Atypical protein kinase C (aPKCζ and aPKCλ) is dispensable for mammalian hematopoietic stem cell activity and blood formation

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The stem-cell pool is considered to be maintained by a balance between symmetric and asymmetric division of stem cells. The cell polarity model proposes that the facultative use of symmetric and asymmetric cell division is orchestrated by a polarity complex consisting of partitioning-defective proteins Par3 and Par6, and atypical protein kinase C (aPKCζ and aPKCλ), which regulates planar symmetry of dividing stem cells with respect to the signaling microenvironment. However, the role of the polarity complex is unexplored in mammalian adult stem-cell functions. Here we report that, in contrast to accepted paradigms, polarization and activity of adult hematopoietic stem cell (HSC) do not depend on either aPKCζ or aPKCλ or both in vivo. Mice, having constitutive and hematopoietic-specific (Vav1-Cre) deletion of aPKCζ and aPKCλ, respectively, have normal hematopoiesis, including normal HSC self-renewal, engraftment, differentiation, and interaction with the bone marrow microenvironment. Furthermore, inducible complete deletion of aPKCλ (Mx1-Cre) in aPKCζ−/[−] HSC does not affect HSC polarization, self-renewal, engraftment, or lineage repopulation. In addition, aPKCζ- and aPKCλ-deficient HSCs elicited a normal pattern of hematopoietic recovery secondary to myeloablative stress. Taken together, the expression of aPKCζ, aPKCλ, or both are dispensable for primitive and adult HSC fate determination in steady-state and stress hematopoiesis, contrary to the hypothesis of a unique, evolutionary conserved aPKCζ/λ-directed cell polarity signaling mechanism in mammalian HSC fate determination.

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Stem cells are characterized by self-renewal and multilineage differentiation potential (1). The choice between these two processes remains a critical issue in stem cell and cancer biology (1–4). It has been proposed that the selection of cell fate determination toward either self-renewal or differentiation of stem cells results from a choice between maintaining contact with the bone marrow (BM) microenvironment to regulate self-renewal, and changing the cell planar polarity to establish an orientation axis for asymmetric distribution (5, 6). Whether stem cell activity is truly dependent on asymmetric cell division remains controversial (1).

Distinct polarity proteins have been shown to act as core components of the cell polarization machinery in animals ranging from Caenorhabditis elegans to humans (6–12). Asymmetric cell division requires polar distribution of a set of proteins that accumulate before mitosis. Classical cell-polarity complex proteins consist of partitioning-defective proteins: Par3, Par6, and atypical protein kinase C (aPKC ζ and aPKC λ) (7). In the hematopoietic system, aPKC signaling has been implicated in the establishment of T-cell polarity during immunological synapse and in the regulation of pathogen-specific $CD8^+$ T-cell activity (12, 13). aPKCs have been shown to play a crucial role in Drosophila neuroblast self-renewal and differentiation (4, 8, 14, 15). Studies based on Drosophila and C. elegans have indicated that spatiotemporal gradients of several polarity aPKC phosphorylated determinants, such as Lgl or Numb, can determine asymmetric stem cell division through microtubule polarization to orient the mitotic spindle (4, 16). However, while the molecules implicated in stem cell polarity are highly conserved in vertebrates, their mechanism of action could be tissue specific, and in some cases differ substantially from that in the invertebrates (1, 5).

Similarly to embryonic and other tissue-specific stem cells, mammalian hematopoietic stem-cell (HSC) fate could be regulated by an active cell-polarity complex (17–21). Using a Notchreporter system, it has been shown that the differential cell fate of paired daughter cells in culture is related to asymmetric acquisition of Numb in vitro (18). A recent RNA-interference (RNAi) based screening has suggested that aPKCζ and Par6 could positively regulate mammalian HSC activity (19). In a separate study, it has also been shown that the pharmacological attenuation of aPKCζ signaling can induce mobilization of murine hematopoietic progenitors (20). However, proper genetic evaluation of the function of aPKCζ and aPKCλ in mammalian HSC fate mapping in vivo has been unexplored because of the unavailability of tissue-specific conditional knock-out animal models.

To elucidate the individual and combined role of aPKCζ and aPKCλ signaling in mammalian HSC activity and hematopoiesis, we generated triple-transgenic, hematopoietic-specific or inducible knock-out mice. We have found that the polarization and activity of mammalian HSC do not depend on aPKCζ and aPKCλ in vivo. Therefore, an alternative tissue-specific and context-dependent signaling complex may be involved in controlling stem-cell fate.

Results

aPKC ζ ^{-/-} Mice Have Normal Steady-State Hematopoiesis. Because aPKCζ has been implicated in determining HSC activity (19, 20), to study the effects of genetic deletion of aPKCζ in mammalian hematopoietic systems in vivo, we analyzed aPKC $\zeta^{-/-}$ mice (22). $a P K C \zeta^{-/-}$ mice are viable and are born following a Mendelian ratio of inheritance (22). We hypothesized that if aPKCζdependent polarity signals regulate HSC and progenitor (HSC/P) activity, we could expect an alteration in the steady-state hematopoiesis of aPKCζ-deficient mice. However, aPKCζ-deficient mice had normal bone marrow (BM) cellularity, similar to their littermates (Fig. 1A). There was no effect of loss of aPKCζ on BM-HSC/P frequency or content (Fig. 1 B–D) or in the frequency of colony-forming progenitors (CFU-C) present in the BM and spleen (Fig. 1 E and F). This finding suggested that aPKC ζ does not regulate steady-state hematopoiesis.

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aPKCζ−/[−] HSCs Have Normal Self-Renewal and Lineage Repopulation Activity. If aPKCζ-dependent polarity signals regulate HSC selfrenewal and differentiation, we would expect an alteration in HSC repopulation ability during serial competitive BM transplantation, the gold-standard assay used to quantitate HSC selfrenewal and differentiation in vivo (23–25). To test this hypothesis, we serially transplanted BM nucleated cells (BMNC) from
CD45.2⁺ WT or CD45.2⁺ aPKC $\zeta^{-/-}$ mice along with CD45.1⁺ B6.SJLPtprca Pep3b/BoyJ competitor cells into lethally irradiated $CD45.1^+$ B6.SJL^{Ptprca Pep3b/BoyJ} primary and secondary recipient mice (Fig. 1G). The donor-derived $CD45.2^+$ chimera level was monitored in the peripheral blood and in the BM in both the primary and secondary recipient mice (Fig. 1H). HSC-repopulation ability did not change in either the primary or secondary recipients because of the deficiency of aPKC ζ (Fig. 1 I and J). There was no change in CD45.2⁺ myeloid (CD11b⁺ graft) or lymphoid (B220⁺ graft and $CD3^+$ graft) cell reconstitution in the recipient BM (Fig. (K) , suggesting that aPKC ζ does not regulate HSC self-renewal and lineage repopulation in vivo.

Because treatment with 5-fluorouracil (5-FU), a potent myeloablative agent, rapidly kills cycling BM cells and brings quiescent HSC into cell division (26, 27), 5-FU administration is used to assess HSC self-renewal and repopulation in vivo (28). Secondary recipient mice, after 16 wk of engraftment, were further challenged with serial administration of 5-FU (on days 0 and 40), and $CD45.2^+$ chimera was monitored over 2 mo (Fig. 1 G and L). There was no significant change in the level of $CD45.2^+$ chimera (Fig. 1L), suggesting that aPKC ζ is dispensable in HSC repopulation activity and stressed hematopoiesis in vivo.

Constitutive and Hematopoietic-Specific (Vav1-Cre) Genetic Deletion of aPKCζ and aPKCλ Does Not Affect Steady-State Hematopoiesis. aPKCζ shares a homologous domain architecture with aPKCλ (29). We wondered whether the loss of function of aPKCζ is

Fig. 1. Deficiency of aPKCζ does not affect HSC activity or steady-state hematopoiesis. (A) Absolute numbers of nuclear cells present in the BM of WT or aPKC $\zeta^{-/-}$ mice (n = 3– 4 mice per group). (B) Representative flow cytometry (FACS) contour diagram showing the frequency of hematopoietic stem cell and progenitors (HSC/P) present in the BM of WT or aPKC $\zeta^{-/-}$ mice. (C) Absolute numbers of HSCs present in the BM of WT or aPKC $\zeta^{-/-}$ mice (n = 3–4 mice per group). (D) Absolute numbers of hematopoietic progenitors (HPC) present in the BM of WT or aPKC $\zeta^{-/-}$ mice (n = 3– 4 mice per group). (E) Absolute numbers of colony-forming progenitors (CFU-C) present in the BM of WT or aPKC $\zeta^{-/-}$ mice ($n = 3-4$ mice per group). (F) Absolute numbers of colony-forming progenitors (CFU-C) present in the spleen of WT or aPKC $\zeta^{-/-}$ mice (n = 3–4 mice per group). (A–F) Error bars represent SD. (G) Experimental set up: 2×10^6 BM cells from CD45.2⁺ WT or aPKC $\zeta^{-/-}$ mice were mixed with 2 \times 10⁶ CD45.1⁺ B6.SJL^{Ptprca Pep3b/BoyJ} WT BM cells and competitively (1:1 ratio) transplanted into lethally irradiated primary, secondary, and tertiary recipient mice and peripheral blood chimera was monitored at different time points. After 16 wk of engraftment, secondary recipient mice were further challenged (intraperitoneally) with two cycles (day 0 and day 40) of 5-FU injections (150 mg/kg body weight) and peripheral blood donor-derived chimera (CD45.2⁺) was monitored for 8 to 10 wk. (H) Representative FACS contour diagram showing competitive chimera of CD45.2⁺ WT or aPKCζ^{-/-} cells serially transplanted into lethally irradiated CD45.1⁺ primary recipient mice from G. (I) Change in donor chimera (CD45.2⁺) in the peripheral blood of primary recipient mice ($n = 6-8$ mice per group). (J) Change in donor chimera (CD45.2⁺) in the peripheral blood of secondary recipient mice ($n = 7$ mice per group). (K) Frequency of lineage-repopulating myeloid (CD45.2⁺CD11b⁺) or B (CD45.2⁺B220⁺) or T (CD45.2⁺CD3⁺) cells present in the BM of the primary recipient mice ($n = 6-8$ mice per group). (L) Change in donor chimera (CD45.2⁺) in the peripheral blood of 5-FU–treated secondary recipient mice $(n = 7$ mice per group). $(l-L)$ Error bars represent SEM.

compensated by aPKCλ in vivo. aPKCλ-deficient mice die at very early embryonic stages before hematopoiesis occurs (12). Mice with aPKC λ ^{f/f} alleles (12) were crossed with Vav1-Cre to generate WT; Vav1-Cre or aPKC $\zeta^{-/-}$; Vav1-Cre or aPKC $\lambda^{\Delta/\Delta}$; Vav1-Cre or $aPKC\zeta^{-/-}\lambda^{A/\Delta}$; Vav1-Cre mice. Expression of Cre recombinase under the promoter of Vav1, a guanine nucleotide exchange factor for Rho-GTPases, induces deletion of aPKCλ alleles, specifically in the fetal and adult hematopoietic system (26, 30).

Genomic DNA PCR revealed complete deletion of aPKCλ
alleles in purified HSCs (Lineage[−]Sca1⁺c-Kit⁺CD34[−]Flk2[−]) from $aPKC\lambda^{A/A}$; Vav1-Cre BM (Fig. 24). We did not detect mRNA expression of aPKCλ in the HSCs isolated from aPKCλ^{Δ/Δ}; Vav1-Cre mice by RT-PCR (Fig. 2B). aPKC $\lambda^{\Delta/\Delta}$; Vav1-Cre and $aPKC\zeta^{-1-\lambda\Delta/\Delta}$; Vav1-Cre mice were born following Mendelian ratio of inheritance [\(Fig. S1](http://www.pnas.org/lookup/suppl/doi:10.1073/pnas.1103132108/-/DCSupplemental/pnas.201103132SI.pdf?targetid=nameddest=SF1)), suggesting that the expression of aPKCζ and aPKCλ is not crucial during fetal hematopoiesis (31).

Because both fetal and adult hematopoiesis use distinct microenvironments and differ in their cell cycle activation (32), we speculated that aPKCζ and aPKCλ may be required for normal HSC activity in adult BM. However, aPKCζ- and aPKCλ-deficient mice had normal BM cellularity, similar to their littermates (Fig. 2C). Loss of aPKCζ and aPKCλ had no detectable effect on BM-HSC/P content (Fig. 2 D and E) or the frequency of BM colonyforming progenitors (Fig. 2F). aPKCζ- and aPKCλ-deficient HSCs showed normal apoptosis (Fig. 2G) and proliferation in vivo (Fig. 2 H and I), similar to WT cells, which suggests that expression of aPKCζ and aPKCλ does not regulate steady-state hematopoiesis.

Deficiency of aPKCζ and aPKCλ Do Not Affect Hematopoietic Recovery from 5-FU–Induced Myeloablative Stress. We examined the hematopoietic response to forced cell cycle activation of quiescent HSC in vivo of WT; Vav1-Cre or aPKCζ−/−; Vav1-Cre or $aPKC\lambda^{\Delta/\Delta}$; Vav1-Cre or $aPKC\zeta^{-/-}\lambda^{\Delta/\Delta}$; Vav1-Cre BM. When challenged with 5-FU, aPKCζ- and aPKCλ-deficient mice elicited

Fig. 2. Constitutive and hematopoietic system-specific loss of aPKCζ/λ does not affect steady-state hematopoiesis. (A) Genomic DNA PCR analysis of WT, floxed, and aPKCλ-deleted alleles in Lineage[−]Sca1⁺c-Kit⁺ (LSK) CD34[−]Flk2[−] cells isolated from the BM of WT; Vav1-Cre or aPKC $\lambda^{\Delta/\Delta}$;Vav1-Cre mice. (B) mRNA expression (RT-PCR) analysis of aPKC λ in LSKCD34⁻⁻Flk2⁻ cells isolated from the BM of WT; Vav1-Cre or aPKCλ^{Δ/Δ}; Vav1-Cre mice. (C) Absolute numbers of nuclear cells present in the BM of WT; Vav1-Cre or aPKCζ^{-/-}; Vav1-Cre or aPKCλ^{Δ/Δ}; Vav1-Cre or aPKC $\zeta^{-/-}$ aPKC $\lambda^{\Delta/\Delta}$;Vav1-Cre mice ($n=3$ –4 mice per group). (D) Absolute numbers of HSCs present in the BM of WT; Vav1-Cre or aPKC $\zeta^{-/-}$ Absolute numbers of HSCs present in the BM of WT; Vav1-Cre or aPKCζ^{-/–};
Vav1-Cre or aPKCλ^{Δ/Δ}; Vav1-Cre or aPKCζ^{-/–} aPKCλ^{Δ/Δ}; Vav1-Cre mice (*n* = 3 mice per group). Error bars represent SD. (E) Absolute numbers of HPC present in the BM of WT; Vav1-Cre or aPKCζ^{-/--};Vav1-Cre or aPKCλ^{Δ/Δ};Vav1-Cre or aPKC $\zeta^{-/-}$ aPKC $\lambda^{\Delta/\Delta}$; Vav1-Cre mice (n = 3 mice per group). (F) Absolute numbers of colony-forming progenitors (CFU-C) present in the BM of WT; Vav1-Cre or aPKCζ^{−/−}; Vav1-Cre or aPKCλ^{∆/∆}; Vav1-Cre or aPKCζ^{−/−} aPKCλ^{∆/∆}; Vav1-Cre mice $(n = 3$ mice per group). (G) Apoptosis (active Caspase 3) of BM-LSK cells in WT; Vav1-Cre or aPKCζ^{−/−}; Vav1-Cre or aPKCλ^{∆/∆}; Vav1-Cre or aPKCζ^{−/−} aPKCλ^{∆/∆}; Vav1-Cre mice in vivo ($n = 3$ mice per group). (H) Representative FACS contour diagram showing BrdU (300 μg i.p. injection, 45-min pulse) incorporation into BM-LSK cells in WT; Vav1-Cre or aPKCζ^{-/−}; Vav1-Cre or aPKCλ^{∆/∆}; Vav1-Cre or aPKC $\zeta^{-/-}$ aPKC $\lambda^{\Delta/\Delta}$; Vav1-Cre mice in vivo. (/) Proliferation of BM-LSK cells in WT; Vav1-Cre or aPKCζ^{−/−}; Vav1-Cre or aPKCλ^{∆/Δ}; Vav1-Cre or aPKCζ^{−/−} aPKCλ^{∆/Δ}; Vav1-Cre mice in vivo ($n = 3$ mice per group). (A– θ) Error bars represent SD.

a normal hematopoietic recovery response similar to littermate control mice [\(Fig. S2](http://www.pnas.org/lookup/suppl/doi:10.1073/pnas.1103132108/-/DCSupplemental/pnas.201103132SI.pdf?targetid=nameddest=SF2)). There were no differences among the groups in the peripheral blood leukocyte, neutrophil, or platelet counts (Fig. $S2$ $A-C$), nor in the evolution of reticulocyte frequency in the peripheral blood (Fig. $S2 D$ and E). No differences were seen in the red blood cell counts in WT and aPKCζand aPKC λ -deficient mice ([Fig. S2](http://www.pnas.org/lookup/suppl/doi:10.1073/pnas.1103132108/-/DCSupplemental/pnas.201103132SI.pdf?targetid=nameddest=SF2)F). These data suggest that aPKCζ and aPKCλ are also dispensable during stressed adult hematopoiesis.

Inducible (Mx1-Cre) Genetic Deletion of aPKCζ and aPKCλ Does Not Affect Steady-State Hematopoiesis. We thought that the constitutive and hematopoietic-specific deletion of aPKCλ, as seen in the Vav1-Cre mice, might provide a selection advantage during development through functional compensation toward normal HSC activity in the aPKCλ-dependent downstream signaling pathway in vivo. We therefore wanted to specifically delete aPKCλ in the adult hematopoietic system using IFN-inducible Mx1-Cre mice (23). Mice with aPKC $\lambda^{f/f}$ alleles (12), were crossed with Mx1-Cre transgenic mice similar to other inducible models of gene deletion (23) and WT; Mx1-Cre or aPKC $\zeta^{-/-}$; Mx1-Cre or aPKC $\lambda^{f/f}$; Mx1-Cre or aPKC $\zeta^{-/-}\lambda^{f/f}$; Mx1-Cre mice were generated ([Fig.](http://www.pnas.org/lookup/suppl/doi:10.1073/pnas.1103132108/-/DCSupplemental/pnas.201103132SI.pdf?targetid=nameddest=SF3) [S3](http://www.pnas.org/lookup/suppl/doi:10.1073/pnas.1103132108/-/DCSupplemental/pnas.201103132SI.pdf?targetid=nameddest=SF3)A). Administration of polyinosinic:polycytidylic acid (pI-pC) resulted in the loss of expression of aPKCλ in the Lineage[−] Sca1+c-Kit⁺CD34⁻Flk2⁻ cells that were isolated from aPKC λ ^{f/f}; Mx1-Cre mice, but not those from WT; Mx1-Cre mice ([Fig. S3](http://www.pnas.org/lookup/suppl/doi:10.1073/pnas.1103132108/-/DCSupplemental/pnas.201103132SI.pdf?targetid=nameddest=SF3)B). Genomic DNA PCR analysis confirmed that there was complete deletion of aPKCλ alleles in the adult hematopoietic system ([Fig. S3](http://www.pnas.org/lookup/suppl/doi:10.1073/pnas.1103132108/-/DCSupplemental/pnas.201103132SI.pdf?targetid=nameddest=SF3)C).

If aPKCζ/λ-dependent polarity signals regulate HSC activity, we should expect an alteration in the steady-state hematopoiesis in aPKCζ- and aPKCλ-deficient mice. However, there was no detectable effect of the loss of aPKCζ and aPKCλ on the content of BM-HSC/P subpopulations when assessed by immunofluores-cence or functional assays ([Fig. S3](http://www.pnas.org/lookup/suppl/doi:10.1073/pnas.1103132108/-/DCSupplemental/pnas.201103132SI.pdf?targetid=nameddest=SF3) $D-F$). Likewise, we did not detect any effect of aPKCζ and aPKCλ deficiency in BM-CD11b⁺ myeloid cell or $B220^+$ B-cell or $CD3^+$ T-cell ([Fig. S3](http://www.pnas.org/lookup/suppl/doi:10.1073/pnas.1103132108/-/DCSupplemental/pnas.201103132SI.pdf?targetid=nameddest=SF3) $G-I$) contents. In addition, aPKCζ- and aPKCλ-deficient HSCs showed normal apoptosis and proliferation (Fig. $S3J$ and K) in vivo, similar to the WT cells, suggesting that the inducible genetic inactivation of aPKCζ and aPKCλ in the adult hematopoietic system does not significantly affect steady state HSC activity or hematopoiesis.

Inducible Deletion of aPKCλ and Constitutive Deficiency of aPKCζ Does Not Affect HSC Self-Renewal and Lineage Repopulation Activity. Despite the normal content, proliferation, and survival of BM-HSC/ Ps in the primary mice, it was possible that the self-renewal ability of long-term HSC (LT-HSC) in the adult hematopoietic system is regulated by aPKCζ and aPKCλ. Compensatory mechanisms during development in Vav1-Cre; aPKC $\zeta^{-/-} \lambda^{f/f}$ mice and the existence of a subtle HSC phenotype in Mx1-Cre; aPKC $\zeta^{-/-}\lambda^{f/f}$ might possible explanations for our findings. To address whether the deficiency of aPKCζ and aPKCλ resulted in a loss of LT-HSC activity, we first analyzed whether aPKCζ and aPKCζ-deficient HSCs are able to reconstitute the hematopoietic system. To test this, we transplanted BM cells from CD45.2+ WT; Mx1-Cre or $aPKC\lambda^{f/f}$; Mx1-Cre or $aPKC\zeta^{-/-}\lambda^{f/f}$; Mx1-Cre mice into lethally irradiated CD45.1⁺ B6.SJLPtprca Pep3b/BoyJ mice. After 4 wk of engraftment, the recipient mice were treated with pI-pC and their peripheral blood $CD45.2^+$ chimera monitored for 16 wk [\(Fig.](http://www.pnas.org/lookup/suppl/doi:10.1073/pnas.1103132108/-/DCSupplemental/pnas.201103132SI.pdf?targetid=nameddest=SF4) [S4](http://www.pnas.org/lookup/suppl/doi:10.1073/pnas.1103132108/-/DCSupplemental/pnas.201103132SI.pdf?targetid=nameddest=SF4)A). pI-pC treatment resulted in the complete excision of donorderived aPKCλ alleles in the transplanted mice [\(Fig. S4](http://www.pnas.org/lookup/suppl/doi:10.1073/pnas.1103132108/-/DCSupplemental/pnas.201103132SI.pdf?targetid=nameddest=SF4)B). However, the deficiency of aPKCζ and aPKCλ did not affect the evolution of myeloid cell $(CD11b²$ cells gated on $CD45.2⁺$ graft) and B-cell (B220⁺ cells gated on $CD45.2^+$ graft) content in the peripheral blood of the recipient mice (Fig. $S4 \dot{C}$ and D). These results indicate that aPKCζ and aPKCλ are not required for hematopoietic reconstitution.

Secondly, to analyze the ability of aPKC-deficient cells to compete with WT cells, we generated mixed chimeric animals by transplanting BMNC from CD45.2⁺ WT; Mx1-Cre or aPKC λ^{if} ; Mx1-Cre or $aPKC\zeta^{-7} \lambda^{f/f}$; Mx1-Cre mice along with CD45.1⁺ B6. SJL Ptprca Pep3b/BoyJ competitor cells into lethally irradiated $CD45.1^+$ B6.SJL^{Ptprca Pep3b/BoyJ} mice (Fig. 3A). After 4 wk of engraftment, the recipient mice were treated with pI-pC (administered on every other day for 14 d) and peripheral blood $CD45.2^+$ chimera was monitored for another 16 wk (Fig. 3A). BMNC from primary recipient mice were serially transplanted into lethally irradiated CD45.1⁺ B6.SJLPtprca Pep3b/BoyJ secondary and tertiary recipients and peripheral blood $CD45.2^+$ cell chimera was determined (Fig. 3A).

The aPKCζ- and aPKCλ-deficient HSCs did not demonstrate a defect in HSC self-renewal (Fig. $3A$ and B) or skewed myeloid/ lymphoid lineage repopulation (Fig. 3 E–H and [Fig. S5\)](http://www.pnas.org/lookup/suppl/doi:10.1073/pnas.1103132108/-/DCSupplemental/pnas.201103132SI.pdf?targetid=nameddest=SF5) when assayed in the setting of serial competitive repopulation. Com-petitively transplanted WT or aPKCζ- and aPLCλ-deficient tertiary recipients showed similar levels of hematopoietic engraftment (Fig. 3I) without bias in the level of gene deletion when compared with primary or secondary recipients [\(Fig. S6](http://www.pnas.org/lookup/suppl/doi:10.1073/pnas.1103132108/-/DCSupplemental/pnas.201103132SI.pdf?targetid=nameddest=SF6) A–C). These data suggest that inducible loss of function of aPKCζ and aPKCλ in the adult hematopoietic system does not affect HSC self-renewal and lineage repopulation activity during serial BM transplantation.

aPKCζ and aPKCλ Are Dispensable in the HSC/P-Interaction with the Hematopoietic Microenvironment. BM-HSC/P retention and mobilization are exquisitely controlled by a delicate network of molecular signals, which are generated by the hematopoietic microenvironment (33, 34). We analyzed whether the loss of function of aPKCζ and aPKCλ could result in defective interaction of HSC/P with the hematopoietic microenvironment and result in impaired HSC/P trafficking. Deficiency of aPKCζ and aPKCλ did not change the content of HSC/Ps present in the peripheral blood (Fig. 4A). aPKCζ- and aPKCλ-deficient mice also showed a normal response to G-CSF–induced HSC/P mobilization similar to the WT mice ([Fig. S7](http://www.pnas.org/lookup/suppl/doi:10.1073/pnas.1103132108/-/DCSupplemental/pnas.201103132SI.pdf?targetid=nameddest=SF7)), suggesting that aPKC ζ and aPKC λ do not regulate HSC/P mobilization.

As mentioned earlier, self-renewal, proliferation, and survival of engrafted aPKCλ or aPKCζ/λ-deficient HSC was not significantly different from their WT counterparts. It is possible, however, that primary deficiency of aPKCζ and aPKCλ results in defective homing/engraftment in the transplanted mice. If this is true, in such case we would expect in the development of a BM failure syndrome. Therefore, $CD45.2^+$ WT or aPKC ζ - and aPKCλ-deficient BM were noncompetitively transplanted into lethally irradiated CD45.1⁺ B6.SJL^{Ptprca Pep3b/BoyJ} recipient mice and peripheral blood lineage repopulation was determined (Fig.

4B). Despite the complete deletion of aPKCλ alleles in the transplanted mice ([Fig. S8\)](http://www.pnas.org/lookup/suppl/doi:10.1073/pnas.1103132108/-/DCSupplemental/pnas.201103132SI.pdf?targetid=nameddest=SF8), we did not see significant changes in the generation of donor-derived myelopoiesis or lymphopoiesis in the peripheral blood in the recipient mice (Fig. $\hat{4}$ C–E). Furthermore, competitive repopulation activities of WT or aPKCζand aPKCλ-deficient HSCs (Fig. 4F) were also similar (Fig. 4G). This finding confirmed that the critical phases of homing, survival, proliferation, and differentiation of aPKCζ- and aPKCλ-deficient HSC/P were preserved.

Finally, adhesion to the C terminus of fibronectin (CH-296), an important constituent of the BM microenvironment and migration toward CXCL12, a physiological chemoattractant, were similar for WT and aPKCζ- and aPKCλ-deficient HSC/Ps (Fig. 4 H and I). Collectively, these results indicate that aPKC ζ and aPKCλ are dispensable for HSC/P trafficking within the hematopoietic microenvironment.

aPKCs Are Not Polarized in HSC and Deficiency of aPKCζ/λ Does Not Impair HSC Polarization In Contact with the Hematopoietic Microenvironment. Our genetic data indicated that aPKCζ/λ does not play a functional role in the critical phases of mammalian HSC engraftment and hematopoiesis. We evaluated whether aPKCζ has a polarized pattern of distribution in the HSCs. In other model systems, asymmetric cell division is preceded by the activation of the aPKCζ-Par6-Par3 polarity complex and the realignment of the microtubule cytoskeleton (7, 11). We analyzed the distribution of aPKCζ and Par6 in WT HSCs in vitro after overnight culture on fibronectin, a major component of the hematopoietic microenvironment, which is responsible for integrin polarization and signaling (35). Although the intracellular locali-

engraftment, and multilineage differentiation. (A) Experimental set up. The 2 \times 10⁶ BM cells from CD45.2⁺ WT; Mx1-Cre or aPKC $\lambda^{\mathsf{f/f}}$; Mx1-Cre or aPKC $\zeta^{-/-}\lambda^{\mathsf{f/f}}$; Mx1-Cre mice were mixed with 2×10^6 CD45.1⁺ B6.SJL^{Ptprca Pep3b/BoyJ} WT BM cells and competitively (1:1 ratio) transplanted into lethally irradiated primary mice. Serial transplantation of 1×10^7 pooled BM cells from primary mice was performed into secondary and tertiary recipient mice. After BM engraftment (4 wk), aPKCλ deletion was induced by administration of pI-pC and CD45.2⁺ chimera was monitored in different time points. (B) Evolution (normalized to baseline) of CD45.2⁺ chimera in peripheral blood of primary recipient mice ($n = 7-9$ mice per group). (C) Evolution (normalized to baseline) of CD45.2⁺ chimera in peripheral blood of secondary recipient mice ($n = 8$ mice per group). (D) Representative FACS contour diagram showing CD45.2⁺CD11b⁺-myeloid and CD45.2+B220+-B-lymphoid cells during HSC engraftment. (E) Evolution (normalized to baseline) of myeloid-cell chimera $(CD45.2⁺$ cells gated on total $CD11b⁺$ graft) in the peripheral blood of primary recipient mice ($n = 7-9$ mice per group). (F) Evolution (normalized to baseline) of myeloid-cell chimera (CD45.2⁺ cells gated on total CD11b⁺ graft) in the peripheral blood of secondary recipient mice ($n = 8$ mice per group). (G) Evolution (normalized to baseline) of B-cell chimera (CD45.2+ cells gated on total B220⁺ graft) in the peripheral blood of primary recipient mice ($n = 7-9$ mice per group). (H) Evolution (normalized to baseline) of B-cell chimera (CD45.2+ cells gated on total B220⁺ graft) in the peripheral blood of secondary recipient mice ($n = 8$ mice per group). (B, C, E-H) Error bars represent SEM. (/) Normalized CD45.2⁺ chimera in peripheral blood of tertiary recipient mice at 8 wk after transplantation. $n = 7-9$ mice per group. Error bars represent SD.

Fig. 3. aPKCζ and aPKCλ does not regulate HSC self-renewal,

Fig. 4. Deficiency of aPKCζ and aPKCλ does not affect interaction of HSC/P within the BM microenvironment. (A) Absolute numbers of CFU-Cs present in the peripheral blood of WT; Vav1-Cre or PKC $\zeta^{-/-}$; Vav1-Cre or aPKC $\lambda^{\Delta\Delta}$; Vav1-Cre or aPKC $\zeta^{-/-}\lambda^{\Delta/\Delta}$; Vav1-Cre mice ($n = 6$ –16 mice per group). Data represent average of two different experiments. (B) Experimental set up. 3 \times 10⁶ BMNCs CD45.2⁺ WT; Vav1-Cre or PKCζ^{-/-}; Vav1-Cre or aPKCλ^{Δ/Δ}; Vav1-Cre or aPKCζ^{-/-}λ^{Δ/λ}; Vav1-Cre mice, were noncompetitively transplanted into
lethally irradiated CD45.1⁺ B6.SJL^{Ptprca Pep3b/BoyJ WT mice. CD45.2⁺ chimera} was monitored in the peripheral blood of the recipient mice. (C) Absolute numbers of CD45.2⁺CD11b⁺-myeloid cells present in the peripheral blood of CD45.1⁺ B6.SJL^{Ptprca Pep3b/BoyJ} mice after 8 wk of transplantation from B (n = 6-9 mice per group). (D) Absolute numbers of CD45.2⁺B220⁺ B-lymphoid cells present in the peripheral blood of CD45.1⁺ B6.SJL^{Ptprca Pep3b/BoyJ} mice after 8 wk of transplantation from $B(n = 6-9$ mice per group). (E) Absolute numbers of CD45.2*CD3* T-lymphoid cells present in the peripheral blood of CD45.1* B6.SJL^{Ptprca Pep3b/BoyJ} mice after 8 wk of transplantation from B ($n = 6-9$ mice per group). (C–E) Error bars represent SD. (F) Experimental set up. The 2×10^6 BM cells from CD45.2⁺ WT; Vav1-Cre or aPKCλ^{∆/Δ}; Vav1-Cre or aPKCζ^{-/-}λ^{∆/Δ}; Vav1-Cre mice were mixed with 2×10^6 CD45.1⁺ B6.SJL^{Ptprca Pep3b/BoyJ} WT BM cells and competitively (1:1 ratio) transplanted into lethally irradiated primary mice. (G) Evolution of CD45.2⁺ chimera in the peripheral blood of recipient mice from $F(n = 7-10$ mice per group). (H) Adhesion (performed in triplicate) of WT; Vav1-Cre, PKCζ¯ ; Vav1-Cre, aPKCλ $^{\Delta\Delta}$; Vav1-Cre or aPKCζ¯ $^{\!\!\!-1}\lambda^{\Delta\Delta}$; Vav1-Cre HSC/Ps to fibronectin (CH-296) in vitro. Error bars represent SEM. (/) Migration (performed in triplicate) of WT; Vav1-Cre, PKCζ^{−/−}; Vav1-Cre, aPKCλ^{∆∆}; Vav1-Cre or aPKC $\zeta^{-/-}\lambda^{\Delta/\Delta}$; Vav1-Cre HSC/Ps toward CXCL12 in vitro. (G, I) Error bars represent SEM.

zation of tubulin, a major constituent of the microtubular network, was highly polarized in isolated HSCs, which were in contact with fibronectin, the distribution of aPKCζ and Par6 was not proportional to the frequency of tubulin polarized HSCs (Fig. 5 A–C). In addition, the deficiency of aPKCζ/λ expression in HSCs did not impair microtubular network polarization (Fig. 5 D and E), indicating that aPKCζ/λ expression is dispensable in HSC polarization in the conditions tested. Together, these data indicate that aPKCζ/λ-downstream cell polarity signaling is dispensable in the determination of mammalian HSC activity and suggests that alternative molecular complexes are involved in stem cell asymmetry.

Discussion

We have explored whether aPKCζ and aPKCλ control HSC polarization in response to developmental and environmental cues, and if HSC polarization is required to produce appropriate numbers of stem cells and differentiated daughter cells. Previous reports using alternative RNA silencing or protein neutralization methods, which are not deprived of potential nontarget effects, suggested that the polarity-determining kinase aPKCζ, in particular, can positively regulate HSC activity (19, 20). To elucidate whether aPKCζ and aPKCλ play a role in HSC activity in vivo, we used combined constitutive, inducible, or tissue-specific gene deletion in vivo. A deficiency of aPKCζ, induced through targeted homologous recombination of embryonic stem cells, did not affect

Fig. 5. aPKCζ and Par6 are not polarized in HSC and aPKCs do not regulate HSC polarization. (A) Immunofluorescence pictures showing aPKCζ (red) along with tubulin (blue) localization in LT-HSCs. (Ai, Aii) aPKCζ is detected diffusely in the cell cytoplasm without any specific asymmetric distribution in most of LT-HSCs. (Aiii, Aiv) Most of LT-HSCs display a highly polarized distribution of the microtubule network. (Scale bar, 5 μ m.) (B) Immunofluorescence pictures showing Par6 (green) along with tubulin (blue) localization in LT-HSCs. (Bi, Bii) Par6 is distributed throughout the cell body and beneath the cell membrane. Par6 localization is not polarized in most of LT-HSCs. (Biii, Biv) LT-HSCs display a highly polarized distribution of the microtubule network. (Scale bar, 5 μm.) (C) Bar diagram showing quantification of immunofluorescence analysis. Fifty to 100 LT-HSCs were singularly analyzed in each experiment ($n = 4$). Error bars represent SEM of the number of tubulin (79.7%), aPKCζ (21.53%), and Par6 (41.24%) polarized cells scored per sample. $*P < 0.05$. (D) Immunofluorescence pictures showing tubulin (blue) localization in LT-HSCs isolated from wild type or aPKCζ- and aPKCλ-deficient mice. Most of LT-HSCs display a highly polarized distribution of the microtubule network ($n = 3-4$ mice per group) (Scale bar, 5 μ m.) (E) Bar diagram showing quantification of immunofluorescence analysis ($n = 30-50$ LT-HSCs from individual mouse). Error bars represent SD.

HSC activity and gross hematopoiesis in vivo, and there was no change in the frequency or content of HSC/P and colony-forming progenitors present in the BM. aPKC $\zeta^{-/-}$ HSCs showed normal survival and proliferation similar to the WT HSC in vivo, and competitive BM transplantation experiments suggested that HSC self-renewal and lineage repopulation ability in primary, secondary, and tertiary recipient mice was not altered because of deficiency of aPKCζ. aPKCζ was also dispensable in 5-FU–induced stressed hematopoiesis in vivo. These results indicate that aPKCζ does not play decisive role in murine HSC fate determination in vivo, contrary to previous reports (19, 20).

An alternative hypothesis was that aPKCλ might compensate for the loss of aPKCζ in vivo. Therefore, we determined the effect of a combined deletion of aPKCζ and aPKCλ in HSC in vivo. Because deletion of aPKCλ results in embryonic lethality (12), we generated mice that had constitutive hematopoietic-specific (Vav1-Cre) (26) and hematopoietic-inducible (Mx1-Cre) (23) deletion of aPKCλ. However, the loss of function of aPKCζ and aPKCλ, using Vav1-Cre and Mx1-Cre mice models, did not affect HSC activity during steady state and 5-FU–induced stressed hematopoiesis. Inducible genetic deletion of aPKCζ and aPKCλ in the adult hematopoietic system also did not affect HSC selfrenewal and lineage repopulation activity. In addition, aPKCζ and aPKCλ were also dispensable for HSC/P homing, retention, adhesion, and migration, which are important parameters for HSC/P interaction within the hematopoietic microenvironment. Furthermore, neither a hematopoietic-specific deficiency of aPKCλ nor a combined deficiency of aPKCζ/λ affected embryonic lethality because of hematopoietic failure, and the aPKCζ/λ-deficient mice were born following a Mendelian ratio of inheritance. This finding is of interest because fetal hematopoiesis requires frequent cell cycling of HSCs and depends on a functionally different hematopoietic microenvironment compared with adult BM hematopoiesis (31, 32). Together, these data indicate that aPKCζ and aPKCλ do not regulate mammalian HSC activity in vivo.

HSC fate may depend on their polarized distribution, particularly when they are in contact within the BM stem-cell microenvironment. Unlike intestinal epithelial cells, which have distinct apical and basolateral cell surfaces, or intrinsically polarized neuronal cells, HSC polarization may not be morphologically appar-

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ent. Interestingly, in other systems, integrin-binding regulates cell polarization and determines cell division (11). In the BM, the vast majority of HSC are quiescent (32). We developed an in vitro assay where fibronectin-dependent forced microtubular network polarization could be efficiently assessed. In these conditions, our genetic models indicate that integrin-dependent binding of HSC to the hematopoietic microenvironment fails to polarize aPKCs and Par6, and a combined deficiency of aPKCζ and aPKCλ did not prevent polarization. This finding suggests that the existence of aPKCζand aPKCλ-dependent stem cell polarity is not a functional determinant for mammalian HSC activity. Conversely, aPKC signaling is implicated in the regulation of mammalian T-cell activity (12, 13), indicating that the contribution of aPKC ζ/λ signaling in mammalian HSCs and in terminally differentiated cells is functionally distinguishable. These data underscore the importance of cell, tissue-microenvironment, and context-dependent regulation of evolutionary conserved signaling pathways.

Given that aPKCζ and aPKCλ are the core kinases within the cell-polarity complex in invertebrates and in other mammalian cells (7, 11, 36), it is an unexpectedly interesting finding that the activity of mammalian HSCs is not dependent on them, and may indicate the existence of an alternative signaling network defining HSC asymmetry with redundancy in mammalian stem cell polarity determination.

Materials and Methods

Information on animals, deletion of aPKCλ, LT-HSC repopulation, 5-FU ad-ministration may be found in [SI Materials and Methods](http://www.pnas.org/lookup/suppl/doi:10.1073/pnas.1103132108/-/DCSupplemental/pnas.201103132SI.pdf?targetid=nameddest=STXT). For HSC/P assays, see [SI Materials and Methods](http://www.pnas.org/lookup/suppl/doi:10.1073/pnas.1103132108/-/DCSupplemental/pnas.201103132SI.pdf?targetid=nameddest=STXT). RT-PCR analysis and statistical analysis are included in [SI Materials and Methods](http://www.pnas.org/lookup/suppl/doi:10.1073/pnas.1103132108/-/DCSupplemental/pnas.201103132SI.pdf?targetid=nameddest=STXT).

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