

# Epidermal Wnt/ $\beta$ -catenin signaling regulates adipocyte differentiation via secretion of adipogenic factors

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Edited by Elaine Fuchs, The Rockefeller University, New York, NY, and approved February 28, 2014 (received for review July 9, 2013)

It has long been recognized that the hair follicle growth cycle and oscillation in the thickness of the underlying adipocyte layer are synchronized. Although factors secreted by adipocytes are known to regulate the hair growth cycle, it is unclear whether the epidermis can regulate adipogenesis. We show that inhibition of epidermal Wnt/ $\beta$ -catenin signaling reduced adipocyte differentiation in developing and adult mouse dermis. Conversely, ectopic activation of epidermal Wnt signaling promoted adipocyte differentiation and hair growth. When the Wnt pathway was activated in the embryonic epidermis, there was a dramatic and premature increase in adipocytes in the absence of hair follicle formation, demonstrating that Wnt activation, rather than mature hair follicles, is required for adipocyte generation. Epidermal and dermal gene expression profiling identified keratinocyte-derived adipogenic factors that are induced by  $\beta$ -catenin activation. Wnt/ $\beta$ -catenin signaling-dependent secreted factors from keratinocytes promoted adipocyte differentiation *in vitro*, and we identified ligands for the bone morphogenetic protein and insulin pathways as proadipogenic factors. Our results indicate epidermal Wnt/ $\beta$ -catenin as a critical initiator of a signaling cascade that induces adipogenesis and highlight the role of epidermal Wnt signaling in synchronizing adipocyte differentiation with the hair growth cycle.

skin | niche cross-talk | stem cells

Mammalian skin is a complex organ composed of a variety of cell and tissue types, including interfollicular epidermis, hair follicles (HFs), melanocytes, nerves, blood vessels, muscles, fibroblasts, and adipocytes. The development and patterning of these cells and tissues are governed by intercellular communication (1–3).

One well-known example of this communication is the link between the HF growth cycle and the oscillation in thickness of the dermal adipocyte layer (4–6). When HFs grow deep into the dermal adipocyte layer in the anagen (growth) phase of the cycle, the adipocyte layer dramatically increases in thickness. This event reflects both increased adipogenesis and hypertrophy of individual adipocytes (7). When HFs regress (catagen phase) and enter the resting (telogen) phase, the adipocyte layer becomes thinner. The growth cycle of rodent HFs is coordinated to form waves of hair growth that traverse the body and the thickness of the skin adipocyte layer oscillates in synchrony with these waves (3).

The synchronized patterns of HF growth and expansion of dermal fat correlate with the activation of the canonical Wnt pathway, which is well established to positively regulate anagen (3, 8). Expression of bone morphogenetic protein 2 (Bmp2) in the mature dermal adipocyte layer is inversely correlated with Wnt activity and inhibits HF growth (3). There is also evidence that immature dermal adipocytes activate HF stem cells to initiate the hair growth cycle (7). These reports suggest that the adipocyte differentiation process is a natural on-off cycling switch for the regulation of hair growth. Although the mechanisms of HF regulation by adipose tissues have been described, it is

unknown whether the HF regulates the morphogenesis, differentiation, and thickness of the dermal adipocyte layer.

In this study, we examined the role of epidermal Wnt/ $\beta$ -catenin signaling in regulating the dermal adipocyte layer and demonstrate that Wnt-dependent epidermal secreted factors promote adipocyte formation.

## Results

**Thickness of the Adipocyte Layer Correlates with Hair Growth in Mouse and Human Skin.** We quantified the volume of the dermal adipocyte layer (hypodermis) in dorsal mouse skin during the first 80 d of postnatal life (Fig. 1A). We confirmed that the adipocyte layer increased in thickness during HF morphogenesis, corresponding to the first anagen [postnatal day (P)10–P15], and decreased dramatically towards the following telogen (P23) (Fig. 1A). During the second anagen (P33), the volume of the adipocyte layer again increased and then decreased with the onset of the second telogen (P68). The increase in adipocyte thickness during the first postnatal HF regeneration happened during the transition from telogen to early anagen, and there was no change during later anagen (Fig. S1 A and B). The thickness of the adipocyte layer correlated with the hair cycle even when hair growth became asynchronous in older mice (Fig. S1 C–E). In addition, the volume of the adipocyte layer in telogen skin increased as mice aged (Fig. S1 E and F).

To examine whether there was also a relationship between HFs and adipocyte layer thickness in human skin, we compared

## Significance

The synchronized patterns of hair follicle growth and expansion of the dermal adipocyte layer have long been recognized. Although factors secreted by adipocytes are known to regulate the hair growth cycle, it is unclear whether, conversely, the epidermis can regulate adipogenesis. Our study now demonstrates that activation of epidermal Wnt/ $\beta$ -catenin signaling stimulates adipocyte differentiation *in vivo* and *in vitro*. The effect can be mediated by secreted factors, including insulin-like growth factor 2 and bone morphogenetic proteins 2 and 6.

Author contributions: G.D. and H.F. designed research; G.D., V.P., B.M.L., and H.F. performed research; G.D., K.N., R.S., H.F., and F.M.W. contributed new reagents/analytic tools; G.D., H.F., and F.M.W. analyzed data; and G.D., H.F., and F.M.W. wrote the paper.

The authors declare no conflict of interest.

This article is a PNAS Direct Submission.

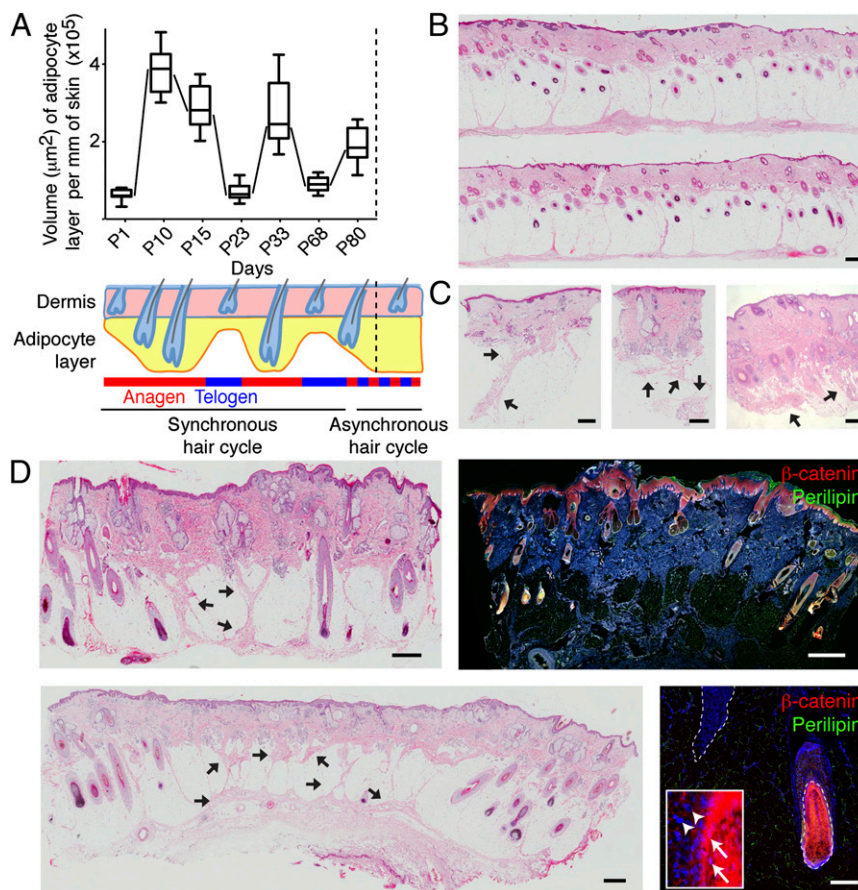
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This article contains supporting information online at [www.pnas.org/lookup/suppl/doi:10.1073/pnas.1312880111/-DCSupplemental](http://www.pnas.org/lookup/suppl/doi:10.1073/pnas.1312880111/-DCSupplemental).



**Fig. 1.** Relationship between hair follicle growth and the thickness of the dermal adipocyte layer. (A) Quantification of adipocyte layer thickness in P1-P80 mice. Diagram shows corresponding status of the hair growth cycle. Data are median and 25th and 75th percentile from three mice per time point. (B–D) H&E-stained human adult scalp skin from healthy (B), alopecia areata (C), and sebaceous nevus (D) patients. (Scale bar, 500  $\mu\text{m}$ .) (D) (Right) Sections of human sebaceous nevus were immunostained for  $\beta$ -catenin (red) and Perilipin (green) with DAPI counterstain (blue). White arrows in the *Inset* indicate nuclear  $\beta$ -catenin, which is absent in the cells indicated by arrowheads. (Scale bars, 500  $\mu\text{m}$ ; except 100  $\mu\text{m}$  in *Lower Right*.)

normal hairy scalp (Fig. 1B) with scalp from three patients with alopecia areata (Fig. 1C) and two with sebaceous nevus (Fig. 1D). In contrast to mice, human dermis is organized into regular lobules of mature adipocytes surrounded by fibroblasts as shown by H&E and Perilipin [a marker of differentiated adipocytes (9)] staining. In pathological samples, these reticular fibroblast networks became more pronounced while the total thickness of the adipocyte layer decreased (Fig. 1C and D, arrows). We quantitated dermal thickness in adjacent affected and unaffected skin in one alopecia areata patient who had not received any treatment for the condition (Fig. 1C, *Right*) and one case of sebaceous nevus (Fig. 1D, *Lower*). In each case, the thickness of the fat layer was decreased by 50% in the regions of hair loss (from  $\sim 2$  to 1 mm). In healthy human anagen HF, as in the mouse, nuclear  $\beta$ -catenin was present in hair bulb keratinocytes, which were surrounded by Perilipin-positive mature adipocytes (Fig. 1D, *Right*). Given the small number of samples analyzed and the complexity of the diseases etiology, we cannot establish a direct link between hair loss and adipocyte thickness in human skin. Nevertheless, regions of hair loss were correlated with reduced adipocyte layer thickness in the samples we examined.

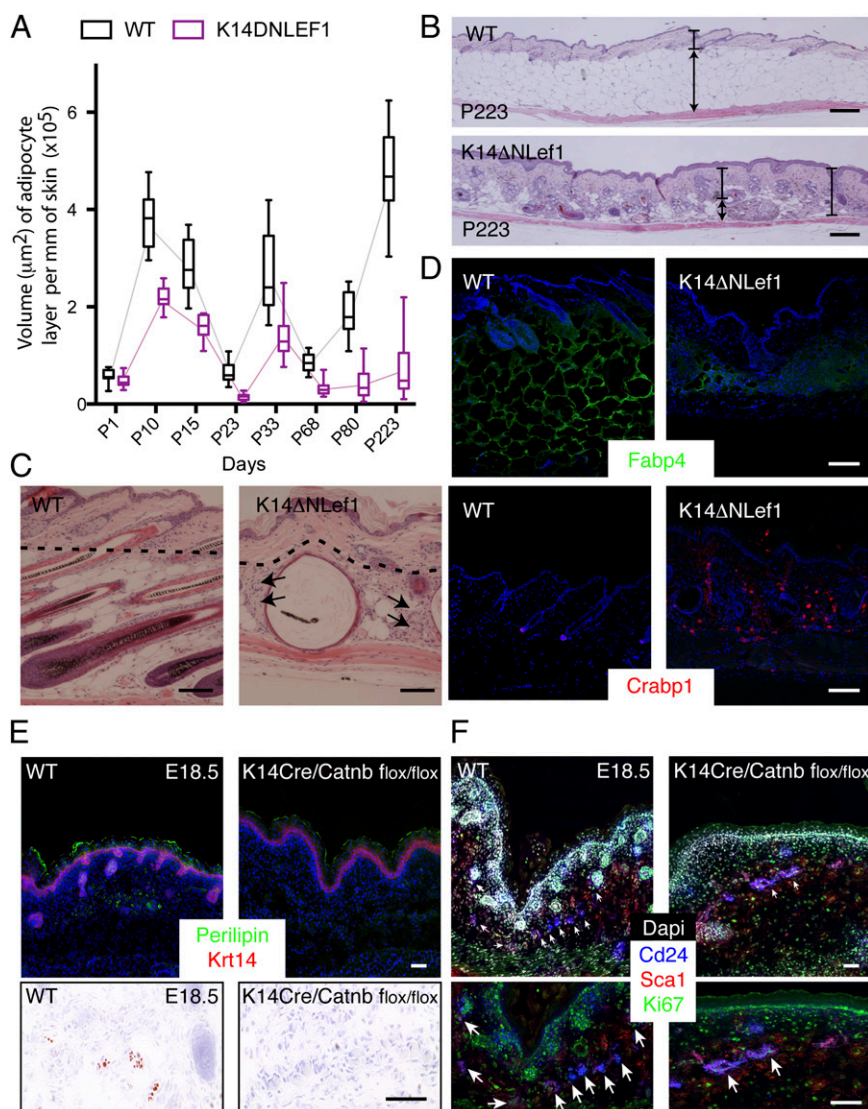
**Inhibition of Epidermal Wnt Signaling Results in a Reduction in the Adipocyte Layer in Adult and Embryonic Skin.** Activation of Wnt signaling stimulates HF to enter anagen and can reprogram cells in adult interfollicular epidermis to form ectopic follicles and differentiate along the HF lineages (10, 11). To investigate whether

Wnt/ $\beta$ -catenin signaling in the epidermis affects the hypodermis, we examined the skin of K14 $\Delta$ NLef1 transgenic mice that express N-terminally deleted Lef1 ( $\Delta$ NLef1), which lacks the  $\beta$ -catenin binding site, under the control of the Keratin 14 promoter (12). The keratin 14 promoter is active in the epidermal basal keratinocyte compartment but not in the dermis.  $\Delta$ NLef1 acts as a dominant-negative inhibitor of Wnt/ $\beta$ -catenin signaling. HF morphogenesis occurs normally in K14 $\Delta$ NLef1 mice, and expression of Wnt target genes is unaffected at P1 (Fig. S2A and B, *Left*). However, the follicles subsequently convert into epidermal cysts with ectopic sebocytes and Wnt signaling is inhibited at the base of the residual hair follicles (Fig. S2B, *Right*) (12).

Taking into account that the onset of the first postnatal anagen is delayed in mutant mice (12), we compared the thickness of the dermal adipose layer of WT and K14 $\Delta$ NLef1 mice during the same hair cycle phase. K14 $\Delta$ NLef1 dermal adipose volume was increased in conjunction with the growth of HF as in WT mice. However, at all time points except P1, the Perilipin-positive adipocyte layer was thinner than in WT mice (Fig. 2A and Fig. S2A). This phenotype became more prominent as mice aged (Fig. 2A and B).

The reduction in thickness of the adipocyte layer corresponded to an increase in the lower, reticular, dermal layer that is rich in fibrillar collagen (Fig. 2B and C). K14 $\Delta$ NLef1 epidermal cysts in the deep dermis were surrounded by reticular dermis rather than mature adipocytes (Fig. 2C, arrows).

The histological observations were confirmed by immunostaining adult mouse skin with antibodies to markers of adipocytes



**Fig. 2.** Inhibition of epidermal Wnt/ $\beta$ -catenin signaling impairs adipogenesis during hair follicle morphogenesis and regeneration. (A) Quantification of adipocyte layer thickness in WT and K14DNLEF-1 skin. Data are median and 25th and 75th percentile from three mice per time point. (B and C) H&E-stained dorsal skin of WT and K14DNLEF-1 mice at P223 (telogen) (B) and P150 (C). Double-headed arrows (B) indicate adipose thickness, whereas bars indicate dermis. In C, arrows show fibroblasts surrounding cysts and dotted lines indicate border between permanent and transient portions of HF. (D) Sections of adult WT and K14DNLEF-1 skin immunostained for Fabp4 (green) and Crabp1 (red), with DAPI counterstain (blue). (E) Sections of embryonic WT and K14Cre/Catnb flox/flox skin immunostained for Perilipin (green) and Krt14 (red), with DAPI counterstain (blue) (Upper) or stained with Oil red O (Lower). (F) Embryonic WT and K14Cre/Catnb flox/flox skin immunostained for Ki67 (green), Sca1 (red), and Cd24 (blue), with DAPI counterstain (white). White arrows indicate cluster of Cd24/Sca1<sup>+</sup> cells. [Scale bars, 300 (B), 100 (C and D), and 25  $\mu$ m (E and F).]

and fibroblasts. There was a striking decrease in Perilipin and Fabp4 [another marker of differentiated adipocytes (13)] positive cells in K14DNLEF1 compared with WT skin (Fig. 2D and Fig. S2A, Right). This decrease correlated with an increase in cellular retinoic acid binding protein 1 (Crabp1) expression. In WT adult skin, Crabp1 expression is restricted to the dermal papilla cells. In contrast, as previously reported (14), Crabp1 was widely expressed in the dermis of K14DNLEF1 skin (Fig. 2D). However, flow cytometric analysis of adipocyte precursor cells (Sca1<sup>+</sup>Cd24<sup>+</sup>) (7) revealed no significant difference between WT and transgenic adult skin (Fig. S2C and D).

To inhibit epidermal Wnt signaling during skin morphogenesis, we ablated  $\beta$ -catenin in embryonic skin via K14Cre. This resulted in a loss of Perilipin-positive, lipid-positive adipocytes (Fig. 2E). In WT skin, small clusters of Sca1/Cd24 double-positive cells were associated with the growing HF at embryonic day

(E)18.5. On removal of epidermal  $\beta$ -catenin, Sca1/Cd24 double-positive cell clusters were still present in the lower dermis (Fig. 2F).

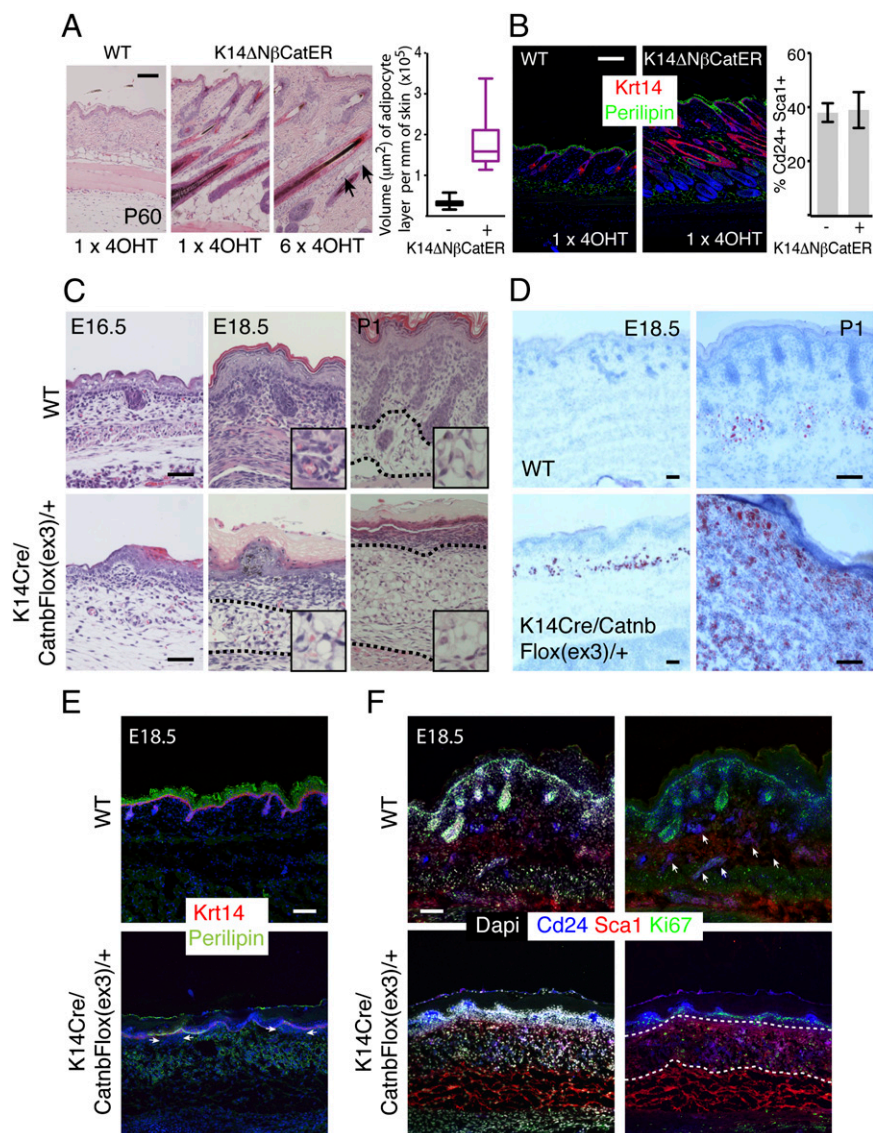
We conclude that epidermal  $\beta$ -catenin signaling is critical for adipocyte differentiation during development and that epidermal inhibition of Wnt/ $\beta$ -catenin activity in adult epidermis decreases the thickness of the adipose layer, due to a decrease in mature adipocytes. Although the increase in thickness of the adipose layer during normal mouse skin aging can have a number of underlying causes (15), our observations on K14DNLEF1 skin suggest an ongoing contribution of Wnt signaling (Fig. 2A).

**Activation of Wnt/ $\beta$ -Catenin Signaling in Adult Epidermis Induces Adipocyte Formation.** To examine whether activation of epidermal  $\beta$ -catenin signaling stimulates adipogenesis, we used K14DN $\beta$ -cateninER transgenic mice, in which stabilized, active  $\beta$ -catenin is fused with the C terminus of a mutant estrogen receptor under the

control of the Keratin 14 promoter (11). Wnt/ $\beta$ -catenin signaling in keratinocytes is activated by topical application of 4-hydroxytamoxifen (4OHT) to mouse dorsal skin. A single application to telogen skin is sufficient to stimulate anagen, whereas six treatments not only stimulate anagen of existing follicles but also induce ectopic HF formation as a result of expansion of the epidermal stem cell compartment and reprogramming cells in the sebaceous gland and interfollicular epidermis to differentiate along the hair follicle lineages (9).

Early anagen WT and K14 $\Delta$ N $\beta$ CateninER skin was treated with two doses of 4OHT skin (at P23 and P25) and analyzed at full anagen (P32 and P38). Although there was a transient increase in Cd24<sup>+</sup>Sca1<sup>+</sup> adipocyte progenitors (P32 only) on transgene activation, the thickness of the adipocyte layer was

unaffected (Fig. S3 A and B). The lack of the effect of the transgene is consistent with the known role of endogenous Wnt activation in promoting anagen (16). To examine whether signals originating from the epidermis can reset the adipocyte oscillation cycle, we next induced precocious activation of epidermal Wnt/ $\beta$ -catenin signaling. Application of one dose of 4OHT to telogen (7–11 wk old) transgenic mice led to induction of anagen and a marked expansion of the adipocyte layer (Fig. 3 A and B). There was no effect on the number of adipocyte precursors (Fig. 3B), but there was an increase in differentiated, Perilipin-positive adipocytes (Fig. 3B). When ectopic HF formation was induced by repeated applications of 4OHT, adipocytes were found in the upper dermis, where they are normally absent (Fig. 3A, arrows).



**Fig. 3.** Activation of epidermal Wnt/ $\beta$ -catenin signaling promotes dermal adipocyte formation. (A) H&E-stained sections of WT or K14 $\Delta$ N $\beta$ -cateninER (K14 $\Delta$ N $\beta$ CatER) dorsal skin treated with one or six doses of 4OHT in telogen (P60). Arrows indicate ectopic adipocyte formation in upper dermis. Graph shows quantification of adipocyte layer thickness after 1 dose of 4OHT. Data are median and 25th and 75th percentile from three mice. (B) Sections of WT or K14 $\Delta$ N $\beta$ -cateninER skin immunostained for Perilipin (green) and Krt14 (red) with DAPI counterstain (blue). (B) (Right) Flow cytometry of Sca1/Cd24 double-positive dermal cells after 1 dose of 4OHT. Data are mean  $\pm$ SD from three mice. (C) H&E-stained dorsal skin of WT and K14Cre/CatnbFlox(ex3)/+ mice. *Insets* show magnified images of lower dermis. (D–F) WT and K14Cre/CatnbFlox(ex3)/+ dorsal skin sections stained for Oil Red O (D), Perilipin (green), and K14 (red), with DAPI counterstain (blue) (E) or Ki67 (green), Sca1 (red), and Cd24 (blue), with DAPI counterstain (white) (F). White arrows indicate dermal condensate (E) or cluster of Cd24/Sca1<sup>+</sup> cells (F). Areas surrounded by black dotted lines are filled by differentiated adipocytes, whereas white dotted lines surround adipocyte precursors. [Scale bars, 100 (A and B) and 50  $\mu$ m (C–F).]

We conclude that activation of Wnt/ $\beta$ -catenin signaling in adult epidermis stimulates adipogenesis in association with induction of anagen and ectopic HF formation.

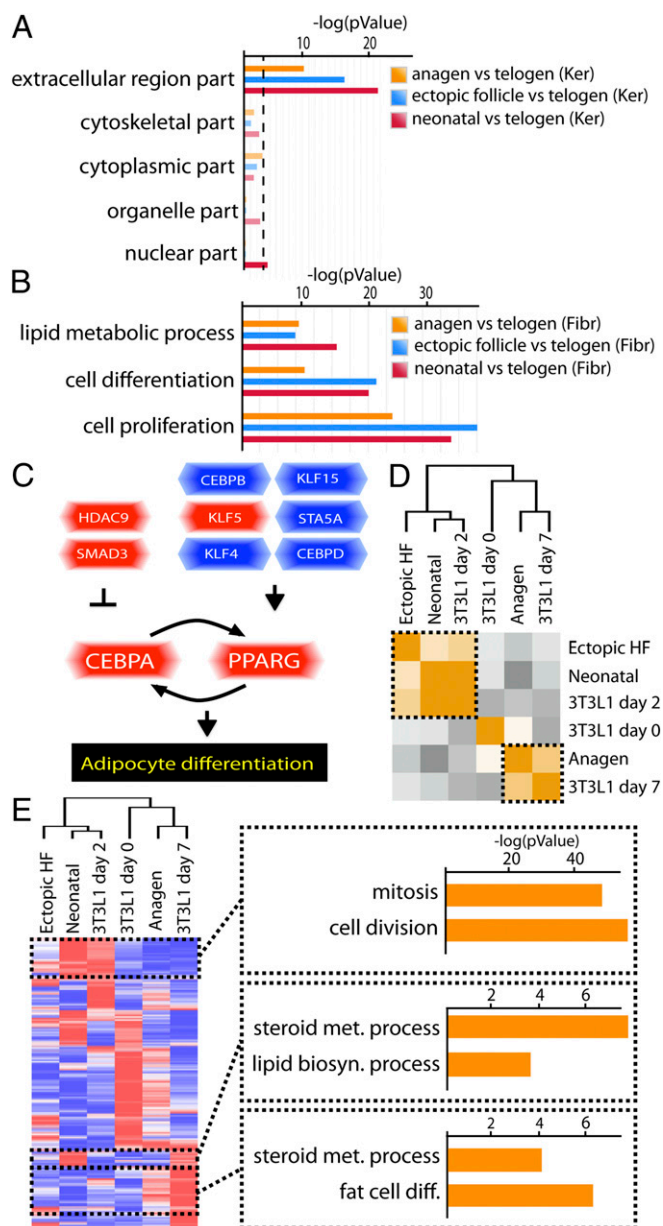
**Activation of Wnt/ $\beta$ -Catenin Signaling in Embryonic Epidermis Induces Adipocyte Differentiation.** To examine whether epidermal Wnt/ $\beta$ -catenin signaling directly stimulated dermal adipocyte formation, rather than indirectly via stimulation of hair growth, we generated K14Cre/CatnbFlox(ex3)/+ mice in which HF morphogenesis fails (17). These mice express stabilized, N-terminally truncated  $\beta$ -catenin in K14-positive epidermal basal keratinocytes. The K14 promoter becomes active in embryonic epidermis by E11.5 (17). During normal dorsal skin development in WT mice, mature adipocytes are readily detected from P1, and some cells with characteristic lipid vacuoles can be seen as early as P1 (E20.5) (areas between dotted lines and *Insets* in Fig. 3C), correlating with HF morphogenesis and activation of epidermal Wnt/ $\beta$ -catenin signaling (18). In K14Cre/CatnbFlox(ex3)/+ mice, cells with lipid vacuoles were already visible by E18.5, and large numbers of adipocytes were present at P1 (areas between dotted lines and *Insets* in Fig. 3C). The embryos died perinatally, allowing us to assess only developmental phenotypes.

To confirm the histological findings, we labeled sections for lipid, Perilipin, Fabp4, and Ppar $\gamma$  (19). K14Cre/CatnbFlox(ex3)/+ mice exhibited adipocyte differentiation in E18.5 skin and a massive increase in the number of adipocytes at P1 (Fig. 3D). Adipocytes were found throughout the dermis (Fig. 3D). In WT skin, expression of Perilipin, Fabp4, and nuclear Ppar $\gamma$  was mainly detectable at P1, whereas in transgenic mice, these markers were widely expressed at E18.5 and P1 (Fig. 3E and Fig. S3C). Sca1, a marker of pre- and mature adipocytes (13), was expressed in a similar pattern to Fabp4 (Fig. S3C). Conversely, in WT P1 skin, many fibroblasts, especially in proximity to the epidermal basement membrane, expressed Crabp1, whereas expression was reduced in K14Cre/CatnbFlox(ex3)/+ mice (Fig. S3C). In K14Cre/CatnbFlox(ex3)/+ skin, there was also a massive increase of Cd24<sup>+</sup>Sca1<sup>+</sup> preadipocytes (area between dotted lines) associated with stimulation of proliferation in the dermis (Fig. 3F).

These results indicate that activation of epidermal Wnt/ $\beta$ -catenin signaling during development induces adipogenesis in the absence of HF morphogenesis.

**Epidermal Wnt/ $\beta$ -Catenin Signaling Promotes Adipocyte Differentiation via Secretion of Ligands for BMP and Insulin Pathways.** Given that the epidermis and dermis are separated by a basement membrane, the inductive effect of epidermal Wnt/ $\beta$ -catenin signaling on adipocyte differentiation is most likely achieved by secreted factors. We searched for candidate factors by analyzing previously published gene expression profiles (9) generated from triplicate age-matched adult K14 $\Delta$ N $\beta$ -cateninER mice that were either treated with vehicle alone (telogen) or received one dose of 4OHT (to induce anagen) or six doses (to induce anagen and ectopic follicles). We did not compare gene expression data from transgenic mice treated with a single dose of 4OHT and anagen WT mice and therefore could not evaluate the extent to which the two situations are similar. Epidermal cells and dermal cells were flow sorted from the skin of each mouse and also compared with cells from triplicate neonatal WT mice (P2) (9). Overrepresentation analysis of gene ontology (GO) terms revealed that the category “extracellular region” was strongly enriched in the sets of genes that were differentially expressed (DEG) in the epidermis on 4OHT-induced activation of the Wnt pathway (Fig. 4A). The presence of a large number of secreted proteins (Table 1) highlights the potential complexity of communication between keratinocytes and neighboring cells.

In the fibroblast dataset, the “lipid metabolic process” and “cell differentiation” GO categories were highly represented in



**Fig. 4.** GO analysis of gene expression profiles of epidermal and dermal cells from K14 $\Delta$ N $\beta$ -cateninER mice. (A) GO overrepresentation analysis of DEG in keratinocytes from K14 $\Delta$ N $\beta$ -cateninER mice (only GO terms related to protein localization). Samples from adult anagen skin (one dose of 4OHT), adult anagen skin with ectopic HF (six doses of 4OHT), and neonatal skin were compared with adult telogen skin (9). (B) GO overrepresentation analysis of DEG in fibroblasts from K14 $\Delta$ N $\beta$ -cateninER mice. (C) Gene expression pattern of transcriptional regulators of adipocyte differentiation. Up- (red) and down- (blue) regulated genes in fibroblasts when epidermal Wnt/ $\beta$ -catenin signaling was activated are shown. (D) Heat map display of hierarchical clustering (Pearson correlation, positive in orange, negative in grey) of fibroblast DEG (neonatal, anagen, and ectopic follicles) and 3T3-L1 adipocyte differentiation markers (25). (E) Z-score heat map representing DEG as in D. Low expressed genes are in blue, high expressed in red. GO analysis of specific clusters is shown.

the DEG, together with the previously reported “cell proliferation” GO term (9) (Fig. 4B). We looked specifically at genes involved in the transcriptional regulation of adipocyte differentiation (20, 21). The master regulators Ppar $\gamma$  and Cebpa were significantly up-regulated. One of their upstream regulators,

**Table 1. Secreted proteins differentially expressed between growing and resting hair follicles**

Gene Categories	Up-regulated in growing hair follicles	Down-regulated in growing hair follicles
Extracellular matrix	Col4a1(A,E,N), Col12a1(A), Col18a1(A,N), Col4a2(A,N), Lamc1(N), Col4a3(E), Col27a1(E,N), Col5a1(E,N), Col14a1(E,N), Col11a1(N), Fbn2(E,N), Col5a2(N), Col2a1(E), Lamb1-1(A,N), Lama2(N), Col6a2(N), Emid1(E,N), Col8a1(E,N), Pcolce(N), Bgn(A,E,N), Fbln1(N), Fmod(N), Tnc(A,E,N), Vcan(N), Npnt(E), Olfml2a(N), Olfml2b(N), Spon1(A,E), Nid1(N), Nid2(N), Bcan(N), Hspg2(N)	Col8a2(A,N), Col27a1(A), Pcolce2(A,E,N), Egfl6(N), Fbln7(A,E,N), Spock2 (A,E,N), Tgfb1(E), Pcolce(E), Ecm1(N)
Peptidases and peptidases inhibitors	Adamts16(A,N), Adamts20(A,E,N), Adamts5(A,N), Adamts1(N), Cpxm1(A,E,N), Itih5(N), Klk7(A,E,N), Klk8(N), Pappa(A,E), Pcsk5(A,N), Pi5(A,E), Prss35(N), Serpine2(E,N), Tmprss11e(E)	Adam23(E), Adamts14(E,N), Angptl4(E,N), C2(A,E), Cpxm2(E), C3(E,N), C4b(E,N), Ctla2a(N), Htra1(A,E), Klk11(A,E), Loc100046035(E), Mmp12(A,E,N), Pm20d1(N), Prss23(A,E,N), Serpinb2(A,E,N), Timp2(N)
Defense response cytokines and chemokines	Ccl12(N), Cxcl14(N), Il1f5(N), Eda(E), Penk(E,N), Lipg(E,N), Pla2g7(A,E,N), Il1rap (N)	Ccl27a(E,N), Ccl8(A,E,N), Ccl9(A,E,N), Cxcl10(N), Cxcl12(N), Cxcl9(A,E,N), Grem1(A,N), Il1b(A,E,N), IL33(A,E,N), IL34(A,E), Xcl1(N), Il7(A,E,N), Pf4(A,E), Lipm(A,E,M), Il1rn(E), Il4ra(E,N), Il15ra(A,E,N), Il22ra2(A,E), Tnfrsf11b(N), Lepr(A,E,N), Tnfrsf18(N), C1qb(A,E), Cd163(A,E,N), C9(A,E,N), Fcgr2b(A,N), Defb1(A,E), B2m(N), Defb6(A,E,N), Smpdl3a(N), Smpdl3b(N)
Growth factors	Artn(N), Bmp6(A,E), Fgf5(A,E,N), Gdf10(A,E), Igf2(N), Igfbp2(N), Igfbp5(N), Ltbp1(E), Mdk(N), Ngf(N), Pdgfa(N), Wnt10b(N), Tgfb3(E), Wnt16(N), Tgfb2(E,N), Wnt4(N), Wnt5b(A,E), Wnt6(E,N), Wnt7a(N), Sfrp2(N), Sema3a(E,N), Hhip(E), Sema3b(E,N), Sema3f(N), Edn1(N), Edn2(A,E)	Fgf18(E,N), Ctgf(A,E,N), Fgfbp1(A,E), Figf(A,E,N), Igfbp3(E,N), Igfbp4(N), Kazald1(A,E,N), Ltbp2(A,E,N), Nrg4(A,E,N), Bmp1(N), Ptn(N), Wnt3(A,E), Dkk3(N), Sema3c(A,E), Sema3d(A,E)
Others	Agrp(A,N), Chi3l1(A), Coch(E), Egfl8(N), Lgi1(N), Egflam(N), Fam20a(A), Fam20c(N), Fstl1(E,N), Gpc3(N), Vash1(N), Vwa2(E), Krt2ap(N), Lepre1(N), Lgals1(N), Lgi2(N), Lox(N), Lypd6(N), Mup1-2-3(A), Kcp(A,E,N), Nppc(A), Nptx2(N), Nts(N), Pcsk5(A,N), Pthlh(E,N), Ptx3(N), S100a9(N), Scrg1(A)	Angptl7(A,E,N), Amy1(N), Bche(N), Cck(A,E,N), Chit1(A,E,N), Chl1(A,N), Cilp(N), Creg1(A,E,N), Cxadr(N), Efemp1(E), Fam132a(N), Lum(A,E), Fam3c(A,E), Gm885(E), Gm94(N), Gpha2(A,E,N), Gsn(N), Xdh(N), Hsd17b13(A,E,N), Sectm1b(E,N), Loc100046120(E,N), Sepp1(N), Sectm1a(A,E,N), Vit(E,N), Acppl(N), Lgals3bp(A,E,N), Myoc(A,E,N), Retnla(A,E,N), Sbsn(N), Npc2(A,E)

A, anagen; E, ectopic HF; N, neonatal.

Klf5, was also increased, whereas the others were decreased (Fig. 4C). These observations indicate that epidermal Wnt signaling stimulates adipocyte differentiation and also indicate some heterogeneity in the dermal cell response.

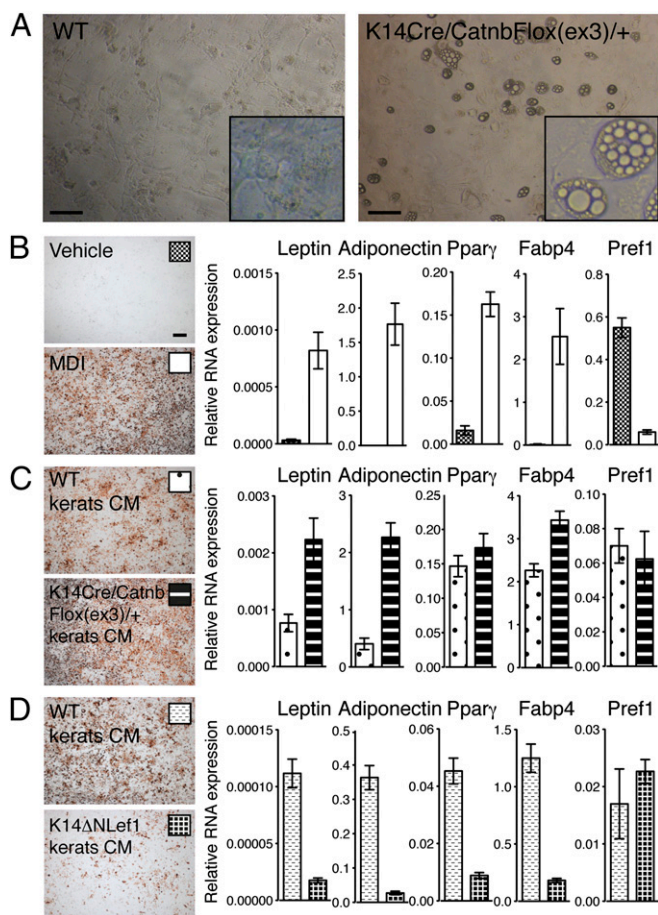
3T3-L1 cells are a well-characterized in vitro model of adipogenesis (22, 23). Differentiation is characterized by an early phase of clonal expansion followed by terminal differentiation (24). To further understand fibroblast responses to epidermal Wnt activation, we compared fibroblast DEG from neonatal, adult anagen, and ectopic follicle-bearing skin with DEG of differentiating 3T3-L1 cells (25). This analysis revealed that fibroblasts from ectopic follicle and neonatal skin were more similar to the early phase of 3T3-L1 differentiation (day 2), whereas adult anagen fibroblasts resembled fully differentiated adipocytes (day 7) (Fig. 4D). To characterize these correlations further, we performed GO analysis on the common gene clusters: adult anagen and 3T3-L1 day 7 cells shared a gene set enriched for “steroids metabolic processes” and “fat cell differentiation,” whereas neonatal fibroblasts and 3T3-L1 day 2 cells shared a gene set enriched for “mitosis” and “cell division” (Fig. 4E). In addition, neonatal fibroblasts and 3T3-L1 day 7 cells were enriched for genes involved in “steroids metabolic process” and “lipid biosynthesis process” (Fig. 4E). These results suggest that neonatal fibroblasts correspond to an earlier stage of adipogenesis, whereas in adult anagen, maturation of adipocytes is taking place.

We routinely culture keratinocytes on an irradiated feeder layer of J2 Swiss mouse 3T3 cells (26). The feeder layer of K14Cre/CatnbFlox(ex3)/+ keratinocytes contained many cells with large lipid vacuoles, whereas that of WT keratinocytes did not (Fig. 5A). To establish experimentally whether  $\beta$ -catenin activation stimulates keratinocytes to secrete adipogenic factors, we cultured 3T3-L1 cells in methylisobutylxanthine, dexamethasone,

and insulin adipocyte differentiation medium (22) and medium conditioned by keratinocytes from WT, K14Cre/CatnbFlox(ex3)/+, or K14 $\Delta$ NLef1 mice. Six days later, we examined lipid production by Oil Red O staining and quantitative RT-PCR (qRT-PCR) of genes that are known to be up- (Leptin, Adiponectin, Pparg, and Fabp4) or down- (Pref-1) regulated during adipocyte differentiation (Fig. 5 B–D).

3T3-L1 cells exposed to K14Cre/catnbFlox(ex3)/+ keratinocyte conditioned medium exhibited greater and more rapid adipogenesis than those exposed to WT keratinocyte medium (Fig. 5C). Conversely, medium conditioned by K14 $\Delta$ NLef1 keratinocytes inhibited adipogenesis (Fig. 5D). When sparse preadipocyte 3T3-L1 cells were treated with conditioned medium, there was no effect on growth rate (Fig. S4). These data suggest that Wnt/ $\beta$ -catenin regulates epidermal production of factors that stimulate adipocyte differentiation.

To identify secreted factors that affect adipocyte differentiation, we tested a panel of 14 candidate factors that were identified from the K14 $\Delta$ N $\beta$ -cateninER expression profile or significantly down-regulated in keratinocytes expressing  $\Delta$ NLef1 or up-regulated in K14Cre/CatnbFlox(ex3)/+ embryonic epidermis. Three factors stimulated 3T3-L1 differentiation: Bmp2, Bmp6, and insulin-like growth factor-2 (Igf2) (Fig. 6 A and B). Consistent with these observations, differentiation of 3T3-L1 cells in the presence of WT keratinocyte conditioned medium was stimulated, in a dose-dependent manner, by exogenous insulin, whereas the effect of insulin was much less pronounced in conditioned medium of K14Cre/CatnbFlox(ex3)/+ keratinocytes (Fig. 6C). In addition, pharmacological inhibition of BMP signaling (27) significantly decreased the adipogenic effect of K14Cre/CatnbFlox(ex3)/+ keratinocyte conditioned medium (Fig. 6D).



**Fig. 5.** Wnt/ $\beta$ -catenin signaling in keratinocytes regulates adipocyte differentiation through secreted factors. (A) Phase contrast images of 3T3-J2 cocultured with WT or K14Cre/CatnbFlox(ex3)/+ keratinocytes for 6 d. Insets are higher magnification views of 3T3-J2 cells. (B–D) Differentiation of 3T3-L1 cells in MDI medium (B), MDI medium conditioned by keratinocytes from WT or K14Cre/CatnbFlox(ex3)/+ mice (C), or K14 $\Delta$ NLeF1 mice (D) for 6 d. Cells were stained with Oil Red O (Left). Gene expression levels of adipocyte markers (Leptin, Adiponectin, Ppar $\gamma$ , and Fabp4) and a preadipocyte marker (Pref1) in 3T3-L1 cells were examined by qRT-PCR. Data are mean  $\pm$  SD from three independent wells. [Scale bars, 100 (A) and 200  $\mu$ m (B–D).]

These *in vitro* results demonstrate that activation of epidermal Wnt/ $\beta$ -catenin signaling can induce adipocyte differentiation by stimulating secretion of ligands for the BMP and insulin signaling pathways.

## Discussion

It has long been recognized that the HF growth cycle and oscillation in the thickness of the skin adipocyte layer are synchronized. Recently, adipocyte-derived factors have been reported to regulate the hair growth cycle (3, 7). Our study now identifies activation of Wnt/ $\beta$ -catenin signaling in keratinocytes as a trigger for adipocyte differentiation, suggesting that periodic activation of epidermal Wnt/ $\beta$ -catenin signaling during the hair cycle contributes to the synchrony.

HF formation was not required for induction and expansion of dermal adipocytes during skin morphogenesis (Fig. 2). In addition, Wnt/ $\beta$ -catenin activation in cultured keratinocytes stimulated differentiation of 3T3-L1 cells into mature adipocytes *in vitro* (Fig. 5). These findings indicate that Wnt/ $\beta$ -catenin activation in keratinocytes, rather than HF formation per se, promotes adipocyte differentiation. Our observations provide an explanation for why adipocytes still form in hairless skin, such as

the paw skin. It remains possible that adipocyte development in embryonic skin and changes in adipocyte thickness in adult skin rely on different signaling pathways and that only the embryonic stage relies directly on epidermal Wnt signaling. *In vitro* coculture of 3T3 cells and keratinocytes may mimic the developmental stage rather than the adult situation.

During mouse skin morphogenesis, activation of Wnt/ $\beta$ -catenin signaling and formation of HFs are observed by E14.5, whereas commitment to the adipocyte lineage occurs at E16.5 (28). This timing suggests that epidermal Wnt/ $\beta$ -catenin signaling is the initial signal that promotes dermal adipogenesis, rather than adipocytes initiating HF formation. When  $\beta$ -catenin was deleted in embryonic epidermis, CD24/Scal<sup>+</sup> adipocyte precursors were still detectable, suggesting that the lack of adipocytes is due to a block in terminal differentiation. When epidermal  $\beta$ -catenin was activated in embryonic and adult skin, there was an expansion of differentiated adipocytes. These results are consistent with recent evidence that epidermal  $\beta$ -catenin activation leads to expansion of both the upper and lower dermal lineages (28).

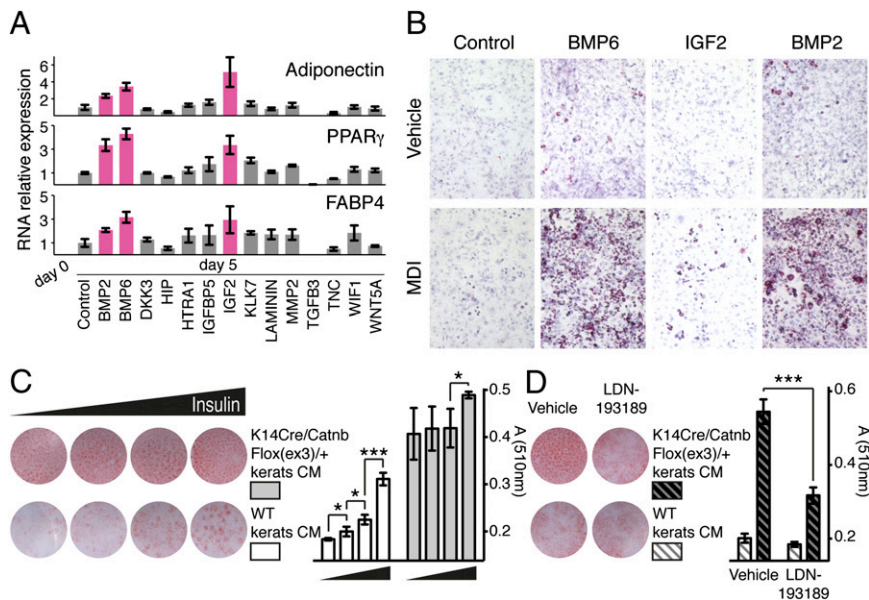
In adult skin, ectopic activation of epidermal Wnt/ $\beta$ -catenin signaling (Fig. 3) or transplantation of adipocyte precursors into the dermis (7) leads to expansion of the adipocyte layer or HF anagen, respectively. These findings suggest that both the epidermis and the adipose layer can initiate the positive feedback loop linking hair cycle progression to adipocyte differentiation. It will be interesting to determine the entity of the clock that regulates the timing and duration of hair growth cycle and other associated cyclic regenerative events in the skin.

Activation of Wnt/ $\beta$ -catenin signaling in keratinocytes led to the expression of a diverse set of secreted factors. Among them, we validated Igf2 and Bmp6 as factors that promote adipogenesis *in vitro* and also confirmed a role for Bmp2 (17, 20, 21). Although we have not determined the sites of expression of these factors during skin morphogenesis and hair follicle cycling, there are indications that they are expressed in the hair follicle (3, 29, 30). In addition, expression of several components of the Igf signaling pathway are altered in skin aging, when the adipocyte layer increases in thickness (15). It is notable that Bmp2, Igf2, and the hair cycle regulator Pdgfa are expressed both by adipocytes and epidermal cells (3, 7, 9), pointing to the complexity of communication between the epidermis and adipocyte layer.

Bmp signaling is known to be important for the functional identity of the dermal papilla (31). The activation of epidermal Wnt/ $\beta$ -catenin signaling during development also induces widespread dermal condensate formation (17) (Fig. 3E, white arrows) and in adulthood leads to expansion of the papillary and reticular dermis (28). Therefore epidermal Wnt affects multiple dermal compartments, indicative of both short and long range signaling.

A balance between pro- and anti-adipogenic signals must be required for spatiotemporal regulation of adipogenesis, and indeed, our gene expression profiles included some antiadipogenic factors (21). Conditioned medium from K14 $\Delta$ NLeF1 keratinocytes inhibited 3T3-L1 adipocyte differentiation, suggesting that they secrete inhibitors of adipogenesis. Because activation of Wnt signaling in keratinocytes influences the behavior of melanocytes, nerves, and fibroblasts (9, 17, 32), some of the antiadipogenic factors may primarily regulate other cell populations in the skin.

In conclusion, our data highlight the role of epidermal Wnt signaling in adipocyte morphogenesis and synchronizing adipocyte differentiation with HF growth. Further studies are necessary to identify the upstream master clock regulators controlling the timing and duration of hair growth cycle and, in particular, whether long-range regulation (neuronal, systemic or hormonal) is involved. There are several physiological and anatomical advantages of synchronizing the HF growth cycle with oscillations in thickness of the adipocyte layer. The combination of a high density of hairs and dermal adipocytes insulates the body from the cold. In addition, the down growth of the HF during



**Fig. 6.** Identification of epidermal adipogenic factors. (A) qRT-PCR of Adiponectin, Ppar $\gamma$ , and Fabp4 expression in 3T3-L1 cells after 5 d of exposure to adipogenic medium containing the indicated recombinant proteins. Data are mean  $\pm$  SD from three independent wells. (B) 3T3-L1 cells stained with Oil Red O after 7 d of exposure to the proteins indicated in the absence (Upper) or presence (Lower) of adipogenic medium. (C) Effect of different insulin concentrations on 3T3-L1 preadipocyte differentiation induced by MDI and conditioned medium from WT or K14Cre/CatnbFlox(ex3)/+ keratinocytes. Oil Red O staining and quantification (absorbance at 510 nm) are shown. Data are mean  $\pm$  SD from four independent wells. (D) Effect of Bmp inhibition during differentiation induced by MDI adipogenic medium and conditioned medium from WT or K14Cre/CatnbFlox(ex3)/+ keratinocytes. Oil Red O staining and quantification are shown. Data are mean  $\pm$  SD from four independent wells. \* $P$  < 0.05; \*\*\* $P$  < 0.0005.

anagen may be facilitated by the presence of an adipocyte layer separating the reticular dermis from the underlying striated muscle, the panniculus carnosus.

## Materials and Methods

**Mice.** All animal procedures were subject to local ethical approval and performed under a UK Government Home Office license and RIKEN's Regulations for the Animal Experiments. K14 $\Delta$ NleF1, K14 $\Delta$ N $\beta$ -cateninER, and Pdgfra-gfp transgenic mice have been described previously (11, 12). K14Cre/CatnbFlox(ex3)/+ and K14Cre/Catnb flox/flox mice were generated by crossing K14Cre (Jackson Laboratory) with CatnbFlox(ex3) mice (generous gift from Makoto Taketo, Department of Pharmacology, Kyoto University, Graduate School of Medicine, Kyoto) (33) and  $\beta$ -catenin lox/lox mice (16), respectively. The K14 $\Delta$ N $\beta$ -cateninER transgene was activated by topical application of 4-hydroxytamoxifen (34). Only female mice were used for this study.

**Human Tissue.** Human tissue samples were obtained with informed consent and processed for research in accordance with the recommendations of the relevant national legislation.

**Histology, Flow Cytometry, and Staining.** Optimal cutting temperature compound or paraffin-embedded tissues were sectioned and stained by conventional methods. Primary antibodies for immunofluorescence were as follows: Perilipin A and Axin2 (Abcam); Keratin14 (Covance); Fabp4 (R&D Systems); Ppar $\gamma$ , Cyclin D1, and Jag1 (Santa Cruz); Sca1 and Cd24 (eBioscience); Ki67 (Novocastra) and Crabp1 (Sigma); and Krt31 (Progen). Images were acquired with a Leica TCS SP5 Tandem Scanner confocal microscope. Images of H&E-stained sections were acquired with an Aperio scan-scope, and quantification of the adipocyte layer was performed with Imagescape. The HF area within the adipocyte layer was excluded from the quantification.

For Oil Red O staining, sections or cultured cells were washed with PBS and fixed in formalin for 10 min at room temperature. Cells were incubated for 30 min at room temperature with 60% filtered Oil Red O stock solution (0.3g/100 mL of isopropanol). Sections/cells were washed with 60% (vol/vol) isopropanol and then water before visualization. Lipid levels were quantified by extracting Oil Red O stained cells with isopropanol and measuring absorbance at 510 nm.

Dermal cells for FACS analysis were isolated as described previously (9) and labeled with the following antibodies: anti-CD24-PE clone: M1/69, anti-Ly-6A/E (Sca1)-FITC or Alexa Fluor 700, anti-Cd29-FITC, anti-Cd34-APC (eBioscience), and lineage antibody mixture V450 (BD Horizon). Pdgfra cells were

sorted with Anti-Cd140a-PE or APC (eBioscience) or Pdgfra-gfp. Labeled cells were analyzed on a BD LSRFortessa cell analyzer. DAPI or LIVE/DEAD Fixable Violet Dead Cell Stain (Life Technologies) was used to exclude dead cells.

**Cell Culture.** Mouse primary dorsal skin keratinocytes were cultured on 3T3-J2 feeders in calcium-free FAD medium (1 part Ham's F12, adenine  $1.8 \times 10^{-4}$  M, 3 parts DMEM) supplemented with 10% FCS, hydrocortisone (0.5  $\mu$ g/mL), insulin (5  $\mu$ g/mL), cholera toxin (8.4 ng/mL), and epidermal growth factor (10 ng/mL) in collagen I-coated culture flasks (35, 36).

3T3-L1 preadipocytes (generous gift from Jaswinder K. Sethi, University of Cambridge Metabolic Research Laboratories, Institute of Metabolic Science, Cambridge, United Kingdom) were maintained in DMEM (10% calf serum). Forty-eight hours after confluence, differentiation was induced with DMEM supplemented with 10% FBS, 1  $\mu$ M dexamethasone, 1  $\mu$ g/mL insulin, and 0.5 mM 3-isobutyl-1-methylxanthine (IBMX) (MDI medium). Two days later, the medium was replaced with DMEM (10% FBS) and was changed every 2 d. The following recombinant proteins were added every second day: Bmp2 (200 ng/mL), Bmp6 (100 ng/mL), Dkk3 (4  $\mu$ g/mL), Hip (10  $\mu$ g/mL), Htra1 (8  $\mu$ g/mL), Igfbp5 (5  $\mu$ g/mL), Igf2 (20 ng/mL), Klk7 (5  $\mu$ g/mL), Laminin (10  $\mu$ g/mL), Mmp2 (5  $\mu$ g/mL), Tgf- $\beta$ 3 (50 ng/mL), Tnc (10  $\mu$ g/mL), Wif1 (1  $\mu$ g/mL), and Wnt5a (1  $\mu$ g/mL). On day 5 after plating, the cells were harvested, and the RNA was extracted. To test the effects of insulin, 100 $\times$  serial dilutions from 1  $\mu$ g/mL were used to supplement the medium. The LDN-193189 Bmp inhibitor was used at final concentration of 0.1  $\mu$ M. Growth rate was determined using the Incucyte real-time video imaging system.

**qRT-PCR.** Total RNA was purified with the RNeasy Mini Kit (Qiagen) with on-column DNaseI digestion according to the manufacturer's instructions. RNA (500 ng) was reverse transcribed with SuperScriptIII (Invitrogen) and random primers. cDNA (10 ng) was used for qRT-PCR with SYBR green super mix (ABI). Data represent mean values  $\pm$  SD. Primers used in this study are listed in Table S1.

**GO and Clustering Analysis.** GO term enrichment analysis was performed using DAVID (<http://david.abcc.ncifcrf.gov>) or GeneGO pathway analysis software (<https://portal.genego.com>) on DEG in keratinocytes and fibroblasts that were previously identified (9). 3T3-L1 DEG at days 0, 2, and 7 from Mikkelsen et al. (25) were compared with fibroblast DEG from Collins et al. (9). Heat maps and clustering analysis were performed with GenePattern 2.0 (37).



**ACKNOWLEDGMENTS.** We acknowledge the core services of the Cancer Research UK Cambridge Research Institute, where the project was initiated, and the assistance of Charlotte Collins, Marta Lesko, Maria Mastrogiannaki, and Kai Kretschmar, who generously provided mice/sections for research. We also thank Asako Nakagawa for technical assistance. G.D. thanks Samuel Woodhouse for daily discussions about this project. This work was supported by grants to F.M.W. from Cancer Research UK, the European Union, the Medical Research Council, and the

Wellcome Trust and to H.F. from RIKEN. B.M.L. was the recipient of a Federation of European Biochemical Societies long-term fellowship. We gratefully acknowledge financial support, in the form of access to the flow facility, from the Department of Health via the National Institute for Health Research comprehensive Biomedical Research Centre award to Guy's & St Thomas' National Health Service Foundation Trust in partnership with King's College London and King's College Hospital NHS Foundation Trust.

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