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Supplemental Information

Improved Human Erythropoiesis and Platelet Formation in Humanized

NSGW41 Mice

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Figure S1:









Figure S1. Analysis of murine and human erythroid progenitors in humanized mice by flow cytometry, Related to Figure 1

(A) Dot plots show the expression of Ter119 and mCD71 on CD45⁻ (human and mouse) bone marrow cells of humanized NSG and NSGW41 mice. (B) Total numbers of murine RBC differentiation stages in 2 femura of non-transplanted and humanized NSG and NSGW41 mice. 6 expts, 2-9 mice each per expt. Depicted are mean \pm SD. (C) Gating strategies to determine the relative contribution of human donor cells to the indicated erythroid progenitor compartments. (D) Dot plots (left) and graph (right) show the presence of CD71⁺ erythroid progenitors within ACK-lysed or CD34-enriched human cord blood (2 expts).

Figure S2. Murine macrophages in humanized NSG and NSGW41 mice, Related to Figure 3

(A) Plot shows the frequency of $CD11b^+$ Gr-1^{lo} macrophages in the blood of humanized or nonhumanized NSG or NSGW41 mice. Mice had received human CD34-enriched CB cells 11-21 weeks before. 3 expts, 2-4 mice each per expt. Depicted are mean \pm SD. (B) Phagocytic activity of murine blood macrophages from indicated mouse strains before and after humanization.

Figure S3. Human thrombocytes in bone marrow after CloLip treatment, Related to Figure 4

NSGW41 mice were injected with 200 μ l clodronate liposomes (CloLip) to deplete macrophages 11 weeks after humanization. (A) Frequency of human platelets within all platelets in bone marrow of CloLip- or PBS-treated NSGW41 mice. (B) Total numbers of human platelets in 2 femura of CloLip- or PBS-treated NSGW41 mice. 1 expt, 4 mice each. Depicted are mean \pm SD.

Supplemental Experimental Procedures

Genotyping of NSGW41 mice

Genotyping of Kit^{W41} allele was performed by PCR on tail-snip DNA (forward: 5²-AAGGAAGGTTAGAACCCCTGG; reverse: 5²-AGCTCCCAGAGGAAAATCCC). The products were sequenced using the forward primer. The Kit^{W41} allele contains a point mutation at position 2519 (G / A) (Nocka et al., 1990).

Antibodies

Human cells were stained using antibodies specific for (clone names given in brackets): CD3 (OKT3), CD19 (HIB19), CD33 (WM-53), CD42a (GR-P), CD45 (HI30), CD235 (HIR2) from eBioscience, CD45 (HI30), CD16 (3G8), CD41 (HIP8), CD61 (VI-PL2) from Biolegend, CD14 (M5E2) (BD Biosciences) and CD71 (MEM-75) from ImmunoTools. Murine cells were stained with the following antibodies: CD11b (M1/70), CD41 (eBioMWReg30), CD45 (30-F11), CD115 (AFS98), Gr-1 (RB6-8C5), Ly-6C (HK1.4), Ter119 (TER-119) (all from eBioscience), CD41 (MWReg30) and CD71 (C2) from BD Biosciences. Blocking reagents used were ChromPure mouse IgG (Jackson ImmunoResearch) for human cells and rat IgG (Dianova) as well as anti-mouse CD16/CD32 (93) (eBioscience) for murine cells. For immunohistochemistry sections were blocked with rat Ig (500 μ g/ml, Jackson ImmunoResearch Laboratories) and Streptavidin/Biotin blocking kit (Vector Laboratories). Antibodies used for immunostaining were hCD45 (HI30), mCD45 (30-F11), hCD42a (GR-P) and mCD41 (MWReg30) (eBioscience). Anti-FITC-Alexa488 (Molecular Probes) and streptavidin-Cy3 (Jackson ImmunoResearch Laboratories) were used for secondary steps. Dapi (2 μ g/ml) was used to stain nuclei.

EPO ELISA

Samples were collected into EDTA containing micro tubes (plasma) or non-coated tubes (serum) and centrifuged at 2000g for 15 minutes at RT. Plasma samples from NSGW41 h*EPO*-tg mice were diluted 2- to 10-fold. All serum samples were diluted 1:2 before use. The optical density was measured on a Sunrise[™] absorbance reader (Tecan). A standard curve was generated using nonlinear regression.

May-Gruenwald-Giemsa staining

May-Gruenwald-Giemsa staining was modified from Waskow et al., 2008. In brief, cytospins were stained in 100% and 50% May-Gruenwald solution (Merck, Darmstadt, Germany) for 3 minutes each. After rinsing in water (pH6.8) for 2 minutes the slides were stained in Giemsa (Merck, Darmstadt, Germany) for 15 minutes. Samples were then washed in water (pH6.8) for 3 minutes and quickly rinsed with ethanol and water to remove excessive dye. Slides were air-dried before microscopy.

Primers for globin chain expression

α-globin (NM_000558.4): f: 5'- CTGGAGAGGATGTTCCTGTCCTTG, r: 5'- CAGCTTAACGG TATTTG GAGGTCAT (t_A =59 °C; 24/26/28/30 cycles)

 β -globin (NM_000518.4): f: 5'- TCCTGAGGAGAAGTCTGCCGTTAT, r: 5'- GAAATTGGACAG CAAGA AAGCGGA (t_a=53 °C; 26/28/30/32 cycles)

 γ -globin (NM_000559.2): f: 5'- ACTCGCTTCTGGAACGTCTGA, r: 5'- GTATCTGGAGGAC AGGGCA CT (t_a=57 °C; 24/26/28/30 cycles)

δ-globin (NM_000519.3): f: 5'-GCAGATTACTGGTGGTCTACCCTTG, r: 5'- GGAAACAGTCC AGGAT CTCAAT GG (t_A =55 °C; 26/28/30/32 cycles)

β-actin (NM_001101.3): f: 5'- AGCGAGCATCCCCCAAAGTT, r: 5'- GGGCACGAAGGCTCAT CATT (t_A =59 °C; 28/30/32/34 cycles)

Phagocytosis assay

3ml defibrinated sheep red blood cells (sRBC, Thermo Scientific) were opsonized in 10ml PBS/1% BSA containing 100 μ g/ml rabbit IgG specific for sRBC (MP Biomedicals) for 30' at 37°C and 30' on ice and subsequently labeled (4*10⁷ sRBCs/ml) with CFSE at a final concentration of 5 μ M. WBCs from lysed blood of non-transplanted and humanized (21 weeks post Tx) NSG and NSGW41 mice were incubated with opsonized CFSE-labeled sheep RBCs in a ratio of 1:100 for 4h at 37°C. Subsequently, sheep RBCs were lysed in ACK buffer for 1min, cells were stained and analyzed by

flow cytometry. Non-opsonized sRBCs were used as control, and non-engulfed sheep RBCs sticking to WBCs were excluded by gating on PE anti-rabbit IgG negative cells.

Supplemental References

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Waskow, C., Liu, K., Darrasse-Jeze, G., Guermonprez, P., Ginhoux, F., Merad, M., Shengelia, T., Yao, K., and Nussenzweig, M. (2008). The receptor tyrosine kinase Flt3 is required for dendritic cell development in peripheral lymphoid tissues. Nature immunology *9*, 676-683.