The role of Hes genes in intestinal development, homeostasis and tumor formation

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SUMMARY
Notch signaling regulates intestinal development, homeostasis and tumorigenesis, but its precise downstream mechanism remains largely unknown. Here we found that inactivation of the Notch effectors Hes1, Hes3 and Hes5, but not Hes1 alone, led to reduced cell proliferation, increased secretory cell formation and altered intestinal structures in adult mice. However, in Apc mutation-induced intestinal tumors, inactivation of Hes1 alone was sufficient for reducing tumor cell proliferation and inducing differentiation of tumor cells into all types of intestinal epithelial cells, but without affecting the homeostasis of normal crypts owing to genetic redundancy. These results indicated that Hes genes cooperatively regulate intestinal development and homeostasis and raised the possibility that Hes1 is a promising target to induce the differentiation of tumor cells.

KEY WORDS: Hes, Notch signaling, Intestinal tumor, Intestine, Stem cell, Mouse

INTRODUCTION
The intestinal epithelium is shaped into crypts, which are invagination structures, and villi, which are finger-like protrusions, and is continuously renewed from stem cells throughout life (Radtke and Clevers, 2005; Cronier et al., 2006; Scoville et al., 2008). Stem cells are located at the bottom of the crypts and produce transit-amplifying progenitors, which then undergo terminal differentiation into four distinct cell types: absorptive cells, mucin-secreting goblet cells, anti-bacterial peptide-secreting Paneth cells, and gut hormone-secreting enteroendocrine cells. The latter three cell types belong to the secretory cell lineage. Postmitotic differentiated cells migrate towards the apex of a villus before being shed into the intestinal lumen. It has been shown that signaling molecules such as those of the Notch signaling pathway are involved in the homeostasis of the intestinal cellular system, including stem cell maintenance, cell fate decision and maturation (Radtke and Clevers, 2005; Cronier et al., 2006; Scoville et al., 2008), but the precise downstream mechanisms and crosstalk of signaling pathways remain to be determined.

It has been shown that the Notch signaling pathway is also involved in the development of colorectal tumors. In ApcMin mice, the Wnt signaling effector β-catenin is stabilized by mutations in the intestinal tumor suppressor gene Apc, and β-catenin forms a complex with Tcf factors in the nucleus, which causes continuous expression of Wnt target genes and the development of adenomas, mainly in the small intestine (Moser et al., 1990; Su et al., 1992; Bienz and Clevers, 2000; van de Wetering et al., 2002). However, blockade of Notch signaling with a γ-secretase inhibitor turned adenoma cells of ApcMin mice into goblet cells (van Es et al., 2005a), whereas activation of Notch signaling in Apc mutant (Apc1638N) mice resulted in at least a 20-fold increase in the number of adenoma cells compared with that in Apc1638N mice (Fre et al., 2009), suggesting that Notch signaling is involved in intestinal tumor formation. Although Notch signaling is a promising target for anti-cancer therapy, all stem/progenitor cells in healthy intestinal regions, as well as the tumor cells, differentiate into goblet cells by inhibition of Notch signaling, which could cause severe side effects. Thus, a strategy to inhibit Notch signaling only in tumor cells without affecting healthy cells is desirable.

Notch signaling is conveyed from cell to cell (Wilson and Radtke, 2006; Kageyama et al., 2007). When the membrane-bound Notch receptor interacts with its ligand anchored in the membrane of neighboring cells, an intrinsic γ-secretase cleaves the receptor, releasing the Notch intracellular domain (NICD). NICD then translocates into the nucleus where it forms an active transcriptional complex with Rbpj, an essential mediator of all Notch receptors, to stimulate the expression of many target genes. Hes1, one of the best-characterized target genes of the Notch signaling pathway, regulates the maintenance of neural and hematopoietic stem cells (Ohtsuka et al., 1999; Kunisato et al., 2003; Shimojo et al., 2008). It has been shown that Hes1 is expressed by proliferating cells, including stem cells in the intestinal crypt, and that it represses the expression of the Math1 (Atoh1) and neurogenin 3 (Ngn3 or Neurog3) genes (Jensen et al., 2000; Yang et al., 2001; Jenny et al., 2002; Kayahara et al., 2003; Fre et al., 2005; van Es et al., 2005a; Suzuki et al., 2005; Riccio et al., 2008; Kopinke et al., 2011). Math1 regulates the development of all types of secretory cells, whereas Ngn3 regulates enteroendocrine cell development (Yang et al., 2001; Jenny et al., 2002). Transgenic mice expressing NICD constitutively upregulate Hes1 expression, increase the number of proliferating cells and lose all secretory cells, whereas Hes1 knockout (KO) mice show overproduction of secretory cells (Jensen et al., 2000; Stanger et al., 2005). These results suggest that the Notch-Hes1 pathway promotes the proliferation of intestinal stem/progenitor cells and inhibits secretory cell development by repressing Math1 and Ngn3 expression. Hes1 is also expressed in human colon cancer cells and in adenoma cells of ApcMin mice (van Es et al., 2005a; Fernández-
Majada et al., 2007), and Hes1 functions as a downstream target of both the Notch and Wnt signaling pathways in Ls174T colon cancer cells (Rodilla et al., 2009), suggesting that crosstalk between Notch and Wnt signaling might take place via Hes1. In the absence of Hes1, Hes3 and Hes5 expression is upregulated in the intestine (Jensen et al., 2000), but it remains unclear whether all Hes genes function similarly or whether Hes1 has a dominant role in intestinal development, homeostasis and tumor formation.

In order to determine the roles of Hes genes in postnatal intestinal development, homeostasis and tumor formation, we analyzed the intestine of Hes1 conditional KO (cKO) and Hes1+/333/555 cKO mice and crossed these mice with ApcMin to determine the roles of Hes genes in intestinal tumors.

MATERIALS AND METHODS

Mice

Villin-Cre mice (el Marjou et al., 2004) and ApcMin mice (Su et al., 1992) were obtained from Jackson Laboratories. Hes1 cKO mice were generated by crossing homozygous Hes1 floxed mice (Imayoshi et al., 2008) with Villin-Cre mice. Hes1/333/555 cKO mice were generated by crossing Hes1floxed/Hes3floxed/Hes5floxed with Villin-CreHes1floxed/Hes3floxed/Hes5floxed. Villin-CreHes1floxed and Villin-CreHes3floxed/Hes5floxed Hes1+/–;Hes3–/–;Hes5–/– mice were normal and used as controls. ApcMin;Hes1 cKO mice were obtained by crossing homozygous Hes1 floxed mice with ApcMin;Villin-Cre;Hes1+/– mice.

Ah-Cre mice were generated as previously reported (Campbell et al., 1996; Ireland et al., 2004). The plasmid pAHIR1 was kindly gifted by Dr C. Roland Wolf (Dundee, UK). Rosa-iacZ mice (Soriano, 1999) were kindly supplied by Dr Philippe Soriano. ApcMin;Ah-Cre;Hes1floxed mice were obtained by crossing homozygous Hes1 floxed mice with ApcMin;Ah-Cre;Hes1floxed mice. For induction of the Ah promoter, mice were treated with 80 μg/g body weight β-naphthoflavone in corn oil by intraperitoneal injection for 5 consecutive days and sacrificed after 3 weeks. All animals were handled in accordance with the Kyoto University Guide for the Care and Use of Laboratory Animals.

Histology and immunohistochemistry

Tissues were fixed overnight in 10% neutral buffered formalin, paraffin embedded and sectioned at 3-5 μm. Sections were stained with Hematoxylin and Eosin or Alcian Blue or subjected to immunohistochemistry with the following primary antibodies: anti-cleaved caspase 3 (1:200 dilution, Cell Signaling), anti-BrdU (1:500, BD), anti-Hes1 (1:100), anti-Hes3 (1:100, Santa Cruz), anti-Hes5 (1:100, Abcam), anti-Ki67 (1:100, DAKO), anti-lysozyme (1:200, DAKO), anti-villin (1:100, Cell Signaling), anti-chromogranin (1:100, Sigma, St Louis, MN, USA), anti-Lgr5 (1:300, MBL International), anti-β-catenin (1:50, Transduction Laboratories), anti-EphB3 (1:50, R&D Systems), anti-phosphorylated Smad1/5/8 (1:200, Cell Signaling), anti-Math1 (1:50, Development Studies Hybriodoma Bank), anti-Ngn3 (1:300, Development Studies Hybriodoma Bank) and anti-Sox9 (1:1000, Chemicon). Antigen retrieval was performed by boiling in 10 mM citrate buffer pH 6.0 for 15 minutes for all antibodies except anti-lysozyme, which was used after treatment with 20 μg/ml proteinase K in TE pH 8.0 (10 minutes at 37°C), and anti-BrdU, which was used after treatment with 2 N HCl (30 minutes at 37°C) and boiling in citrate buffer as above. Immunoperoxidase labeling was performed with the VECTASTAIN ABC Kit (Vector Laboratories). Slides were developed using DAB and counterstained with Hematoxylin.

For X-gal staining, perfusion with 4% paraformaldehyde in 0.1 M phosphate buffer was performed, and frozen sections (8 μm) were prepared. These sections were stained with X-gal as previously described (Imayoshi et al., 2006). For single BrdU labeling, mice were injected intraperitoneally with 100 μg/kg BrdU 2, 48 or 96 hours before sacrifice. For multiple BrdU labeling, mice were injected intraperitoneally with 100 mg/kg BrdU four times at 2-hour intervals and sacrificed after 9 days.

For quantification, the number of each cell type per section was counted in at least five sections of three animals per genotype. Statistical differences were determined by Student’s t-test.

RESULTS

Accelerated differentiation of secretory cells in the small intestine of Hes1 cKO mice at a postnatal stage

Because Hes1 KO mice die soon after birth (Ishibashi et al., 1995), the postnatal role of Hes1 was not known. To understand the role of Hes1 in postnatal development and homeostasis of the intestine, we generated Hes1 cKO mice by crossing Hes1 flox mice (Imayoshi et al., 2008) with Villin-Cre mice, which induced a Cre-dependent recombination in the intestinal epithelium by embryonic day (E) 12.5 (Madison et al., 2002; el Marjou et al., 2004). In Villin-Cre;Hes1flox mice, Hes1 was almost completely ablated and Hes1 expression was not detectable in the intestinal epithelium (supplementary material Fig. S1-A-F, S-T). Therefore, these mutants were used as Hes1 KO mice. Villin-Cre;Hes1flox mice were apparently normal and were used as controls.

Hes1 cKO mice were born at the expected frequency and survived throughout adulthood. On postnatal day (P) 2.5, the crypt-villus structure of the small intestine was apparently normal (Fig. 1-A,B) and the numbers of proliferating cells (Ki67+) and absorptive cells (villin+) were unchanged in Hes1 KO mice compared with controls (Fig. 1-C-F). However, the numbers of apoptotic cells (cleaved caspase 3), goblet cells (Alcian Blue+) and enteroendocrine cells (chromogranin+) were noticeably increased in Hes1 KO mice (Fig. 1-G-L, O-Q). Normally, Paneth cells (lysozyme) appear 2 weeks after birth, and they were not detectable in control mice at this stage (Fig. 1-M). In Hes1 cKO mice, Paneth cells were largely absent at this stage (Fig. 1-N), although a very small number were prematurely differentiated (Suzuki et al., 2005). Thus, Hes1 seems to be important for generation of the correct number of intestinal cells and for their survival. However, at 2 months of age, these defects were not observed in Hes1 cKO mice (supplementary material Fig. S2), suggesting that Hes1 deficiency was mostly compensated in the adult.

Accelerated differentiation of secretory cells in the small intestine of Hes1/333/555 cKO mice at postnatal and adult stages

It was previously shown that, in the absence of Hes1, the related genes Hes3 and Hes5 are upregulated in the embryonic intestine (Jensen et al., 2000). This was confirmed in the postnatal Hes1 cKO mice (supplementary material Fig. S10-R), suggesting that Hes3 and Hes5 compensate for a Hes1 deficiency. To examine this possibility, we generated Villin-Cre;Hes1flox;Hes3flox–/–;Hes5flox–/– (Hes1/333/555 cKO) mice (supplementary material Fig. S1-G-N). Hes3–/–;Hes5–/– (Hes3/5 KO) mice (Hatakeyama et al., 2004) and Villin-Cre;Hes1flox–/–;Hes3–/–;Hes5–/– mice were apparently normal and were used as controls. Hes1 is expressed by proliferating cells but not by Paneth cells in controls (supplementary material Fig. S1-D, red arrows), whereas Hes5 is normally expressed by Paneth cells (van Es et al., 2005a). However, Hes1 was expressed by Paneth cells in Hes3/5 KO mice (supplementary material Fig. S1-H, red arrows), suggesting that Hes1 compensates for Hes3 and Hes5 deficiency.

In Hes1/333/555 cKO mice, the histological structure of the small intestine was apparently normal (Fig. 2-A,B) and absorptive cells (villin+) were not affected at P2.5 (Fig. 2-C,D). However, the numbers of goblet cells (Alcian Blue+) and enteroendocrine cells (chromogranin+) were increased in Hes1/333/555 KO mice at P2.5 (Fig. 2-E-H, P-Q), as observed in Hes1 cKO mice. Furthermore,
Paneth cells (lysozyme+) were prematurely differentiated in Hes1/3/5 cKO mice at P2.5, whereas they were not present in controls (Fig. 2J,R). Interestingly, some of these prematurely differentiated Paneth cells did not remain at the intervillus pocket but moved upwards along the crypt-villus axis (Fig. 2J). Furthermore, the number of apoptotic cells was increased in Hes1/3/5 cKO mice (Fig. 2K,L, arrows, Fig. 2S). Thus, Hes genes are required for the proper differentiation of secretory cells and for the survival of intestinal cells.

Proliferating cells were present at the intervillus pocket as well as in the upper region in controls, whereas they were absent from the intervillus pocket in Hes1/3/5 cKO mice (Fig. 2M,N). Quantification indicated that the number of proliferating cells at P2.5 was reduced in Hes1/3/5 cKO mice compared with controls (Fig. 2O). Similar defects were observed in Hes1/3/5 cKO mice at 2 months of age (supplementary material Fig. S3). At this age, crypt base columnar (CBC) cells, which constitute actively cycling Ki67+ Lgr5+ stem cells (Barker et al., 2007), resided...
Fig. 2. Defects in the small intestine of Hes1/3/5 cKO mice at P2.5. The small intestines of Hes1/3/5 cKO and control mice were examined at P2.5. (A,B) Hematoxylin-Eosin staining showing that the gross structure of the small intestine of Hes1/3/5 cKO mice (B) was unaffected compared with controls (A). (C,D) The development of villin+ absorptive cells was not disturbed in Hes1/3/5 cKO mice. (E-H) In Hes1/3/5 cKO mice, the numbers of goblet cells (E,F) and enteroendocrine cells (G,H) were increased. (I,J) In control mice, Paneth cells were not observed (A,I), but in Hes1/3/5 cKO mice Paneth cells were present (B,J, arrows). Some Paneth cells did not remain in the intervillus pockets but migrated towards the top of the villi. (K,L) The number of apoptotic cells (arrows) was increased in Hes1/3/5 cKO mice. (M,N) The number of Ki67+ proliferating cells was decreased in Hes1/3/5 cKO mice, and these cells were absent from the intervillus pocket (N), although they were present at the intervillus pocket in control mice (M). Scale bars: 100 μm. (O) The number of proliferating cells per intervillus region was reduced in Hes1/3/5 cKO mice. (P-S) Quantification of the number of goblet (P), enteroendocrine (Q), Paneth (R) and apoptotic (S) cells. Error bars indicate s.d.
between Paneth cells at the crypt bottom in controls (supplementary material Fig. S3I,K, black and red arrows). In Hes1/3/5 cKO mice, however, CBC cells were not present at the crypt bottom, but were located away from this region, although their number was not noticeably changed (supplementary material Fig. S3J,L,N, arrows). These results suggest that, in the absence of Hes genes, cell proliferation is reduced, although stem cells are well maintained.

The above data also indicate that whereas differentiation of secretory cells was exclusively inhibited by Hes1 at a neonatal stage, it was cooperatively inhibited by Hes1, Hes3 and Hes5 in order to maintain adult homeostasis. It has been shown that differentiation of secretory cells is regulated by Math1 and Ngn3 (Yang et al., 2001; Jenny et al., 2002), so we examined Math1 and Ngn3 expression in Hes1/3/5 cKO mice. Immunohistochemical analysis showed that Math1 and Ngn3 expression was upregulated in the small intestine of Hes1/3/5 cKO mice (supplementary material Fig. S4, black and red arrows). Thus, we conclude that Hes genes inhibit the differentiation of secretory cells by repressing Math1 and Ngn3 expression.

In control mice, Paneth cells remained at the crypt bottom and expressed nuclear β-catenin, an active Wnt signaling effector (supplementary material Fig. S5C,D, arrows). These cells also expressed EphB3, a downstream target of Wnt signaling, which determines the position of Paneth cells (supplementary material Fig. S5E, arrows) (Batlle et al., 2002). In Hes1/3/5 cKO adult mice, however, many Paneth cells moved away from the crypt bottom (supplementary material Fig. S5F, red arrows), and these mispositioned Paneth cells expressed neither nuclear β-catenin nor EphB3 (supplementary material Fig. S5F-H, red arrows), suggesting that these Paneth cells failed to undergo Wnt-dependent maturation.

Abnormal structures and enhanced secretory cell differentiation in the large intestine of Hes mutant mice

We next examined the large intestine of Hes1 cKO and Hes1/3/5 cKO mice. Compared with control mice, the crypts of the large intestine of Hes1 cKO mice were greatly expanded, and proliferating cells were mispositioned in Hes1 cKO mice at P16 (supplementary material Fig. S6A-D). Furthermore, although Paneth cells (lysozyme+) were not observed in the large intestine of control mice (I), but many were present in Hes1/3/5 cKO mice (J, arrow in inset). Scale bars: 100 μm. (O,P) Quantification of the number of enteroendocrine (O) and Paneth (P) cells. Error bars indicate s.d.
expressed nuclear β-catenin similarly in control and Hes1/3/5 cKO mice (Fig. 3K,L, arrowheads in insets). However, the number of enteroendocrine cells was increased in Hes1/3/5 cKO mice (Fig. 3G,H,O). Furthermore, Paneth cells were formed ectopically in Hes1/3/5 cKO mice at 2 months of age (Fig. 3I,J, arrows, Fig. 3P).

**Abnormal cell proliferation and migration in the large intestine of Hes mutant mice**

We next examined the proliferation and migration of cells of the large intestine in mutant and control mice at 2 months of age by administration of BrdU and performing immunohistochemistry for Ki67. Two hours after BrdU administration, all BrdU+ cells were present within the bottom two-thirds of the crypt in controls (Fig. 4A), and Ki67+ cells were also found in the same region (Fig. 4B,F,J, arrows), indicating that cell proliferation occurs within the bottom two-thirds of the crypt. Lgr5+ stem cells also resided at the crypt bottom (supplementary material Fig. S7A, arrows). After 48 hours, BrdU+ cells that had migrated to the luminal side disappeared, whereas those that had migrated to the bottom side still remained in crypts (K, arrows). (M,N) The turnover of differentiated cells of control (M) and Hes1/3/5 cKO (N) mice was examined by multiple BrdU injections. Nine days after multiple BrdU injections, no BrdU+ cells were detectable in the large intestine of control mice (M), whereas many BrdU+ cells were observed in the cystic crypt bottom (N, below dotted red line, arrows in yellow inset) but not in the luminal side (arrows in blue inset). Scale bars: 100 μm. (O) Quantification of the number of proliferating cells. Error bars indicate s.d.
Large intestine at one year of age

**Control**

**Hes1/3/5 cKO**

**Control**

**Hes1/3/5 cKO**

4O). Lgr5+ stem cells were also observed in the middle region of the crypts in Hes1/3/5 cKO mice, although the number of Lgr5+ stem cells was not noticeably different from that in control mice (supplementary material Fig. S7B, arrow). In these mutant mice, BrdU+ cells migrated not only towards the luminal side but also towards the bottom (Fig. 4G, arrows). Moreover, 96 hours later, BrdU+ cells that migrated upwards had disappeared, whereas those that migrated to the bottom still remained in the crypts (Fig. 4K, arrows). Apoptosis (cleaved caspase 3+ cells) occurred only at the flat surface of the large intestine in control mice (supplementary material Fig. S7C, arrows), but occurred at both the flat surface and the crypt base region (red arrows) in the large intestine of cKO mice (supplementary material Fig. S7D, arrow). In these mutant mice, survival of differentiated cells is enhanced in the large intestine of Hes1/3/5 cKO mice.

**Progressive alteration in the large intestine of Hes1/3/5 cKO mice**

In the large intestine of 1-year-old Hes1/3/5 cKO mice, the number of villin+ absorptive cells was noticeably increased and they expanded into the luminal side, forming villus-like structures with a morphology reminiscent of the small intestine (Fig. 4M, bottom two-thirds of the crypts, L, arrows). Proliferating cells were present within the bottom two-thirds of the crypts (K, arrows), whereas in Hes1/3/5 cKO mice they were present around the middle region of the crypts (L, arrows). Immunohistochemical analysis for β-catenin, phosphorylated Smad1/5/8 and Sox9. All three were similarly expressed (arrows and arrowheads) in Hes1/3/5 cKO mice and controls. Boxed regions are enlarged to the right. Scale bars: 100 μm.

**Fig. 5.** Defects in the large intestine of Hes1/3/5 cKO mice at 1 year of age. The large intestines of Hes1/3/5 cKO and control mice were examined at 1 year of age. (A,B) Hematoxylin-Eosin staining. Crypts of the large intestine were greatly expanded and formed villus-like structures in Hes1/3/5 cKO mice (B) as compared with controls (A). (C,D) Absorptive cells (villin+) were present at the flat surface in controls (C, above dotted line). However, in Hes1/3/5 cKO mice, the absorptive cell population was remarkably expanded in the inner half of crypts (D, above dotted line, arrows). (E,F) Goblet cells (Alcian Blue+) formed expanded crypts in their bottom region (below dashed line). (G,H) The number of enteroendocrine cells (chromogranin+) was increased mainly in the luminal half of the crypts of Hes1/3/5 cKO mice (H, arrows), as compared with the control (G). (I,J) Paneth cells (lysozyme+) were not present in the large intestine of control mice (I), but were abundant in Hes1/3/5 cKO mice (J). Proliferating cells were present within the bottom half of the crypts of Hes1/3/5 cKO mice (J). Goblet cells (Alcian Blue+) were also noticeably increased in number, but they were present in the bottom side of Hes1/3/5 cKO mice (Fig. 5E,F). Enteroendocrine cells (chromogranin+) were overproduced in the luminal side, and Paneth cells (lysozyme+) were ectopically present in the bottom side of the large intestine of Hes1/3/5 cKO mice (Fig. 5G-J). Proliferating cells were present in the bottom half of the large intestine of control mice (Fig. 5K, arrows), whereas they were mispositioned around the middle region of the crypt in Hes1/3/5 cKO mice (Fig. 5L, arrows), as observed at 2 months of age. These results suggest that the large intestine of Hes1/3/5 cKO mice was progressively altered in morphology and became somewhat similar to small intestine, although their detailed structures were different.

**DEVELOPMENT**
Because the distribution of differentiated epithelial cells was altered in Hes1/3/5 cKO mice, we analyzed the Wnt and BMP signaling pathways, which antagonistically regulate the lineage commitment of intestinal stem cells (Scoville et al., 2008). The distributions of nuclear β-catenin, a Wnt signaling effector, and of phosphorylated Smad1/5/8, a BMP signaling effector, were very similar in control and Hes1/3/5 cKO mice (Fig. 5M,N, arrows; Fig. 5O,P, arrowheads), suggesting that no significant alteration of the Wnt and BMP pathways had occurred in the mutants. Because the large intestine also shows the small intestine-like morphology in the absence of Sox9 (Bastide et al., 2007), we examined Sox9 expression. Sox9 was similarly expressed in the bottom regions of both control and Hes1/3/5 cKO mice at 2 months (Fig. 5M,N, arrows, and supplementary material Fig. S5A,B) and 1 year (Fig. 5Q,R, arrows) of age. However, some of the mispositioned proliferating cells in Hes1/3/5 cKO mice, which were located at the luminal side of the large intestine (Fig. 5L), seemed to lose Sox9 expression, suggesting that these Sox9-negative proliferating cells might contribute to the small intestine-like morphology.

Decreased proliferation and increased differentiation of tumor cells in the absence of Hes1
Finally, to determine the role of Hes1 in intestinal tumor development, we crossed ApcMin mice with Hes1 cKO mice. Tumors developed in the intestine of ApcMin mice by loss-of-function mutation of Apc (supplementary material Fig. S8A-E), and nuclear β-catenin was stabilized in these tumor cells (Fig. 6A,J,M,P). In ApcMin;Hes1 cKO mice (Fig. 6E,F and supplementary material Fig. S8F,G), tumors also developed and nuclear β-catenin was stabilized in such tumor cells (Fig. 6B,K,N). However, whereas many tumor cells proliferated (Ki67+) in ApcMin mice (Fig. 6G and supplementary material Fig. S8J), only subsets of them were positive for Ki67 in ApcMin;Hes1 cKO mice (Fig. 6H and supplementary material Fig. S8K), indicating that many tumor cells exited the cell cycle in the absence of Hes1. Furthermore, there were many differentiated goblet cells, enteroendocrine cells, absorptive cells and Paneth cells
in the tumor regions of Apc\textsuperscript{Min};Hes1 cKO mice (Fig. 6Q,T,W,Z, arrows, and supplementary material Fig. S8I) as observed in non-tumor regions (Fig. 6R,U,X,AA), whereas there were virtually no such differentiated cells in tumors of Apc\textsuperscript{Min} mice (Fig. 6P,S,V,Y and supplementary material Fig. S8H). Math1 and Ngn3, which induce secretory and enteroendocrine cell development, respectively, were also upregulated in the tumor regions of Apc\textsuperscript{Min};Hes1 cKO mice as compared with Apc\textsuperscript{Min} mice (supplementary material Fig. S9). These results suggest that, in the absence of Hes1, tumor cells do not continue to proliferate but instead differentiate into postmitotic intestinal cells.

To clarify this idea, we examined the proliferation activity of tumor cells 24 hours after BrdU administration. In Apc\textsuperscript{Min} mice, almost all BrdU\textsuperscript{+} tumor cells were positive for Ki67 even 24 hours after BrdU administration (Fig. 7A,E,I), suggesting that these tumor cells continued to proliferate. By contrast, in Apc\textsuperscript{Min};Hes1 cKO mice, the numbers of BrdU\textsuperscript{+} or Ki67\textsuperscript{+} tumor cells were almost all BrdU\textsuperscript{+} tumor cells did not express Ki67 (supplementary material Fig. S10F). However, in Apc\textsuperscript{Min};Hes1 cKO mice, the numbers of BrdU\textsuperscript{+} or Ki67\textsuperscript{+} tumor cells continued to proliferate. By contrast, most of the BrdU\textsuperscript{+} cells in Apc\textsuperscript{Min};Hes1 cKO mice did not express Ki67 (F,G,J,K), indicating that many tumor cells had exited the cell cycle. Note that Ki67\textsuperscript{+} (G, arrows) and BrdU\textsuperscript{+} (K, arrows) cells were mostly distinct from each other. Scale bars: 100 \( \mu \text{m} \).

**DISCUSSION**

**Hes genes promote cell proliferation and negatively regulate intestinal secretory cell development**

In this study, we examined the roles of Hes genes in the regulation of intestinal development and homeostasis by analyzing Hes KO mice. In previous reports, the small intestine of Hes1 KO neonatal mice showed overproduction of secretory cells (Jensen et al., 2000; Suzuki et al., 2005). However, it was unclear whether Hes1 is similarly important for regulating postnatal development and homeostasis of intestinal cells, as Hes1 KO mice die soon after birth. It was previously shown that, in adult Rhpph cKO mice and adult wild-type mice treated with a \( \gamma \)-secretase inhibitor, all proliferating cells exit the cell cycle and differentiate into goblet cells (van Es et al., 2005a; Riccio et al., 2008). We found that Hes1 cKO mice only displayed an increase in the number of secretory cells during a neonatal stage, whereas Hes1/3/5 cKO mice displayed a reduction in the number of proliferating cells as well as an increase in the number of secretory cells. In these mutant mice, in addition to goblet cells, the number of enteroendocrine cells was noticeably increased. Furthermore, Paneth cells were observed in neither the neonatal small intestine nor the large intestine, whereas they appeared in these regions of Hes1/3/5 cKO.
Hes genes keep stem cells in the correct position on the crypt-villus axis

Intestinal stem cells reside at the crypt bottom (Barker et al., 2007; Barker et al., 2008; Sangiorgi and Capecchi, 2008); stem cells exist between and just above Paneth cells in the small intestine and between progenitors of goblet cells in the large intestine. In Hes1/3/5 cKO mice, proliferating cells were reduced in number and their position was changed. This defect was particularly evident in the large intestine of Hes1/3/5 cKO mice, in which the lower parts of the crypts were gradually expanded. Although normal progeny cells only migrated towards the lumen surface, the progeny of mutant stem/progenitor cells moved along the crypt-villus axis not only towards the surface lumen but also towards the crypt bottom in the large intestine. In addition, the cells that migrated abnormally to the crypt base displayed a lower turnover than normal intestinal cells, which probably led to expansion of the mutant crypts. It is likely that the abnormal migration and expansion of crypts lead to the mislocalization of stem/progenitor cells.

Mislocalization of stem/progenitor cells could in turn lead to abnormal cell differentiation. In the intestine, signaling factors such as Wnt and BMP form gradients along the crypt-villus axis to regulate stem/progenitor cell differentiation (Radtke and Clevers, 2005; Crosnier et al., 2006; Scoville et al., 2008). Wnt signaling activity is highest at the bottom of crypts and gradually decreases along the crypt-villus axis, whereas the activity of BMP signaling shows the opposite gradient (Kosinski et al., 2007; Scoville et al., 2008), and BMP signaling and Wnt signaling antagonize each other (He et al., 2004). Normally, Paneth cells reside at the crypt bottom, receive Wnt signaling and express EphB3, a downstream target for Wnt signaling, which keeps Paneth cells at the crypt bottom (Batlle et al., 2002). In the small intestine of Hes1/3/5 cKO mice, however, some Paneth cells did not remain at the crypt bottom but migrated towards the luminal side along the crypt-villus axis. Transgenic mice expressing the secreted Wnt inhibitor Dkk1 (Villin-Dkk1), and intestine-specific cKO mice of the Wnt receptor frizzled 5 (Fzd5), display mislocalization of Paneth cells resulting from a decrease in Wnt signaling activity (Pinto et al., 2003; van Es et al., 2005b). In Hes1/3/5 cKO mice, Paneth cells remaining at the crypt bottom showed nuclear β-catenin expression and also expressed EphB3, whereas mispositioned Paneth cells showed neither nuclear β-catenin staining nor EphB3 expression, similar to the defects observed in Villin-Dkk1 and Fzd5 cKO mice. It is likely that some differentiating Paneth cells cannot receive proper Wnt signaling owing to changes in the position of the stem/progenitor cells in Hes1/3/5 cKO mice and consequently lose EphB3 expression, which further enhances the mislocalization of Paneth cells.

The large intestine also showed abnormal cell differentiation in Hes1/3/5 cKO mice 1 year after birth. In the mutant large intestine, stem cells were located around the middle region of the crypt.
Absorptive cells and enteroendocrine cells were distributed above the stem cells, whereas goblet cells and Paneth cells were located below them. In the adult intestine, Wnt signaling regulates the development of goblet cells and Paneth cells via Sox9 (Ireland et al., 2004; Bastide et al., 2007; Mori-Akiyama et al., 2007; Wang et al., 2007), whereas BMP signaling mainly controls the development of enteroendocrine cells (Auclair et al., 2007). Therefore, it is likely that, in the large intestine of Hes1/3/5 cKO mice, progenitors migrating to the lumen lose Sox9 expression and differentiate into enteroendocrine cells as well as absorptive cells, whereas secretory progenitors moving to the crypt base differentiate into Paneth cells or goblet cells under the control of Wnt signaling via Sox9.

Thus, Hes genes not only control the size of the proliferating cell population, but also keep stem cells in the correct position on the crypt-villus axis, thereby allowing stem cells to receive signaling molecules and to differentiate properly into epithelial cells.

**Hes1 is required for maintenance of the undifferentiated state of tumor cells**

We also showed that Hes1 plays an important role in maintaining the undifferentiated state of intestinal tumor cells. In the absence of Hes1, many tumor cells exited the cell cycle and differentiated into postmitotic epithelial cells, even though these cells showed stabilization of nuclear β-catenin expression. A recent study showed that Hes1 expression is directly regulated by Notch and Wnt/β-catenin signaling (Peignon et al., 2011), suggesting that these pathways converge on Hes1.

It was previously shown that inhibition of Notch signaling by a γ-secretase inhibitor induced differentiation of adenoma cells into goblet cells, suggesting that Notch signaling might be a promising target for colorectal cancer therapy. However, γ-secretase inhibitors also caused all proliferating cells within the crypts of healthy intestinal regions to differentiate into goblet cells throughout the small and large intestines (Wong et al., 2004; van Es et al., 2005a), which could cause severe diarrhea and malabsorption. By contrast, inactivation of Hes1 had no apparent effect on cell proliferation and differentiation in the crypts of non-tumor regions, probably because other factors compensate for Hes1 deficiency. Thus, Hes1 would be a better target to induce the differentiation of tumor cells. Recently, selective anti-Notch receptor antibodies were developed and shown to inhibit the growth of human colon cancer cells in xenotransplantation models (Wu et al., 2010). However, it remains to be investigated whether each Notch-specific antibody can repress tumor growth in Apcmin mice, as Notch1 and Notch2 are functionally redundant (Riccio et al., 2008) and because Hes1 expression remains after treatment with Notch1- or Notch2-specific antibody in the adult intestine (Wu et al., 2010).

Our study revealed the roles of Hes genes in the regulation of intestinal development and homeostasis. Hes genes control not only the proliferation and differentiation of stem/progenitor cells but also their positions, so that these cells properly receive and respond to signaling molecules. We also showed that inactivation of Hes1 alone is sufficient to reduce tumor cell proliferation and induce the differentiation of tumor cells into all types of intestinal epithelial cells. Importantly, inactivation of Hes1 does not affect the proliferation or differentiation of cells within normal crypts in adult animals, raising the possibility that Hes1 might be a good target for a therapy that aims to induce tumor cell differentiation.

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**Competing interests statement**

The authors declare no competing financial interests.

**Supplementary material**

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