

aggregate was DiI-negative, although some single CM-DiI<sup>-</sup>/Kit<sup>+</sup> cells were observed (data not shown). It is possible that not all allantoic cells were CM-DiI-tagged and that non-tagged cells gave rise to single Kit<sup>+</sup> cells, given the technical difficulty of tagging all cells. We injected CM-DiI into the basal part of allantois, implying that HC clusters are originated from this part. It is also possible that chorionic cells per se may give rise to Kit<sup>+</sup> cells: chorion reportedly has a potential to generate myeloid and definitive erythroid cells (Corbel et al., 2007; Zeigler et al., 2006). Thus, although some HC clusters may have been derived from chorion, it is more likely that the mouse placenta does autonomously generate HSCs and that the allantois is at least a major source of placental HSCs. As shown in Fig. 4A, HC clusters first form cell aggregates. Although several reports suggest that HC clusters in the AGM region are derived from endothelial cells expressing VE-cadherin (Dzierzak and Speck, 2008), the HC clusters in the placenta probably did not originate from endothelial cells. Interestingly, Fraser et al. demonstrated that VE-cadherin is also expressed in HC clusters in the AGM region, indicating that VE-cadherin is not a specific marker of endothelial cells (Fraser et al., 2003). It may be further necessary to evaluate the origin of HC clusters both in the AGM region and the placenta in the future.

### Niche regulation of placental HSCs

HSCs are regulated by niche cells surrounding HSCs. However, it remains unclear how embryonic HSCs are regulated by niche cells. In the bone marrow, expression of niche cell markers such as N-cadherin and CXCL12 enables their isolation by flow cytometry and has contributed greatly to an understanding of niche regulation (Arai and Suda, 2007; Sugiyama et al., 2006). Conversely, investigation of the placental niche has been impeded by a lack of markers for placental niche cells. To address this issue, we isolated niche cells surrounding HC clusters in placenta by LCM. Using this system, we obtained niche cells despite the lack of markers. HC clusters were found inside blood vessels, suggesting that niche cells are mostly composed of endothelial cells. In addition, we sorted out both endothelial and mesenchymal cells, and performed real-time PCR with SCF gene (Fig. 6). Our gene expression analysis revealed that SCF is predominantly expressed in niche cells, and protein expression analysis suggested that SCF is predominantly expressed in niche endothelial cells. In agreement, we found that SCF is predominantly expressed in endothelial cells, in particular cells surrounding HC clusters by immunostaining. Interestingly, SCF was expressed in clusters as well as in endothelial cells, implying an autocrine mechanism. It would be of interest to investigate whether SCF plays a role in specification as well as niche development. To understand the role of the SCF/Kit signal in regulating placental HSCs, we performed a loss-of-function experiment in vivo to inhibit SCF/Kit signaling in the mouse placenta using a WEC system with 10.25 dpc embryos – a stage suitable for manipulation. SCL is not required for HSC development once commitment to hematopoietic lineages has occurred (D'Souza et al., 2005; Mikkola et al., 2003b; Robb et al., 1995; Shivdasani et al., 1995). However, Gata2 is crucial for definitive hematopoiesis and functions in the generation and expansion of HSCs in the AGM region (Ling et al., 2004; Lugus et al., 2007; Tsai et al., 1994). Our study confirmed that expression of *Runx1*, *Myb* and *Gata2* was significantly downregulated compared with control samples in Kit loss-of-function analyses but *SCL* expression was not altered. Kit receptor activation plays a major role in regulating survival, proliferation and self-renewal of HSC phenotypes (Kent et al., 2008), but how SCF/Kit signal regulates

*Runx1*, *Myb* and *Gata2* remains unclear. In addition to SCF/Kit signaling, other signals may regulate HC clusters. SCF secreted by niche cells may modulate proliferation of CD31<sup>+</sup>/CD34<sup>+</sup>/Kit<sup>+</sup> cells between 10.5 and 12.5 dpc, as shown in Fig. 5, although this proliferation might be due to an accumulation of the hematopoietic cells in the placental vasculature as this organ increases in size. Decrease of Ki-67 positive cells might be due to the downregulation of SCF by niche cells.

IL3 reportedly increases the number of HSCs in the AGM region (Robin et al., 2006). However, these authors demonstrated that IL3 has no effect on HSC activity in the placenta at 10.5 dpc, an observation compatible with our data showing that IL3 is not expressed in placental niche cells (Fig. 6A). Hedgehog, BMP4, bFGF and VEGF signals from the surrounding micro-environment are required for mesodermal cells to commit to hematopoietic cells (Dzierzak and Speck, 2008). In the AGM region, location plays a role in regulating HSC generation: ventral tissues induce AGM HSCs, whereas dorsal tissues suppress them (Peeters et al., 2009). Hedgehog protein(s) have been identified as positive effectors that increase the number of AGM HSCs (Peeters et al., 2009). Moreover, there is greater expansion of placental HSCs from 11.5 dpc to 12.5 dpc than of hematopoietic progenitors at this site, suggesting that other signals in the placental niche probably inhibit HSC differentiation (Gekas et al., 2005).

This is the first report to identify and examine the function of cytokine signals regulating HSCs in the mouse placenta. Our study is also evidence that LCM is a useful tool with which to study molecular mechanisms in specific cell aggregates. Recently, it was demonstrated that human placenta contains HSCs and that stromal cells (derived from human placenta) could support hematopoiesis (Robin et al., 2009). Clarifying how the niche regulates HSCs in the placenta could lead to an understanding of how to manipulate HSC generation from ES/iPS cells and, thus, be applicable to future clinical applications.

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### Competing interests statement

The authors declare no competing financial interests.

### Supplementary material

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