Fig. 5). Subsequently, we infected melan-a cells with AdSimMITF to knockdown the endogenous MITF. According to western blot analysis, the expression of MITF, tyrosinase, and TRP1 was dramatically decreased in the cells infected with AdSimMITF (Fig. 6). Co-infection of melan-a cells with AdSimMITF and AdWnt3a demonstrated that Wnt3a rescued the expression of MITF, tyrosinase and TRP1 (Fig. 6).

These results suggested that Wnt3a contributed to increase melanin synthesis through upregulation of the expression of MITF and its downstream genes, tyrosinase and TRP1.
The Wnt signaling pathway is critical for regulating multiple aspects of basic cell functions, including proliferation, polarity, differentiation, and migration of cells (10-12). Despite the well-known role of Wnt3a in the development of melanocytes, the role of Wnt3a in mature melanocytes remains unknown. Therefore, this study was conducted to address this question.

The melan-a melanocytes were originally derived from normal epidermal melanoblasts from embryos of inbred C57BL/6J mice. When the cell line was established, the primary culture was unpigmented melanoblasts, which then matured to pigmented melanocytes in the incubation (21). We thus used melan-a as an in vitro cell model to investigate the effect of Wnt3a in mature melanocytes.

Wnt/β-catenin signaling is mediated by binding to Frizzled (Fzd) receptors and low density lipoprotein-related protein (LRP)5/6, and induces various cellular events. Fzd1, Fzd3, Fzd8, and LRP6 have been demonstrated to act as receptors for Wnt3a (28-32). Our RT-PCR result showed that melan-a cells expressed Wnt receptors, Fzd (Fzd1-10), and the co-receptors LRP5 and LRP6, suggesting that melan-a cells could respond to Wnt3a. In this study, we infected melan-a cells with AdWnt3a and demonstrated that AdWnt3a expressed Wnt3a protein and efficiently activated Wnt/β-catenin signaling in melan-a cells. Rabbani et al. (33) co-cultured Wnt10b-transfected epithelial cells with melan-a cells and found that Wnt10b secreted by epithelial cells could activate Wnt/β-catenin signaling in melan-a cells. We treated melan-a cells with the conditioned medium from JB6 mouse keratinocytes infected with AdWnt3a for 2 days. Wnt/β-catenin signaling in melan-a cells was also activated by testing TOP-flash system (data not shown). These results suggest that melan-a cells may respond to Wnt3a secreted by itself in an autocrine manner or secreted by keratinocytes in a paracrine manner in vitro.

Wnt signaling has been associated with proliferation of cells. Previous studies showed that Wnt3a promoted the proliferation of epidermal stem cells, mesenchymal stem cells, and fibroblast cells, and inhibited the proliferation of B-ALL cell lines (33-37). As for the melanocyte cell lineage, Dunn et al. (16) reported that Wnt3a did not stimulate proliferation of melanoblasts in vitro, and Delmas et al. (38) reported that stabilized β-catenin reduced the number of melanoblasts in vivo. Similar to findings in melanoblasts, Wnt3a failed to induce and, in fact, inhibited mature melanocyte proliferation in this study. It is important to bear in mind that Wnt3a may play distinct roles in different cell types, developmental stages, and organismal origin.

In mammalian melanocytes, melanin biosynthesis is catalyzed by three melanocyte-specific enzymes: tyrosinase, tyrosinase-related protein-1 (TRP1) and TRP2 (6). Tyrosinase is the rate-limiting enzyme in melanogenesis while TRP1 and TRP2 function as downstream of enzymes in the melanin biosynthetic pathway. To investigate the influence of Wnt3a on melanogenesis of mouse melanocytes, we analyzed tyrosinase activity and melanin content in melan-a cells. The data showed that Wnt3a significantly increased tyrosinase activity and melanin synthesis in melan-a cells. In support of our findings, previous studies have reported that inhibition of Wnt/β-catenin signaling strongly inhibited melanin synthesis (20,39).

Melanin synthesis is stimulated by a large number of effectors, including cAMP-elevating agents, cholera toxin, UV light and so on. MITF is the master regulator of melanogenesis that regulates the expression of the melanogenic enzymes, tyrosinase, TRP1, and TRP2 as well as other pigmentation factors (25,40). In this study, we detected the expression of MITF and its downstream target genes, tyrosinase, TRP1, and TRP2 as well as other pigmentation factors (25,40). In this study, we detected the expression of MITF and its downstream target genes, tyrosinase, TRP1, and TRP2, and demonstrated that Wnt3a upregulated the expression of MITF, tyrosinase, and TRP1 in melan-a cells at the protein level. Furthermore, Wnt3a rescued the expression of MITF and its downstream target genes following reduction of endogenous MITF in melan-a cells by AdSimMITF. With the same melan-a cells, Takeda et al. (41) found that Wnt3a protein induced MITF mRNA expression and activated the MITF promoter by recruiting LEF-1 and β-catenin to the LEF-1-binding site. Our data were consistent with the result of Takeda et al. (41) which demonstrated that MITF is a target gene of Wnt3a signaling.

**Figure 6.** Effect of Wnt3a on the expression of the melanogenic enzymes in AdSimMITF-infected melan-a cells. (A) Melan-a cells were infected with AdGFP, AdSimMITF, or co-infected with AdSimMITF and AdWnt3a, respectively. After 72 h, western blot analyses were performed with antibodies specific for MITF, tyrosinase, TRP1, TRP2 and GAPDH. (B) The relative protein expression level. The data are representative results of three independent experiments.

*P<0.05; **P<0.01.
In conclusion, the study demonstrates that Wnt3a reduces the proliferation of melan-a cells while simultaneously increases the melanin synthesis through the upregulation of MITF and its downstream genes, tyrosinase and TRP1.

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