c-myc in the hematopoietic lineage is crucial for its angiogenic function in the mouse embryo

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The c-myc proto-oncogene, which is crucial for the progression of many human cancers, has been implicated in key cellular processes in diverse cell types, including endothelial cells that line the blood vessels and are critical for angiogenesis. The de novo differentiation of endothelial cells is known as vasculogenesis, whereas the growth of new blood vessels from pre-existing vessels is known as angiogenesis. To ascertain the function of c-myc in vascular development, we deleted c-myc in selected cell lineages. Embryos lacking c-myc in endothelial and hematopoietic lineages phenocopied those lacking c-myc in the entire embryo proper. At embryonic day (E) 10.5, both mutant embryos were grossly normal, had initiated primitive hematopoiesis, and both survived until E11.5-12.5, longer than the complete null. However, they progressively developed defective hematopoiesis and angiogenesis. The vast majority of embryos lacking c-myc specifically in hematopoietic cells phenocopied those lacking c-myc in endothelial and hematopoietic lineages, with impaired definitive hematopoiesis as well as angiogenic remodeling. c-myc is required for embryonic hematopoietic stem cell differentiation, through a cell-autonomous mechanism. Surprisingly, c-myc is not required for vasculogenesis in the embryo. c-myc deletion in endothelial cells does not abrogate endothelial proliferation, survival, migration or capillary formation. Embryos lacking c-myc in a majority of endothelial cells can survive beyond E12.5. Our findings reveal that hematopoiesis is a major function of c-myc in embryos and support the notion that c-myc functions in selected cell lineages rather than in a ubiquitous manner in mammalian development.

KEY WORDS: c-myc, Angiogenesis, Vasculogenesis, Mouse, Hematopoiesis, Myc, Vascular development

INTRODUCTION

The c-myc (Myc – Mouse Genome Informatics) proto-oncogene encodes a basic helix-loop-helix/leucine zipper transcription factor that is short-lived but rapidly induced upon serum stimulation. Its deregulation is associated with a wide range of human cancers (Adhikary and Eilers, 2005; Evan et al., 2005; Grandori et al., 2000; Pelengaris et al., 2002), and its overexpression is highly tumorigenic in many types of tissues in animals. A large body of work performed in cultured cells shows that c-Myc is expressed broadly and functions as a central regulator of normal cellular programs, including cell proliferation, differentiation, growth, survival and migration in many cell types (Grandori et al., 2000). Although these studies have made tremendous contributions to our understanding of the cellular mechanisms underlying the oncogenic effects of c-Myc, the physiological function of this protein remains largely unknown.

c-myc-null embryos exhibit severe developmental abnormalities in a wide range of organs and die early in gestation before 10.5 days post coitum (dpc) (Davis et al., 1993; Trumpp et al., 2001), supporting the notion that c-myc is essential for a broad range of organ development. Recent studies suggest that c-Myc is required for the proliferation of progenitor cells and the self-renewal of stem cells (Murphy et al., 2005). In the intestine, c-Myc is expressed in the proliferative zone of intestinal crypts, where putative intestinal stem cells reside, and is essential for the formation of these crypts (Bettess et al., 2005; Muncan et al., 2006). Similarly, in the skin epidermis, c-Myc is expressed in the proliferative basal layer and bulge region, where stem and progenitor cells are located (Bull et al., 2001). These new in vivo findings suggest that c-myc is uniquely required in the stem and progenitor cell compartments.

The role of c-myc in the development of the vascular system is of particular interest because it is crucial not only for all aspects of normal tissue function but also for pathological tumor growth and survival. Endothelial cells (ECs) line blood vessels and are the primary cell type responsible for blood vessel function and regeneration. Hematopoietic cells (HCs) give rise to the blood cells of the circulatory system. Differentiation of these two lineages first occurs in yolk sac blood islands, where ECs and HCs may arise from a common mesoderm-derived precursor: the hemangioblast (Cumano and Godin, 2007; Eichmann et al., 2002; Ema and Rossant, 2003). During vascular morphogenesis, ECs coalesce to assemble a primitive vascular network composed of a capillary plexus with uniform caliber and honeycomb appearance. This formation of blood vessels by de novo EC differentiation is known as vasculogenesis (Adams and Alitalo, 2007; Carmeliet, 2005). The primitive capillary plexus subsequently undergoes growth and remodeling to shape the mature vascular tree. Angiogenesis is the process of new blood vessel growth from existing vessels (Folkman, 2006; Hanahan and Folkman, 1996; Thurston et al., 2007). c-Myc has been shown to regulate angiogenesis by promoting the expression of pro-angiogenic factors such as VEGF in stromal cells.
while inhibiting the expression of the anti-angiogenic factor thrombospondin 1 (Baudino et al., 2002; Dews et al., 2006; Knies-Bamforth et al., 2004; Mezquita et al., 2005; Shchors et al., 2006; Watnick et al., 2003). c-Myc is reportedly required for vasculogenesis during development, as c-myc-null embryos have no detectable blood vessels (Baudino et al., 2002). However, whether c-Myc in ECs plays an essential role in vasculogenesis or angiogenesis is currently unknown.

To ascertain the cell type-specific role of c-Myc during vascular development, we generated conditional knockouts (CKs) of the c-myc gene in c-myc<sup>fl<sub>ox</sub></sup> mice using cell lineage-specific Cre lines. We were surprised to find that c-myc was not required for vasculogenesis, and that deleting c-myc in a majority of ECs was compatible with early embryo survival. By contrast, c-myc deletion was detrimental to hematopoietic lineages during development, and c-myc deletion in these lineages was sufficient to cause vascular developmental defects.

**MATERIALS AND METHODS**

**Mice**

Tie2-Cre, Tie1-Cre, Sox2-Cre, Vav-Cre, Tie1-GFP, c-myc<sup>fl<sub>ox</sub></sup> and c-myc<sup>−/−</sup> mice were previously described (Braren et al., 2006; de Boer et al., 2003; Gustafsson et al., 2001; Hayashi et al., 2002; Iijin et al., 2002; Trumpf et al., 2001). All animals were treated in accordance with the guidelines of the University of California San Francisco (UCSF) Institutional Animal Care and Use Committee.

**Materials**

Details of antibodies and real-time PCR primers used can be provided on request.

**Imaging of embryos, whole-mount immunofluorescence staining, and EC isolation and culture**

Previously established procedures (Braren et al., 2006) were followed.

**Whole-mount lacZ staining**

lacZ staining of embryos and yolk sacs was as previously described (Carpenter et al., 2005). Samples were fixed in 4% paraformaldehyde (PFA) overnight after lacZ staining. Specimens were then embedded in paraffin and sectioned at 5 μm. Sections were stained with Eosin and visualized using a Zeiss Axioskop 2 Plus microscope (Zeiss, Thornwood, NY). Images were captured using a DC 300 camera and IM50 software (Leica, San Jose, CA).

**Cell proliferation and TUNEL assays**

Cell proliferation and apoptosis were evaluated as previously described (Braren et al., 2006). TUNEL ('TdT-mediated dUTP nick-end labeling') was performed according to the manufacturer's instructions (Santa Cruz, Santa Monica, CA). For TUNEL analysis, embryos were fixed in 4% PFA for 2 hours and washed three times in PBS. They were then incubated in 10 μg/ml of terminal deoxynucleotidyl transferase (TdT) and 50 μg/ml of biotinylated dUTP in 0.1x PBS for 90 minutes. The reaction was terminated by washing the embryos three times in PBS. The embryos were then treated with 3% H<sub>2</sub>O<sub>2</sub>-in PBS for 5 minutes to block endogenous peroxidase activity. The embryos were washed in PBS and incubated in 10% normal goat serum in PBS for 1 hour to block non-specific binding. The embryos were then incubated in the primary antibody solution (1:500 in PBS-T-0.5% BSA) at 4°C overnight. After washing to remove unbound antibody, the embryos were incubated in the secondary antibody solution (1:200 in PBS-T-0.5% BSA) at room temperature for 1 hour. The embryos were then washed to remove unbound antibody and incubated in the avidin-biotin-peroxidase complex (ABC) solution (1:500 in PBS-T-0.5% BSA) at room temperature for 1 hour. The embryos were then washed and incubated in the DAB solution (1:100 in PBS-T-0.5% BSA) at room temperature for 10 minutes. The embryos were then washed and mounted with Vectorshield containing DAPI (4′,6-diamidino-2-phenylindole; Vector Laboratories, Burlingame, CA). The slides were then observed using a Zeiss Axioskop 2 Plus microscope equipped with a Sensicam CCD camera and Slidebook software (Intelligent Imaging Innovations, Santa Monica, CA).

**Allantoic explants**

Allantoic explant was performed as described (Braren et al., 2006). Allantois were isolated at E8.0 at the 6- to 8-somite stage and cultured for 24-48 hours in FN-coated dishes. Cultures were stained with anti-CF31 as described above.

**RESULTS**

**c-myc is essential for angiogenesis but not vasculogenesis**

Gross abnormalities and developmental retardation by 9.5 dpc and death by 10.5 dpc have been previously observed in c-myc<sup>−/−</sup> embryos (Davis et al., 1993; Trumpf et al., 2001). Multi-organ failure, including circulatory defects, is believed to underlie this early developmental arrest, although the specific cell lineages affected are largely unknown. We therefore analyzed c-myc<sup>−/−</sup> embryos specifically for vascular defects, using whole-mount immunostaining against CD31, an EC-specific marker. As previously reported, these mutant embryos were developmentally retarded and displayed major developmental defects. However, we were surprised to find that major vessels such as the dorsal aorta had developed by 8.75 dpc. Smaller inter-somitic and cranial vessels had also formed in the embryo proper, although they were underdeveloped compared with wild-type vessels (Fig. 1A,B). In the yolk sac, primitive vascular networks had formed, although the mutant yolk sac contained more vascular plexuses and less organized hierarchical branches than the control (Fig. 1C,D). At 10.25 dpc, although the control vasculature had matured to an elaborate and well-organized system (Fig. 1E,G), the c-myc<sup>−/−</sup> embryos retained a primitive vasculature in the head (Fig. 1F) and the yolk sac (Fig. 1H). Nonetheless, embryo cross-sections revealed that, like controls, c-myc<sup>−/−</sup> embryos had ECs lining their dorsal aortae, common cardinal veins and other blood vessels (Fig. 1I,J). These results indicate that, in contrast to a previous report (Baudino et al., 2002), c-myc<sup>−/−</sup> embryos contained differentiated ECs that were capable of assembling a primitive vasculature. Subsequent vascular remodeling, however, was defective in the mutants.

Because the detection of ECs in the c-myc<sup>−/−</sup> mutant was surprising, we examined the vasculature in another mutant embryo, in which c-myc is deleted in the entire embryo proper but retained in the placenta. We created a CK using the Sox2-Cre mouse line, in which the antibody-cell suspension was incubated on ice for 30 minutes. A BD LSRII FACS Machine (BD Biosciences, San Diego, CA) was used to perform flow-cytometry, and FlowJo software was used for data analysis. Propidium iodide (1 μg/ml) staining was used to exclude dead cells.

**Collection of embryonic peripheral blood cells**

Embryonic peripheral blood (PB) was isolated by opening the embryonic vitelline vessels, dorsal aortae and the heart to release blood cells completely. The cells were passed through a 40 μm nylon mesh before use.

**Immunofluorescence staining of sections and isolated cells**

Rehydrated paraffin sections were blocked with 5% donkey serum in PBS for 2 hours at room temperature. They were then incubated with primary antibody at 4°C overnight, washed three times with PBS, followed by 1 hour secondary antibody incubation at 4°C in blocking solution (2% BSA, 3% normal donkey serum, 0.01% Triton X100, 1xPBS). Samples were washed with PBS and mounted with Vectorshield containing DAPI (4′,6-diamidino-2-phenylindole; Vector Laboratories, Burlingame, CA). Cells grown on six-well plates were fixed in 4% PFA/PBS for 20 minutes and permeabilized in 0.1% Triton X-100/1xPBS/2%BSA for 10 minutes at room temperature before blocking (2% BSA/0.01% of Triton X100/1xPBS). Images were captured using either air lenses or a 63x Anachroplan water immersion lens, and a Zeiss Axiosvert2 Plus microscope equipped with a Sensicam CCD camera and Slidebook software (Intelligent Imaging Innovations, Santa Monica, CA).
which Cre is active in all cells of epiblast origin, including the entire embryo proper, the yolk sac mesoderm, the amnion membrane and the embryonic vessels in the placenta, but not the visceral endoderm or the extra-embryonic ectoderm (Vincent and Robertson, 2003). We verified Cre expression throughout the entire embryo by Cre reporter assay (data not shown). The Sox2-Cre; c-myc\textsuperscript{floxed/–} mutant progeny displayed growth retardation by 11.5 dpc and died between 11.5 and 12.5 dpc. These embryos, unlike the null mutants, appeared grossly normal without major organ defects before E10.5 (data not shown). However, at E11.5 they were anemic (Fig. 1L,N) and displayed abnormal vasculature (Fig. 1P). Anti-CD31 staining revealed abundant ECs in mutant yolk sacs and embryos proper (Fig. 1O-T), demonstrating that the mutant embryos were not defective in EC differentiation, although capillary remodeling was defective. Mutant yolk sac microvessels were more primitive, comprising disorganized capillary plexuses with larger intercapillary spaces than in the controls. Major vitelline vessels were narrower and underdeveloped in the mutant (Fig. 1O,P). Similarly, head capillaries were more primitive. The carotid arteries were less elaborate and the primitive head veins were narrower compared with controls (Fig. 1Q-T). These findings confirmed that c-myc is not required for the onset of vasculogenesis in embryos. However, it is essential for subsequent vascular morphogenesis.

The Sox2-Cre; c-myc\textsuperscript{floxed/–} mutant embryos that maintained c-myc in their visceral endoderm and placenta died 2 days later than c-myc\textsuperscript{–/–} embryos, suggesting that expression of c-myc in these extra-embryonic tissues is essential to the survival of the embryos at this stage. Consistent with this notion, c-myc is expressed at high levels in the ectoplacental cone starting at 6.5 dpc (Downs et al., 1989), and c-Myc promotes trophoblast proliferation (Erlebacher et al., 2004). We found that by 8.25 dpc, c-myc\textsuperscript{–/–} placentas contained fewer trophoblasts, exhibiting defective placentation morphology and cellular composition lacking trophoblast integrity. In addition, the mutant chorionic plate was thinner, and the ectoplacental cone was not integrated (see Fig. S1 in the supplementary material). These findings suggest that c-myc is required for placental development, and that the cellular function of c-myc in this organ requires future investigation. In this report, we have focused on the function of c-myc in the embryo proper.

**Loss of c-myc in endothelial and hematopoietic cells induces similar vascular defects as loss of c-myc in the entire embryo proper**

To determine whether eliminating c-myc specifically from the circulatory system would result in developmental and vascular defects, we crossed c-myc\textsuperscript{floxed/–} mice with Tie2-Cre; c-myc\textsuperscript{–/–} mice,
in which Cre is active in EC and HC lineages starting as early as 7.5 dpc in the common progenitor of these two lineages (Braren et al., 2006). To assess the efficiency of Tie2-Cre-mediated c-myc deletion, we measured nuclear-specific c-Myc expression in isolated ECs by immunostaining (see Fig. S3 in the supplementary material). Quantitative analysis showed that virtually no (0.62%) mutant ECs and 91.2% of control ECs had c-Myc staining by 10.5 dpc (Fig. 2M). These experiments demonstrate the success of c-myc deletion. These embryos, like Sox2-Cre CKs, appeared grossly normal without major organ defects before E10.5 (data not shown). At 9.5 dpc, the Tie2-Cre; c-myc<sup>flox/–</sup> embryos were similar to controls, with all embryos showing blood-filled vasculatures, although some were slightly paler (Fig. 2A,B). However, at 10.5 dpc, the mutants appeared anemic (Fig. 2D). At 11.5 dpc, the mutant yolk sacs and embryos were completely white, whereas control yolk sacs displayed vessels filled with red blood cells (Fig. 2E,F). Mutant embryos were smaller than controls (Fig. 2F). No mutant embryos survived beyond 12.5 dpc (see Table S1 in the supplementary material). The gross abnormalities and the stage at which the phenotype occurred in Tie2-Cre CK closely resembled that of the Sox2-Cre CK embryos. Anti-CD31 staining on 11.5 dpc Tie2-Cre CK embryos also revealed a very similar vascular phenotype to the Sox2-Cre CK, with abundant ECs but more primitive capillaries and smaller major vessels in mutant yolk sacs and heads compared with the controls (Fig. 2G-L). Quantitative analysis of the yolk sac vascular defects is summarized in Fig. S2 (see supplementary material). These findings from Tie2-Cre CK embryos confirm that vasculogenesis occurred without c-myc, but further capillary remodeling was defective. Additionally, the allantois is vascularized by vasculogenesis, and we did not detect any apparent vascular defects in the Tie2-Cre CK allantoic explants (Fig. 3A,B), further suggesting that vasculogenesis occurs in the absence of c-myc.

Expression of N-Myc (Mycn – Mouse Genome Informatics), another Myc family member, driven by the c-myc promoter can functionally replace c-Myc activity in vivo (Malynn et al., 2000). To
a nearly complete absence of blood cells by 11.5 dpc (data not shown). Hemogenic ECs that reside in the ventral side of the dorsal aorta at around 10.5 dpc are thought to give rise to HSCs (de Bruijn et al., 2002; Taoudi and Medvinsky, 2007). We examined serial cross-sections of dorsal aortae from four pairs of embryos at 10.5 dpc. Cells located at the ventral wall of the aorta and morphologically resembling hemogenic ECs were seen in all controls but in none of the mutants (Fig. 3E,F). These findings suggest that removing c-myc from ECs and HCs is sufficient to induce the hematopoietic, angiogenic and survival defects observed in embryos harboring a global c-myc deletion.

**c-myc deletion in HCs is sufficient to induce vascular defects and embryonic lethality**

To delineate the effect of c-myc deletion in hematopoietic lineages on vascular development, we examined embryos in which c-myc had been deleted specifically in HC lineages using Vav-iCre. Vav-iCre has been shown to mediate gene excision in adult HCs (de Boer et al., 2003). We thus analyzed Vav-iCre activity in embryos using a Rosa26R-lacZ reporter according to our established method (Braren et al., 2006). At 11.5 dpc, we found that Vav-iCre was active almost exclusively in fetal liver HCs (Fig. S4A,C in the supplementary material) and in some circulating blood cells in the yolk sac (Fig. S4B in the supplementary material), but not in the endothelium or any other tissues (see Fig. S4D in the supplementary material). This result indicates that Vav-iCre mediated c-myc deletion in HCs without affecting ECs or any other cell types. We assessed Vav-iCre activity in individual embryos by FACS analysis in 11.5 dpc fetal liver HCs using the Rosa26YFP reporter (Srinivas et al., 2001). The fraction of HCs expressing Cre varied among individual embryos in all three HC lineages tested, with averages around 50% (see Fig. S4K in the supplementary material).

About 60% of Vav-iCre;c-myc flox/–; mutants (27/43) appeared anemic at 11.5 dpc (Fig. 4B), with paler and smaller fetal livers compared with controls (see Fig. S4L in the supplementary material). A similar % of mutants died around 12.5dpc, a stage similar to the Tie2-Cre and Sox2-Cre CKs. About 16% of mutants (7/43) showed various degrees of hemorrhaging in the embryo proper (see Fig. S4E,G in the supplementary material). About 21% of mutants (9/43) survived through 12.5 dpc but developed anemia and died by 15.5 dpc. None of the Vav-iCre mutants survived to birth. It is likely that non-uniform Cre activity among individuals resulted in these variable phenotypes (see Fig. S4K in the supplementary material). Nonetheless, deletion of c-myc solely in HCs resulted in complete embryonic lethality of all embryos examined.

To visualize the vasculature of the Vav-iCre;c-myc flox/– mutant embryos, we performed whole-mount anti-CD31 staining on 11.5 dpc embryos. We found that the vascular defects in these anemic embryos resembled those in Tie2-Cre- and Sox2-Cre CKs (see Fig. 1P,R,T, Fig. 2H,I,J,L). Although the overall vascular patterning in Vav-iCre;c-myc flox/– embryos proper and yolk sacs was in place, the vessels were narrower and the vascular network appeared underdeveloped and primitive compared with controls (Fig. 4D,F,H). Taken together, these results suggest that c-myc deletion in HC lineages alone likely accounts for the anemia, embryonic lethality and vascular developmental defects induced by c-myc deletion in ECs and HCs combined.

To quantify the hematopoietic defects in 11.5 dpc Vav-iCre CKs, we performed HC counts in peripheral blood (Fig. 4I) and fetal liver cells (Table 1). We found a several fold decrease of cells (7.6 fold in Ter119+ cells, 4.3 fold in CD45+ and 4.6 fold in CD11b+ cells) in the mutant peripheral blood. The mutants’ fetal livers had cytopenia and
contained 12.5-fold fewer cells than their littermate controls. In addition, we used Lin markers, containing CD3e, CD11b, CD45R/B220, Ter119, Ly-6G and Ly-6C to label the committed hematopoietic lineages, which include T lymphocytes, B lymphocytes, monocytes/macrophages, NK cells, erythrocytes and granulocytes. We found the proportion of committed (Lin+) cells was reduced while that of uncommitted cells (Lin-) was increased in Vav-iCre;c-myc\textsuperscript{flox/-} embryos. Among the Lin- cells, the proportion of KLS-HSCs (Kit\textsuperscript{hi}, Lin\textsuperscript{-}, Sca1\textsuperscript{+}) (Ivanova et al., 2002) in the Vav-iCre;c-myc\textsuperscript{flox/-} fetal liver was increased, while the proportion of Kit\textsuperscript{lo}, Lin- cells was significantly decreased. In summary, these results demonstrate that elimination of c-myc in HCs by Vav-iCre compromised definitive hematopoiesis.

c-myc deletion in the majority of ECs and a subset of HCs results in partial survival

As c-myc deletion in HCs alone using Vav-iCre resulted in defects similar to c-myc deletion from both HCs and ECs (with Sox2-Cre or Tie2-Cre), we investigated the effect of endothelial c-myc expression on vascular development. For this experiment, we needed a Cre line that would remove c-myc in the ECs but not the HCs; however, such a reagent is not currently available. We therefore chose the Tie1-Cre line, with Cre active in the majority of ECs and a subset of HCs by 10.5 dpc (Gustafsson et al., 2001) (see Fig. S5 in the supplementary material). We verified Tie1-Cre activity using the Rosa26R reporter. Cre was active in ~54% of myeloid cells, 26% of erythroid cells and 15% of lymphocyte precursor cells at 11.5 dpc (Fig. S6 in the supplementary material). These results suggest that Tie1-Cre mediated c-myc deletion in the majority of ECs and a subset of HCs.

By 12.5 dpc, we did not detect any defects in Tie1-Cre;c-myc\textsuperscript{flox/-} mutants. By 17.5 dpc we observed anemic, dying mutant embryos. Remarkably, half of the mutants survived to birth and a third to post-weaning (Table 2). Surviving mutant adults appeared normal. These results demonstrate that c-myc deletion in a majority of ECs is compatible with embryo survival.

To examine the HC development in these mutants, we performed HC counts from the peripheral blood (Fig. 5) and fetal livers (Table 3) of 11.5 dpc Tie1-Cre CKs. We found a less than twofold decrease in peripheral blood cell lineages (Ter119\textsuperscript{+}, CD45\textsuperscript{+}, CD11b\textsuperscript{+}) in the mutants compared with controls. *P<0.01 and **P<0.05 by Student's t-test.

c-Myc-depleted ECs show no significant cell biological defects

Most Tie1-Cre CKs examined at 11.5 dpc showed no detectable abnormalities, suggesting that c-myc expression in the majority of ECs is not essential for vascular development. To

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<tr>
<th>CD45\textsuperscript{-}</th>
<th>Ter119\textsuperscript{-}</th>
<th>CD11b\textsuperscript{-}</th>
<th>Lin\textsuperscript{-}</th>
<th>Lin\textsuperscript{hi/-}</th>
<th>KLS within Lin\textsuperscript{-}</th>
<th>Kit\textsuperscript{hi} within Lin\textsuperscript{-}</th>
<th>Kit\textsuperscript{lo} within Lin\textsuperscript{-}</th>
<th>Live cells/ fetal liver</th>
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<tr>
<td>Controls</td>
<td>14.3 (1.1)*</td>
<td>22.5 (3.0)*</td>
<td>3.8 (0.5)*</td>
<td>44.0 (2)*</td>
<td>56.0 (1.9)*</td>
<td>0.08 (0.02)*</td>
<td>19.7 (2.2)*</td>
<td>5.1 (0.4)*</td>
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<tr>
<td>Mutants</td>
<td>3.5 (1.1)*</td>
<td>14.5 (2.4)*</td>
<td>1.8 (0.5)*</td>
<td>25.0 (4.6)*</td>
<td>75.0 (4.7)*</td>
<td>0.2 (0.1)*</td>
<td>5.1 (1.6)*</td>
<td>11.8 (2.8)*</td>
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Table 1. Impaired differentiation and proliferation of hematopoietic cells in Vav-iCre-mediated c-myc deletion in 11.5 dpc fetal livers

Data represent the mean percentage of live cells (as e.m.) and the mean of live cell numbers per fetal liver (as e.m.) from 10 mutant fetuses and 20 control fetuses. Differences between mutant and control fetuses were significant at *P<0.01 and **P<0.05, or not significant at ‡P=0.112. Controls were Vav-iCre;c-myc\textsuperscript{+/-} and c-myc\textsuperscript{flox/-}; mutants were Vav-iCre;c-myc\textsuperscript{flox/-}.
verify whether c-myc is required in ECs for blood vessel formation, we examined the in vivo and in vitro behaviors of Tie2-Cre CK ECs.

We analyzed EC proliferation in vivo by BrdU incorporation and did not observe any significant difference in proliferation rates of aortic endothelium between control and mutant embryos (Fig. 6A). We also did not detect an obvious proliferative difference in isolated ECs from BrdU-labeled embryos following maternal BrdU injection (data not shown).

To determine whether loss of c-myc affected EC survival in vivo, we performed TUNEL assays on cryosections of yolk sacs. We found a significant increase in TUNEL staining in vivo, we performed TUNEL assays on cryosections of yolk (data not shown).

We also did not detect an obvious proliferative difference in isolated c-myc Cre ECs, we isolated ECs from 10.5 dpc embryos and cultured on we performed time-lapse video-microscopy of actin cytoskeleton between control and c-myc Cre ECs, we isolated ECs from 10.5 dpc embryos and cultured on fibronectin. The paths of both mutant and control ECs were random and indistinguishable from one another, as shown by our measurements of net path length and average speed (Fig. 6E,F,G). These results demonstrate that c-myc-null ECs are not defective in cell motility in vitro.

To further address the angiogenic potential of c-myc-null ECs, we tested their ability to form endothelial tubes in vitro (Fig. 6H,I). Because this assay requires a large number of ECs, we isolated ECs from the vena cava of adult c-mycfl/fl mice and removed their c-myc gene using an adenovirus-mediated Cre (AdCreGFP) deletion system. FACS analysis of GFP expression in the cultured ECs showed that the efficiency of adenovirus infection was 98% (Fig. 6J), and PCR analysis of genomic DNA confirmed that the c-myc gene was excised in the majority of the cells (Fig. 6K). However, we found no statistically significant difference in the number of branch points (data not shown) or the lengths of tubes formed by mutant and control ECs on the Matrigel surface (Fig. 6L). These findings suggest that c-myc-null ECs are not defective in cell migration or capillary morphogenesis.

Deletion of c-myc in HCs leads to reduction of proangiogenic factors crucial for vascular morphogenesis

HCs have been increasingly recognized as significantly contributing to angiogenesis by modulating the production of proangiogenic factors (Tordjman et al., 2001; Kopf et al., 2006; Shojaei et al., 2007). We therefore examined the mRNA levels of various proangiogenic factors, including Pdgfa, Mmp2 and Il1b, in whole embryos, using quantitative-PCR analysis. We found that mRNA levels of these genes were significantly decreased in the anemic Vav-iCre; c-mycfl/fl-embryos compared with their control

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<th>Table 2.</th>
<th>Tie1-Cre mediated c-myc deletion resulted in partial lethality</th>
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<tr>
<td>Survival after weaning (48 mice from seven litters)</td>
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<tr>
<td>Ratio</td>
<td>Mutants*</td>
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<tr>
<td>Expected§ 25%</td>
<td>75%</td>
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<tr>
<td>Experimental§ 8%</td>
<td>92%</td>
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<td>Expmt/Expt¶ 32%</td>
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Newborn survival (56 mice from eight litters)

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<th>Ratio</th>
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<th>Controls†</th>
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<tr>
<td>Expected§ 25%</td>
<td>75%</td>
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<tr>
<td>Experimental§ 12.5%</td>
<td>87.5%</td>
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<tr>
<td>Expmt/Expt¶ 50.0%</td>
<td>116.6%</td>
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*Mutants include Tie1-Cre; c-mycfl/fl and Tie1-Cre; c-mycfl/+.
†Controls include c-mycfl/fl, c-mycfl/+, c-myc+/-, c-mycnull.
§Expected indicates the expected Mendelian genetic ratio of genotypes.
¶Expmt/Expt indicates the ratio of experimental over expected.

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<th>Table 3.</th>
<th>Tie1-Cre mediated c-myc deletion led to mild defects in hematopoiesis in 11.5 dpc fetal livers</th>
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<tbody>
<tr>
<td>Controls</td>
<td>Mutants</td>
</tr>
<tr>
<td>CD45*</td>
<td>15.4 (1.3)</td>
</tr>
<tr>
<td>Ter119*</td>
<td>24.0 (4.7)</td>
</tr>
<tr>
<td>CD11b*</td>
<td>4.2 (0.5)</td>
</tr>
<tr>
<td>Lin*</td>
<td>42 (4.6)</td>
</tr>
<tr>
<td>Linhi*</td>
<td>51 (4.6)</td>
</tr>
<tr>
<td>KLS within Lin*</td>
<td>51 (4.6)</td>
</tr>
<tr>
<td>Kitlo within Lin*</td>
<td>51 (4.6)</td>
</tr>
<tr>
<td>Kithi within Lin*</td>
<td>21.0 (2.8)</td>
</tr>
<tr>
<td>Live cells/fetal liver</td>
<td>3.5 (0.3)</td>
</tr>
<tr>
<td>Controls</td>
<td>412,591 (47807)*</td>
</tr>
<tr>
<td>Mutants</td>
<td>195,350 (43767)*</td>
</tr>
</tbody>
</table>

Data represent the mean percentage of live cells (as e.m.) and the mean of live cell numbers per fetal liver (as e.m.) from seven mutant fetuses and 24 control fetuses. Differences between mutant and control fetuses were significant at *P<0.01, †P<0.05. Controls were Tie1-Cre;c-myc+/–, Tie1-Cre;c-mycfl/fl and c-mycnull. Mutants were Tie1-Cre;c-mycfl/fl.
littermates (Fig. 7A). By contrast, levels of VEGF transcripts were dramatically increased, indicating a hypoxic response in the embryos suffering anemia (see Fig. 4A,B). Similarly, elevated VEGF protein was found in Tie2-Cre;c-myc^{floxed} embryos (Fig. 7B). These results demonstrate that c-Myc deficiency-induced hematopoietic defects may have reduced expression of certain proangiogenic factors, thereby hindering normal vascular morphogenesis.

**DISCUSSION**

To ascertain the physiological functions of c-Myc in development, we examined embryos lacking c-myc completely, in the embryo proper but not in extra-embryonic tissues, in hematopoietic and endothelial lineages, and in hematopoietic cells specifically. Our findings demonstrate that c-myc is not required for vasculogenesis in the embryo but can indirectly control angiogenesis through its vital role in hematopoiesis. Deletion of c-myc in HCs alone is lethal and sufficient to elicit both hematopoietic and vascular defects.

**c-Myc plays an essential role in embryonic hematopoiesis**

Fetal hematopoiesis begins with primitive hematopoietic differentiation in blood islands of the yolk sac at 7.5 dpc and lasts until 10.5 dpc in mice. Definitive hematopoiesis, which generates enucleated erythrocytes among other hematopoietic lineages like HSCs, starts at 10.5 dpc in the aorta-gonad-mesonephros region. Concurrently, HSCs colonize the developing fetal liver. The murine placenta also harbors HSCs during midgestation. Around birth, hematopoiesis translocates to the bone marrow (BM). Adult and fetal hematopoiesis differ in the types of niches in which they occur, as well as in the capacity of adult versus fetal HSCs to renew, proliferate and differentiate (Cumano and Godin, 2007; Mikkola and Orkin, 2006; Wilson and Trumpp, 2006). The role of c-Myc in adult hematopoiesis in BM has been reported (Wilson et al., 2004). Our experiments demonstrate the essential function of c-Myc in fetal hematopoiesis.

Our data suggest that HCs undergo primitive differentiation in the absence of c-myc. At 9.5 dpc, prior to definitive hematopoiesis, the majority of Tie2-Cre CKs were indistinguishable from the controls, with blood-filled circulatory systems. This phenotype is in sharp contrast to that of embryos lacking Scl, a gene required for the
The absence of c-Myc in definitive hematopoiesis seems to affect the differentiation of HSCs, resembling c-myc-deficient adult hematopoiesis in BM (Wilson et al., 2004). The proportion of uncommitted HCs, KLS-HSCs and Kitlow progenitor cells increased in the Vav-iCre fetal liver. However, the proportion of Kithi progenitor cells decreased. This reduction could be a genuine decrease of Kithi progenitors in mutants or a loss of cell surface Kit expression on otherwise functional HSCs, as has been observed following myeloid ablation (Randall and Weissman, 1997). However, the proportion of the committed HCs, including Ter119+, CD45+ and CD11b+ lineages, decreased significantly in Vav-iCre CKs. In addition, the total number of isolated fetal liver cells, which are composed primarily of HCs, was decreased. These data suggest that the mutant HSCs and progenitor cells can survive and divide but subsequent differentiation into HCs are defective. Our findings provide crucial evidence that c-Myc is required in a cell-autonomous fashion for HSC differentiation. This finding is complementary to the finding in Sox2-2 Cre CKs (Dubois et al., 2008).

The fact that mutant embryos in which c-myc is deleted in the entire embryo proper but not the placenta survived 2 additional days beyond the survival of the complete null embryo shows that c-myc is essential for placental development. It is currently unknown which specific cell lineage(s) c-Myc may function in and what precise role c-Myc may play in the placenta. However, this finding is intriguing in light of recent discoveries that the placenta is an active site for HSC development (Gekas et al., 2005; Mikkola et al., 2005). Given the crucial function of c-Myc in both fetal and adult hematopoiesis, it is plausible that c-Myc may function in placental HSCs. The c-Myc CKs described here may serve as an excellent model to elucidate the molecular control of placental hematopoiesis.

**c-Myc is not required for embryonic vasculogenesis**

Previous reports suggest that c-myc is required for vasculogenesis (Baudino et al., 2002) and downregulation of c-myc in cultured ECs leads to cellular senescence (Guney and Sedivy, 2006). By contrast, we show here that vasculogenesis occurs in the absence of c-myc. Our method of gene excision is efficient, leading to the deletion of the entire coding region for c-Myc (Trumpp et al., 2001). We therefore respectfully disagree with this earlier conclusion. Gene disruption in the two studies was achieved by targeting a similar region of c-myc, so the reason for the discrepancy between our observations and those of Baudino et al. are currently unclear. However, we confirmed our results in three independent c-myc-deficient mutants (c-myc null, Sox2-2 Cre and Tie2-2 Cre CKs), and the presence of ECs in c-myc deleted embryos was also verified in a different laboratory (Dubois et al., 2008). We conclude that c-myc is not required for vasculogenesis in embryos.

At the cellular level, c-myc-deficient ECs did not exhibit detectable defects in cell proliferation, survival, migration or even capillary morphogenesis. These results are in contrast to the report that c-Myc is essential for EC proliferation in culture (Guney and Sedivy, 2006). We performed proliferation assays using primary ECs to closely mimic in vivo conditions. In addition, neither N- nor L-Myc compensated for the loss of c-Myc. We also show that about one third of Tie1-1 Cre CKs, in which c-Myc was deleted in the majority of ECs, survived into adulthood without apparent abnormalities. This result suggests that widespread deletion of c-myc in the endothelium is compatible with survival. Taken together, our observations suggest that abrogating c-Myc in ECs may not disrupt angiogenesis, and c-Myc likely regulates angiogenesis through a non-cell-autonomous fashion.
Hematopoietic abnormalities caused by c-myc deletion lead to defects in angiogenesis

Although a primitive vascular network formed in the absence of c-myc, its angiogenic remodeling into a complex vascular tree was abnormal. Because c-Myc-deficient ECs appear to function normally, we propose that defective HCs cause the vascular defects observed in our c-myc CK embryos. Supporting this notion, vascular defects in the Vav-iCre CKs, where c-myc is deleted specifically in the hematopoietic lineage, resembled those in the Tie2-Cre CK embryos.

HCs are known to affect angiogenesis through hemodynamic influence (Lucitti et al., 2007) and oxidative stress, such as hypoxia (Jones et al., 2004; Ramirez-Bergeron et al., 2006). The reduced hematocrit in the mutants likely changes the viscosity of the blood, and hence alters the hemodynamic forces required for growth and maintenance of vessel size (Lucitti et al., 2007). Moreover, both Vav-iCre and Tie2-Cre CK embryos were anemic by 11.5 dpc. Hypoxia was evident by elevated VEGF levels, a common consequence of embryos in hypoxic conditions. Hypoxia causes pan-tissue damage via apoptosis (Graven et al., 1993). Therefore, both low hematocrit and the hypoxia-mediated apoptosis could contribute to the vascular defects observed in the c-myc mutants.

However, lack of hemodynamic force and increased hypoxic stress are not the only explanation for the absence of vascular remodeling in c-myc mutants. HCs secrete factors capable of promoting angiogenesis in a paracrine manner (Okamoto et al., 2005; Okuda et al., 1996). Lack of proangiogenic factors from HCs are responsible for angiogenic defects in Arnt (Ramirez-Bergeron et al., 2006) and Aml1 (Takakura et al., 2000) mutants. As Tie2-Cre;c-mycflac embryos have hematopoietic defects, they might also lack HC-derived proangiogenic factors, which could explain their angiogenesis defects.

Consistent with this hypothesis, pale Vav-iCre;c-mycflac embryos exhibited a significant decrease of Il1b and Mmp2 mRNA at 11.5 dpc. IL1β is secreted primarily from monocytes and macrophages. In the anemic Vav-iCre;c-mycflac embryos, the significant decrease of CD11b⁺ myeloid cells in fetal livers and peripheral blood likely explains the decrease of IL1β expression and secretion. IL1β and MMPs were recently found to form an axis to regulate the bioavailability of VEGF in angiogenesis (Shchors et al., 2006). IL1β mobilizes VEGF from the extracellular matrix (ECM) to ECs during active angiogenesis, via its ability to promote expression and proteolytic activation of stromal MMPs (Mountain et al., 2007; Shchors et al., 2006). MMPs not only modulate the ECM but also cleave the ECM binding domain of VEGF and release isoforms of VEGF to ECs (Bergers et al., 2000). Depletion of these and other paracrine factors originating from hematopoietic cells is likely to contribute to the impaired angiogenesis in the mutant, despite the elevated VEGF mRNA levels.

We therefore suggest that a combination of defects including reduction in hemodynamic stress and hypoxia-induced apoptosis with a shortage in proangiogenic factors contributes to the vascular defects in the mid-gestation c-myc mutant embryos. If these vascular defects resulted exclusively from the loss of HCs, then preserving c-myc in HCs while deleting it in ECs should allow the mutant embryos to survive beyond midgestation and develop a normal vasculature. Indeed, when we deleted c-myc in the majority of ECs but only a subset of HCs using the Tie1-Cre line, all of these mutants survived to late gestation, well past the lethality of Vav-iCre;c-mycflac embryos. These results suggest that c-Myc regulates angiogenesis through its control over hematopoiesis and the production of paracrine factors.

The physiological function of c-myc may be restricted to hematopoietic lineages in the embryo and placenta

Retention of c-myc in the visceral endoderm and the extra-embryonic ectoderm of Sox2-cre mutants prevented the gross organ abnormalities seen in c-myc⁻/⁻ embryos and extended the embryo survival, demonstrating that c-myc plays an essential function in these tissues. In the embryo proper, the severe Vav-iCre CKs phenocopied the Tie2-Cre and Sox2-Cre CKs, suggesting the possibility that c-myc in the HCs is most crucial for the development and survival of the embryo at this stage.

Therefore, our genetic evidence suggests the possibility that c-myc functions restrictively in placenta and HCs but less so in other tissues. Supporting this notion, we found no significant cell autonomous requirement for c-myc in ECs. Other studies also indicate that c-Myc is dispensable for the homeostasis of the adult intestinal epithelium (Benitah et al., 2005; Oskarsson et al., 2006), postnatal hepatocyte proliferation (Baena et al., 2005) and liver regeneration (Li et al., 2006). Although further investigation is required to delineate the precise physiological function of c-Myc, our data and the published findings raise the hypothesis that c-myc may be uniquely required in the hematopoietic lineage and placenta, playing a less crucial role in other cell lineages in vivo.

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Supplementary material

Supplementary material for this article is available at http://dev.biologists.org/cgi/content/full/135/14/2467/DC1

References


