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Development of Mature and Functional Human Myeloid Subsets in Hematopoietic Stem Cell-Engrafted NOD/SCID/IL2rγKO Mice

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Although physiological development of human lymphoid subsets has become well documented in humanized mice, in vivo development of human myeloid subsets in a xenotransplantation setting has remained unverified. Therefore, we investigated in vivo differentiation and function of human myeloid subsets in NOD/SCID/IL2rγnull (NSG) mouse recipients transplanted with purified lineage−CD34+CD38− cord blood hematopoietic stem cells. At 4–6 mo posttransplantation, we identified the development of human neutrophils, basophils, mast cells, monocytes, and conventional and plasmacytoid dendritic cells in the recipient hematopoietic organs. The tissue distribution and morphology of these human myeloid cells were similar to those identified in humans. After cytokine stimulation in vitro, phosphorylation of STAT molecules was observed in neutrophils and monocytes. In vivo administration of human G-CSF resulted in the recruitment of human myeloid cells into the recipient circulation. Flow cytometry and confocal imaging demonstrated that human bone marrow monocytes and alveolar macrophages in the recipients displayed intact phagocytic function. Human bone marrow-derived monocytes/macrophages were further confirmed to exhibit phagocytosis and killing of Salmo nella typhimurium upon IFN-γ stimulation. These findings demonstrate the development of mature and functionally intact human myeloid subsets in vivo in the NSG recipients. In vivo human myelopoiesis established in the NSG humanized mouse system may facilitate the investigation of human myeloid cell biology including in vivo analyses of infectious diseases and therapeutic interventions. The Journal of Immunology, 2012, 188: 6145–6155.

The use of genetically modified mice has led to significant advances in biomedical research by providing insights into the role of individual genes both in normal physiology and in disease pathogenesis. However, translation of these findings into effective therapies for human diseases has been limited by the gap that exists between murine and human biology. The availability of human samples (e.g., cells and tissues), although supporting successful translation from bench research to clinical medicine, is limited by both logistic and ethical concerns.

Therefore, mice repopulated with human hematopoietic cells through xenogeneic transplantation were developed directly to investigate the human immuno-hematopoietic system in vivo. Based on pioneering work using the Hu-PBL-SCID (1) and SCID-hu (2) systems, investigators have aimed to improve xenotransplantation models using immunodeficient strains of mice as recipients of human hematopoietic stem cells (HSCs) to achieve long-term engraftment of multiple human hematopoietic and immune subsets (3). NOD/SCID mice, established by back-crossing the scid mutation onto the NOD strain background, are characterized by partially impaired innate immunity and deficient complement-dependent cytotoxicity and are the gold standard for stable human hematopoietic stem/progenitor cell engraftment (3, 4). The ability of NOD-scid mice to support human HSC engraftment is associated with a human-like polymorphism in the IgV domain of the signal regulatory protein-α (SIRPa) allele in the NOD strain background resulting in the expression of SIRPa with enhanced binding of human CD47 (5). The interaction between SIRPa and CD47 has previously been implicated in the negative regulation of phagocytosis by macrophages through a “do-not-eat-me signal” (6).

The introduction of IL-2γ common γ-chain mutations onto the NOD/SCID and BALB/c-Rag2KO backgrounds led to the generation of the NOD/SCID/IL2rγnull (NOG) (7, 8) strain with a truncated IL-2γR, the NOD/SCID/IL2rγnull (NSG) (9, 10) strain with a complete IL-2γR-null mutation, and BALB/c-Rag2KO/γcnull mice (11), resulting in more profound defects in innate immunity. In vivo human hematopoietic repopulation through transplantation...
of human CD34+ or CD34+CD38− HSCs in these severely immunocompromised recipients accelerated human stem cell and immunology research by allowing higher levels of human HSC engraftment, differentiation of human T cells in the murine thymus and secondary lymphoid organs, enhanced maturation of human B cells, and human immune function in vivo (7–12). Despite these advances, one of the remaining issues to be clarified in humanized mouse research has been the development of human myeloid lineage cells in the host mouse tissues. To date, we and others reported the development of human CD33+ myeloid cells and myeloid subsets in NOG mice, BALB/c-Rag2KO/− mice, and secondary lymphoid organs, enhanced maturation of human CD33+ myeloid cells and myeloid function in vivo (7–12). In this study, by using NSG newborns as recipients, we present in vivo differentiation of human myeloid subsets in the bone marrow (BM), spleen, and respiratory tract of NSG mice engrafted with purified lineage−CD34+CD38− human cord blood (CB) HSCs. Human granulocytes (neutrophils, basophils, and mast cells) and APCs (monocytes/macrophages, conventional dendritic cells [cDCs] and plasmacytoid dendritic cells [pDCs]) developing in NSG mice exhibited characteristics of human myeloid cells including morphological features and expression of surface molecules known to be associated with the myeloid cell subsets. Moreover, human myeloid cells developing in the NSG recipients displayed functionality such as responsiveness to cytokine or TLR adjuvant and phagocytic function. The in vivo system supporting the development of mature human myeloid cells with intact function will facilitate the evaluation of human myeloid development from hematopoietic stem/progenitor cells, advance in vivo investigation of human myeloid cell-mediated immune responses against pathogens and malignancies, and will support studies of therapeutic agents.

Materials and Methods

Mice

NOD.Cg-PrkdcsidIl2rgtm1Wjl/J (NSG) mice were developed at The Jackson Laboratory by back-crossing a complete null mutation at the Il2rg locus onto the NOD.Cg-PrkdcsidIl2rgtm1Wjl (NOD/SCID) strain (9, 10). Mice were bred and maintained under defined flora with irradiated food at the animal facility of RIKEN and at The Jackson Laboratory according to guidelines established by the institutional animal committees at each respective institution.

Purification of human HSCs and xenogeneic transplantation

All experiments were performed with authorization from the Institutional Review Board for Human Research at RIKEN Research Center for Allergy and Immunology. CB samples were first separated for mononuclear cells (MNCs) by LSR lymphocyte separation medium (MP Biomedicals). CB MNCs were then enriched for human CD34+ cells by using anti-human CD34 microbeads (Miltenyi Biotec) and sorted for 7AAD− cells (98%) with purity above 98% by analysis of forward scatter-height/forward scatter-width and side scatter-height/side scatter-width. The purity of HSCs was higher than 98% after sorting. Newborn (within 2 d of birth) recipients received 150 cGy total body irradiation using a 137Cs source irradiator, followed by i.v. injection of 1 × 10^5–3 × 10^5 sorted HSCs via the facial vein (14). The recipient peripheral blood (PB) harvested from the retro-orbital plexus was evaluated for human hematopoietic engraftment every 3 to 4 wk starting at 6 wk posttransplantation. At 4–6 mo posttransplantation, recipient mice were euthanized for analysis.

Flow cytometry

Erythrocytes in the PB were lysed with Phos Lyse (BD Biosciences). Single-cell suspensions were prepared by BM and spleen using standard procedures. To isolate MNCs from the lung, lung tissues were carefully removed by gentle dissection. MNCs were then enriched for human CD34+ cells by using anti-human CD34 microbeads (Miltenyi Biotec) and sorted for 7AAD− cells out of total 7AAD− cells with purity above 98% by analysis of forward scatter-height/forward scatter-width and side scatter-height/side scatter-width. The purity of HSCs was higher than 98% after sorting. Newborn (within 2 d of birth) recipients received 150 cGy total body irradiation using a 137Cs-source irradiator, followed by i.v. injection of 1 × 10^5–3 × 10^5 sorted HSCs via the facial vein (14). The recipient peripheral blood (PB) harvested from the retro-orbital plexus was evaluated for human hematopoietic engraftment every 3 to 4 wk starting at 6 wk posttransplantation. At 4–6 mo posttransplantation, recipient mice were euthanized for analysis.

Flow cytometry

Erythrocytes in the PB were lysed with Phos Lyse (BD Biosciences). Single-cell suspensions were prepared by BM and spleen using standard procedures. To isolate MNCs from the lung, lung tissues were carefully excised, teased apart, and dissociated using collagenase (Wako) (15). The following mAbs were used for identifying engraftment of human hematopoietic cells in NSG recipients: anti-human CD34 V450 (clone UCHT1) and PE–Cy5 (H17A3), anti-human CD4 PE–Cy5 (RPA-T4), anti-human CD8 PE–Cy5 (RPA-T8), anti-hCD11b/Mac-1 Pacific blue (ICRF44), anti-hCD11c allophycocyanin (B-ly6), anti-hCD14 Alexa700 (M5E2), allophycocyanin–H7 and V450 (M6p9), anti-hCD15 allophycocyanin (H98) and V450 (MMA), anti-hCD19 PerCP–Cy5.5, PE–Cy5 and PE–Cy7 (SJ25C1), anti-hCD33 PE and PE–Cy7 (p67.6), anti-hCD34 PE–Cy7 (8G12), anti-hCD38 FITC and allophycocyanin (HB7), anti-hCD45, V450 and V500 (H30), anti-hCD45 AmCyan and allophycocyanin–Cy7 (2D1), anti-hCD56 FITC (NCAM16.2) and PE–Cy5 (B159), anti-hCD114/G-CSFR PE (LM741), anti-hCD116/G-CSFR FITC (hGMCSFR-M1), anti-hCD117c/Ki-PerCP–Cy5.5 (104D2), anti-hCD119/IFN–γR PE (GIR-208), anti-hCD123/IL-3R PE and PerCP–Cy5.5 (7G3), anti-hCD28/TLR2 Alexa647 (11G7), anti–H–L–A–DR allophycocyanin–H7 (L243), anti-mouse CD45 PerCP–Cy5.5 and allophycocyanin–Cy7 (30-F11), all from BD Biosciences; anti-human CD1c/BDCDA–1 FITC (ADS–8E7), anti-hCD141/ BDCDA–3 FITC, PE and allophycocyanin (AD5–14H12), anti-hCD303/ BDCA–2 PE (AC144) from Miltenyi; anti-human CD115/M-CSFR PE (9-4D2–1E4), anti-hCD203/e–NPF3 PE (NP4D6), anti-hCD284/TLR4 PE (HTA125), anti-hFcεRI FITC (AER–37), anti-mouse CD45 Alexa700 (30- F11) from BioLegend. The labeled cells were analyzed using FACScan or FACScan (BD Biosciences).
HSC-engrafted NSG recipients at 4–6 mo posttransplantation were injected i.v. with LPS at 15 μg/mouse. After LPS injection, plasma was collected from 0 to 4 h. Human cytokines IL-1β, IL-6, IL-8, IL-10, IL-12p70, and TNF were measured by cytometric bead array (BD Biosciences).

**IFN-γ–induced Salmonella killing activity by humanized mouse-derived monocytes/macrophages**

*Salmonella typhimurium* PhoPc strain transformed with the pKKGFP plasmid was kindly provided by J.P. Kraehenbuhl (17). *S. typhimurium* was grown shaking at 180 rpm overnight in Luria–Bertani broth supplemented with 100 μg/ml ampicillin at 37°C. BM monocytes/macrophages were purified by FACSAria (BD) based on the phenotypic characterization of lineage (CD3, CD7, CD16, CD19, CD56, CD123, CD235a)-negative, mouse CD45 and Ter119-negative, human CD45+CD11b+. Aliquots of (10⁴) human monocytes/macrophages derived from humanized mouse BM were cultured on collagen type I-coated 96-well plates (BD) for 24 h in either the presence or the absence of 1000 U/ml recombinant human IFN-γ (BD). Then, cells were infected with *S. typhimurium* at multiplicity of infection (MOI) of 20 at 37°C for 2 h, and the infection was confirmed by fluorescence microscopy (Biorevo BZ-9000; Keyence). For intracellular CFU determination, cells were washed twice with PBS and lysed in 0.2% Triton X-100 in PBS for 2 min, and lysates were diluted and plated onto Luria–Bertani broth agar plates containing 100 μg/ml ampicillin for colony enumeration.

**Statistical analysis**

The numerical data are presented as means ± SEM unless otherwise noted. Where noted, two-tailed t tests were performed, and the differences with the p value <0.05 were deemed statistically significant (GraphPad Prism; GraphPad).

**Results**

**Human myeloid lineage cells develop in NSG mice transplanted with human CB HSCs**

Recent advances in our knowledge of innate immunity reemphasize the importance of myeloid cells for sensing, capturing, and processing Ags for the initiation of innate and acquired phases of immune responses. The development of human myeloid cells in HSC-engrafted NSG mice has not previously been studied in detail. To evaluate in vivo differentiation and function of human myeloid lineage cell populations, we transplanted 1 × 10⁵–3 × 10⁶ purified human lineage CD34⁺CD38⁻ CB HSCs intravenously into sublethally irradiated newborn NSG mice. At 4–6 mo post-transplantation, we confirmed high levels of human hematopoietic chimerism and multilineage differentiation of human immune subsets as evidenced by flow cytometry (Fig. 1A, 1B). In addition to the reconstitution of human adaptive immunity (CD3⁺ T cells and CD19⁺ B cells), we identified the development of human innate immune cell subsets such as CD56⁺ NK cells and CD33⁺ myeloid cells in the recipient mice. The frequency of human myeloid lineage cells was higher in the recipient BM (30.7 ± 3.9%, n = 11) compared with that in the spleen (6.2 ± 1.2%, n = 11, p < 0.0001 by paired two-tailed t test) and PB (7.1 ± 1.3%, n = 11, p < 0.0003) (Fig. 1B).

We then identified the subsets of human myeloid cells present in the humanized NSG recipient mice through flow cytometry. Human myeloid subsets are classified into HLA class II-negative granulocytes and class II-positive APCs. In the BM and spleen...
of NSG recipients at 4–6 mo posttransplantation, we observed differentiation of both human granulocytes and APCs. Among the granulocyte lineage, human CD15^+CD33^lowHLA-DR^+ neutrophils, CD117^+CD123^+CD203c^+ basophils, and CD117^+CD203c^+HLA-DR^+ mast cells were observed in the recipient BM and spleen. Analyses of APC populations found that CD14^+CD33^+HLA-DR^+ monocytes, cDCs, mast cells, basophils, and pDCs in the BM and spleen of NSG recipients are summarized (n = 10). (C) In the humanized mouse BM and spleen, two distinct subsets of DCs, BDCA-1^+ DCs and BDCA-3^+ DCs, were identified in HLA-DR^+CD33^+CD11c^+ conventional DCs. Frequencies of the two DC subsets within BM and spleen hCD45^+CD33^+ cells are shown (BM, n = 9, *p = 0.007, significant differences between cDCs; spleen, n = 6, **p = 0.046). (D) Human myeloid cells isolated by cell sorting of recipient BM demonstrate characteristic morphological features on May–Grünwald–Giemsa stain. Baso, Basophils; Mast, mast cells; Mo, monocytes; Neu, neutrophils. Scale bars, 10 μm.

FIGURE 2. Development of human myeloid lineages in NSG recipients. (A) Representative flow cytometry contour plots demonstrating differentiation of human HLA-DR^+ granulocytes and HLA-DR^+ APCs in the BM and spleen of an NSG recipient. (B) The frequencies of human neutrophils, monocytes, cDCs, mast cells, basophils, and pDCs in the BM and spleen of NSG recipients are summarized (n = 10). (C) In the humanized mouse BM and spleen, two distinct subsets of DCs, BDCA-1^+ DCs and BDCA-3^+ DCs, were identified in HLA-DR^+CD33^+CD11c^+ conventional DCs. Frequencies of the two DC subsets within BM and spleen hCD45^+CD33^+ cells are shown (BM, n = 9, *p = 0.007, significant differences between cDCs; spleen, n = 6, **p = 0.046). (D) Human myeloid cells isolated by cell sorting of recipient BM demonstrate characteristic morphological features on May–Grünwald–Giemsa stain. Baso, Basophils; Mast, mast cells; Mo, monocytes; Neu, neutrophils. Scale bars, 10 μm.
BDCA-1⁺ BDCA-3⁺ monocytes, CD14⁺ CD33⁺ HLA-DR⁺ BDCA-1⁺ or BDCA-3⁺ cDCs, and CD123⁺ BDCA-2⁺ HLA-DR⁺ pDCs developed in the recipients BM and spleen (Fig. 2A, 2B). The frequency of CD15⁺ CD33⁺ HLA-DR⁻ neutrophils within human CD45⁺ CD33⁺ myeloid cells were present at the highest level in the BM (35.0 ± 5.4%, n = 10), whereas CD117⁺ CD203c⁺ FcɛRIlow HLA-DR⁻ mast cells developed at a higher efficiency in the recipient spleen (43.3 ± 4.0% within CD45⁺ CD33⁺, n = 10) (Fig. 2B). Among human APCs developing in the NSG recipients, monocytes accounted for the highest frequency in total myeloid cells both in the BM (32.6 ± 3.1%, n = 10) and spleen (25.2 ± 4.0%, n = 10). cDCs are divided into two subsets according to the expression of BDCA-1⁺ and BDCA-3⁺. Within human CD45⁺ CD33⁺ myeloid cells, the frequencies of BDCA-1⁺ DCs accounted for 6.4 ± 1.2% in BM (n = 9) and 6.7 ± 1.7% in spleen (n = 6) and were significantly higher than those of BDCA-3⁺ DCs (2.4 ± 0.3% and 2.8 ± 0.4%, respectively) (Fig. 2C). We then performed flow cytometric analysis using the same mAbs to determine the frequencies of each myeloid subset in primary BM MNCs. Although we could not directly compare human neutrophil development, the proportion of human monocytes, BDCA1⁺ cDCs, BDCA3⁺ cDCs, and pDCs was similar between primary human BM and humanized mouse BM (Supplemental Fig. 1). In addition to the expression analysis of cell surface molecules, we performed May–Gru¨ nwald–Giemsa staining to identify the morphology of the myeloid lineage cells developing in the NSG recipients. Human myeloid cells purified from NSG recipient BM exhibited characteristic morphological features (Fig. 2D).

**Human myeloid lineage cells developing in NSG recipients demonstrate intact functional responses to human cytokines in vitro and in vivo**

We confirmed the development of various human myeloid subsets in the BM and spleen of NSG recipients and next examined the expression of human cytokine receptors including IFN-γR, G-CSFR, GM-CSFR, and M-CSFR compared with that in human CB (Fig. 3A–C). By using human CD45⁺ CD33⁺ CB myeloid cells as control, we confirmed that human CD45⁺ CD33⁺ cells in the recipient BM expressed comparable levels of IFN-γR, G-CSFR, GM-CSFR and M-CSFR (n = 5, no significant difference between humanized mouse BM and CB, p = 0.6444, p = 0.0985, p = 0.3879, and p = 0.5816, respectively) (Fig. 3D).

To demonstrate functional responses to human cytokine stimulation at a cellular level, we examined the phosphorylation of STAT molecules using flow cytometry. Recent studies have revealed that hematopoietic cytokine receptor signaling is largely mediated by JAK kinases and STAT molecules known as the downstream transcription factors (18). BM cells from NSG recipients (n = 3) were

**FIGURE 3.** Expression of cytokine receptors on human myeloid cells in NSG recipients. (A–C) Representative flow cytometry contour plots demonstrating the expression of IFN-γR, G-CSFR, GM-CSFR, and M-CSFR by CB hCD45⁺ CD33⁺ myeloid cells (top) and by hCD45⁺ CD33⁺ myeloid cells derived from humanized NSG BM (bottom). Contour plots for isotype control Ig are also shown. (D) Expression of each cytokine receptor within hCD45⁺ CD33⁺ cells is summarized (CB, n = 5; humanized NSG BM, n = 5).
stimulated with rhIFN-γ, rhG-CSF, rhGM-CSF, or rhM-CSF in vitro for 15 min at 37˚C. In neutrophils and monocytes, rhGM-CSF specifically induced STAT5 phosphorylation, but not irrelevant STATs (e.g., STAT4 and STAT6) (Fig. 4A, 4C). Additionally, rhIFN-γ and rhG-CSF induced optimal STAT phosphorylation (Fig. 4B, 4D, 4E). Indeed, rhIFN-γ stimulation resulted in intracellular STAT1, STAT3, and STAT5 phosphorylation, and rh-G-CSF stimulation induced STAT3 and STAT5 phosphorylation, respectively (Fig. 4E). These results indicate that intact molecular events occur in human neutrophils and monocytes in response to recombinant human cytokines in vivo.

We next investigated in vivo cytokine response by human myeloid cells in the NSG humanized mice. Stimulation with rhG-CSF in vivo is known to induce proliferation of myeloid precursors and mobilization of myeloid subsets from BM (19). After in vivo treatment of humanized mice by rhG-CSF for 5 d, the frequencies of hCD45+CD15+CD33low fraction (human neutrophils) and hCD45+CD15−/lowCD33+ fraction (human monocytes and DCs) increased in the PB (three out of three recipients) (Fig. 4F). These findings indicate that human myeloid cells developing in the humanized NSG recipients demonstrate functional cytokine response both in vivo and in vitro.

**Human inflammatory response via TLR signaling**

Along with the role of cytokine receptor signaling in development and function of myeloid cells, signaling via TLRs serves fundamental roles in evoking systemic inflammatory response by myeloid cells (20). We therefore analyzed the expression of TLRs in human myeloid cells developed in the engrafted NSG recipients by flow cytometry. We examined the surface expression of TLR2 and TLR4 in the human myeloid cells developed in the humanized mouse BM. TLR2 is specifically expressed in human monocytes and BDCA1+ DCs, and TLR4 is expressed in the four distinct myeloid subsets, neutrophils, monocytes, BDCA1+ cDCs, and BDCA3+ cDCs (Fig. 5A, 5B). The expression of TLR4 was also confirmed in humanized mouse BM-derived monocytes and other myeloid subsets, which has led us to investigate the in vivo response of human innate immunity against LPS, a potent TLR4 agonist.
ligand and endotoxin. To this end, we have administered 15 μg LPS to NSG humanized mice followed by detection of human inflammatory cytokines by flow cytometry. Bead-attached Abs for human cytokines did not detect 5000 pg/ml of mouse cytokines demonstrating that these Abs and analyses are human-specific (Supplemental Fig. 2). Of the cytokines examined, we have seen the significant elevation of plasma levels of human IL-6, human IL-8, and human TNF (Fig. 5C). Time-dependent kinetics showed that the prompt response of human innate cells to the LPS was achieved between 30 min and 1 h after injection. Consequently, humanized mice could be used to examine human innate immune response against infectious organisms and to predict inflammatory response provoked by the TLR ligands.

Human myeloid cells present in NSG recipient lung exhibit functional phagocytosis

In the human immune system, myeloid cells serve an important role in immune surveillance not only in the systemic immune compartments but also in the mucosal tissues, especially the respiratory compartment of lung protected by both mucosal and systemic immune systems (21, 22). To examine whether functional reconstruction of human myeloid cells occurs in the lung, we evaluated the differentiation and function of human myeloid lineage cells isolated from the lungs of NSG recipients. Among human CD45+ cells present in the recipient lungs, myeloid lineage cells constituted 20.3 ± 3.8% (n = 8; a representative set of flow cytometry plots shown in Fig. 6A). The majority of human myeloid lineage cells residing in the recipient lungs were CD33+/H-2DR+ monocytes/macrophages (60.9 ± 5.1% within huCD45+CD33+, n = 8) (Fig. 6B).

The respiratory tract represents a major port of entry for inhaled pathogenic organisms, and resident alveolar monocytes/macrophages play a major role in surveillance and immune defense. To confirm the phagocytic function of human monocytes/macrophages present in the lungs of NSG recipients, we performed in vitro phagocytosis assay using yellow-green fluorescent beads by flow cytometry and confocal microscopy imaging. After in vitro incubation of NSG recipient-derived human CD45+ cells with 1 and 2 μm fluorescent beads at 37˚C, uptake of beads was noted in 9.0 and 7.9% of hCD45+CD33+ human myeloid cells, respectively (Fig. 6C). It should be noted that uptake of fluorescent beads was observed only in hCD45+CD33+ myeloid cells, but not in hCD45−CD33− lymphoid cells (Fig. 6C). This demonstrates that the fluorescent bead uptake specifically represents phagocytic function by human lung myeloid cells, not nonspecific uptake of the beads or binding or coating of the cells by the beads. The efficiency of uptake was 24.4 ± 3.0% in the lung-derived hCD45+ cells (n = 6, p = 0.001 compared with 4˚C incubation by two-tailed t test), equivalent to that in BM-derived hCD45+ cells (16.6 ± 2.7%, n = 4, p = 0.01 compared with 4˚C incubation by two-tailed t test) (Fig. 6C, 6D).

Next, phagocytosis of fluorescent beads by human myeloid cells was confirmed by direct visualization by confocal microscopy. Three-dimensional confocal imaging demonstrated intracellular localization of the fluorescent bead signal in sorted fluorescent bead (YG signal)+hCD45+CD33+ human myeloid cells, confirming the
internalization of microparticles by human monocytes/macrophages (Fig. 6E). Taken together, these findings demonstrate the presence of human innate immunity with intact phagocytic function in the NSG recipient lung.

Humanized mouse BM-derived monocytes/macrophages exhibit IFN-γ-induced phagocytosis and killing against Salmonella typhimurium

Myeloid subsets serve essential roles in host defense against various infectious microorganisms as a part of innate immunity. Of the various myeloid subsets discussed in the current study, monocytes and macrophages display excellent phagocytic potential by phagolysosome formation, by the effects of oxidative and nitrosative stress, and by antimicrobial cationic peptides and enzymes (23). To evaluate future application of the humanized mouse system in infectious disease research, we examined the phagocytic function of human monocytes/macrophages derived from humanized NSG BM against S. typhimurium. We purified mCD45^−TER119^−hCD45^+Lin^−CD11b^+ cells as monocytes/macrophages from the recipient BM (Fig. 7A) and cultured 10,000 purified human monocytes/macrophages with S. typhimurium at an MOI of 20 with or without prestimulation of human recombinant IFN-γ at 1000 U/ml. In the five in vitro experiments, stimulation of human monocytes/macrophages with rhIFN-γ resulted in the significantly potentiated phagocytosis and killing of Salmonella by the humanized mouse-derived monocytes/macrophages as evidenced by the decreased numbers of colony formation by S. typhimurium (at 3 h postinfection p = 0.023, at 12 h postinfection p = 0.091 [n.s.] compared with control versus IFN-γ stimulation by two-tailed t test) (Fig. 7B). Taken together, human monocytes that develop in the humanized NSG mice possess phagocytic activity against microbeads and bacteria and kill phagocytized bacteria presumably via signaling through cytokine receptors and TLRs.

Discussion

In vivo reconstitution of mature and functional human myeloid cells not only facilitates in vivo examination of human innate immunity but also offers a promising platform for translational research in the areas of infectious immunity and drug development. In the current study, we have aimed to clarify how functional human myeloid cells develop in NSG humanized mice.

In the NSG recipients, we found distinct levels of reconstitution of myeloid subsets in the BM and spleen. The differential myeloid reconstitution in the humanized hematopoietic organs is comparable to that seen in the human tissues, reflecting the distinct physiological roles of each hematopoietic organ in mammals. BM acts an essential reservoir of short-lived neutrophils and monocytes that readily migrate into sites of infection and inflammation. In addition, BM neutrophils function as paracrine regulators for mobi-
The development of human myeloid lineages is regulated by various cytokine signals (18, 27). In the current study, we directly compared the frequencies of human myeloid subsets using humanized mouse BM and primary human BM MNCs. As to the development of human APCs, humanized mouse BM recapitulates physiological development of human monocytes and two different subtypes of cDCs. However, we could not directly compare the frequencies of human neutrophils between humanized mouse BM and primary human BM, as we have used frozen BM MNCs. According to the previous reports, the frequency of human neutrophils in the humanized mouse BM is lower than that in the primary human BM (28, 29).

Human myelopoiesis within the mouse microenvironment may occur through multiple cooperative mechanisms. First, mouse cytokines such as stem cell factor, FLT3 ligand, G-CSF, and thrombopoietin may directly stimulate human myelopoiesis by cross-reacting with their respective receptors on human hematopoietic stem and/or myeloid progenitor cells. These human myeloid cells in turn produce cytokines such as GM-CSF and IL-3, resulting in the differentiation, maturation, and maintenance of human granulocytes, monocytes, and DCs. At the same time, the cytokine milieu within the NSG recipient repopulated with human hematopoietic cells may not be completely sufficient, to support human hematopoiesis as evidenced by the relative paucity of human neutrophils in the recipient BM, spleen, and circulation that might suggest the requirement of human cytokine or adhesion molecules in the hematopoietic tissues of the recipients. Recent studies suggested that the induced expression of human cytokines in the mouse environment may lead to enhanced differentiation and maturation of human myeloid subsets including neutrophils (30–33). In the current study, however, human monocytes develop in NSG recipients despite the fact that M-CSF is exclusively produced in non-hematopoietic cells and that murine M-CSF does not cross-react with human M-CSFR. This may be attributable to the redundancy among cytokines such as M-CSF, GM-CSF, and IL-3 as demonstrated in previous studies using M-CSF–deficient mice (34).

As a measure of human myeloid cell function, we investigated cytokine responses in human neutrophils and monocytes developing in the NSG recipients. Consistent with the expression of cytokine receptors identified on the human myeloid cells, neutrophils and monocytes showed intact responses to human cytokines both in vivo and in vitro. Phosphorylation of STAT molecules represents a molecular event downstream of cytokine receptor activation. STAT1 is a key mediator of IFN-γ activation of cells and an indispensable component of IFN-γ–dependent innate defense mechanisms against infections (35). The STAT3 signaling pathway is essential for G-CSF–mediated granulopoiesis (36). Specific phosphorylation of STAT5 may be an essential molecular event enabling generation of granulocytes from myeloid progenitors and proliferation and survival of mature neutrophils (37). STAT4 and STAT6 are essential for mediating IL-12 and IL-4 signaling in Th cells (38, 39). Human myeloid cells developing in humanized NSG recipients responded to human cytokines in a specific manner, as determined by the selective activation of JAK–STAT signaling pathways to corresponding cytokines.

Similar to the analysis of the expression of cytokine receptors and signaling, we showed that human myeloid subsets developing in the NSG humanized mice expressed various TLRs at the protein level. In the analysis of TLR expression in humanized mouse BM–derived cells, specific expression of TLR2 was observed in human monocytes and BDCA1+cDCs rather than neutrophils or BDCA3+cDCs. Consistent with the expression of TLR4 in human myeloid subsets developing in humanized NSG recipients responded to human cytokines in a specific manner, as determined by the selective activation of JAK–STAT signaling pathways to corresponding cytokines.
immunological homeostasis in the respiratory tracts (21, 22). In the recipient lung, unlike the BM or spleen, CD33+CD14+HLA-DR+ macrophages were the predominant myeloid population. Frequencies of human B cells, T cells, and myeloid cells in the recipient lung were distinct from those in the recipient PB, excluding the possibility that the human myeloid cells isolated from the recipient lung are contaminating PB myeloid cells. Importantly, macrophages, the predominant human myeloid subset in the recipient lung, demonstrated intact phagocytic function. Macrophages in the NSG recipients will be compared with the recently reported hGM-CSF and hIL-3 knock-in Rag2KO/IL2rgKO humanized mice showing abundant human macrophages in bronchoalveolar lavage (32). Establishment of an in vivo model of human pulmonary mucosal immunity may enable investigation of in vivo immune surveillance in the respiratory tract and in allergic pulmonary disorders and may allow evaluation of vaccines at preclinical stages (43, 44).

In this study, the reconstitution of both systemic and mucosal human innate immunity was observed in the NSG humanized mice. We performed phenotypic characterization and functional evaluation of human myeloid cells developing in the recipients, including granulocytes and APCs. Humanized mice reconstituted with both lymphoid and myeloid human lineages would facilitate in vivo investigation of interactions between the lymphoid and the myeloid compartments, allowing the dissection of the coordinated human immune response at the level of the whole organism.

Disclosures

The authors have no financial conflicts of interest.

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