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Figure legends

Figure 1. Experimental design for clonal and quantitative evaluation of self-renewal (A) A mouse is reconstituted with a single HSC. BM cells isolated from this mouse are transplanted into multiple mice. Upon secondary transplantation, a population-type assay (B) and a limiting dilution-type assay (C), both based on competitive repopulation, are performed. The number of RU in defined numbers of BM cells (such as 10^6 cells) can be determined from % chimerism obtained by FACS analysis of peripheral blood cells (B). The number of CRU in the BM can be calculated from the frequency of CRU in defined numbers of BM cells (C). The amount of RU as well as the number of CRU regenerated in BM by 1 CRU can be finally estimated.

Hiro-sensei, this doesn't make sense to me. How is "the amount of RU" different from "the number of RU"? Should this read "The number of RU (and the number of CRU) that 1 CRU regenerates in BM can thus be estimated"?

The MAS, an average RU per CRU, indicates the quality of regenerated CRU. Thus, RU, CRU, and MAS can be used as parameters for measurement of self-renewal activity in an individual HSC. Black circles indicate CRUs. a, cells derived from test cells; b, cells derived from competitor cells; c, cells derived from a host mouse; d, limiting dose of cells.

Figure 2. Increased frequency of HSCs in BM cells of Lnk-deficient mice
Graded numbers of BM cells from wild-type (WT) or Lnk-deficient (Lnk-KO) mice were transplanted together with 2×10^5 competitor cells. The frequency of CRUs in BM cells of Lnk-deficient or wild-type mice was estimated to be one in 1,900 cells or one in 31,000 cells.

Figure 3. Expansion of HSC populations in Lnk-deficient mice
(A) Representative FACS profiles show Sca-1 and c-Kit expression in CD34⁺Lin⁻ BM cells (upper panels) and the SP in Lin⁻ BM cells (lower panels). The proportions of CD34⁺KSL cells or Lin⁻SP cells were remarkably increased in Lnk-deficient mice (right panels) as compared with those in wild-type mice (left panels). The percentages of gated cells among total BM cells are shown.
(B) The kinetics of development of CD34⁺KSL cells (upper panel) or Lin⁻SP cells (lower panel) are shown, along with mouse age.

Figure 4. Multilineage reconstitution by single CD34⁺KSL cells from wild-type or Lnk-deficient mice
A single CD34⁺KSL cell from wild-type (WT) or Lnk-deficient (Lnk KO) mice was mixed with 2×10^5 BM cells and transplanted into a lethally irradiated mouse. All myeloid and B- and T-lymphoid lineages were reconstituted with a single test donor cell 4 months after transplantation in 24 of 92 (26%) mice receiving wild-type cells and in 17 of 76 (22%) mice receiving Lnk-deficient cells. (A) The list of RU per cell calculated based on % chimerism for each CRU. The MAS (2.76 ± 3.84 , $n=17$) in Lnk-deficient repopulating cells was significantly higher than that (0.64 ± 0.80 , $n=24$) in wild-type repopulating cells ($p < 0.05$, unpaired t-test with Welch correction). (B) Data are graphically presented with myeloid and B- and T-lymphoid lineage constituents (M-RU, B-RU, and T-RU).

Figure 5. In vitro maintenance of Lnk-deficient HSCs
CD34⁺KSL cells (10-cell equivalent) from 8-week-old wild-type or Lnk-deficient mice were transplanted into a group of lethally irradiated mice along with 2×10^5 competitor cells (indicated as "Before"). Such cells were also co-cultured with OP-9 cells for 7 days, followed by transplantation (indicated as "After"). Recipient mice were analyzed 3 months after transplantation. The mean of RU per 10 CD34⁺KSL cells from Lnk-deficient mice (5.04 ± 5.34 , $n=9$) was significantly greater than that from wild-type mice (0.48 ± 0.32 , $n=9$) on Mann-Whitney testing ($P=0.0002$). 1 of 9 mice that

had received cultured wild-type cells showed reconstitution with test donor cells (0.04 RU/10 cells, n=1). 9 of 9 mice that had received cultured Lnk-deficient cells were reconstituted (3.40 ± 2.55 RU/10 cells, n=9).

Table 1. Increases of RU and CRU in BM of Lnk-deficient mice.

Mice	BM cells ($\times 10^7$)	RU/ 10^5 BM cells	CRU/ 10^5 BM cells (95% CI)	MAS (RU/cell)
WT B6	6.4 \pm 1.2 (n=8)*	1.0	3.2 (1.8-5.6)	0.3
Lnk KO B6	6.6 \pm 1.9 (n=5)	63.3 \pm 33.2 (n=5)	52.6 (31.3-83.3)	1.2

Wild-type (WT) and Lnk-deficient (Lnk-KO) B6 mice aged 8 weeks were compared. The numbers of bone marrow (BM) cells in the hindlimbs (two femora and two tibiae) and RU per 10^5 BM cells are expressed as mean \pm S.D. CRU per 10^5 BM cells is expressed as mean (95% confidence interval). Mean activity per stem cell (MAS) was calculated using the averages of RU and CRU. *, previously published data (Sudo et al., 2000).

Table 2. Self-renewal activity in single HSCs

Clones		Primary transplantation		Secondary transplantation				
Donor mice	No.	RU/cell	BM cells (x10 ⁶) /mouse	RU/10 ⁶ cells	RU/BM	CRU/10 ⁶ cells	CRU/BM	MAS (RU/cell)
WT	1	3.26	400	0.16	64	2.44	976	0.07
	2	1.57	245	0.25	61	3.03	742	0.08
	3	0.82	325	0.19	62	2.50	813	0.08
	4	0.69	285	0.18	51	2.17	619	0.08
	5	0.47	280	0.04	11	1.25	350	0.03
	6	0.28	275	0.14	39	1.56	429	0.09
Lnk	1	7.62	385	1.00	385	2.50	963	0.40
KO	2	3.38	545	1.60	872	5.56	3,030	0.29

The number of RU/cell was obtained from analysis of mice that were primary recipients of single CD34⁺KSL cells. The numbers of RU and CRU/10⁶ BM cells were obtained from analysis of mice that were secondary recipients of BM cells reconstituted with single CD34⁺KSL cells (clones) from wild-type or Lnk-deficient mice. The number of BM cells in a primary recipient mouse (BM cells/mouse) was calculated based on the minimal assumption that 20% of all BM cells are present in the femora and tibiae. The total numbers of secondary RU and CRU in BM per mouse (RU/BM and CRU/BM) were obtained by calculation of (BM cells)(RU/10⁶ cells)/10⁶ and (BM cells)(CRU/10⁶)/10⁶. The MAS is the number of RU divided by the number of CRU.

Supplemental Figure Legends

Supplemental Figure S1 *Lnk* is expressed in various hematopoietic lineages

Normalized cDNAs obtained from various populations in BM, spleen, and thymus of B6 mice underwent PCR analysis for *Lnk* sequences. Expression of *Lnk* was detected in stem and progenitor cell populations represented by CD34⁻KSL and CD34⁺KLS cells or KSL cells in combination. *Lnk* and *Hprt* were amplified for 35 cycles using *Lnk* primers (5'-TCT CTC AGG CAC CAG GTT C-3' and 5'-ATT CAC ACG TCT GCC TCT CT-3') and *Hprt* primers (5'-GCT GGT GAA AAG GAC CTC T-3' and 5'-CAC AGG ACT AGA ACA CCT GC-3').

Supplemental Figure S2 Cell cycle analysis of CD34⁻KSL cells from wild-type or *Lnk*-deficient mice

(A) Cell cycle analysis was performed on CD34⁻KSL cells isolated from wild-type or *Lnk*-deficient mice at 8 weeks of age after stained with PI. (B) Cell cycle kinetics was analyzed. Mice were fed water containing BrdU for 2 weeks. 10,000 or more CD34⁻KSL cells or CD34⁺KSL cells were stained with PI and anti-BrdU antibody. At least 5,000 events were collected on FACS analysis. Data from one of the two experiments are shown.

Supplemental Figure S3 Relationship between RU per cell and secondary CRU
Using data in Table 2, the linear regression between RU/cell and secondary CRU was analyzed. RU/cell is positively correlated with secondary CRU ($p < 0.05$). The correlation coefficient is 0.09. Dotted lines show 95% confidence interval.

Supplemental Figure S4 In vitro survival and division of *Lnk*-deficient CD34⁻KSL cells

(A) Single CD34⁻KSL cells isolated from wild-type (WT) or *Lnk*-deficient (*Lnk* KO) mice were cultured under serum-free conditions at graded concentrations of KL for 5 days. Subsequently, all cells were cultured in the presence of 10% FCS with 100 ng/ml of KL, 10 ng/ml of thrombopoietin, 10 ng/ml of interleukin-3, and 1 U/ml of erythropoietin for 10 days more. Colonies consisting of 50 or more cells were counted. There was no significant difference between the numbers of colonies formed by WT cells and those formed by *Lnk*-deficient cells at any concentration of KL examined, indicating that the survival rates by day 5 of culture did not differ between WT and *Lnk*-deficient CD34⁻KSL cells. (B) Single CD34⁻KSL cells from WT or *Lnk*-deficient

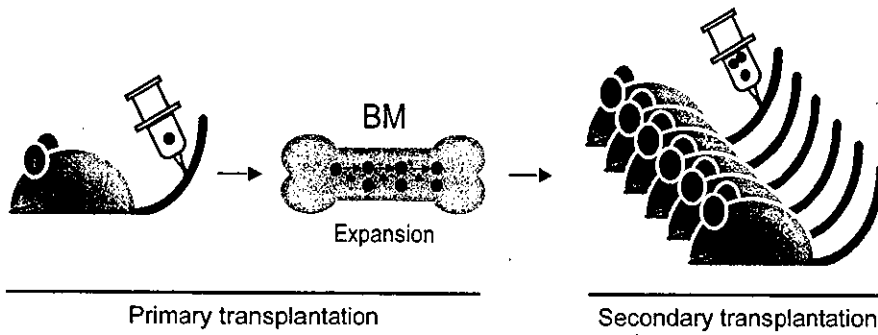
mice were cultured with 10 ng/ml of KL for 7 days. The number of cells produced by single cells was monitored. Data show clones (black circles) which gave rise to 2 or more cells. The number of cells produced by single cells did not differ significantly between WT and Lnk-deficient mice.

Supplemental Procedures

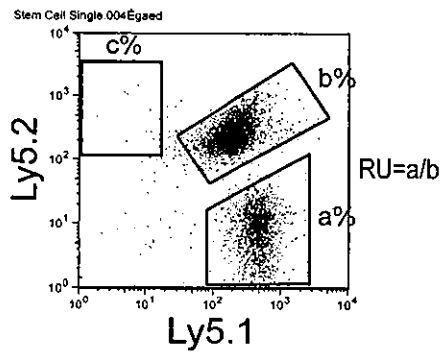
BrdU labeling and analysis

A stock solution of (+)-5-bromo-2'-deoxyuridine (Sigma) was prepared in sterile water at a concentration of 10 mg/ml and stored at -20 °C. Mice were fed water containing 0.5 mg/ml of BrdU for 2 weeks; the water bottle containing BrdU was protected from light and was changed twice a week. CD34⁺KSL cells or CD34⁻KSL cells were sorted into an Eppendorf tube containing one ml of 10% FCS in α -MEM. Cells were centrifuged at 5,000 rpm for 3 min. After the supernatant was discarded, cells were resuspended in 150 μ l of PBS, followed by addition of 350 μ l of ethanol at -20 °C. Cells were fixed at 4 °C overnight. After two washes with PBS, cells were gently vortex-resuspended in 500 μ l of 4N HCl and 20% Tween 20 in water and were incubated at room temperature (RT) for 30 min. After being spun down, cells were incubated in borate-buffered saline (pH 9.0) at RT for 5 min. Cells were washed twice with 500 μ l of 0.5 % Tween 20 in PBS and allowed to react with a titrated amount of FITC-conjugated anti-BrdU antibody (BD Biosciences) at RT for 30 min. After being washed with PBS, the cells were resuspended in 300 μ l of PBS containing 15 μ g/ml of RNase A (Sigma) and 5 μ g/ml of PI, and were incubated at RT for 30 min. FACSCalibur was used for cell cycle analysis.

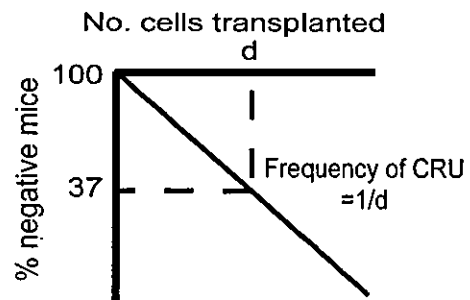
A. Single cell reconstituted mouse



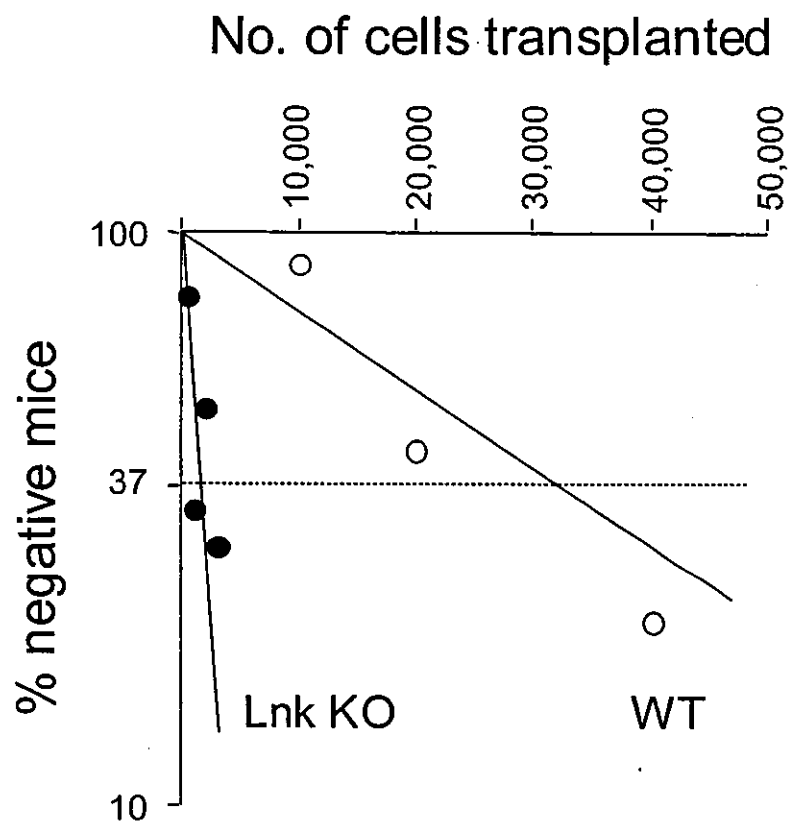
B. Population type CR

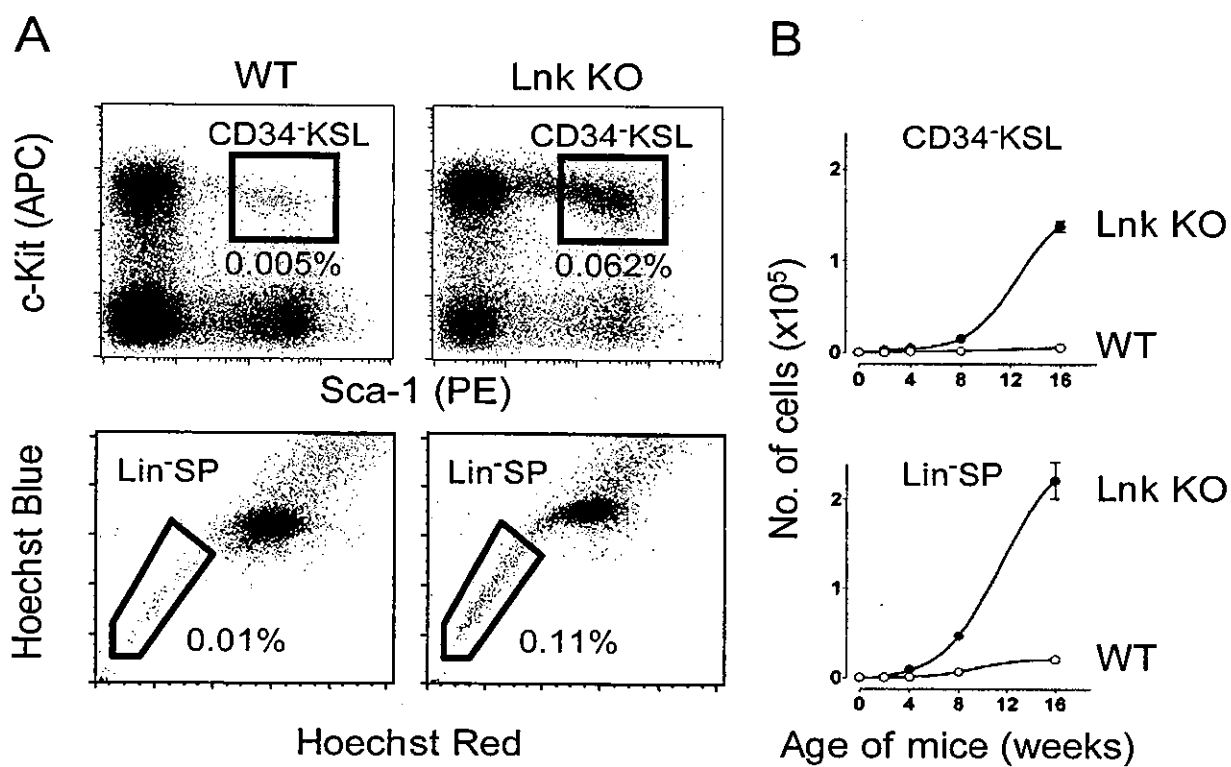


C. Limiting dilution type CR



Ema et al. Fig. 2





Ema et al. Fig. 4

A

WT		Lnk KO	
% Chimerism	RU/cell	% Chimerism	RU/cell
62.0	3.26	87.8	14.39
52.6	2.22	79.2	7.62
44.0	1.57	76.7	6.58
39.7	1.32	69.8	4.62
37.7	1.21	67.4	4.13
34.3	1.04	62.8	3.38
29.0	0.82	50.4	2.03
25.7	0.69	36.9	1.17
19.0	0.47	36.6	1.15
16.8	0.40	26.6	0.72
12.9	0.30	18.2	0.44
12.4	0.28	10.3	0.23
12.1	0.28	7.1	0.15
10.9	0.24	6.8	0.15
9.4	0.21	4.5	0.09
7.0	0.15	4.5	0.09
6.2	0.13	3.0	0.06
5.9	0.13		
5.9	0.13	MAS	2.76 ± 3.84 (n=17)
5.9	0.12		
5.0	0.11		
3.8	0.08		
3.1	0.06		
2.5	0.05		
MAS	0.64 ± 0.80 (n=24)		

P < 0.05

B

