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Ptk7 deficient mice have decreased Hematopoietic Stem Cell pools as a result of deregulated proliferation and migration

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Running title: Role of the polarity protein PTK7 in murine haematopoiesis.

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Abstract

Hematopoietic Stem Cells (HSCs) located in adult bone marrow or fetal liver in mammals produce all cells from the blood system. At the top of the hierarchy are long-term HSCs endowed with lifelong self-renewal and differentiation properties. These features are controlled through key microenvironmental cues and regulatory pathways such as Wnt signaling. We have previously shown that PTK7, a tyrosine kinase receptor involved in planar cell polarity (PCP), plays a role in epithelial Wnt signaling. However, its function in hematopoiesis has remained unexplored. Here we show that PTK7 is expressed by hematopoietic stem and progenitor cells (HSPCs) with the highest protein expression level found on HSCs. Taking advantage of a *Ptk7* deficient mouse strain, we demonstrate that loss of *Ptk7* leads to a diminished pool of HSCs, but does not affect *in vitro* or *in vivo* hematopoietic cell differentiation. This is correlated with increased quiescence and reduced homing capacities of *Ptk7* deficient HSPCs, unravelling novel and unexpected functions for planar cell polarity pathways in HSC fate.

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41 **Introduction**

42 Hematopoiesis is a biological process consisting on the production of all blood cell
43 types from hematopoietic stem cells (HSCs) located in the bone marrow (BM) of adult
44 mammals. Fetal hematopoiesis occurs in different sites, particularly in the liver, during its
45 earliest phases. Long term hematopoietic stem cells (LT-HSCs) are characterized by their
46 capacity to provide lifelong reconstitution of all blood cell lineages after transplantation
47 into lethally irradiated recipients, whereas short term HSCs (ST-HSCs) do so for only 8-10
48 weeks.

49 In the mouse, LT-HSCs belong to the LSK cell compartment defined by the
50 Lineage$^{-/low}$ Sca1^{+}c-Kit^{+} phenotype irrespectively of their fetal liver (FL) or adult bone
51 marrow (BM) origin. Additional markers known as SLAM (Signal Lymphocytes Activating
52 Molecules) can further subdivide this compartment (1)(2). Indeed, CD150^{+}CD48^{-} LSK have
53 been shown to contain around 40% of HSCs with long-term reconstitution potential in BM
54 as well as in FL (3).

55 Hematopoiesis is a highly controlled multi-step process that relies on complex
56 interaction networks involving cell surface receptors, growth factors, and adhesion
57 molecules expressed by HSCs and their environment (4)(5)(6)(7)(8). Because HSCs
58 generate mature hematopoietic cells, including immune cells, their replacement must be
59 adjusted to homeostatic or stress conditions such as infections, inflammation or blood loss,
60 and their expansion must be controlled to avoid exhaustion. This control is made possible
61 by the coordinated regulation of quiescence, self-renewal, and differentiation through
62 appropriate signals delivered by functional BM niches (9). HSCs are retained in these BM
63 niches by cell surface molecules endowed with adhesive and/or signalling functions
64 expressed by HSCs. These receptors belong to different protein families: integrins (VLA-4)
65 (10), Immunoglobulin superfamily (Ig Sf) adhesion molecules (6) (11), G Protein Coupled

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Receptors (CXCR4)(12), or tyrosine kinase receptors (TIE2, c-Kit)(13)(14)(15) interacting with ligands present within the BM microenvironment (16). Discovery of novel molecules implicated in this multi-step program is of particular importance to embrace its complexity, develop new strategies aiming at the regeneration of damaged hematopoietic tissues and to understand hematopoietic diseases.

PTK7 is a planar cell polarity (PCP) receptor belonging to the Immunoglobulin Superfamily (Ig Sf) and playing important roles during development (17)(18). PCP is controlled by a non-canonical Wnt pathway, organizes polarization of many epithelial tissues and organs within the plane and drives cell migration and cell intercalation of non-epithelial cells such as mesenchymal cells (19)(20). *Ptk7* deficient mouse embryos show all signs of typical PCP defects such as severe tube neural and abdomen closure defect with secondary development abnormalities (21). PTK7 is a type I protein composed of seven Ig loops in its extracellular region, a transmembrane domain, an intracellular region with a tyrosine kinase domain (17)(18)(22)(23). While PTK7 is catalytically inactive, the cytoplasmic domain containing the tyrosine kinase domain is mandatory for receptor function *in vitro* and *in vivo* (24)(25)(26)(27). Both extracellular and intracellular domains of PTK7 can be released from the plasma membrane following the sequential cleavage of the receptor by membrane type 1-matrix metalloproteinase (MT1-MMP)/ADAM-17, and γ-secretase, interfering with its PCP and promigratory activities (24)(28)(29).

We have recently found that PTK7 expression is not restricted to epithelial cells. Indeed, PTK7 is expressed at the cell surface of human hematopoietic progenitors and acute myeloid leukemia (AML) patient blast cells. Overexpression of the PCP protein leads to increased cell survival and cell migration of leukemic cells and confers a poor prognosis to patients independently of other risk factors (27). Similar conclusions were recently drawn

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from studies done in chronic lymphocytic leukemia in which a set of PCP proteins is upregulated (30).

Although functions of canonical and non-canonical Wnt pathways have been extensively studied in hematopoiesis (for review see (31)(32)(33)), the role of PTK7 has never been explored in this process. In the present study, we have thus addressed the function of PTK7 in physiological hematopoiesis using a mouse model deficient for *ptk7* expression. Using flow cytometry, we first mapped expression of PTK7 on HSCs and progenitors. As PTK7 deficient mice die perinatally, we then studied function of PTK7 in fetal hematopoiesis and revealed a decreased number of HSPCs in FL of *Ptk7* deficient mice. Although HSC differentiation was not affected, we found a dramatic decrease in proliferation and migration properties of *Ptk7* deficient HSPCs that reveal novel unexpected functions for this planar cell polarity molecule.

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Methods

Mice

C57BL/6 and C57BL/6-CD45.1 were purchased from Janvier Laboratory and Charles River Laboratories, respectively. All mice used for transplantation experiments are 6 to 12 weeks old. Lethal irradiation was done with an X-ray irradiator (X-RAD® 160 X-Ray, PXi) using a single dose of 8Gy.

PTK7 gene-trap mice were obtained by injection of gene-trap ES XST087 (Bays Genomics, USA) in E3.5 blastocyst from C57BL/6 mouse. Chimeric mice were back-crossed with B6 mice for seven generations to obtain homogenous genetic background for experiments. All mice were housed and maintained under specific pathogen-free condition. E14.5 and 15.5 were used in all FL experiments. All animal experiments were performed in agreement with the French Guidelines for Animal Handling.

Genotyping

Genotyping of adult mice and embryos was performed by genomic PCR using the following pairs of primers: 5'- ACGTCACTGGAGAGGAAGTCCGCAG-3' and 5' GATCGAAGCAGGTCCTGTGGTCCT-3' for wild type allele and 5'-TTGTGAAGTGGAGGCTCCGGACC-3' and 5'-CGATCTTCCTGAGGCCGATACTGTC-3' for gene trap insertion allele.

Flow cytometry and cell sorting

For the analysis of hematopoietic stem/progenitor cells in FLs, FLs were dissected from E14.5 embryos and single cell suspension was made in phosphate-buffered saline (PBS)-2% Fetal Calf Serum (FCS).

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127 For analysis of hematopoietic stem/progenitor cells in BM, thymus or blood cells obtained
128 from adult mice were suspended in PBS-2% FCS. For BM and thymus, cells were filtered
129 using 70 µm cell strainer (BDFalcon) and lysis of red blood cells was done using ACK lysis
130 buffer (0.15 M NH4Cl, 10mM KHCO3, 0.1mM Na2EDTA, Life Technology) for 2 min at
131 room temperature. Cells were spun and resuspended in PBS-2%FCS +/- 0.125mM EDTA
132 for staining. For blood samples, red blood cell lysis was done after staining using BD FACS
133 Lysing buffer solution (BD Biosciences) following manufacturer instructions.
134 For flow cytometry analysis and cell sorting, cells were stained with monoclonal antibodies
135 purchased from Ebiosciences or Biolegend against the following molecules: B220 (clone
136 6B2), CD3 (145-2C11), CD4 (RM4-5), CD8 (53-6.7), CD11b/Mac-1 (M1/70), CD11c
137 (N418), CD16/32 (93), CD19 (6D5), CD24 (M1/69), CD34 (RAM34), CD45 (30F11),
138 CD45.1/Ly5.1 (A20), CD45.2/Ly5.2 (104), CD48 (HM48-1), CD49b (DX5), CD49d (R1-
139 2), CD117/c-Kit (2B8 and/or ACK2), CD135 (A2F10), CD150 (TC15-12F12.2), F4/80
140 (BM8), Gr1 (RB6-8C5), Sca-1 (D7), Ter-119. Antibody human PTK7 and glycophorin A
141 (11E4B-7-6) were purchased from Miltenyi biotech and Beckman Coulter, respectively.
142 All experiments were done by multicolour parametrics analysis using FITC, PE, APC, PE-
143 Cy7, APC-Cy7, PerCPCy5.5, AlexaFluor700, AlexaFluor594, Pacific blue, PE-Cy5, Q-dot
144 605 fluorochroms or spectral equivalents. Commercial antibodies were used either directly
145 conjugated or biotinylated. For mouse PTK7 expression, polyclonal rabbit antibody raised
146 against recombinant PTK7-Fc was used in the experiments. For detection of interesting
147 antigen, PE or APC-conjugated goat-anti–rabbit (Jackson Immunoresearch Laboratories)
148 were used as secondary antibodies. To exclude lineage positive cells, streptavidin
149 conjugated with Alexa A594 (Invitrogen) or Pacific blue (Ebiosciences) were used.
150 Irrelevant isotype-matched antibodies were used as controls. Cells were stained 20 minutes
151 at 4°C and rinsed in PBS-2% FCS solution. Viability marker (vivid dye, Invitrogen) was

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added in last staining in PBS solution. Stained cells were analysed by LSR II SORP (laser 405, 488, 561, and 633; BD Bioscience) or sorted by ARIA III (BD Bioscience). LSK cell populations were purified by cell sorting using a FACS-ARIA with > 95% purity. Results were analyzed using BD-DIVA Version 6.1.2 software (BD Biosciences) or FlowJo Version 7.6.2 (TreeStar) software.

Homing experiments

E14.5 embryos were sacrificed and were immediately genotyped. FL cells from wild type or gene trap mice were suspended, counted and stained with cell tracker calcein AM and calcein red-orange (Invitrogen), respectively, according manufacturer instructions. Cells were mixed at 1/1 ratio and 4.10^6 total cells were injected through retro-orbital vein in previously irradiated (8 Gy) mice. 12h after injection, mice were sacrificed and blood, bone marrow, spleen, liver and thymus samples were analysed for fluorescents cells.

Transplantation assays

For LSK sorting, FL cells (CD45.2+) were labelled for lineage positive (lin+) cells with following biotinylated primary rat antibodies (lineage mix): CD3, CD4, CD8, CD19, B220, DX5, TER-119, GR-1, CD11c. Lin+ cells were depleted by magnetic separation using anti-rat Ig coated magnetic beads (DynaBeads, Lifetechnology). Lin^{Neg} cells were also stained and sorted by FACS according to the phenotype Lin^{Neg}, $CD45.2^+$, Kit^+ and $Sca\text{-}1^+$. CD45.1 mice were lethally irradiated and grafted with 1000 or 400 $CD45.2^+$ LSK cells through the retro-orbital vein along with 1.10^5 total BM recipient cells to allow short term reconstitution and mice survival. Recipient mice survival was followed and BM engraftment was examined by monitoring donor ($CD45.2^+$ cells) and host cell ($CD45.1^+$ cells) frequency in

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the blood. After lethal irradiation, mice were maintained on water containing neomycin sulfate 1g/500mL for 2 weeks.

Serial transplantation assay

Lethally irradiated (8 Gy) mice (C57Bl/6, CD45.1) received 3.10^6 total fetal liver cells (first transplantation) or BM total cells (next transplantations). BM engraftment was monitored as described above.

Competition assay

Lethally irradiated (8 Gy) mice (C57Bl/6, CD45.1) received 2.10^6 total FL cells from CD45.2 PTK7 $^{-/-}$ mutant embryos and 2.10^6 total FL cells from $CD45.1^+CD45.2^+$ $PTK7^{+/+}$ embryos, issues from crossing of CD57Bl/6, CD45.1 and CD57Bl/6, CD45.2. Plugs are obtained the same day. BM engraftment was examined by monitoring frequency of $CD45.2^+$ cells from PTK7 mutant donors and $CD45.1^+CD45.2^+$ cells from wild type donor, excluding $CD45.1^+$ cells from host cells, in the blood. After lethal irradiation, mice were maintained on water containing neomycin sulfate 1g/500mL for 2 weeks.

Cell cycle analysis and apoptosis analysis

FL progenitors were enriched as previously described and stained with cKit–APC-eFluor750, CD150-APC, Sca-1-PerCP-Cy5.5, and CD48-PECy7. The biotin-conjugated lineage combination was revealed with AlexaFluor 594–streptavidin (Invitrogen). For cell cycle analysis, after extracellular staining, cells were permeabilized, fixed, and stained with anti–Ki67-FITC MoAb (BD Biosciences) and DAPI as previously described (34). For apoptosis analysis, cells were stained with PE-annexine V to evaluate annexine V externalization (BD biosciences) and DAPI.

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In vitro methylcellulose assays

For methylcellulose colony assays, FL or BM cells suspensions were prepared in Iscove modified Dulbecco medium (IMDM) supplemented with 2% FBS. 3.10^4 cells of FL or 1.10^5 cells of BM were seeded in Methocult M3434 (containing rmSCF 50 ng/mL, rmIL-3 10ng/mL, rhIL-6 10ng/mL, and EPO 3U/mL) (from Stem Cell Technologies, Grenoble, France) in 35 mm dishes accordingly to manufacturer's instructions. All assays were done in triplicated. Colonies (>30 cells) were counted after 13 days of incubation at 37°C, 5% CO_2 and humidified atmosphere.

Adhesion assays

MS5 stromal cells were plated on 96-well flat bottom plates, at a concentration of 5.10^3 cells/well 1 day before the assay. MS5 is a stromal cell line that has been generated by irradiating adherent cells in long-term BM culture. Lin^{Neg} FL cells (5.10^5) obtained as previously described were cocultured for 2 hours with MS5 stromal cells. Plates were washed 3 times with PBS and adherent cells were removed by strong pipetting in PBS 1.25mM EDTA 2% FCS. The phenotype of adherent cells was determined using anti-CD117, Sca-1, CD150, CD48, and FITC-lineage mix to exclude any residual mature cells contaminating the Lin^{Neg} preparation. Absolute numbers were measured using the HTS coupled to the BD-LSR2 SORP (BD Bioscience).

Migration assays

Lin^{Neg} FL cells were enriched as previously described and suspended in Iscove modified Dulbecco medium (IMDM) containing 2% of bovine serum albumine. About 2.10^5 cells in 75 µl were loaded in 5-µm pore Transwell system (Corning Life Sciences). The lower

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chamber was filled with Serum free Media supplemented with the chemoattracting murine stromal-derived factor-1 (mSDF-1α, 100 ng/mL, Peprotech). The same number of cells was plated in distinct wells to be used as control for migration. Plates were then incubated 4 hours at 37°C, 5% CO_2. All experiments were performed in duplicates. Cells were also stained using anti-CD117, Sca-1 and streptavidin coupled to Pacific Blue as described above. Absolute numbers were measured using the HTS coupled to the BD-LSR2 SORP (BD Bioscience) and the migration ratio was defined as the number of migrating cells between migration wells and control wells.

TF1 cell differentiation

Growth of the human erythroid/macrophage progenitor cell line TF1 is dependent on GM-CSF or IL3 and can be induced to differentiate in the presence of growth factors. TF1 cells were grown in IMDM medium supplemented with 5 ng/ml GM-CSF and supplemented with EPO (2 UI/ml) for erythroid differentiation. TF1 cells were infected by viral particles containing shRNA against PTK7 as previously described (27). A scramble non-targeting shRNA was used as a control. shRNA efficiency for PTK7 downregulation was controlled by flow cytometry before and at the end of experiments. TFI cells were then differentiated in IMDM with 2 IU/ml EPO and without GM-CSF for a total of seven days as previously described (35). Erythroblastic differentiation was evaluated by glycophorin A expression.

Statistical analysis

Statistical significance was determined with nonparametric Mann-Withney *U* test using Prism Version 5 software (Graphpad Software Inc).

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Results

PTK7 is expressed on HSCs, myeloid progenitors, lymphoid and thymic progenitors

In a first approach, we studied PTK7 pattern of expression in adult murine hematopoietic cells. To this end, a rabbit polyclonal antibody generated against the extracellular region of PTK7 was used. This antibody recognizes mouse PTK7 endogenously expressed on endothelial and hematopoietic cells by FACS analysis in wild type but not *ptk7* deficient mice (Supplementary Fig. S1). Analysis of PTK7 expression in peripheral blood and in splenocytes revealed that none of the mature mouse leukocytes subsets (granulocytes, monocytes, B cells and T cells) expressed PTK7 to detectable level (Fig. 1A and data not shown). We next analysed PTK7 expression in murine BM hematopoietic progenitors using multiparametric flow cytometry and gating strategies described in Figure 1B. When LSK subsets were examined, PTK7 expression was found to be inversely correlated to differentiation stages with the highest expression on HSCs and multipotent progenitors (MPP) type 1 HSC-MPP1 (RFI=40), and intermediate expression on MPP2 (RFI=20) and MPP3-MPP4 (RFI=15) (Fig. 1C). More committed progenitors including common myeloid progenitors (CMP) and common lymphoid progenitors (CLP) expressed PTK7 at lower levels while the protein was almost undetectable on the surface of granulocyte-macrophage progenitor (GMP), and megakaryocyte–erythroid progenitor (MEP) (Fig. 1D). A similar phenotypic analysis was conducted on differentiating T cells isolated from the thymus. T cell progenitors do not express CD4 and CD8 markers and are called double negative (DN) cells before differentiation into double positive (DP) and single positive (SP) cells. Within the DN compartment, four thymic progenitor populations can be identified according to CD44 and CD25 expression from the most immature DN1 to the most mature DN4 subsets. PTK7 is expressed by the CD44+CD25- (DN1) population and expression is conserved

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during thymic differentiation within the DN2, DN3 and DN4 subpopulations (Supplementary Fig. S2). PTK7 expression is also detected on DP thymocytes before becoming progressively lost at the SP stage. Altogether, these phenotypic analyses reveal that PTK7 expression is inversely correlated to hematopoietic differentiation with the highest expression found in HSCs. This suggests that PTK7 may play a role at this early stage of hematopoietic differentiation.

Ptk7$^{-/-}$ embryos have a quantitative defect of hematopoietic stem cells

We next questioned the function of PTK7 in hematopoiesis. To this end, we generated a *Ptk7* deficient mouse strain using a gene-trapped ES cell available through the International Gene Trap Consortium. As expected from the previous description of *Ptk7* deficient strains, we observed a perinatal lethality due to profound developmental defects (21). *Ptk7* $^{-/-}$ embryos harbour a severe phenotype with cranial neural tube closure defect, gastroschisis, kidney and forelimb abnormalities (Fig. 2A, 2B). Considering this dramatic phenotype, it was impossible to study adult hematopoiesis in these mice and we thus decided to study FL hematopoiesis. In FL, as well as in the BM, HSCs are comprised within the LSK cell compartment. Using the same gating strategy as in Figure 1, we analysed the consequences of PTK7 deficiency in the different LSK cell populations (Fig. 2C). We observed a 25% decrease of total cellularity at E14.5 in PTK7 deficient FL compared to wild type FL (Fig. 2D). However proportion of leucocytes (e.g. CD45+ cells) is equivalent between deficient and wild type FL (Fig. 2E) suggesting that the decrease in FL cellularity in *Ptk7* $^{-/-}$ embryos is likely due to a global effect of *ptk7* deficiency on FL size. We next examined the hematopoietic progenitor compartment. A decrease of about 50% of LSK cell frequency was observed in *Ptk7* deficient FL compared to control FL (1.33% *Ptk7* $^{+/+}$ and $^{+/-}$ LSK cells versus 0.67% *Ptk7* -/- LSK cells) (Fig. 2F). In the different LSK cell populations, the

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most affected one was the HSC-MPP1 subset (Fig. 2C and 2G). No significant difference in the frequency of MMP2 and MPP3-MPP4 subsets was observed in *Ptk7* deficient FL compared to wild type or heterozygous FL. When we examined the absolute number of the different subsets of hematopoietic progenitors, we noticed that all LSK subsets were affected by PTK7 deficiency. Indeed, there was a 68% decrease in the LSK cell absolute number in *Ptk7* deficient embryos compared to control embryos (Fig. 2H). The most dramatic decrease was observed in the HSC compartment. Indeed, there was up to 75% decrease in HSC number in *Ptk7* $^{-/-}$ embryos compared to control one whereas there was around a 50% decrease in MPP2 and MPP3/4 cell numbers (Fig. 2I)

To assess the frequency of functional HSC in FL, we transplanted 1000 and 400 sorted LSK cells from *Ptk7* $^{+/+}$ or *Ptk7* $^{-/-}$ FL bearing the CD45.2 allele into two groups of lethally irradiated mice bearing the CD45.1 allele. As expected, all mice included in the group injected with 1000 LSK cells survived. However, with 400 injected cells, *Ptk7* $^{+/+}$ FL LSK cells were able to reconstitute recipient mice between 2 and 12 weeks after injection whereas all mice except one injected with *Ptk7* $^{-/-}$ LSK cells died within 6 weeks (Fig. 2J). We concluded that loss of PTK7 leads to a defect in functional HSC frequency. *In vitro* clonogenicity assays confirmed the defect in HSCs and early precursor frequency as we observed a twofold reduction in colony-forming units starting from 3.10^4 total FL cells. (Fig. 2K).

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PTK7 deficient fetal liver cells are more quiescent than wild type fetal liver cells

To determine if the absence of PTK7 could alter HSC quiescence, we performed cell cycle phenotypic analysis of LSK FL cells using an antibody directed against Ki-67, a nuclear marker to measure cell cycle, and DAPI to evaluate the DNA content (Fig. 3A). We found that a greater proportion of HSCs and early progenitors were in G0 phase in *Ptk7* deficient embryos as compared to control. Indeed, 56% of HSC-MPP1 cells were quiescent in *Ptk7* deficient embryos with respect to 46% in control embryos (Fig. 3B). This significant increase in quiescent cells was maintained throughout hematopoietic differentiation with respectively 46% versus 39% and 19% versus 14% of the cells in G0 phase at the MMP2 and MPP3-MPP4 stages (Fig. 3B). To exclude that these unbalanced proportion of quiescent cells was not due to selective apoptosis of cycling *Ptk7* deficient cells, we analyzed the percentage of apoptotic cells in *Ptk7* deficient and control FL cells. We did not observed a difference of apoptosis as measured by Annexin V staining between wild type and mutant mice in total FL cells and LSK cells (Fig. 3C). These results demonstrate that PTK7 delivers proliferative signals to HSCs but does not affect HSC survival.

PTK7 deficient cells are able to reconstitute definitive hematopoiesis in recipient mice

To investigate the functional capacity of *Ptk7* deficient HSCs, we generated chimeric mice using FL cell transplantation. CD45.1 C57Bl/6 mice were lethally irradiated (8 Gy) and then reconstituted with 3.10^6 total FL cells from *Ptk7* deficient or wild type FL embryos (CD45.2). Reconstitution and chimerism were followed over time by monitoring the frequency of donor cells (CD45.2) versus recipient cells (CD45.1) in the peripheral blood of recipient mice (Fig. 4A). As expected, all recipient mice injected with control FL cells were reconstituted between 2 and 12 weeks. The same results were obtained with PTK7 deficient FL cells demonstrating that in absence of any competition, *Ptk7* deficient HSCs

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are able to reconstitute all mature hematopoietic lineages of lethally irradiated recipients (Fig. 4B-C). To investigate whether *Ptk7* deficiency could lead to more subtle hematopoietic defects, BM of chimeric mice was analyzed after twelve weeks. No significant difference was found in frequencies of mature and progenitor cells in the BM (Fig. 4E-F). In agreement with reconstitution experiments, in *vitro* differentiation assays showed that *Ptk7* deficient FL cells are able to give rise to all subtypes of colony forming units in the same proportion than control FL cells (Fig. 4G, left panel and supplementary Fig.S4). However, we observed fewer colonies generated by *Ptk7* deficient HSPC than by control cells (Fig. 4G, right panel). This is likely due to decreased cell cycle activity of *Ptk7* deficient HSPCs while the potential of differentiation is not affected. Since quiescence of HSCs could be correlated to loss of self-renewal (36), we then tested self-renewal capacities of *Ptk7* deficient HSCs by serial transplantation assays. Recipient CD45.1 mice were lethally irradiated and reconstituted with CD45.2 donor cells from wild type and mutant mice. The first injection was performed with 3.10^6 total FL cells. After 12 weeks, BM cells were harvested and 3.10^6 cells were then injected to a second lethally irradiated recipient. The same protocol was applied for the following transplantations. We monitored chimerism by flow cytometry analysis and obtained a very good reconstitution with about 95 to 100% of donor cells in BM for all transplantations (Fig. 4H). We also investigated erythroblastic reconstitution in mice and did not find any differences in term of hemoglobin level and red blood cell count (Supplementary Fig. S3A and S3B). *In vitro* differentiation using TF1 cells that are able to differentiate in response to EPO did not reveal any difference in the presence or absence of PTK7 (Supplementary Fig. S3C and S3D). We thus concluded that PTK7 is not required for hematopoietic differentiation.

To get further insights into PTK7 function in early hematopoiesis, we performed mixed chimera competition experiments. CD45.1 C57Bl/6 mice were lethally irradiated (8 Gy)

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and then reconstituted with 2.10^6 total FL cells from *Ptk7* deficient (CD45.2) and 2.10^6 total FL cells from wild type FL embryos (CD45.2xCD45.1). Reconstitution and chimerism were followed over time by monitoring the frequency of *PTK7*$^{-/-}$ cells (CD45.2) and *PTK7*$^{+/+}$ (CD45.2$^+$CD45.1$^+$) versus recipient cells (CD45.1) in the peripheral blood of recipient mice. We did not observe significant differences at D30 and D60, but observed an advantage of *PTK7*$^{-/-}$ cells at long term (>120 jours) (Fig. 5A). Since maintenance of quiescence is a stem cell property, we tested cell cycle activity of adult HSC in chimeric mice. Recipient CD45.1 mice were lethally irradiated and reconstituted with CD45.2 donor cells from wild type and mutant FL mice as previously described. We performed cell cycle analysis on hematopoietic progenitors more than 6 months post transplantation, when stable hematopoietic reconstitution is obtained. We confirmed that hematopoietic progenitors isolated from mice reconstituted with PTK7 deficient cells are more quiescent than wild type counterparts (Fig. 5B and 5C). Clonogenic experiments confirmed that fewer colonies are generated by *Ptk7* deficient HSPC than by control from chimeric mice (Fig. 5D). These results suggest that PTK7 is essential for quiescence independently of the microenvironment.

Homing and migration defects of Ptk7 deficient cells

Since we have previously described a pro-migratory function of PTK7 in leukemic cell lines and because increased HSC quiescence may not be the unique cause of the functional defect of *Ptk7*$^{-/-}$ HSCs (Fig. 2J), we decided to investigate the role of PTK7 in hematopoietic homing. FL cells from several *Ptk7*$^{-/-}$ or *Ptk7*$^{+/+}$ embryos were pooled and stained separately with two different fluorescent trackers. Cells were injected according to a 1/1 ratio and mice were sacrificed 12h after injection. Analysis of homing was performed by flow cytometry (Fig. 6A-C). Although we observed comparable levels of residual

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circulating cells regardless the PTK7 status in peripheral blood, *Ptk7* deficient cells had a very poor capacity to home to the BM, spleen, liver, and thymus as compared to control (Fig. 6D). Since homing defect can be explained by loss of cell adhesion to BM stromal cells and/or by a defect of cell migration toward a chemoattractant, we tested these two hypotheses. Adhesion of *Ptk7* deficient LSK cells on the murine MS5 stromal cell line was not significantly different from control LSK cells (Fig. 7A). In contrast, we found a decrease in spontaneous transmigration of *Ptk7* deficient FL cells as compared to control cells in Boyden chamber assay (8% versus 5%, Fig. 7B). No effect was observed with heterozygous FL cells. We also observed significant differences when the chemokine SDF1α was used as a chemoattractive molecule. Indeed, whereas 12% of control FL cells migrated toward SDF1α, only 8% of PTK7 deficient FL cells were able to do so (Fig. 7C). Without chemoattractant, less than 5% of LSK cells migrated across the transwell regardless the PTK7 status (Fig. 7D). When SDF1α was used as a chemoattrative molecule, a twofold decrease in LSK transmigration was observed for *Ptk7* deficient cells as compared to control cells (20% versus 40%, Fig. 7E).

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412 **Discussion**

413 Discovery of novel molecular pathways implicated in HSC maintenance is of
414 particular interest to develop new strategies aiming at the regeneration of damaged
415 hematopoietic tissues and to understand hematopoietic diseases such as leukemia. We have
416 previously identified the cell polarity protein, PTK7, as a novel tyrosine kinase receptor
417 overexpressed in acute myeloid leukemia (AML). High expression of PTK7 is correlated to
418 poor prognosis and resistance to chemotherapeutic regimens (27). Recently, another study
419 revealed that PTK7 is also upregulated in T acute lymphocytic leukemia (T-ALL)(37). In
420 humans, we and others have evidenced that PTK7 expression is mostly restricted to normal
421 early progenitors committed to myeloid and T lymphoid lineages.

422 In the present study, we show that PTK7 expression is similar between mouse and
423 human systems. PTK7 is absent in mature blood cells and its expression in BM
424 hematopoietic progenitors is inversely correlated with differentiation. The highest
425 expression of PTK7 is found in HSC-MPP1 compartment. PTK7 expression is also
426 conserved between mice and human when the thymic compartment is considered. PTK7 is
427 easily detectable at the cell surface of CLP and immature DN and DP populations whereas
428 its expression is lost in single positive CD4 or CD8 lymphocytes. These data correlate with
429 results from other groups that demonstrated expression of PTK7 by recent thymic emigrants
430 cells and thymocytes (38)(37).

431 In order to study the role of PTK7 in hematopoiesis, we generated *Ptk7* deficient mice using
432 available gene-trapped ES cells. Our *Ptk7* deficient strain exhibited expected planar cell
433 polarity defects as compared to another strain of mice issued from the same ES clone (21).
434 Since these mutant mice cannot reach adulthood, functions of PTK7 were investigated in
435 FL hematopoietic cells. The major hematopoietic phenotype of *Ptk7* deficient mice consists
436 in increased quiescence of HSPCs and shrinkage of the FL HSPC pool. Although these

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phenotypes may sound logical if one considers that less proliferation produces less cells, they are difficult to reconciliate with loss of functional reconstitution observed when recipient mice are engrafted with limited numbers of LSK cells (Fig. 2J). Indeed, PTK7 is implicated in canonical and non-canonical Wnt pathway (17)(39), which are thought to be the guardians of the balance between HSC self-renewal and HSC differentiation (32). We have recently shown that PTK7 plays a role in canonical pathway downstream of Wnt3a suggesting that deficiency in PTK7 may mirror deficiency in Wnt3a (25). This would be consistent with results showing that Wnt3a deficiency leads to a reduction in the numbers of HSPCs in the FL but would not fit with the reduced reconstitution capacity observed in secondary transplantation assays of Wnt3a deficient cells which is not observed with *Ptk7* deficient cells (40). Alternatively, it may well be that PTK7 controls non-canonical pathway through interaction with Wnt ligands (41) (42). In this way, Wnt5a antagonizes Wnt3a-mediated canonical signaling and increases quiescence of HSCs (43). This occurs through activation of the small Rho GTPase Cdc42 that increases HSC polarity (44)(45). Such a function of PTK7 in Wnt5a-mediated non-canonical pathway would fit with the planar cell polarity function of PTK7 and with altered homing and migrating properties of *Ptk7* deficient cells that likely depend on small Rho GTPase activation. Some studies show that PTK7 regulates the balance between Wnt canonical and non-canonical pathways by interacting with Dishevelled (26)(46), ROR2 (42)(47), β-catenin (25) and LRP6 (39). We unsuccessfully tried to identify downstream pathways that could be regulated by Ptk7 in HSCPs using a candidate gene/protein approach. We also looked at gene expression differences between Ptk7 knock-out and control hematopoietic cells using unbiased gene expression profiling. Although we identified differences (data not shown), we could not find obvious Gene Ontology enrichments or pathways that could be experimentally tested. Further studies will be needed to unravel how Ptk7 functions in HSPCs.

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462 In our model of FL hematopoietic cells, we confirmed the role of PTK7 in cell
463 migration and found a new role of this protein in cell homing to hematopoietic organs and
464 in the control of cell proliferation. This observation confirms our previous data obtained
465 from human leukemia samples (27) can be linked with already described functions of PTK7
466 in cell migration and in metastatic dissemination. Indeed, we and others have found that
467 PTK7 expression on various cancer cells is linked with higher metastasis potential
468 (27)(29)(48)(49). Recently, we also described a poor prognosis of PTK7 expression in
469 colon cancer cells linked with higher metastatic events (50). In addition, our data show for
470 the first time that PTK7 is involved in the regulation of hematopoietic and stem and
471 progenitors cell (HSPC) cycle. Lack of PTK7 expression increases quiescence and
472 enhances the long term repopulating potential after six months, while a trend toward
473 decreased chimerism is observed after one or two months. This could reflect the dual
474 function of PTK7 loss of expression in decreasing short term homing to the bone marrow
475 and increasing long term establishment of hematopoiesis. Alternatively, it may well be that
476 homing assay performed with total liver cells reveals a constitutive promigratory function
477 of PTK7, in contrast to long term engraftment reflecting bone marrow micro-environmental
478 signaling through PTK7 in HSCs. Similar complex mechanisms of cell cycle regulation and
479 differential homing of mature cells versus HSPCs homing to the bone marrow have also
480 been attributed to CXCR4 (51) (52) (53) (54). Whether PTK7 and CXCR4 signaling share
481 more in common remains to be addressed. In conclusion, PTK7 overexpression in
482 immature blasts cells of acute myeloid leukemia has been associated to bad prognosis, early
483 relapse and chemotherapeutic resistance and a recent study shows that PTK7 can be used as
484 a stem cell marker of colon cancer stem cells (27)(55). Taken together, all these data form a
485 set of arguments to involve PTK7 expression in stem cells including cancer stem cells but
486 also to prove a functional role of PTK7 in stem cell biology. We think that these new data

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487 on PTK7 involvement on cell cycle are very promising to understand molecular
488 mechanisms of leukemia/cancer development.
489 Altogether, our results report a novel unexpected function for the planar cell polarity
490 molecule PTK7 in the crosstalk between expansion of HSPC compartment and HSPC
491 migration. This opens new avenue for manipulation of HSC fate using agonists or
492 antagonists of PTK7.
493

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Author contribution

A.-C.L designed experiments, wrote the paper, performed all experiments and generated deficient mice; M.D.G, M.G., F.L., F.B., S.M. helped to perform experiments; J.-C.O. generated deficient mice; M-L.A., M.A-L, J-P.B. designed experiments and wrote the paper.

Disclosure of potential conflicts of interest

All authors declare no conflict of interest.

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694 **Figure Legends:**
695

696 **Figure 1: PTK7 expression in wild type bone marrow.** (A) Expression of PTK7 in blood
697 mature cells. Granulocytes are defined as $Gr1^{+}CD11b^{+}$ cells. Monocytes are defined as
698 $F4/80^{+}$ $CD11b^{+}CD115^{+}$ cells. Lymphocytes are gated using B220, CD19, CD3, CD4 and
699 CD8 combination. MFI ratios are indicated in dots plots. (B) Strategy of bone marrow
700 progenitors gating. First panel: Dot plot showing the LSK^{+} gate within Lin^{-} bone marrow
701 cells. Second panel: dot plot showing early progenitors gating using SLAM markers.
702 Hematopoietic stem cells (HSC) are defined as $CD150^{+}CD48^{-}$ cells. More mature
703 multipotents progenitors (MPP) are defined as $CD150^{+}CD48^{+}$ (MPP2) and then $CD150^{-}$
704 $CD48^{+}$ (MPP3/4). Third panel: dot plot showing engaged progenitors within $c\text{-}Kit^{+}Sca\text{-}1^{-}$
705 Lin^{-} cells. CMP (Common Myeloid Progenitor) is defined as $CD16/32^{lo}CD34^{+}$. GMP
706 (Granulocyte-Macrophage Progenitor) is defined as $CD34^{+}CD16/32^{hi}$. MEP
707 (Megakaryocyte-Erythroid Progenitor) is defined as $CD16/32^{-}CD34^{-}$. Fourth panel: Gating
708 of CLP (Common Lymphoid Progenitor) defined as $Lin^{-}IL7Ra^{+}B220^{-}CD24^{lo}CD117^{lo}$. (C)
709 Overlays showing expression of PTK7 during differentiation. Black line: PTK7, gray
710 histogram: isotypic control (pre-immune rabbit antiserum). (D) Quantitative comparison of
711 expression levels between different progenitor populations: normalization of PTK7 mean of
712 fluorescence (MFO) with CD117 MFO on LSK subpopulations and progenitors
713 ($CD117^{+}Sca\text{-}1^{-}$ cells). Statistical significance was calculated by nonparametric Mann-
714 Whitney test. All data are shown as the mean ± s.d. *$p<0.05$. **$p<0.01$.***$p<0.001$.
715

716 **Figure 2: Phenotype of PTK7 deficient mouse and analysis of fetal liver at E14.5.** (A)
717 PTK7 mutants embryos showing severe planar cell polarity and developmental
718 abnormalities. (B) Western blot of murine embryonic fibroblasts (MEFs) obtained from a

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719 litter of our PTK7 gene trapped established mouse line. (C) Dot plots showing LSK gate
720 within lin$^-$ FL cells and SLAM gating within LSK cells. (D) Quantification of total number
721 fetal liver cells. Data are pooled from 6 litters: *Ptk7*$^{-/-}$ N=12; *Ptk7*$^{+/-}$ N=19; *Ptk7*$^{+/+}$ N=12
722 (E) Percentage of CD45$^+$ cells within fetal liver. Data are pooled from 8 litters: *Ptk7*$^{-/-}$
723 N=15*; Ptk7*$^{+/-}$ N=26*; Ptk7* $^{+/+}$ N=9**.** (F) Percentage of LSK cells within CD45$^+$ FL cells. (G)
724 Percentage of LSK subpopulation using SLAM markers. (H) Absolute number of LSK cells
725 in E14.5 FL. (I) Absolute number of LSK subpopulations using SLAM markers. From (F)
726 to (I) data are pooled from 10 litters: Ptk7$^{-/-}$ N=19; Ptk7 $^{+/-}$ N=29; Ptk7$^{+/+}$ N=10. (J) Survival
727 curves of CD45.1 C57Bl/6 mice engrafted with 400 LSK cells sorted from CD45.2 FL
728 together with 1.10^{5}congenic CD45.1 total bone marrow cells. Black line: engraftment with
729 *Ptk7*$^{+/+}$ FL cells; Dashed line: engraftment with *Ptk7*$^{-/-}$ FL cells. Shown experiment is
730 representative from 3 separate experiments. N=6 mice per group. (K) Clonogenic capacity
731 of FL total cells. 3.10^4 FL cells are seeded in M3434 methylcellulose. Shown experiment is
732 representative from 3 separate experiments. *Ptk7*$^{-/-}$ N=6; *Ptk7*$^{+/+}$ N=4. All statistical
733 significance was calculated by nonparametric Mann-Whitney test. All data are shown as the
734 mean ± s.d. *p<0.05. **p<0.01.***p<0.001.
735
736 **Figure 3: Cell cycle and apoptosis analysis of LSK FL cells.** (A) Representative dot plots
737 of DNA content (DAPI) plotted versus Ki-67 nuclear antigen staining on LSK cells. The
738 cell-cycle phases were defined as G0 (Ki-67$^-$ and *2n* DNA), G1(Ki-67$^+$ and *2n* DNA) and S-
739 G2-M (Ki-67$^+$ and DNA >*2n*). (B) Percentage of LSK FL cells in cell-cycle phases using
740 SLAM markers. Data are pooled from 3 separate experiments and embryos (Ptk7$^{-/-}$ N=10;
741 Ptk7$^{+/+}$ N=11) from 7 different litters. *p<0.05. **p<0.01. (C) Determination of percentage
742 of apoptotic cells by staining with annexin V and DAPI. Apoptotic cells were defined as
743 annexin V$^+$ DAPI$^-$. Results are representative of 8 *Ptk7*$^{-/-}$ and 6 WT embryos from 5

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different litters. All statistical significance was calculated by nonparametric Mann-Whitney test. All data are shown as the mean ± s.d. *p<0.05. **p<0.01.***p<0.001.

Figure 4: Hematological capacity of reconstitution and differentiation. Lethally irradiated CD45.1 mice (8 Gy) were reconstituted with 3.10^6 CD45.2 total FL cells isolated from *Ptk*$^{+/+}$ and *Ptk*$^{-/-}$ embryos. Percentage of each hematological populations from donor (CD45.2) was monitored by flow cytometry (A). Percentage of matures cells in recipient blood (B) and in spleen (C). Percentage of BM mature cells (D) and BM progenitors (E) in recipient mice. (F) Percentage of LSK cells and LSK subpopulation in recipient BM. White histograms depict mice reconstituted with *Ptk*$^{-/-}$ FL. Black histograms depict mice reconstituted with *Ptk*$^{+/+}$ FL. N=6 mice per group. (G) *In vitro* differentiation capacity of FL cells. 3.10^4 FL cells were seeded in methylcellulose. Left panel: Quantification and characterization of CFU-C. Right panel: Proportion of CFU. (H) Serial transplantation. First transplantation: Lethally irradiated mice were injected with 3.10^6 of total E14.5 FL cells (T1: transplantation 1). Next transplantations were done by injection of 3.10^6 total bone marrow cells from previously reconstituted mice in irradiated C57Bl/6 CD45.1 mice (T2 to T5). Percentage of donor cells (CD45.2) *versus* recipient cells (CD45.1) was monitored. Each data represents chimerism of definitive hematopoiesis 3 months after irradiation. Data are the mean ± s.d. N=6 mice per group.

Figure 5: Competition experiments and cell cycle in adult mice. (A) 2.10^6 cells from Ptk7$^{+/+}$ (CD45.1xCD45.2) and 2.10^6 cells from Ptk7$^{-/-}$ (CD45.2) are injected in recipient (CD45.1) mice. Chimerism is followed in blood samples. Data are representative of 4 independent experiments (N=6 mice per group). (B-D) Recipient CD45.1 mice reconstituted with CD45.2 donor cells from wild type and mutant FL mice (B)

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769 Representative dot plots of DNA contents (DAPI) plotted versus Ki-67 nuclear antigen
770 staining on LSK cells. (C) Percentage of LSK cells in cell cycle phases using SLAM
771 markers. (D) Clonogenic capacity of BM total cells. 1.10^5 BM cells are seeded in M3434
772 methylcellulose. Data are representative of 2 independent experiments (N=6 mice per
773 group). All statistical significance was calculated by nonparametric Mann-Whitney test.
774 *p<0.05. **p<0.01.***p<0.001.
775
776 **Figure 6: Homing capacities of deficient PTK7 cells.** (A) 2.10^6 of total fetal liver cells
777 from *Ptk7*$^{-/-}$ or *Ptk7*$^{+/+}$ embryos were stained with cell tracker calcein AM and calcein red-
778 orange respectively, pooled in 1/1 ratio and injected in recipient mice. (B-D) 12h after
779 injection, mice were sacrificed and biological samples were removed and analyzed to
780 evaluate homing capacities of each cells . (B) Bone marrow of uninjected mice was used as
781 control for staining specificity. (C) Dot plots of stained cells removed from blood, BM,
782 spleen, liver and thymus. (D) Quantification and statistical analysis. Data are pooled from 3
783 independent experiments or representative (for spleen panel). N=6 mice per group. All
784 statistical significance was calculated by nonparametric Mann-Wihtney test. *p<0.05.
785 **p<0.01.***p<0.001.
786
787 **Figure 7: Adhesion and migration assays** (A). Adhesion assay on MS5 stromal cells.
788 Percentage of adherent FL cells on MS5 stromal bone marrow cell line after 2 hours of
789 incubation. Left panel: total FL cells. Right panel: LSK cells. Data are representative of 3
790 independent experiments. *Ptk*$^{-/-}$ N=4; *Ptk*$^{+/-}$ N=6; *Ptk*$^{+/+}$ N=3. (B to E) Migration in Boyden
791 chamber assay. Number of migrating cells are normalized to the number of input cells for
792 each sample (duplicate wells). (B) Migration of lin^{Neg} enriched FL cells without
793 chemoattractant. (C) Migration of lin^{Neg} enrichied FL cells with SDF1a as chemoattracting

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794 reagent. (D). Migration of LSK FL cells without chemoattractant. (E). Migration of LSK
795 FL cells with SDF1α as chemoattracting reagent. All statistical significance was calculated
796 by nonparametric Mann-Wihtney test. *$p<0.05$. **$p<0.01$.***$p<0.001$.

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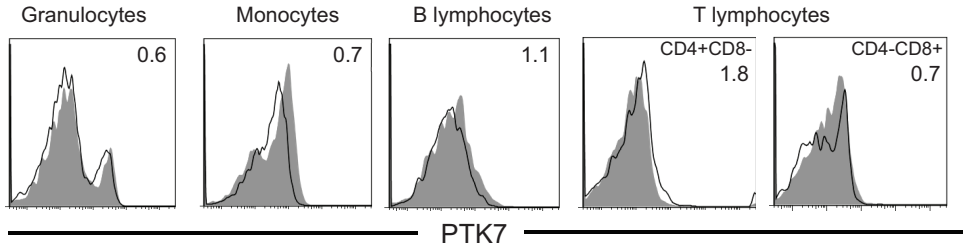

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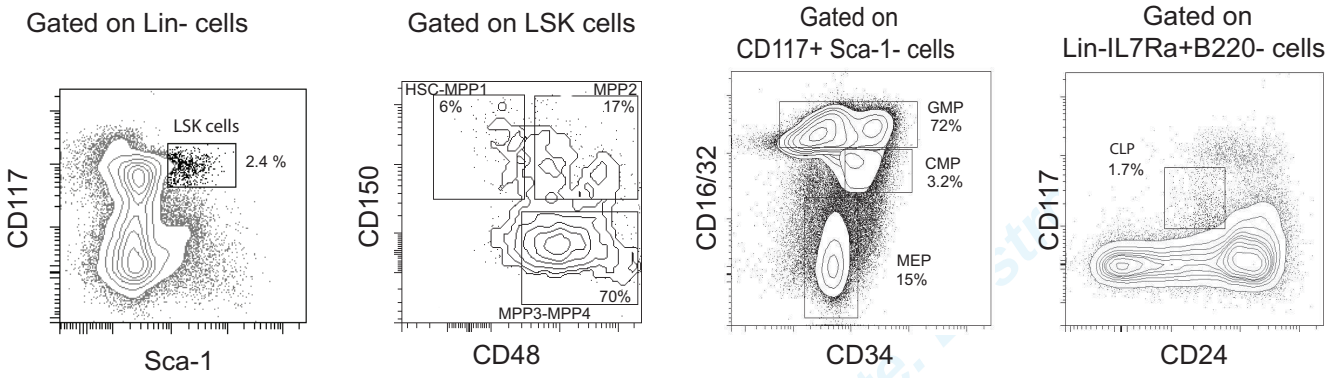

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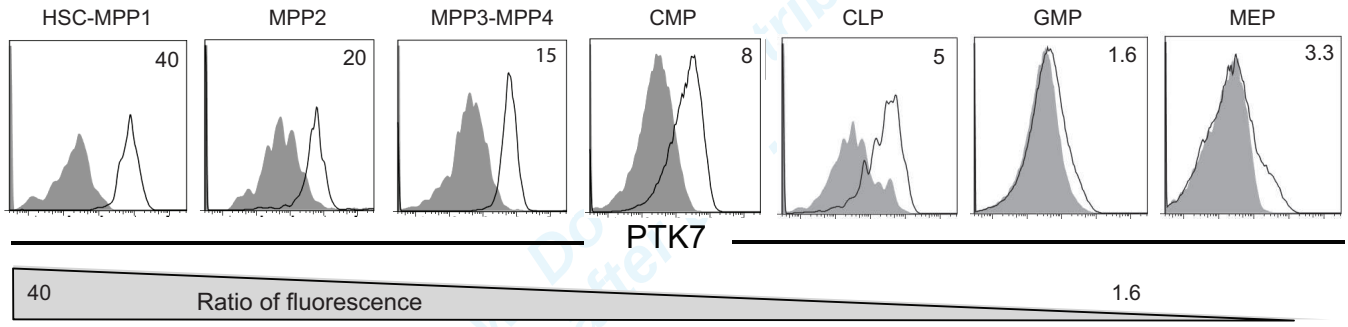

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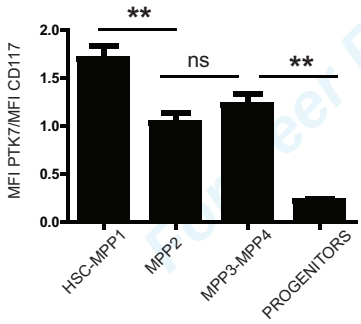
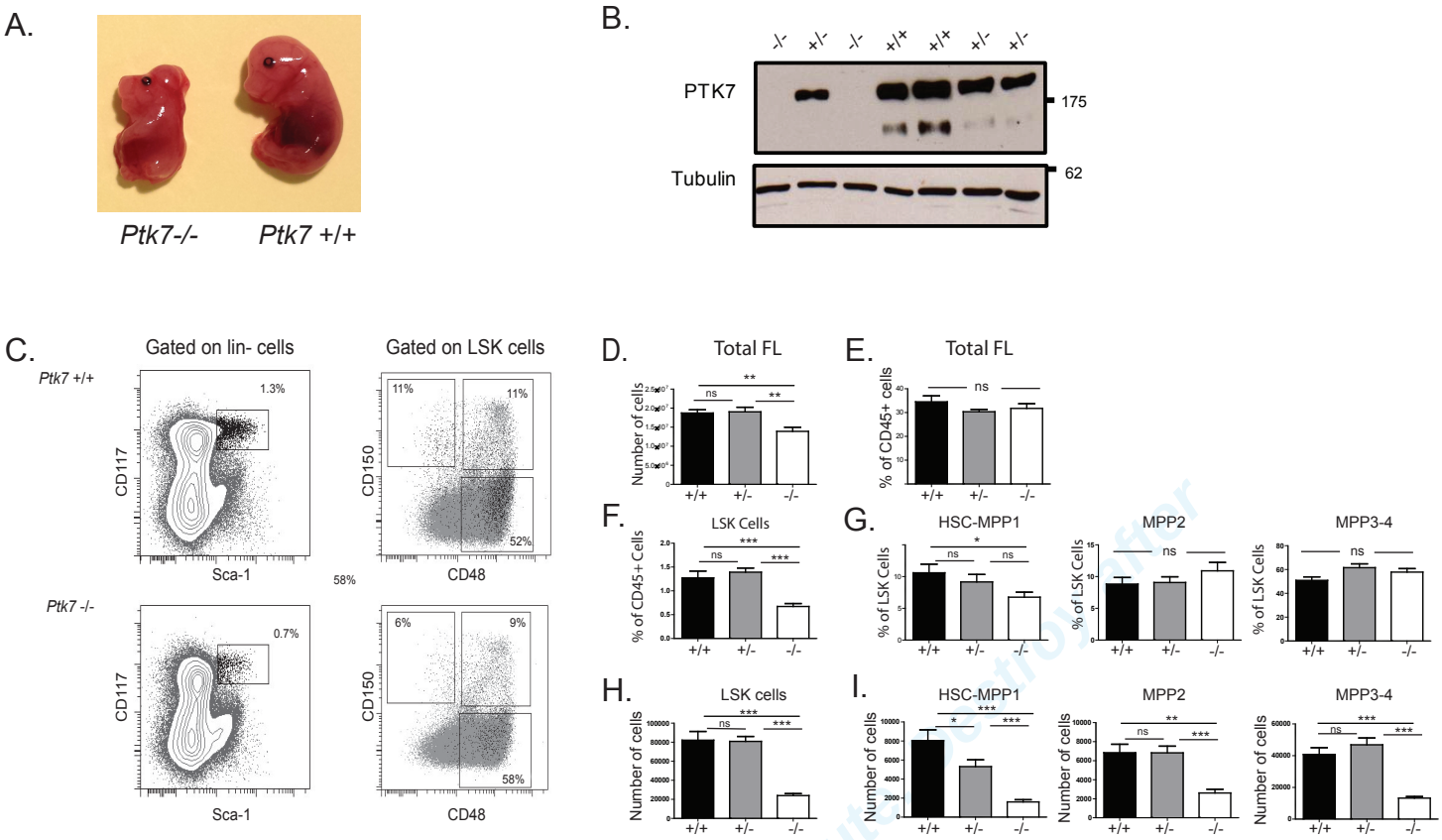
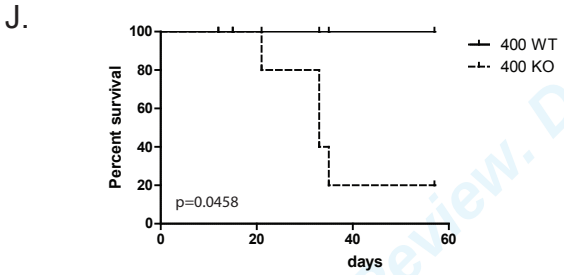


Figure 1





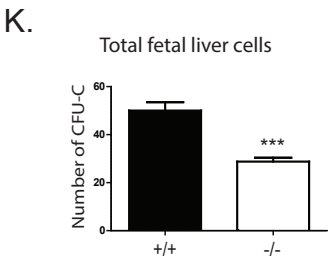


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Figure 2

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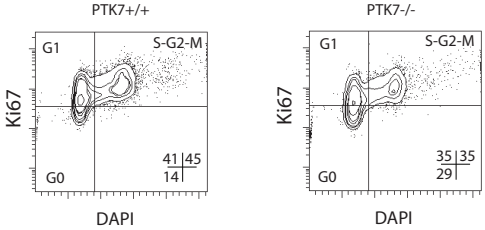

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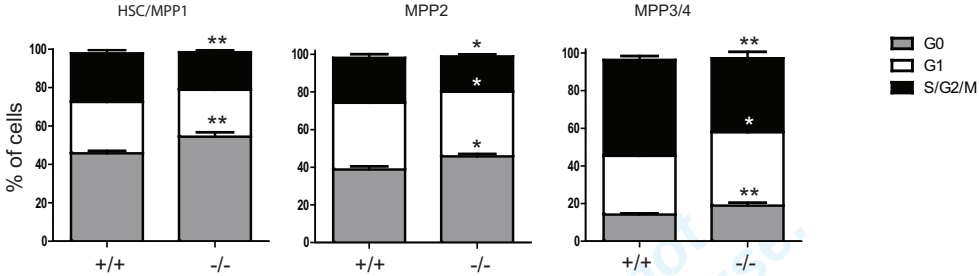

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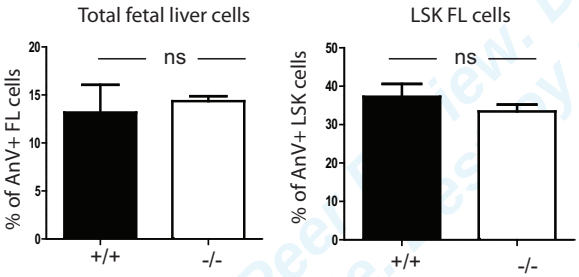

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Figure 3

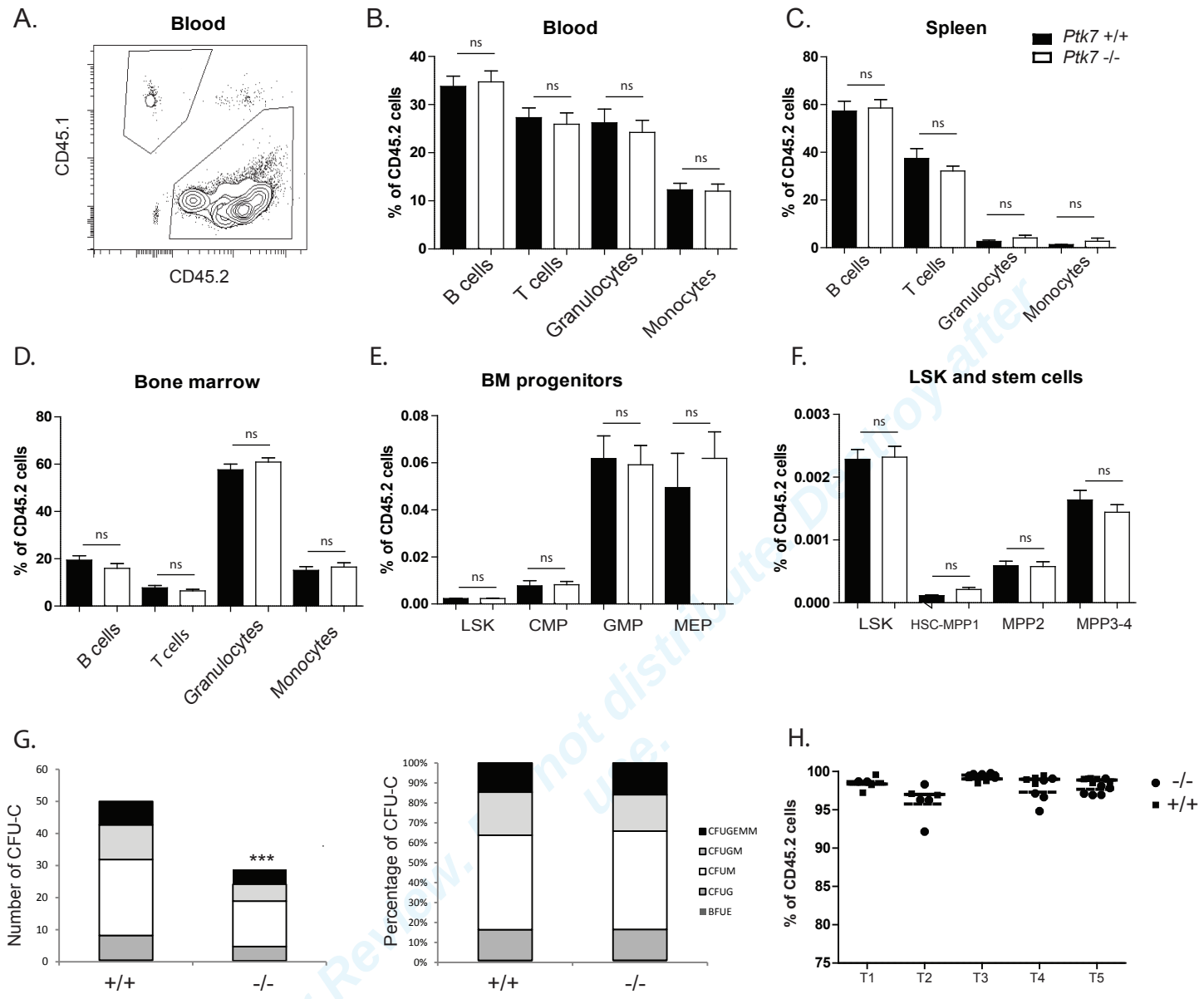

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Figure 4

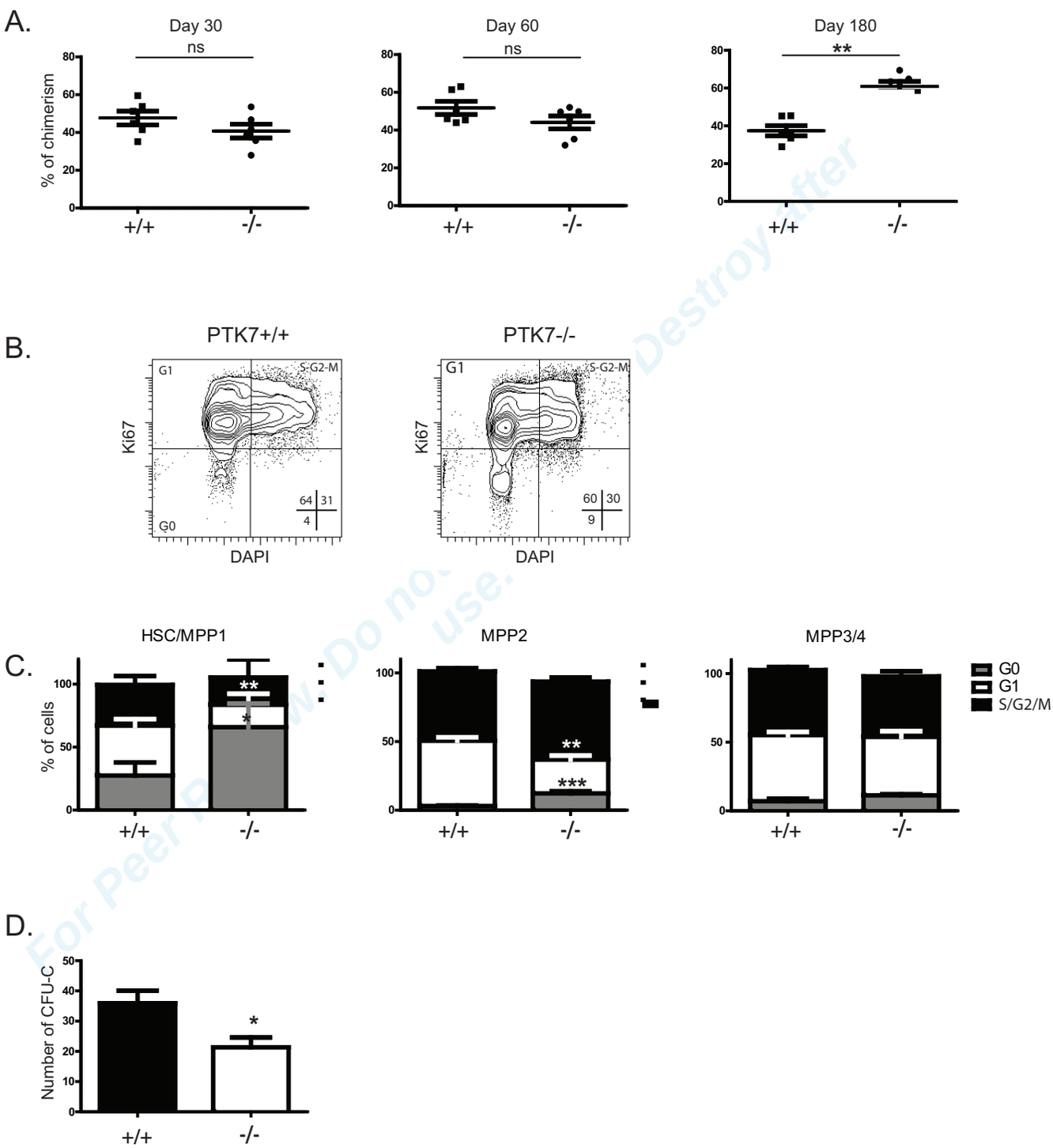
A.
Day 30
ns
Day 60
ns
Day 180
**
% of chimerism
+/+
-/-
B.
PTK7+/+
PTK7-/-
G1
S-G2-M
G0
Ki67
DAPI
64
31
4
60
30
9
C.
HSC/MPP1
MPP2
MPP3/4
% of cells
G0
G1
S/G2/M
D.
Number of CFU-C

Figure 5

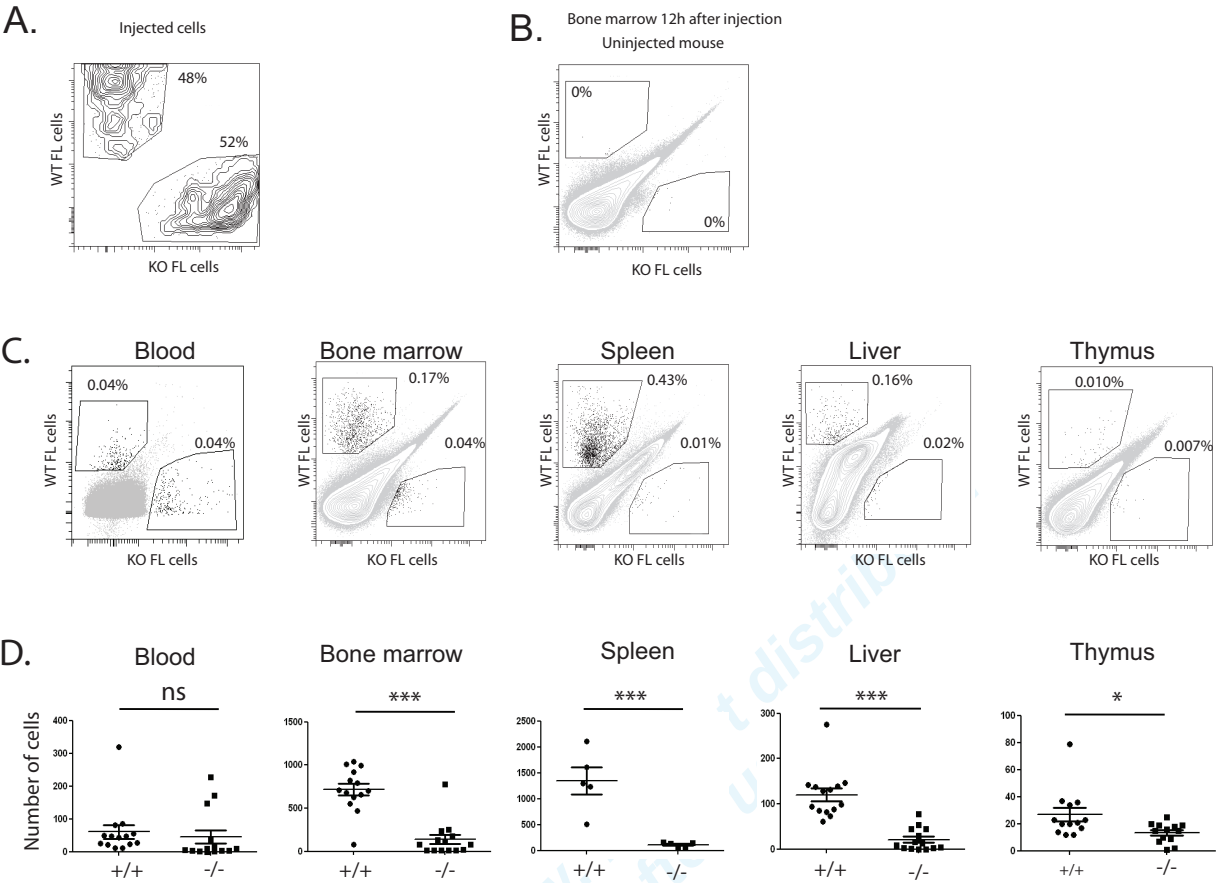

Figure 6

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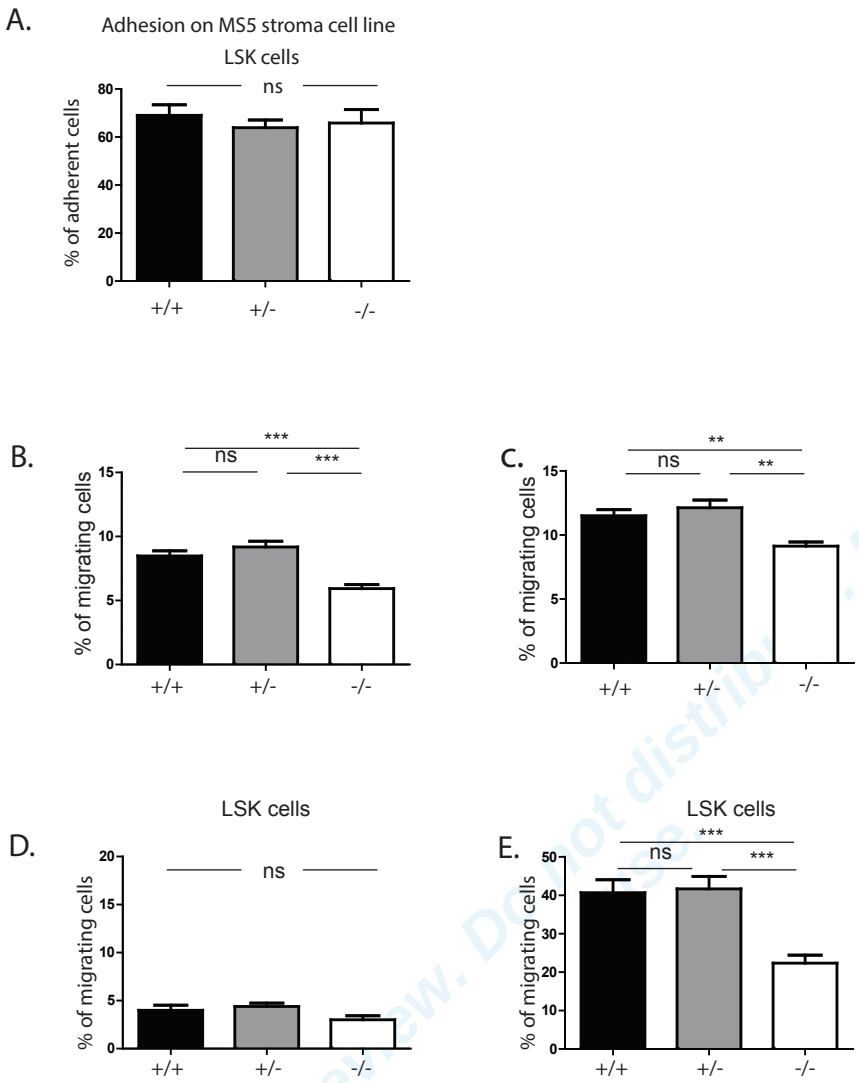

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Figure 7