

## Chronic exposure to IFN $\alpha$ drives medullar lymphopoiesis towards T-cell differentiation in mice

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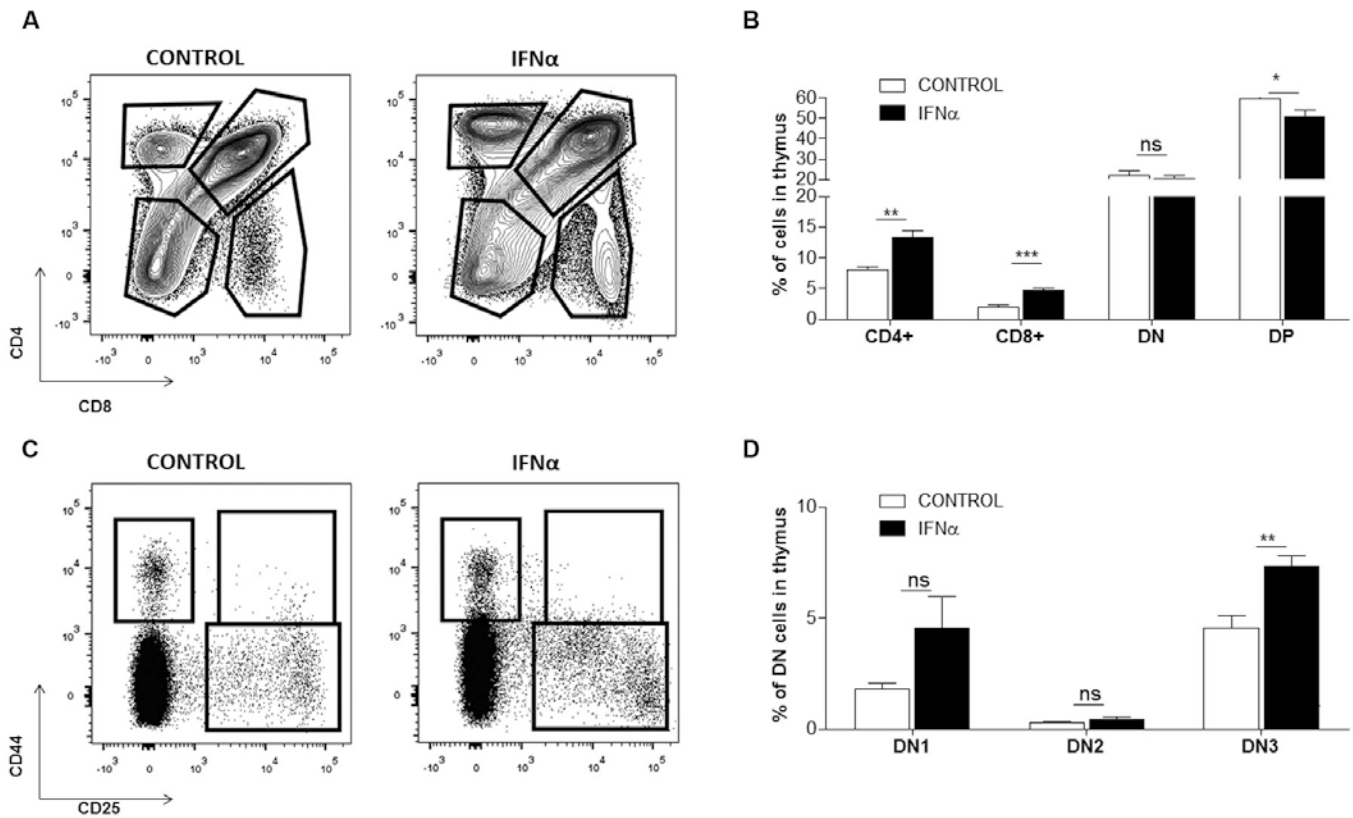
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The online version of this article has a Supplementary Appendix.

Manuscript received on August 8, 2014. Manuscript accepted on February 12, 2015.

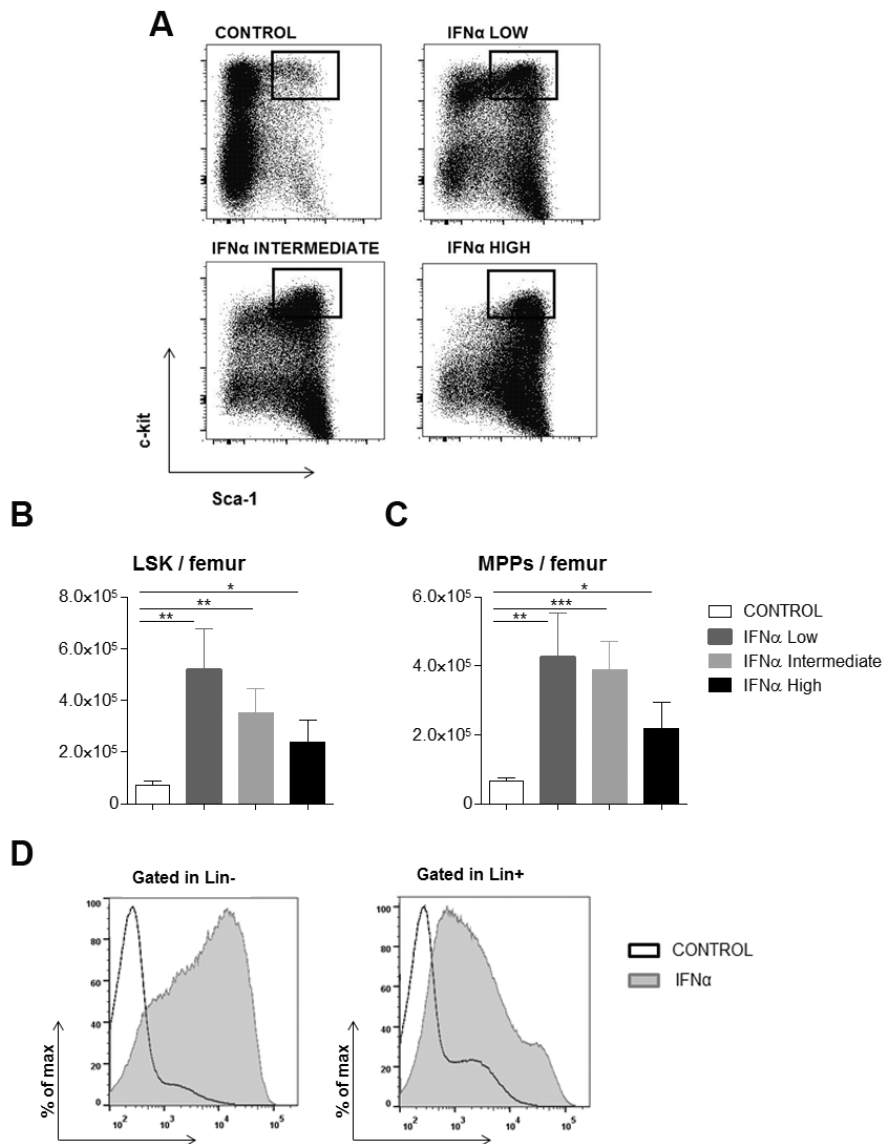
Correspondence: ggasegui@unav.es

## SUPPLEMENTARY FIGURE S1



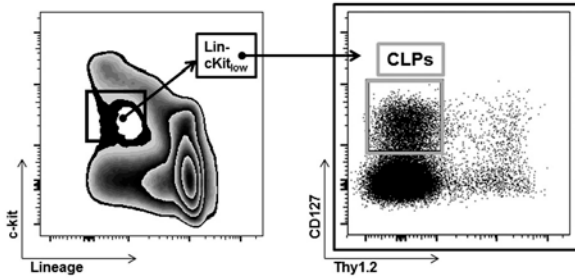
**Chronic IFN $\alpha$  expression affects the cellular composition of the thymus.** Three weeks after AAV-IFN $\alpha$  or AAV-Luc injection (5-6 mice/group) the cellular composition of the thymus was analysed by flow cytometry analysis. (A) Representative flow cytometry analysis of thymic cells labeled with anti-CD4 and anti-CD8 antibodies. (B) Percentage of CD4+, CD8+, double-negative (DN) or double-positive (DP) cells in thymi. (C) Representative flow cytometry analysis of thymus cells labeled with anti-CD44 and anti-CD25 cells gate on DN cells to differentiate among DN1, DN2 and DN3 subsets. (D) Percentage of DN1 (CD8-CD4-CD44+CD25-) or DN2 (CD8-CD4-CD44+CD25+) or DN3 (CD8-CD4-CD44-CD25+) in thymi. Results are expressed as the mean +/- SD. Statistical significance was determined by Student's t-test (\*,  $P \leq 0.05$ ; \*\*,  $P \leq 0.01$ ; \*\*\*,  $P \leq 0.001$ ).

## SUPPLEMENTARY FIGURE S2



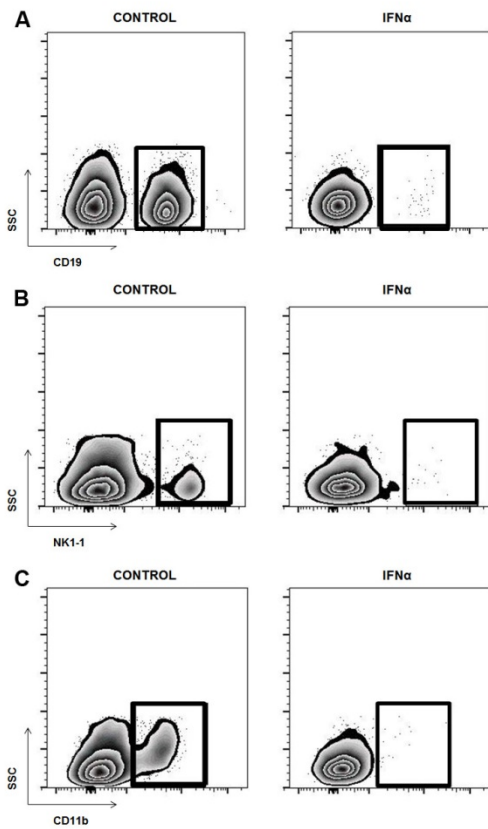
**IFN $\alpha$  induces an aberrant expression of Sca-1 marker.** (A) Representative flow cytometry analysis of Lineage negative (Lin<sup>-</sup>) cKit<sup>+</sup> and Sca-1<sup>+</sup> cells (LSK) of BM cells obtained from mice treated with different doses of AAV-IFN $\alpha$  or AAV-Luc (high dose). The analysis was performed three weeks after vector injection. (B-C) Bar graphs represent the numbers of LSK (B) and MPPs (C) per femur (5-6 mice/group). Results are expressed as the mean  $\pm$  SD. Statistical significance was determined by Student's t-test (\*,  $P \leq 0.05$ ; \*\*,  $P \leq 0.01$ ; \*\*\*,  $P \leq 0.001$ ). (D) Representative flow cytometry histogram analysis of Sca-1<sup>+</sup> cells in the BM of AAV-treated mice gated in Lin<sup>-</sup> (left) or Lin<sup>+</sup> (right) population.

### SUPPLEMENTARY FIGURE S3



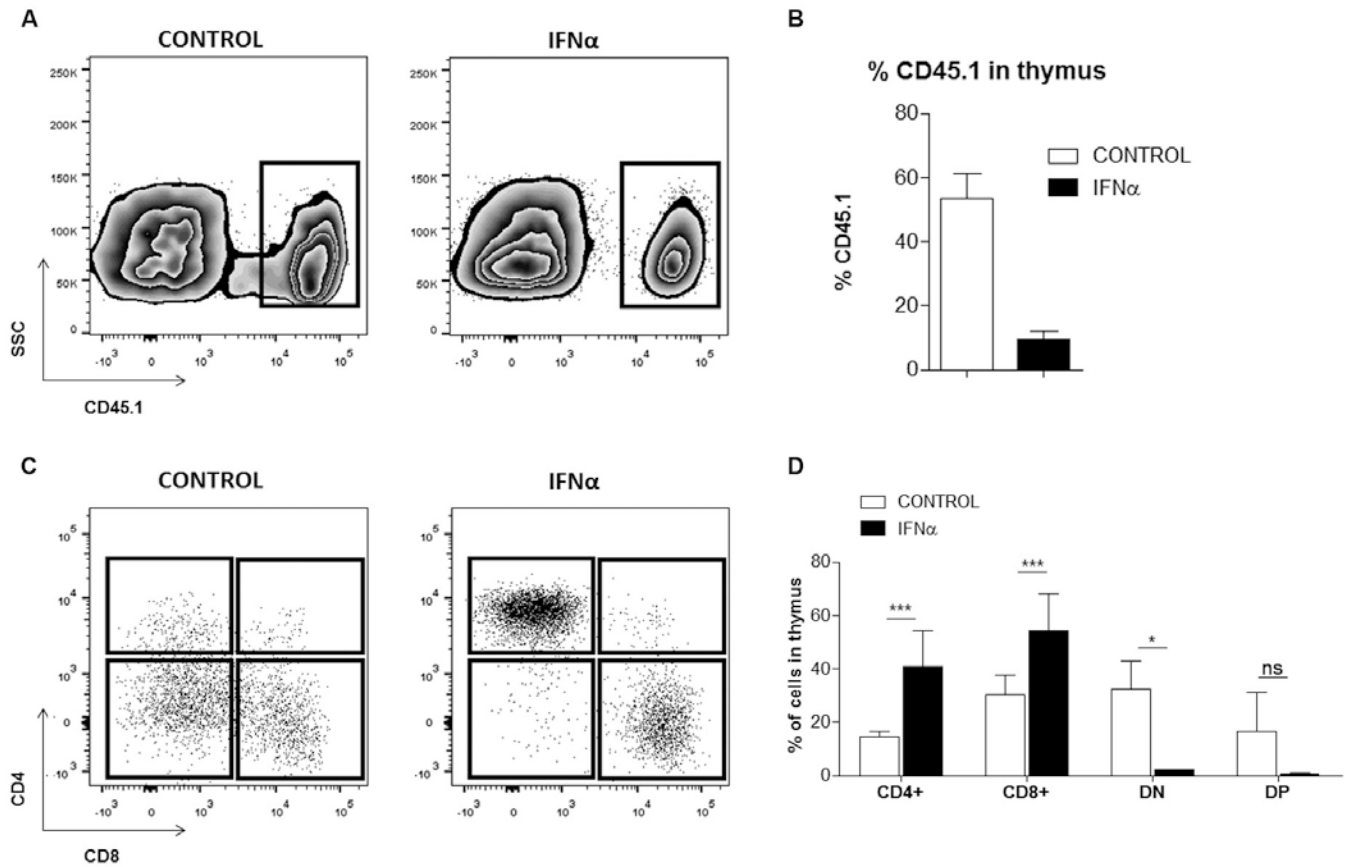
**Gating strategy for the identification of CLPs population.** CLPs were defined as Lin- ckit<sup>low</sup> CD127+ Thy- cells.

### SUPPLEMENTARY FIGURE S4



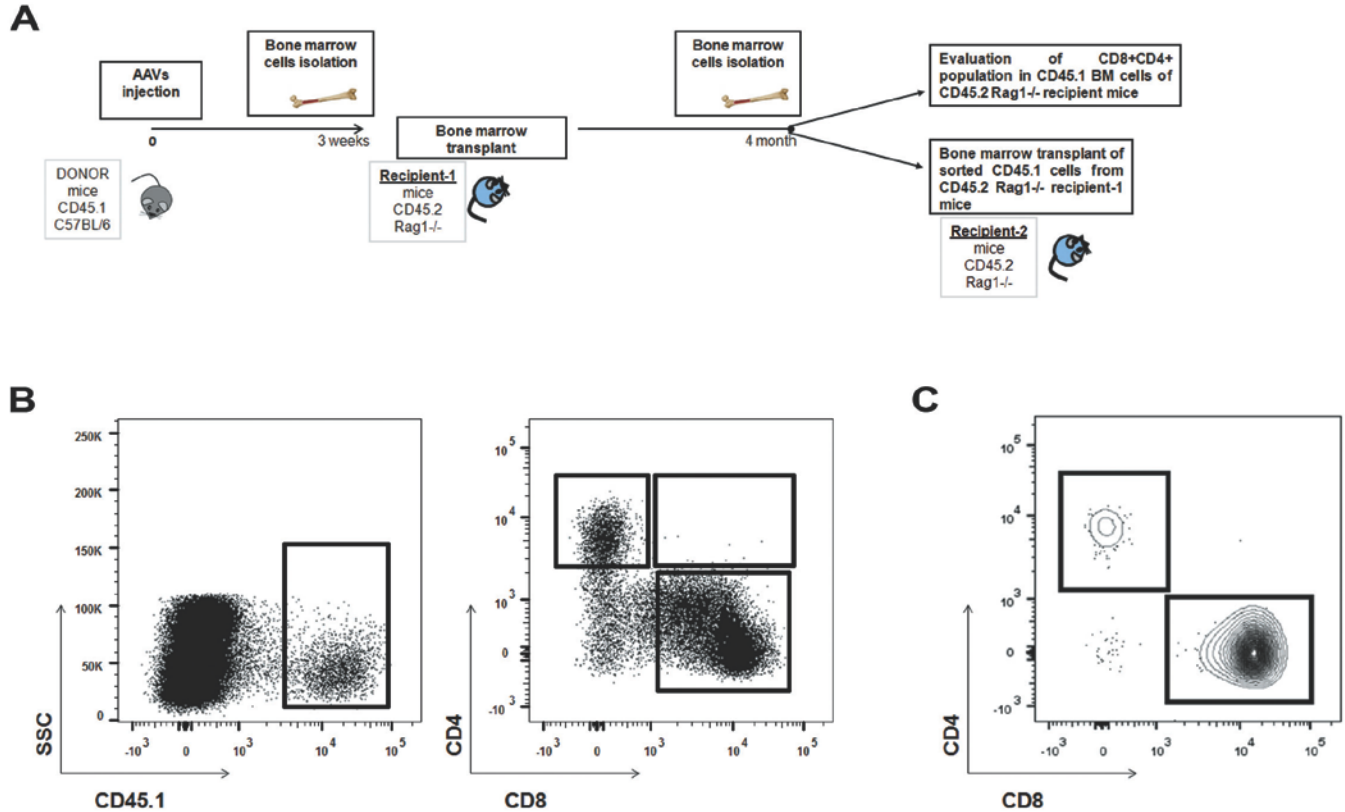
**IFN $\alpha$  expression affects the repopulating ability of hematopoietic cells.** (A-C) Representative flow cytometry analysis of CD45.1+ CD19+ (A), CD45.1+NK1.1+ (B), CD45.1+CD11b+ (C) donor-cells from AAV- IFN $\alpha$  or AAV-Luc treated mice in blood of CD45.2 recipient mice (4-5 mice/group).

**SUPPLEMENTARY FIGURE S5**



**Analysis of cellular composition of the thymus in RAGB6 CD45.2+ recipient animal transplanted with CD45.1+ BM cells from AAV-IFN $\alpha$  or AAV-Luc treated mice.** (A) Representative flow cytometry analysis of CD45.1 donor-cells obtained from the thymus of RAGB6 mice transplanted with bone marrow cells from AAV-IFN $\alpha$  or AAV-Luc treated mice (5 mice/group). (B) Mean percentage of CD45.1+ cells in thymus of recipient animals. (C) Representative flow cytometry analysis of the CD45.1+ CD4+ and/or CD8+ cells in thymi obtained from recipient mice 120 days after BM transplantation. (D) Mean percentage of CD45.1+ CD4+ and/or CD8+ cells in thymi of recipient animals.

## SUPPLEMENTARY FIGURE S6



**Sustained IFN $\alpha$  expression drives medullar lymphopoiesis to T cell development and it is not due to BM-DP population present in the AAV-IFN $\alpha$  treated mice.** (A) Schematic representation of the experiment. Three weeks after AAV-IFN $\alpha$  injection BM cells from C57/BL6 CD45.1 were isolated and transplanted by iv injection into RAG-1-deficient (RAG $^{-/-}$ ) CD45.2 recipient mice (Recipient-1). Four months after transplantation CD45.1 were isolated from the BM of Recipient-1 and re-transplanted by iv injection into RAG-1-deficient (RAG $^{-/-}$ ) CD45.2 (Recipient-2). (B) Representative flow cytometry analysis of CD45.1 $^{+}$  CD4 $^{+}$  and/or CD8 $^{+}$  cells in the bone marrow of Recipient-1 mice (5 mice/group). (C) Analysis of CD45.1 $^{+}$  CD4 $^{+}$  and/or CD8 $^{+}$  cells in the peripheral blood of Recipient-2 mice (5 mice/group).

**Supplementary Table 1.** List of antibodies used for flow cytometry of blood cells

<b>Antibody</b>	<b>Conjugate</b>	<b>Clone</b>	<b>Conc.</b>	<b>Dilution</b>	<b>Supplier</b>
CD8a	FITC	53-6.7	0.5 mg/ml	1/200	eBiosciences
CD8a	Pacific Blue	53-6.7	0.5 mg/ml	1/200	Biolegend
CD4	APC/Cy7	GK1.5	0.2 mg/ml	1/1000	Biolegend
CD19	APC/Cy7	PeCa1		1/100	ImmunoTools
CD3	PerCP/Cy5.5	17A2	0.2 mg/ml	1/200	Biolegend
NK1.1	APC	PK136	0.2 mg/ml	1/200	Biolegend
CD11b	FITC	M1/70.15		1/100	ImmunoTools
CD11c	APC	HL3		1/100	ImmunoTools
CD45.1	PE	A20	0.2 mg/ml	1/150	Biolegend
CD45R/B220	FITC	RA3-6B2	0.5 mg/ml	1/200	BD Pharmingen
CD44	APC/Cy7	IM7	0.2 mg/ml	1/400	Biolegend
CD25	BV421	PC61	12 µg/mL	1/200	Biolegend

**Supplementary Table 2.** List of antibodies used for flow cytometry of bone marrow cells

Antibody	Conjugate	Clone	Conc.	Dilution	Supplier
Lineage cocktail	APC			1/150	BD Pharmingen
Lineage cocktail	FITC			1/150	Biologend
CD117(C-KIT)	PE	2B8	0.2 mg/ml	1/150	Biologend
CD117(C-KIT)	APC/Cy7	2B8	0.2 mg/ml	1/200	Biologend
Ly-6A/E (Sca-1)	Pacific Blue	D7	0.5 mg/ml	1/200	Biologend
CD48	FITC	HM48-1	0.5 mg/ml	1/500	Biologend
CD150	PE/Cy7	TC15-12F12.2	0.2 mg/ml	1/150	Biologend
CD34	PE	RAM34	0.2 mg/ml	1/100	Biologend
CD34	BV421	MEC14.7	0.1 mg/ml	1/100	Biologend
CD34	FITC	RAM34	0.5 mg/ml	1/100	Biologend
CD16/32	PE/Cy7	93	0.2 mg/ml	1/300	Biologend
CD90.2	BV510	53-2.1	0.1 mg/ml	1/500	Biologend
CD127	APC	SB/199	0.2 mg/ml	1/400	Biologend
CD135	PE/Cy5	A2F10	0.2 mg/ml	1/100	Biologend

CD8a	FITC	53-6.7	0.5 mg/ml	1/200	eBiosciences
CD8a	Pacific Blue	53-6.7	0.5 mg/ml	1/200	Biologend
CD4	APC/Cy7	GK1.5	0.2 mg/ml	1/1000	Biologend
CD19	APC/Cy7	PeCa1		1/100	ImmunoTools
CD3	PerCP/Cy5.5	17A2	0.2 mg/ml	1/200	Biologend
NK1.1	APC	PK136	0.2 mg/ml	1/200	Biologend
CD11b	FITC	M1/70.15		1/100	ImmunoTools
CD11c	APC	HL3		1/100	ImmunoTools
CD45.1	PE	A20	0.2 mg/ml	1/150	Biologend
CD45R/B220	FITC	RA3-6B2	0.5 mg/ml	1/200	BD Pharmingen
Ter-119	APC	TER-119	0.2 mg/ml	1/200	BD Pharmingen
CD41	PE	MW/Reg30	0.2 mg/ml	1/100	BD Pharmingen



**Supplementary Table 3:** quantitative PCR primer list

<b>Gene name</b>	<b>Forward</b>	<b>Reverse</b>
IL-7 RECEPTOR ALPHA	CTGCAGTCCCAGTCATCATGA	GTGGCACTCAGATGATGTGACA
NOTCH-1	CGTGATGACCTAGGCAAGT	CAGTCTCATAGCTGCCCTCA
GATA 3	CTTATCAAGCCCAAGCGAAG	CCCATTAGCGTTCCCTCCTC
DLL-4	TGTCTCCACGCCGGTATTG	AGGTCGTCTCCCGGTGTGT
DELTEX-1 (DTX-1)	GAGGATGTGGTTCGGAGGTA	CAGGCAGAGCAGGTGATACA
RUNX1	GCTTTCAAGGTGGTGGCAC	CTCGCTACCTGGTTCTTCATG
GATA1	CCAGTTTGTGGATTCTGCCC	GTAGGCCAGTGCTGATGCTGC
MEIS-1	GCATGCAGCCAGGTCCAT	TAAAGCGTCATTGACCGAGGA
SOCS1	CACCTTCTTGGTGCGCG	AAGCCATCTTCACGCTGAGC
E2A	CTTTGACCCTAGCCGGACATAC	GTGCCAACACTGGTGTCTCTC
EBF1	AGATTGAGAGGACGGCCTTTGT	TCTGTCCGTATCCCATTGCTG
PAX5	AATCGCTGAGTACAAACGCCAA	TCCGAATGATCCTGTTGATGGA
FOXO1	TCGTACGCCGACCTCATCA	TCCTTGAAGTAGGGCACGCTC

## **SUPPLEMENTARY METHODS**

### **Determination of murine IFN $\alpha$**

Concentration of murine IFN $\alpha$  was determined by VeriKine Mouse Interferon-Alpha ELISA Kit (PBL interferon source).

### **Hemogram**

Blood samples were analyzed using an automated veterinary hematological analyzer with a preprogrammed murine calibration mode (Hemavet 950FS; Drew Scientific, Waterbury, CT).

### **Cell isolation**

For bone marrow cell isolation bones were flushed using a 23-gauge (femurs and tibias) or 26-gauge needle (iliac crests) and the bones discarded. The cells were obtained by mechanical disruption and washed by centrifuging at 1300rpm for 5 minutes at 4°C, resuspended in PBS, and then filtered through a 70- $\mu$ m filter. Red blood cells were removed using a lysis Buffer. Cell concentrations were determined with an automatic cell counter (Z1 Coulter Particle Counter, Beckman Coulter).

For thymi cell isolation thymocytes were harvested by homogenizing thymic lobes with a syringe followed by centrifugation through a 40- $\mu$ m Nylon mesh. Cells were washed in FACS-Buffer (PBS with 0.5% BSA and 0.05% NaN<sub>3</sub>), counted and stained using various antibody combinations.

### **Blood cells analysis**

Blood single-cell suspensions were pretreated with FcR-Block (anti-CD16/32 clone 2.4G2; BD Bioscience-Pharmigen). Afterward, cells were stained with different combinations of antibodies. A list of antibodies is provided in supplemental Table 1.

### **Cell cycle analysis of CD8+ and CD4+ T cells.**

Cell cycle analysis of CD8+ and CD4+ T cells were performed using Click-it EdU Flow cytometer assay kit according to the manufacturer's instructions (Invitrogen).

### **Hematopoietic Stem cell (HSC) staining**

Whole bone marrow were isolated and stained on ice with various antibody cocktails to identify each progenitor compartment. A list of antibodies used is provided in supplemental Table 2.

### **Cell sorting**

Different cell fractions were stained with antibodies and further purified on a FACSARIA cell sorter (BD Biosciences, San Diego, CA).

### **In vitro treatment**

$3 \times 10^5$  LK cells were cultivated in StemSpa Serum-Free Expansion Medium (SFEM) medium (StemCell technologies) in presence or absence of 0,5U/ $\mu$ L of recombinant IFN $\alpha$ . LK were treated for 8h or 24h and harvested to analyse the TF by RT-PCR.

### **Bone marrow re-transplant**

CD45.1+ cells derived from AAV-treated mice were purified on a FACSARIA cell sorter from BM of CD45.2 Rag-/- (recipient-1) mice. Then these CD45.1+ cells were re-transferred by retroorbital injection into other group of RAG1-/- mice.

### **Statistical analysis of RT-PCR data**

RT-PCR data were preprocessed and analyzed with R/Bioconductor.<sup>23</sup> The Delta Ct (DCt) method was used to quantify relative expression of each gene with respect to the housekeeping gene control GAPDH. The  $2^{-\Delta\Delta C_t}$  formula was used to calculate the differential gene expression. Data are shown as log2 of the ratio between IFN $\alpha$ -exposed and control cells. LIMMA (Linear Models for Microarray Data)<sup>24</sup> was used to find out the

differentially expressed genes between the samples treated with IFN $\alpha$  and the controls. A False Discovery Rate (FDR) of 0.05 was established as selection criteria.<sup>25</sup>