Adenosine signaling promotes hematopoietic stem and progenitor cell emergence

Lili Jing,1,2 Owen J. Tamplin,1,2 Michael J. Chen,1,2 Qing Deng,5,6 Shenia Patterson,7 Peter G. Kim,1,2,3 Ellen M. Durand,1,2 Julie M. Green,5,6 Shinobu Matsuura,7 Julien Ablain,1,2 Margot K. Brandt,1,2 Thorsten M. Schaüger,1,2 Anna Huttenlocher,5,6 George Q. Daley,1,2,3 Katya Ravid,7 and Leonard I. Zon1,2,3,4

1Stem Cell Program, Division of Hematology/Oncology, Boston Children’s Hospital and Dana-Farber Cancer Institute, Boston, MA 02115
2Harvard Stem Cell Institute, 3Howard Hughes Medical Institute, and 4Department of Stem Cell and Regenerative Biology, Harvard University, Cambridge, MA 02138
5Department of Medical Microbiology and Immunology and 6Department of Pediatrics, University of Wisconsin-Madison, Madison, WI 53706
7Department of Medicine, Boston University School of Medicine and the Evans Center for Interdisciplinary Biomedical Research, Boston, MA 02118

Hematopoietic stem cells (HSCs) emerge from aortic endothelium via the endothelial-to-hematopoietic transition (EHT). The molecular mechanisms that initiate and regulate EHT remain poorly understood. Here, we show that adenosine signaling regulates hematopoietic stem and progenitor cell (HSPC) development in zebrafish embryos. The adenosine receptor A2b is expressed in the vascular endothelium before HSPC emergence. Elevated adenosine levels increased runx1+/cmyb+ HSPCs in the dorsal aorta, whereas blocking the adenosine pathway decreased HSPCs. Knockdown of A2b adenosine receptor disrupted scl+ hemogenic vascular endothelium and the subsequent EHT process. A2b adenosine receptor activation induced CXCL8 via cAMP–protein kinase A (PKA) and mediated hematopoiesis. We further show that adenosine increased multipotent progenitors in a mouse embryonic stem cell colony-forming assay and in embryonic day 10.5 aorta–gonad–mesonephros explants. Our results demonstrate that adenosine signaling plays an evolutionary conserved role in the first steps of HSPC formation in vertebrates.

Hematopoietic stem cells (HSCs) are a reservoir of rare, multipotent stem cells that provide a continuous supply of various hematopoietic lineages circulating in the blood (Orkin and Zon, 2008). It is of great therapeutic interest to generate transplantable HSCs from human embryonic stem cells (ESCs) or induced pluripotent stem cells. Despite many years of studies, such in vitro bona-fide HSC generation has proven difficult, which is partly the result of our incomplete understanding of the pathways that regulate HSC formation during development.

In the embryo, HSCs are first specified in the aorta–gonad–mesonephros (AGM) region (Medvinsky and Dzierzak, 1996). HSCs derive directly from a unique population of aorta endothelial cells termed hemogenic endothelium (HE; Yoshimoto and Yoder, 2009). By in vivo time-lapse confocal imaging, recent studies have captured the emergence of HSCs from the ventral aorta endothelium through a process known as the endothelial-to-hematopoietic transition (EHT; Bertrand et al., 2010; Boisset et al., 2010; Kissa and Herbomel, 2010). During this process, hemogenic endothelial cells bend, round up to transform to HSCs, and release from the aorta ventral wall to the vascular lumen. Previous studies have identified pivotal transcription factors that regulate this process. runx1, a central player expressed in HE and nascent HSCs, is essential for the transformation of hemogenic endothelial cells to HSCs (Okuda et al., 1996; Wang et al., 1996; Chen et al., 2009). Additionally, scl,
Adenosine-regulated HSPC formation is primarily through $A_2b$ adenosine receptor

Adenosine signaling exerts its effect by activating G protein–coupled receptors on target cells. Four adenosine receptors, $A_1$, $A_2a$, $A_2b$, and $A_3$, are expressed in hematopoietic and endothelial cells, with $A_2b$ being the most abundant in both cell types. Adenosine signaling through these receptors modulates hematopoietic progenitor cell development and function.

In the context of hematopoietic stem cell (HSC) development, adenosine signaling plays a crucial role in regulating HSPC formation and differentiation. The $A_2b$ receptor is particularly important, as it promotes the proliferation and survival of hematopoietic progenitors. This receptor-mediated effect is thought to be mediated by the adenosine A2b receptor, which is highly expressed in hematopoietic progenitors and is involved in regulating HSC development in the ventral wall of the dorsal aorta.

The effect of adenosine signaling on HSC development is mediated through a cAMP-dependent pathway. Adenosine receptor agonists increase the production of cAMP, which activates protein kinase A (PKA), leading to the differentiation of HSPCs. In contrast, adenosine receptor antagonists, such as NECA, inhibit cAMP production and block HSPC differentiation.

In conclusion, adenosine signaling through the $A_2b$ receptor is an essential regulator of HSC development and function. Understanding the molecular mechanisms underlying this process is crucial for advancing our knowledge of hematopoietic stem cell biology and for developing strategies to manipulate HSC development in disease contexts.
A1, A2a, A2b, and A3, have been identified. Each receptor has a distinct cellular and tissue distribution and modulates different biological functions (Haskó et al., 2008). We focused on A2a and A2b adenosine receptors (denoted here as A2a and A2b) because of their enrichment in endothelial and hematopoietic cells (Adair, 2005; Rossi et al., 2012) and because of the

**Figure 1.** An adenosine analogue regulates definitive hematopoiesis. (A–C’) In situ hybridization of runx1/cmyb at 36 hpf. Representative images of wild-type embryos treated with DMSO (A and A’), adenosine receptor agonist NECA (20 µM; B and B’), and antagonist (10 µM CGS15943; C and C’). Prime panels depict the AGM region of the embryos in A–C (red arrows). The numbers at the right corner (here and following) indicate embryos with altered HSPC staining over the total number of embryos examined. (D) Bright view of Tg(cmyb:GFP, flk1:mcherry) embryos. The square indicates the live-imaging position. (E–G’) Confocal images of double-transgenic embryos at 36 hpf. The embryos were treated with DMSO (E and E’), NECA (F and F’), or CGS15943 (G and G’). The white arrows point to the double-positive HSPCs in the aorta region. The blue arrowheads point to the multiciliate cells of the pronephros. Dashed lines indicate the somite boundaries. (H) Quantification of the number of HSPCs per somite in Tg(cmyb:GFP, flk1:mcherry) embryos at 36 hpf. The results are presented as mean ± SE (Student’s t-test: *, P < 0.05; n = 5–8 per group). Bars, 100 µm.
During embryo development, A2b is predominantly expressed in the vasculature within the AGM and CHT (Fig. 2, A and B), which is consistent with the previous findings (Boehmler et al., 2009). Real-time quantitative PCR (qPCR) above-described effect of NECA (an A2-type adenosine receptor agonist). We designed morpholinos (MOs) to knock down the two receptors and identified A2b as an important player in HSC development.

Figure 2. Adenosine functions through A2b to regulate HSPC formation. (A and B) In situ hybridization of A2b at 24 hpf (A) and 32 hpf (B). A2b is expressed in the vasculature of the AGM (red arrowheads) and the CHT (blue arrowheads). Insets are enlarged images of the AGM. (C) qPCR analysis of A2b expression in flk1:GFP+ FACS-sorted cells with flk1:GFP− cells at 24, 27, and 32 hpf. Results are shown as fold change of GFP+ to GFP− cells and normalized to expression of β-actin mRNA (Student’s t test: *, P < 0.05; n = 3 samples). (D) qPCR analysis of A2b expression in flk1:GFP+ cells from control or sih morphant embryos at the indicated stages: 16 and 23 hpf (before circulation) and 28 hpf (after circulation starts). Results are shown as fold change relative to A2b level at 16 hpf and normalized to expression of β-actin mRNA (Student’s t test: *, P < 0.05; n = 3 samples). (E) RT-PCR on uninjected or A2b splicing MO (A2b SP MO, here and following)—injected embryos with A2b- or actin-specific primers. Note the absence of correctly spliced A2b transcript after MO injection. (F–I) Expression of runx1/cmyb at 36 hpf in the AGM of control embryos (F), embryos injected with A2b MO (G), and hA2b mRNA–injected (H) and A2b MO- and hA2b mRNA–co-injected embryos (I). The expressions in each panel are combined from multiple experiments. (J and K) Expression of runx1/cmyb at 36 hpf in control embryos (J) and embryos treated with 10 µM BAY 60-6583 (K). (L–O) Confocal images of Tg(cmyb:GFP; flk1:mcherry) embryos at 36 hpf. Control embryos (P) and embryos injected with A2b MO (P′), and hA2b mRNA–injected (P″) are shown. White arrowheads mark the double-positive HSPCs. Blue arrowheads point to the pronephros. White dashed lines mark somite boundaries. (S) Quantification of the experiments in G and I. The results are presented as the mean percentage of embryos with restored runx1/cmyb staining as in I ± SE (Student’s t test: *, P < 0.05; n = 3 independent experiments, 20–25 embryos per experiment). (T) Quantification of number of HSPCs per somite in Tg(cmyb:GFP; flk1:mcherry) embryos. The results are presented as mean ± SE (Student’s t test: *, P < 0.05; **, P < 0.01; n = 5–8 embryos per group). Bars, 100 µm.
analysis confirmed that A2b is enriched in FACS-sorted flk1:GFP+ endothelial cells from 20-somite stage to 40 hpf (Fig. 2 C and not depicted). We found that A2b expression in endothelial cells is elevated shortly before the circulation starts and HSPCs emerge and that A2b expression is strongly reduced in embryos injected with silent heart (silb) MOs that lack a heartbeat and blood circulation (Fig. 2 D). These results suggest that circulation is required to maintain A2b expression, but not for induction of expression.

We used a translation–blocking MO and a splice MO to block the A2b receptor. We confirmed that the splicing MO inhibits A2b splicing (Fig. 2 E). Knockdown of A2b, using either MO did not affect the general development of the embryos (not depicted) but caused a strong reduction of runx1/cmyb in the AGM (Fig. 2 F, G, L, and M). Injection of zebrafish A2b mRNA rescued the expression of runx1/cmyb in A2b splicing MO–injected embryos (Fig. 2, N and O), and injection of human A2b mRNA partially rescued runx1/cmyb in the embryos injected with A2b translation–blocking MO (Fig. 2, H, I, and S), which supports the specificity of MO targeting. In addition, we injected A2b MO into p53−/− mutant embryos to attenuate the off-target effect (Robu et al., 2007), and we saw similar phenotypes in p53 mutant embryos (not depicted), which also supports that the observed HSPC defects are caused by specific loss of A2b function. The robust phenotype after blocking A2b function prompted us to examine the effect of an A2b receptor selective agonist (BAY 60-6583; Koupenova et al., 2012) in zebrafish embryos. As expected, BAY 60-6583 significantly increased the expression of runx1/cmyb (Fig. 2, J and K).

Consistent with the runx1/cmyb staining, knockdown of A2b in Tg(amyb:GFP, flk1:mcherry) reduced the double fluorescent-labeled HSPCs (1.1 ± 0.2/somite vs. 3.6 ± 0.6/somite [conj]). BAY 60-6583 treatment increased the number of HSPCs in the Tg(amyb:GFP, flk1:mcherry) embryos (4.6 ± 0.3/somite; Fig. 2, P–R) and T). Alterations of adenosine signaling with either treatment did not have an effect on the GFP-labeled cells in the pronephros (Fig. 2, P–R).

Compared with amyb, runx1 is more specifically expressed in nascent HSCs. We examined embryos using runx1 probe alone and found that knockdown of A2b strongly reduced runx1+ cells (Fig. 3, A and B). We also studied hematopoiesis and subsequent lineage differentiation at later time points after A2b inhibition. A2b morphants exhibit reduced runx1+/amyb+ HSPCs in the CHT at 60 hpf (Fig. 3, C and D), reduced amyb+ multipotent progenitor cells in the CHT at 4.5 d post-fertilization (dpf; Fig. 3, E and F), and reduced rag1+ thymic T cells (Fig. 3, G and H). Together, our results indicate that A2b regulates HSPC development.

To determine the specificity of loss of A2b-induced HSPC defects, we examined the development of other tissues. A2b morphant embryos demonstrated normal heart beating and circulation (not depicted). Those embryos developed normal vasculature differentiation, as shown by the normal expression of flk1:mcherry (Fig. 2, P–Q′), the vessel marker flk1, the artery marker EphB2, and the normal intersegment vessels (flk1 and flk1:mcherry), pronephros (cd11h17), and somite (myoD; Fig. 3, I–P). In addition, A2b MO–injected embryos showed normal primitive erythroid cells (gata1) and angioblasts (sl; Fig. 3, Q–T). The transient expression of amyb and runx1 in early somite–stage embryos was not affected (Fig. 3, U–X). These results support that the defects in HSPC specification in the absence of A2b are highly specific, not caused by defects from the nearby tissues.

Adenosine promotes hematopoietic development in mESC culture and in mouse AGM explant

To test whether adenosine signaling would have an effect on mammalian hematopoiesis, we assessed the effect of adenosine on the development of mouse HSPC populations. We first used an mESC hematopoietic differentiation system, an accessible in vitro model which recapitulates mouse embryonic hematopoiesis (Kim et al., 2013). In this model, mESCs are aggregated into embryoid bodies (EBs) and differentiated into primitive streak–like mesoderm on day 2, hemangioblasts on day 3.25, and hematopoietic precursors after day 5. A2b receptor is present between days 0 and 6 of EB differentiation (not depicted). Analysis of gene expression data in sorted populations from day 6 EBs (Kim et al., 2013) showed that A2b is high in the VE-cadherin ‘CD41’ endothelial cells and low in the CD41+ populations (Fig. 4 A), indicating that A2b likely regulates VE-cadherin+ cell intermediates, a fraction from which HSCs emerge. Methylcellulose–based colony–forming assay was performed to measure both hematopoietic progenitor types and numbers. Addition of A2b selective agonist BAY 60-6583 to EBs between days 4 and 6 significantly increased the number of hematopoietic colonies, including definitive erythroid (CFU-E), granulocyte/monocyte (CFU-GM), and multipotent granulocyte/erythrocyte/macrophage (CFU-E/GM/GEMM) colonies, at a concentration of 0.5–2.5 µM (Fig. 4 B). Consistently, qPCR analysis revealed that many hematopoietic genes, including sl, lmo2, runx1, gata1, bH1, and Bmajor (adult globin), were up–regulated on day 6 after BAY 60-6583 treatment (Fig. 4 C). In contrast, BAY 60-6583 did not increase the expression of the mesoderm marker Cerberus, the expression of endothelial gene fli1 (Fig. 4 D). Interestingly, the expression of flk1 was decreased after BAY 60-6583 treatment (Fig. 4 D), consistent with the previous study showing the antagonist effect of runx1 on flk1 (Swiers et al., 2013). These data support a positive role of adenosine and its A2b receptor in regulating mESC hematopoietic differentiation.

Next, we examined whether BAY 60-6583 promotes HSCs in the mouse AGM. A recent study performed expression profiling of the AGM HE at the onset of circulation (Swiers et al., 2013). We analyzed the microarray data and found that A2b expression is high in HE and low in non–HE and hematopoietic progenitors (Fig. 4 E), similar to A2b expression from day 6 EBs. We confirmed A2b expression in VE-Cadherin+ cells from the E10.5 AGM by RT–PCR (Fig. 4 F). To test the effect of adenosine signaling on AGM HSPCs, AGM explants from E10.5 embryos were cultured with DMSO or BAY 60-6583, and HSC activity was measured by CFU...
culture (CFU-C) assay. As shown in Fig. 4 G, the addition of BAY 60-6583 significantly increased the numbers of erythroid, granulocyte/macrophage (GM), and multipotent progenitor (GEMM) colonies. Thus, adenosine promotes AGM HSCs in mouse.

Adenosine is required for the EHT

It is known that HSCs originate from HE via EHT. Recent time-lapse imaging studies have directly captured the emergence of HSPCs from flk1:GFP+ aortic endothelial cells in live zebrafish embryos (Bertrand et al., 2010; Kissa and Herbomel, 2010). To examine whether adenosine regulates the EHT, we injected A2b MO into Tg(flk1:GFP) embryos and used time-lapse imaging to track the formation of HSPCs between 32 and 60 hpf, as previously described (Kissa and Herbomel, 2010). As shown in Fig. 5 and Videos 1 and 2, in A2b-deficient embryos, flk1:GFP+ endothelial cells initiated bending, and sometimes contraction, normally, albeit more rarely than in control embryos. We noticed that as the flk1:GFP+ cells rounded up to become HSCs, a subset of these cells burst into small cellular pieces (Fig. 5, A–C). The bursting is very similar to the events seen in runx1-deficient embryos, in which the EHT event of flk1:GFP+ cells abort into cell death (Kissa and Herbomel, 2010). These results suggested that adenosine signaling is required to achieve successful EHT.

To explore the relationship between adenosine signaling and runx1, we used an inducible runx1 Tg line (hsp70:runx1). After heat shock induction, runx1 was expressed throughout the embryo (Fig. 5 D) and slightly increased the definitive hematopoietic marker cmyb in control embryos (Fig. 5, E and F). In A2b
MO–injected embryos, overexpression of runx1 showed a partial, but statistically significant rescue of cmyb staining (Fig. 5, G–I). Thus, overexpressing runx1 rescued HSPC formation in the absence of A₂b, suggesting that runx1 acts downstream of A₂b.

Adenosine signaling regulates scl⁺ hemogenic vascular endothelium

The abortive EHT of flk1:GFP cells could result from a lack of hemogenic potential in these cells. We then examined whether adenosine regulates HE specification, the onset of EHT. The master hematopoietic transcriptional regulator scl is critical for establishing the hemogenic potential in aorta endothelial cells and plays its role before runx1 during definitive hematopoiesis. In zebrafish, an N-terminal truncated scl⁻ isoform is essential for HSPC emergence. The scl reporter line Tg(scl⁻:d2eGFP) marks the hemogenic vascular endothelium, which is later transformed to HSPCs via EHT (Zhen et al., 2013). We altered adenosine signaling in Tg(scl⁻:d2eGFP) embryos. In the absence of A₂b, scl⁻:GFP⁺ HE cell population was strongly reduced (1.6 ± 0.2/somite vs. 3.5 ± 0.3/somite [con]; Student’s t test: *, P < 0.05; **, P < 0.01; n = 3–5 per group). Addition of A₂b receptor agonist BAY 60-6583 to wild-type embryos increased the number of scl⁻:GFP⁺ in the AGM (4.6 ± 0.7/somite vs. 3.5 ± 0.5/somite [con]; Student’s t test: *, P < 0.05; **, P < 0.01; n = 3 per group). Overexpression of scl mRNA in zebrafish embryos partially rescued runx1⁺/cmyb⁺ HSPCs in A₂b-deficient embryos (Fig. 6, D–G and I). These results indicate that adenosine signaling mediates scl⁺ hemogenic vascular endothelium. Together, our data suggest that adenosine mediates HSPC formation via regulation of hematopoietic commitment of endothelial cells.
Figure 5. Adenosine signaling is required for EHT in zebrafish. (A and B) Still images from in vivo time-lapse confocal imaging of Tg(flk1:GFP) embryos between 32 and 48 hpf. Embryos are either uninjected (A) or injected with A\(_{2b}\) MO (B). Numbers indicate recording time in hours and minutes. Arrows in A point to three aorta cells undergoing successful EHT. Cells 2a and 2b are the daughter cells of cell 2. The two dorsal aorta cells 1' and 2' in B initiate EHT and burst into pieces (marked by circles). (C) Quantification of the number of flk1:GFP\(^+\) cells per somite undergoing burst during the 16-h time-lapse period. The results are presented as mean ± SE (Student's t test; **, P < 0.01; n = 6–8 embryos per group). (D) Tg(hsp70:runx1) embryos stained for runx1 without heat shock (HS) induction or after heat shock induction. (E–H) Expression of cmyb at 36 hpf in Tg(hsp70:runx1) embryos. Embryos were either uninjected (E and F) or injected with A\(_{2b}\) MO (G and H). Embryos either received no heat shock treatment (E and G) or received heat shock induction...
Adenosine regulates CXCL8 production and mediates hematopoiesis via CXCL8

In many cell types, adenosine promotes the production of cytokines, chemokines, and growth factors and often exerts its cellular effects through or in association with these factors (Adair, 2005). In vascular endothelial cells, A2b activation modulates the production of several angiocrine factors (Feoktistov et al., 2002). One of the major factors is CXCL8 (also known as IL-8). CXCL8 is known as a potent stem cell—mobilizing agent (Laterveer et al., 1995), and it also increases the proliferation of hematopoietic progenitor cells in vitro (Hermouet et al., 2000). Human aortic endothelial cells (HAECs) express the A2b receptor (Iwamoto et al., 1994), and we confirmed A2b expression by RT-PCR (not depicted). We treated HAECs with BAY 60-6583, which increased CXCL8 protein production (Fig. 7 A). To demonstrate that A2b regulates CXCL8 in zebrafish embryos, we examined cxcl8 transcripts. Knockdown of A2b strongly decreased cxcl8 and runx1 expression in flk1;GFP+ endothelial cells, but did not affect the expression of endothelial marker fli1 (Fig. 7 B). We next examined the function of CXCL8 in HSC development. Inhibition of cxcl8 with MO (Stoll et al., 2011) in zebrafish embryos strongly reduced runx1/cmyb+ HSPCs in the AGM (Fig. 7, C–F) and the CHT (Fig. 7, I and J). Loss of CXCL8 did not have an effect on the development of the dorsal aorta at the time when HSPCs start to emerge (Fig. 7, K and L) and did not interfere with primitive hematopoiesis (not depicted), which supports a specific role of cxcl8 in HSPC formation. In addition, we generated a CXCL8 mutation by a transcription activator–like effector nuclease (TALEN) approach. This mutation
Adenosine regulates HSC formation via cAMP–PKA

A<sub>2b</sub> receptor is commonly coupled to adenyl cyclase via G protein subunit Gs and increases intracellular cAMP that activates PKA (Fredholm, 2007). We treated HAECs with BAY 60-6583 and performed chemiluminescence assay for cAMP activity. BAY 60-6583 induced cAMP production (Fig. 8A).

In addition, treatment of HAECs with H89 reduced BAY 60-6583–induced CXCL8 production (Fig. 8B). To examine the role of cAMP in A<sub>2b</sub>-mediated hematopoietic induction in embryos, we applied H89 to zebrafish embryos. H89 at 1 µM decreased runx1<sup>+</sup>/cmyb<sup>+</sup> HSPCs in the wild-type embryos and abolished the enhancement effect of NECA on leads to the deletion of two amino acids in the conserved CXC domain (Fig. 7 M; Fernandez and Lolis, 2002), which has been shown to be important for its tertiary structure. We found that the HSPC staining is reduced in CXCL8 mutant embryos compared with those in control siblings (Fig. 7, N and O), in accordance with the defects we observed in CXCL8 MO knockdown embryos. Similar to loss of A<sub>2b</sub>, loss of CXCL8 also reduced scl<sup>+</sup> hemogenic vascular endothelium (Fig. 7, P–R).

More importantly, cxcl8 mRNA restored HSCs and progenitors in the embryos that lack the A<sub>2b</sub> (Fig. 7, G and H). These results suggest that adenosine signaling regulates CXCL8 production to mediate HSPC development.

Adenosine mediates HSPC development by regulating CXCL8. (A) BAY 60-6583 induces CXCL8 secretion in HAECs (Student’s t test: *, P < 0.05; n = 3). (B) Relative mRNA expression of cxcl8, runx1, and flt1 in FACs-sorted flk1:GFP<sup>+</sup> cells in A<sub>2b</sub> MO–injected compared with control MO–injected embryos (Student’s t test: **, P < 0.01; n = 3). (C–H) Expression of runx1/cmyb at 36 hpf. Control embryos (C) and embryos injected with cxcl8 MO (D), injected with cxcl8 mRNA (E), co-injected with cxcl8 MO and cxcl8 mRNA (F), injected with A<sub>2b</sub> MO (G), and co-injected with A<sub>2b</sub> MO and cxcl8 mRNA (H) are shown. (I and J) Expression of runx1/cmyb in the CHT of control embryos or embryos injected with cxcl8 MO. (K and L) Control or cxcl8 MO–injected embryos stained for dorsal aorta (ephrinB2). (M) A CXCL8 TALEN mutation leads to the deletion of two amino acids (in yellow) in the conserved CXC domain. (N and O) Expression of cmyb at 36 hpf in control sibling embryos and CXCL8 mutant embryos. htz, heterozygous; hom, homozygous. (P–Q) Confocal imaging of Tg(sclβ:z2eGFP, flk1:mcherry) embryos at 30 hpf. Control embryos (P and Q) and embryos injected with cxcl8 MO (Q and Q<sup>′</sup>) are shown. Arrowheads indicate the hemogenic endothelial cells marked by sclβ:GFP<sup>+</sup>. Dashed lines mark the somite boundaries. (R) Summary of the number of sclβ:GFP<sup>+</sup> hemogenic endothelial cells per somite (Student’s t test: **, P < 0.01; n = 5 per group). The results are presented as mean ± SE. Bars, 100 µm.
scl+ hemogenic vascular endothelium (Fig. 8, K–M). These data support that cAMP–PKA acts downstream of A2b in promoting HSPC emergence.

Overall, these data indicate that adenosine signaling through A2b in endothelial cells activates the cAMP–PKA pathway, which up-regulates the production of cytokine CXCL8.
turn, CXCL8 promotes the emergence of HSPCs from the endothelium that is mediated by a transcriptional program including \(sl\) and \(nunx1\).

**DISCUSSION**

Adenosine signaling via its receptors has been studied in tissue regeneration. For example, adenosine promotes the regeneration of pancreatic \(\beta\) cells (Andersson et al., 2012). In the hematopoietic system, adenosine enhances hematopoietic cell repair in adult mice after irradiation (Pospíšil et al., 1993). Although components of adenosine signaling are expressed in the early embryo (Massé and Dale, 2012), the role of adenosine during embryogenesis, particularly in progenitor/stem cell development, has not been characterized previously. Our work has discovered an unrecognized role of adenosine in regulating HSPC development. We demonstrate that adenosine promotes HSPC emergence in zebrafish embryos and promotes hematopoietic differentiation in mESCs and in E10.5 AGM explants.

In the developing embryo, HSPCs derive from HE, a crucial intermediate population generated from endothelial cells. However, the signaling pathways that induce the hemogenic commitment of endothelial cells remain largely unknown. \(sl\) is one of the early molecular determinants for HE (Lancrin et al., 2009). But little is known regarding how an \(sl\) event is regulated. Our experiments demonstrate that adenosine \(A_2b\) is expressed in vascular endothelial cells in zebrafish embryos and in VE-cadherin\(^+/+\) +23runx1-GFP\(^+\) HE in the mouse AGM. Adenosine signaling modulates \(sl\)-marked HE, and ectopic expression of \(sl\) partially rescues HSPC development in the absence of \(A_2b\). These findings support that adenosine through \(A_2b\) regulates \(sl\)-mediated hematopoietic commitment from aortic endothelium.

It is well known that adenosine stimulates the production of multiple growth factors and cytokines (Feoktistov and Biaggoni, 2011). In the AGM, endothelial cells are a probable source of the environmental factors that could affect HSC specification. We show that adenosine regulates CXCL8 production in the vascular endothelial cells. CXCL8 is often associated with neutrophil chemotaxis (Baggiolini and Clark-Lewis, 1992). It also has other cellular functions such as inducing rapid mobilization of HSCs in rodents (Laterveer et al., 1995). But its role during developmental hematopoiesis remains unexplored. We demonstrate that CXCL8 regulates \(sl\)\(^+\) HE and subsequent HSPC development during embryogenesis, similar to adenosine \(A_2b\). Moreover, \(cxl\) restores HSPC formation in \(A_2b\)-deficient embryos. Although adenosine may induce hematopoiesis through additional growth factors, our data suggest that adenosine acts at least partly through stimulating CXCL8 to promote HE and HSPC specification.

In addition, we show that cAMP–PKA is a downstream effector of adenosine signaling, and it underlies adenosine-induced CXCL8 production in endothelial cells, which is consistent with a previous report that the CXCL8 gene promoter has a cAMP-responsive element (CRE)–like site (Iourgenko et al., 2003). An accompanying study in this issue by Kim et al. demonstrated that shear stress activates the PKA–CREB pathway in VE-cadherin\(^+\) cells, which promotes AGM HSC emergence. Recent studies indicate that shear stress can lead to the release of ATP, which is quickly converted to adenosine (Yegutkin et al., 2000; Wen et al., 2011). In our study, we found that the expression of \(A_2b\) in vascular endothelial cells is maintained by blood flow. Therefore, it is possible that during HSC development, shear stress induces adenosine release, and its effect via \(A_2b\) contributes to the activation of the cAMP–PKA pathway in endothelial cells. We propose that cAMP–PKA–CREB regulates the production of growth factors and cytokines, which in turn promotes HSPC development controlled by a transcriptional program. It is also possible that cAMP–PKA–CREB may directly regulate \(sl\), and CXCL8 and other factors may act to reinforce the transcriptional program. In any case, our experiments together highlight the critical role of the cAMP–PKA signaling pathway during vertebrate HSPC emergence.

It remains unknown how CXCL8 regulates HSPC emergence. Since it has been suggested that CXCL8 enhances cell growth and survival in different types of cells (Hermoust et al., 2000), it is possible that CXCL8 production in the AGM microenvironment affects the survival and/or proliferation of newly formed \(sl\)\(^+\) hemogenic endothelial cells and HSPCs. We found that CXCL8 receptors \(cxr1\) and \(cxr2\) are expressed in \(flk1\)\(^+\) vascular endothelial cells in zebrafish embryos before HSPC emergence (unpublished data), which supports an autocrine signaling of CXCL8. The precise mechanism underlying CXCL8-mediated HSPC induction awaits further study.

In addition to acting in the formation of HE, our time-lapse imaging experiments suggest that adenosine might also regulate the transition of endothelial cells to hematopoietic cells. \(nunx1\) is a central factor critical for the generation of HSCs from HE. It has been suggested that \(sl\) might activate \(nunx1\), directly or indirectly, to promote the conversion of endothelial cells to hematopoietic cells. In zebrafish embryos that lack \(nunx1\), \(sl\)\(^+\) HE remains intact (Zhen et al., 2013). In \(A_2b\)- and CXCL8-deficient embryos, \(nunx1\) expression was decreased, and \(sl\)-marked hemogenic endothelial cells were also reduced. Our data support that the \(A_2b\)-CXCL8 axis acts upstream of \(sl\) rather than \(nunx1\). It is intriguing that we found that overexpression of \(nunx1\) restored HSPCs in \(A_2b\)-deficient embryos. This finding supports the idea that overexpression of \(nunx1\) may bypass the requirement of \(sl\)-induced hematopoietic potential in endothelium.

Collectively, our experiments have discovered that adenosine regulates the generation of HSCPs in the early embryo. We provide evidence that it acts through \(A_2b\) in endothelial cells to regulate HE and subsequent HSPC emergence, partly through up-regulation of CXCL8. These results provide novel insights into the signaling pathways that will be needed to derive HSCs from ESCs or pluripotent cells.

**MATERIALS AND METHODS**

**Animals.** Mice and zebrafish were maintained at Boston Children’s Hospital according to institutional animal care and use committee protocols. The zebrafish wild-type Tubingen and transgenic lines \(flk1\):\(GFP\), \(flk1\):\(mcherry\), cMyb: \(GFP\), \(cMyb\):\(cMyb\)\_\(GFP\) (Z. Wen, Hong Kong University of Science and Technology, Hong Kong, China), and \(p53\)\(^{−/−}\) were used in this study.
Embryo chemical treatment. Wild-type and transgene embryos were treated from 5-somite to 36 hpf. 20–50 µM NECA, 10–20 µM CGS19755, and 5–10 µM BAY 60-6853 were obtained from Tocris Bioscience.

Confocal imaging. Live embryos were anesthetized in 0.04 mg/ml Tricaine, mounted in 1% low-melting agarose, and imaged with Leica confocal microscopes. GFP+/β′′/β′′′ cells were counted in projections of z-stack images (n = 5–10 embryos per group).

In situ hybridization. In situ hybridization was performed as previously described (Thiese and Thisse, 2008). The images were taken with a Leica stereoscope and processed using Adobe Photoshop.

FACS sort cells and qPCR analysis. β′′/GFP embryos were manually dissociated with 0.9% PBS containing Liberase (Sigma–Aldrich). GFP+ cells were sorted into LS–TRIzol (Sigma–Aldrich). Total mRNA was extracted according to the manufacturer’s protocol. Genomic DNA was removed using TURBO DNA-free kit (Life Technologies). cDNA was synthesized using SuperScript First-Strand synthesis system (Life Technologies). qPCR was performed using SYBR green supermix on a CFX96 real-time PCR detection system (Bio-Rad Laboratories) with gene-specific primers listed in Table S1. Relative expression level was determined.

MO and mRNA injection. MOs were ordered from Gene Tools and injected into 1–2 cell-stage embryos. Aβ, translation blocking MO (5′-CAATGGCGATGTGAGCGAATCCAT-3′, 5 ng) was injected at 1–2 cell-stage. Aβp, splice-blocking MO (5′-AAAGCGGAGAAGACTCCTCGAGG-3′, 2 ng) was designed against the splice donor site of ex on. For RT-PCR analysis, cDNA templates were synthesized from five 24-hpf embryos. PCR primers were 5′-TCTTCAATCTCTCGCTGTCC-3′ (forward) and 5′-TCTTCAATCTGTCGCAATC-3′ (reverse). IL-8 MO targeting exon 2–intronic 2 junction sequence (5′-CTGATGCTGGTGAACCTCAGTA-3′ [Stoll et al., 2011]) was injected at 0.5 ng/embryo. Standard control MO was purchased from Gene Tools.

mRNA was in vitro transcribed from linearized constructs using SP6 mMESSAGE mMACHINE kit (Ambion) and injected into the embryos at 1-cell stage. Zebrafish Aβp mRNA was injected at 50 pg/embryo. Human Aβp mRNA was injected at 400 pg/embryo. scl mRNA was injected at 150 pg/embryo, and cdx8 mRNA was injected at 200 pg/embryo.

Methylocellulose colony formation. mESC maintenance and differentiation were modified from Nostro et al. (2008). The Bry–GFP cells were maintained in serum-free culture system for 3 days and then dispensed and aggregated to EBs using aggreWell (STEMCELL Technologies). After 24 h, EBs were grown in 6-well ultra-low attachment plate (Costar) with serum-free medium containing 2 ng/ml Activin, 3 ng/ml Wnt3a, 3 ng/ml BMP4, and 1 mM GSK269662A for 48 h. After that, EBs were switched to serum-free medium containing 5 ng/ml VEGF and 3 ng/ml BMP4 for 72 h. BAY 60-6853 was added between day 4 and day 6 during EB expansion. On day 6, EBs were dissociated to single cells and plated on M3434 methylocellulose. Colonies were counted on days 7–9 by morphology. For qPCR analysis, total RNA was extracted from day 6 EBs, and after cDNA synthesis, qPCR was performed using gene-specific primers as in Table S1.

AGM explant. Whole AGM region from day 10.5 embryos were cultured in IMDM (Invitrogen) containing 2% fetal calf serum and 50 µU/ml penicillin/streptomycin overnight in the presence of 0.1% DMSO or 0.5 µM BAY 60-6853. Tissues were maintained on a shaker in 5% CO2 at 37°C in a humidified incubator. After 14-h (overnight) culture, AGM tissues were dissociated in 0.1% collagenase at 37°C for 1 h and plated using M3434 medium (STEMCELL Technologies) according to the manufacturer’s instructions. Colonies were counted on days 7–9 by morphology.

Time-lapse video. Confocal microscopy was performed according to Kissi and Herbomel (2010). Embryos were mounted in 1% LMP agarose with E3 media and tricaine as described previously (Bertrand et al., 2010). Embryos were imaged in MatTek glass-bottom multi-well plates (no. 1.5 coverslip). Time-lapse recording was performed at 28°C. Microscopy was performed using a Yokogawa spinning disk confocal and Nikon inverted Ti microscope. The objective was a Nikon 20x air Plan-Apo differential interference contrast NA 0.75. Images were acquired with dual Andor Xon x3 EMCCD cameras (512 × 512 pixels) and NIS-elements software. Confocal z-stacks were acquired every 5 min over 16 h (between 32 hpf to 48 hpf) in 17 planes spaced by 2 µm. Multiple stage-matched embryos were imaged in parallel using a moving XY stage.

Heat shock of Tg(hsp70:runx1). The embryos of hsp70:runx1, either uninject or injected with Aβp MO, were heat-shocked twice at 38.5°C for 30 min at 3-somite stage and at 22 hpf. Embryos were then fixed at 36 hpf for analysis.

Cell culture. Primary HAEcs were cultured in EBM-2 medium–supplemented EGM-2 medium containing hEGF, hydrocortisone, hFGF-2, VEGF-R3-IGF-1, acidic acid, Heparin, fetal bovine serum, and Gentamicin/Ampicillin–B [GA] (Lonza). The cells were plated on gelatin-coated plates and kept under a humidified atmosphere of air/CO2 (19:1) at 37°C.

Measurement of CXCL8 secretion. After HAEcs were cultured to confluence, cells were serum starved overnight and suspended in serum-free medium containing 1 U/ml adenosine deaminase. Cells were treated with chemicals at the desired concentration and incubated for 6 h. At the end of incubation, the culture media were collected, and the cells were lysed for total protein concentration measurement (DC protein assay; Bio-Rad Laboratories). CXCL8 concentration was measured using an ELISA kit (BioLegend). cAMP luminescence assay. After cultured to confluence, HAEcs were serum starved overnight. The cells were resuspended in fresh serum-free medium containing 1 U/ml adenosine deaminase and incubated at 37°C for 10 min. Before chemical treatment, cells were treated with 200 µM papaverine and incubated at 37°C for 10 min to block phosphodiesterase. Chemicals were added at the desired concentrations, and cells were incubated at 37°C for 15 min. At the end of incubation, cells were lysed for cAMP ELISA assay according to the manufacturer’s instructions (Enzo Life Sciences). A small portion of cell lysates was used to measure protein concentration.

Generation of CXCL8 mutant. The cxcl8 mutant was generated with TALENs using backbones containing heterodimer endonucleases pCS2– AL3RR and pCS2TAL3DD (Dahlem et al., 2012) and the Golden Gate TALEN kit (Cermak et al., 2011). The two TALEN arms are designed to target Danio rerio CXCL8 (NCBI Gene ID: 100002946) at 5′-TGAGGGTTCTGGTCTAGAT-3′ and 5′-TGAACAGAAAGCGGACCCG-3′. CXCL8 mutant fish were identified by high-resolution melt analysis (Roche 490963101). Primers used were forward, 5′-GAATAGAGCCTTG- CAGGAGTCTGGCTG-3′; reverse, 5′-GGAGGGAAGACTCCA- CATCTTTAT-3′. To sequence individual F0 fish, genomic DNA was isolated from a fin clip using 50 mM NaOH. A PCR was run using the above HRMA primers, and the product was cloned into pCR-4-TOPO (Life Technologies) and then transformed into DH5α competent cells. Six to eight colonies were grown up and sequenced using the M13R primer (5′-CAGGAACACGCTATGAC-3′) to identify any mutations or deletions in the fish. The F2 heterozygous and homozygous mutant embryos were genotyped using the following primer pair: forward, 5′-GAACATGCA- AAACCATATACTGC-3′; and reverse, 5′-AGGGGTCGACAGATC- TCC-3′. The PCR product was then sequenced to confirm the deletion of TCGTGT at positions 214 to 219.

Online supplemental material. Table S1 provides the primers sequences used in this study. Videos 1 and 2 are the time-lapse imaging of a control wild-type zebrafish embryo and an embryo in which Aβp expression was knocked down; the two videos demonstrate that in Aβp-deficient embryos, fewer
flk1:GFP+ cells initiate EHT, and the process ends with an explosion. Online supplemental material is available at http://www.jem.org/cgi/content/full/jem.20141528/DC1.

We would like to thank Dr. Zilong Wen for providing the scfβ2-GFP line, Ronald Mathiess from Boston Children’s Hospital flow cytometry laboratory for assistance with FACS, Dr. Divashit Sah for shi MO, members of the Zhang lab for comments, and Dr. David Wiley, Julie Perlin, and Eva Fast for critical reading of the manuscript. This work was supported by grants from the National Heart, Lung, and Blood Institute (NHLBI; R01 HL04880-21 to Li. Zon, P01 HL10001-05; and H33149 to K. Ravid), National Institute of Diabetes and Digestive and Kidney Diseases (5F30 DK49216, R24 DK092760, and 5RO1 DK52598), and Howard Hughes Medical Institute to Li. Zon. S. Patterson was supported by a Cardiovascular Training Grant from the NHLBI (HL037969).

Li. Zon is a founder and stockholder of Fate Therapeutics Inc. and Scholar Rock.

Submitted: 10 August 2014
Accepted: 20 March 2015

REFERENCES


Anwar, S.S., A.A. Wolf, S. Stainier, et al. 2015. Cardiac training grant from the NHLBI (HL007969).


B978-0-12-385526-8.00005-9


Dr. David Wiley, Julie Perlin, and Eva Fast for critical reading of the manuscript.


