Essential Role of Signal Transducer and Activator of Transcription (Stat)5a but Not Stat5b for Flt3-dependent Signaling

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Abstract

The receptor tyrosine kinase Flt3 plays an important role in proliferation and survival of hematopoietic stem and progenitor cells. Although some post-receptor signaling events of Flt3 have been characterized, the involvement of the Janus kinase/signal transducer and activator of transcription (Jak/Stat) pathway in Flt3 signaling has not been thoroughly evaluated. To this aim, we examined whether Flt3 activates the Jak/Stat pathway in Baf3/Flt3 cells, a line stably expressing human Flt3 receptor. Stat5a, but not Stats 1–4, 5b, or 6, was potently activated by Flt3 ligand (FL) stimulation. Interestingly, FL did not activate any Jaks. Activation of Stat5a required the kinase activity of Flt3. A selective role for Stat5a in the proliferative response of primary hematopoietic progenitor cells to FL was documented, as FL did not act on progenitors from marrows of Stat5a^{-/-} mice, but did stimulate/costimulate proliferation of these cells from Stat5a^{+/+}, Stat5b^{-/-}, and Stat5b^{+/+} mice. Thus, Stat5a is essential for at least certain effects of FL. Moreover, our data confirm that Stat5a and Stat5b are not redundant, but rather are at least partially distinctive in their function.

Key words: Flt3 • signal transducer and activator of transfection 5 • signal transduction • hematopoiesis • gene knockout

Introduction

Flt3 ligand (FL)¹ is a potent cytokine that acts synergistically with a wide range of CSFs and ILs to stimulate proliferation and differentiation of hematopoietic stem and progenitor cells (1–4). It also enhances survival of progenitor cells (5, 6). Its receptor, Flt3, belongs to the type III receptor tyrosine kinases (RTKs) that also include receptors for CSF-1, Steel (or stem cell) factor (SLF), and platelet-derived growth factor (PDGF; reference 7). This family has five immunoglobulin-like domains in their extracellular region and an intracellular tyrosine kinase made up of an ATP-binding loop and a catalytic domain separated by a kinase

insert domain. In vivo and in vitro studies have shown that FL and Flt3 play important roles in proliferation and differentiation of both multipotent stem and lymphoid progenitor cells (1–4, 8–10). Mice lacking Flt3 receptor have normal mature hematopoietic populations; however, they exhibit reduced numbers of primitive B lymphoid progenitors and multipotent stem cells (11).

Before the cloning of Flt3 ligand, the capacity of the Flt3 receptor to transmit a signal was studied by constructing chimeric receptors between the cytoplasmic and transmembrane domains of Flt3 and the extracellular domains of either the c-kit (12) or CSF-1 receptor (13–15). Using this strategy, some early signaling events and substrate specificities of Flt3 receptor were characterized (13, 15, 16). The chimeric receptor transduced mitogenic signals through association with and/or phosphorylation of a number of cytoplasmic proteins, including RAS GTPase–activating protein, phospholipase C- γ , Vav, Grb2, Shc, and the p85 subunit of phosphatidylinositol 3 kinase. p85 has been

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¹Abbreviations used in this paper: FL, Flt3 ligand; Jak, Janus kinase; PDGF, platelet-derived growth factor; RTK, receptor tyrosine kinase; SLF, Steel factor; Stat, signal transducer and activator of kinase.

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shown to bind tyrosine 958 (YQNM) in the COOH-terminal tail of murine Flt3 (15, 16). Recently, we found that p85 does not bind directly to human Flt3, which lacks a potential p85-binding site in the COOH terminus (17). In addition, FL stimulation of the human Flt3 receptor resulted in phosphorylation of Src homology 2 (SH2)-containing inositol-5-phosphatase (SHIP) and a 100-kD protein in monocytic THP-1 cells (17); phosphorylation of Shc and Cbl in myeloid cells (18); and SH2-containing tyrosine phosphatase, SHIP, and Cbl-b in pro-B cells (17, 19, 20). However, whether Janus kinase/signal transducer and activator of transcription (Jak/Stat) proteins are involved in Flt3 signaling has not been clearly studied.

In this study, we examined whether Flt3 activates the Jak/Stat pathway in a murine IL-3–dependent hematopoietic cell line Baf3, which stably expresses full-length human Flt3 receptor. We found that, unlike most RTKs, Flt3 potently activates only Stat5a without activation of any Jaks. This activation of Stat5a by Flt3 was also observed in COS-7 cells cotransfected with Flt3 receptor and Stat5 expression vectors. Moreover, FL did not stimulate/costimulate proliferation of primary hematopoietic progenitor cells from marrows of mice functionally deleted in Stat5a (-/-) but it did act normally on Stat5a littermate control (+/+), Stat5b (-/-), and Stat5b (+/+) progenitor cells. These results demonstrate that Stat5a, but not Stat5b, plays a critical role in mediating FL effects.

Materials and Methods

Cytokines and Antibodies. Recombinant human FL and murine SLF were provided by Immunex. Recombinant preparations of murine IL-3, GM-CSF, M-CSF, and G-CSF were purchased from R&D Systems. Rabbit polyclonal anti-Flt3, anti-Stat5a, anti-Stat5b, anti-Stat6, and anti-Tyk2 antibodies were purchased from Santa Cruz Biotechnology, Inc. Rabbit polyclonal anti-Stat5a, anti-Stat5b, anti-Jak1, anti-Jak2, and anti-Jak3 antibodies and antiphosphotyrosine mAb (4G10) were obtained from Upstate Biotechnology. Anti-Stat1, anti-Stat2, and anti-Stat4 antibodies were from Transduction Laboratories. AG490 and AG1296 were from Calbiochem-Novabiochem.

Cell Culture, DNA Constructs, and Transfection. The murine IL-3–dependent hematopoietic cell line Baf3/Flt3, a subline of Baf3 transduced with the human Flt3 receptor gene, was cultured as previously described (17). Before cytokine stimulation, Baf3/Flt3 cells were washed with PBS and starved in serum-free RPMI 1640 overnight. COS-7 and HEK293 cells were cultured in DMEM containing 10% fetal bovine serum and 100 U/ml penicillin and streptomycin.

The cDNA encoding human Flt3 (provided by Dr. Steward Lyman, Immunex, Seattle, WA) was subcloned into the expression vector pCDNA3 (Invitrogen). Mutants of Flt3 that lack either COOH terminus (Flt3- Δ CT; amino acids 950–993) or both COOH terminus and the second kinase domain (Flt3- Δ CT/TKII; amino acids 782–993) were constructed by PCR-based mutagenesis. The cDNAs encoding murine Stat5a and Stat5b were obtained from Dr. X.-Y. Fu (Yale University, New Haven, CT). Transfection of COS-7 and 293 cells was performed using SuperFect reagent (Qiagen) according to the manufacturer's protocol. For cytokine stimulation experiments, COS-7 or 293 cells

were serum-starved for 24 h and then stimulated with FL (100 ng/ml).

Immunoprecipitation and Immunoblotting. After starvation, cells were suspended in serum-free media $(1-2 \times 10^7 \text{ cells/ml})$ and stimulated at 37°C with human FL or murine IL-3 in the absence of any serum. Cell extracts were prepared and subjected to immunoprecipitation and immunoblotting with the antibodies indicated as previously described (20).

Electrophoretic Mobility Shift Assay. Nuclear extracts from untreated and FL-treated cells were prepared as described elsewhere (21). For the EMSA, 20,000 cpm of 32 P-labeled γ -activated sequence motif corresponding to the β -casein gene promoter (5'-AGATTTCTAGGAATTCAATCC-3') was incubated with 20 µg of nuclear extract proteins in 20 µl of binding buffer containing 10 mM Tris, pH 7.5, 50 mM NaCl, 1 mM dithiothreitol, 1 mM EDTA, 0.05% NP-40, 2 µg poly(dI·dC):poly(dI·dC), and 10% glycerol at room temperature for 30 min and then resolved on 4% polyacrylamide gels containing $0.5 \times$ TBE (1 \times TBE is 89 mmol/liter Tris borate and 1 mmol/liter EDTA, pH 8) and 2.5% glycerol. Gels were run at 4°C in $0.5 \times$ TBE at 20 V/cm, dried, and autoradiographed. Oligonucleotide competition was performed by preincubating nuclear extracts with the cold probe (50-fold excess) and poly(dI·dC):poly(dI·dC) for 30 min at 4°C before the addition of labeled probe. For supershifts, samples were preincubated for 1 h at 4°C with the indicated antibodies.

Mice. Mice functionally deleted (-/-) in Stat5a (22) and Stat5b (23) were from fifth generation backcrosses to the C57BL/6 background. These mice were generated from heterozygous matings. Some of the characteristics of immune cells from these mice have been described previously (24, 25). As controls, cells from age- and sex-matched littermate Stat5a^{+/+} and Stat5b^{+/+} mice were used.

Hematopoietic Progenitor Cell Assay. Unseparated bone marrow (5 × 10⁴ cells/ml) obtained from femurs of Stat5a^{-/-}, Stat5a^{+/+}, Stat5b^{-/-}, and Stat5b^{+/+} mice were plated in 0.3% agar (Difco) culture medium in the presence of 10% heat-inactivated fetal bovine serum (Hyclone) in the absence and presence of varying concentrations of human FL; murine SLF; murine GM-CSF; murine M-CSF; human G-CSF; the combination of SLF with GM-CSF, M-CSF, or G-CSF; or the combination of FL with SLF, GM-CSF, M-CSF, or G-CSF. Plates were incubated at 5% CO₂ and lowered (5%) O₂ for 7 d in a humidified atmosphere and scored for colonies deriving from granulocyte and/or macrophage progenitor cells (26). Three plates were scored per point per experiment.

Flow Cytometric Analysis. After lysis of red blood cells, mouse bone marrow cells were subjected to negative selection using lineage-specific antibodies (anti-Gr1, anti-CD11b, anti-CD3, and anti-B220 antibodies from PharMingen) and complement to remove granulocytes, macrophages, and mature T and B cells. Then the cells were stained with fluorochrome-conjugated antibodies (anti-Flt3-PE and anti-c-kit-FITC from PharMingen) and analyzed using FACScan[®] with CELLQuest software (Becton Dickinson).

Results

FL Preferentially Activates Stat5a over Stat5b in Baf3/Flt3 Cells. To test whether any Stat proteins are involved in Flt3 signaling, we used Baf3/Flt3 cells transduced with fulllength Flt3 cDNA. This subline stably expresses Flt3 receptor on the cell surface and proliferates in response to human FL (20). Among the Stat proteins we examined (Stat1, 2, 3, 4, 5, and 6), only Stat5 was tyrosine phosphorylated by FL stimulation, while IL-3 induced tyrosine phosphorylation of both Stat3 and Stat5 (Fig. 1 A). Tyrosine phosphorylation of Stat5 induced by FL was transient and reached a maximal level at 5 min (Fig. 1 B). It was weak compared with IL-3 stimulation. Fig. 1 C shows the dose–response of tyrosine phosphorylation of Flt3 and Stat5 induced by FL stimulation. We also checked Stat1, 2, 4, and 6. Although they are expressed in Baf3/Flt3 cells, none of them were tyrosine phosphorylated by FL stimulation (data not shown). These results demonstrate that unlike most RTKs, Flt3 selectively activates Stat5.

Two different Stat5 genes, encoding Stat5a and Stat5b and sharing >90% identity at the amino acid level, have been identified (27–30). Both Stat5a and Stat5b are activated by many cytokines and growth factors. The targeted knockout of the individual genes in mice has suggested that they play essential and at least partially overlapping roles in the physiological responses associated with several cytokines (22–25, 31). Despite their homology, there is evidence suggesting that Stat5a and Stat5b may be differentially activated (32). Since the anti-Stat5 antibody we used in Fig. 1 recognizes both Stat5a and Stat5b, we assessed whether FL activated both or only one of the Stat5 proteins. By using specific anti-Stat5a and -Stat5b antibodies, we found that FL induced tyrosine phosphorylation of Stat5a but not Stat5b (Fig. 2 A). IL-3 induced tyrosine phosphorylation of both Stat5a and Stat5b in Baf3/Flt3 cells (data not shown).

We next investigated if FL could stimulate Stat5a DNA binding activity by EMSAs using a γ -activated sequence motif from the β -casein gene promoter, which is known to bind to both Stat5a and Stat5b. As shown in Fig. 2 B, FL stimulation induced a DNA binding complex that was clearly supershifted by anti-Stat5a antibody, but anti-Stat5b antibody had a much smaller effect. This is consistent with the tyrosine phosphorylation pattern of Stat5 and suggests that gel shift assays tend to be more sensitive than phosphotyrosine blots. These results demonstrate that FL preferentially activates Stat5a in Baf3/Flt3 cells.

FL Does Not Activate Jaks. Since Stat5a is activated by FL stimulation, we tested whether Jaks are activated by FL. Jak1, Jak2, Jak3, and Tyk2 were immunoprecipitated from cells treated with either FL or IL-3, separated by SDS-PAGE, transferred to polyvinylidene difluoride membranes, and blotted with antiphosphotyrosine antibody. IL-3 has been reported to activate both Jak1 and Jak2 (33–35). As shown in Fig. 3, although IL-3 induced rapid tyrosine phosphorylation of Jak1 and Jak2, none of the Jaks were activated by FL. These results show that FL does not activate Jaks and suggests that activation of Stat5a is likely mediated by other tyrosine kinases, possibly the Flt3 receptor.





rosine antibody. (B) Stat5 is transiently tyrosine phosphorylated by FL stimulation. Growth factor-starved Baf3/Flt3 cells were stimulated with FL for various periods of time or with IL-3 for 5 min. Stat5 was immunoprecipitated from cell lysates and immunoblotted with antiphosphotyrosine antibody. (C) Dose-response of tyrosine phosphorylation of Flt3 and Stat5. Growth factor-starved Baf3/Flt3 cells were stimulated with different dose of FL for 5 min. Flt3 and Stat5 were immunoprecipitated from cell lysates and immunoblotted with antiphosphotyrosine antibody. The same membranes were stripped and reblotted with different antibodies as shown. Results shown are from one representative of two to three experiments. pY, antiphosphotyrosine antibody.

Stat5

IB: a-Stat5



(A) Growth factor-starved Baf3/Flt3 cells. (A) Growth factor-starved Baf3/ Flt3 cells were stimulated with FL for 5 min. Stat5a and Stat5b were immunoprecipitated from

cell lysates and immunoblotted with anti-phosphotyrosine antibody (pY). The same membrane was stripped and reblotted with anti-Stat5 antibody that recognizes both forms of Stat5. Results shown are from one representative of three experiments. (B) FL activates Stat5a DNA binding activity. Growth factor-starved Baf3/Flt3 cells were either untreated (control) or stimulated with FL for 10 min. Nuclear extracts from these cells were incubated with ³²P-labeled probe and subjected to EMSA. Oligonucleotide competition was performed by preincubating nuclear extracts with the cold probe. For supershifts, nuclear extracts were preincubated with the indicated anti-Stat5a or anti-Stat5b antibodies. The arrow indicates the DNA/Stat5 complex. Results shown are from one representative of two experiments.

Activation of Stat5a Requires the Kinase Activity of Flt3. Since Stat5a is tyrosine phosphorylated after FL stimulation and none of the Jaks is activated by FL stimulation, it is possible that Flt3 may directly phosphorylate Stat5a. To test this hypothesis, we first used type III receptor tyrosine kinase inhibitor AG1296, which has been shown to specifically inhibit SLF and PDGF signaling (36). As a control, we

IP: Jak1 IP: Jak2 IP: Jak3 IP: Tyk2 С FL IL-3 FL IL-3 C C FL C FL Jak1 - Jak2 Blot: a-pY Blot: α-pY Blot: a-pY Blot: a-pY - Jakl - Jak3 Tvk2 Jak2 Blot: a-Jakl Blot: a-Tyk2 Blot: a-Jak2 Blot: a-Jak3

used AG490, which inhibits the kinase activity of Jak2 and Jak3 (37, 38). As shown in Fig. 4 A, AG1296 at 50 μ M diminished tyrosine phosphorylation of Flt3 to the control level, whereas AG490 did not have an effect. After Baf3/ Flt3 cells were pretreated with these two inhibitors, Stat5 was immunoprecipitated from cells treated with FL and tyrosine phosphorylation of Stat5 was determined by Western blot analysis with antiphosphotyrosine antibody. As shown in Fig. 4 B, tyrosine phosphorylation of Stat5a was decreased close to control levels by AG1296, but not by AG490. This is consistent with the results that Jaks were not activated by FL stimulation and suggests that the tyrosine kinase activity of Flt3 is required for activating Stat5a.

Next, we transiently transfected Flt3 cDNA with either Stat5a or Stat5b cDNA into COS-7 and HEK293 cells and examined whether Flt3 could induce tyrosine phosphorylation and DNA binding activity of Stat5. After serum-starvation for 24 h, the transfected cells were stimulated with human FL (100 ng/ml) for 5 min. Stat5a and Stat5b were immunoprecipitated from cell lysates and tyrosine phosphorylation was analyzed by immunoblotting with antiphosphotyrosine antibody. As shown in Fig. 5 A, no phosphorylation of either Stat5a or Stat5b was detected in COS-7 cells that were mock transfected. Cotransfection of Flt3 and Stat5a cDNA resulted in strong phosphorylation of Stat5a. Addition of FL slightly increased phosphorylation of Stat5a. In contrast, the phosphorylation of Stat5b was very weak, and could only be detected after longer exposure when coexpressed with Flt3. Control immunoblots showed that equivalent levels of Flt3 were expressed. Similar results were obtained when using 293 cells (data not shown). We then evaluated the DNA binding activity of Stat5 by EMSAs. As shown in Fig. 5 B, no DNA binding activities were detected in COS-7 cells that were transfected with vector control plus Stat5a or Stat5b alone. Cotransfection of Flt3 cDNA with Stat5a resulted in high levels of a DNA complex that was supershifted by anti-Stat5a antibody. Although Stat5a and Stat5b were expressed at similar levels (data not shown), cotransfection of Flt3 with Stat5b yielded only a weaker DNA binding activ-

Figure 3. Jaks are not activated by FL stimulation in Baf3/Flt3 cells. Growth factor–starved Baf3/Flt3 cells were stimulated with FL (100 ng/ml) or IL-3 (10 ng/ml) for 5 min. Jak1, Jak2, Jak3, and Tyk2 were immunoprecipitated from cell lysates and immunoblotted with antiphosphotyrosine antibody (pY). The same membranes were stripped and reblotted with different antibodies as shown in the figure. Results shown are from one representative of three experiments.

722 Flt3 Ligand Preferentially Activates STAT5A in a JAK-independent Manner



Figure 4. Inhibition of tyrosine phosphorylation of Flt3 and Stat5a by AG1296, but not by AG490. Baf3/Flt3 cells were pretreated with either control diluent DMSO or AG1296 and AG490 for 1 h at 37°C and washed once with PBS. Then the cells were treated with FL for 5 min. Flt3 (A) and Stat5a (B) were immunoprecipitated from cell lysates and immunoblotted with antiphosphotyrosine antibody (pY). The same membranes were stripped and reblotted with different antibodies as shown in the figure. In B, AG1296 and AG490 were used at 50 μ M. Results shown are from one representative of two experiments.

ity. These results confirmed that Flt3 preferentially activated Stat5a.

Although expression of wild-type Flt3 in COS-7 cells led to phosphorylation of Stat5a, mutant Flt3 lacking the second kinase domain did not induce phosphorylation of Stat5a, and mutant Flt3 lacking the COOH terminus induced less phosphorylation of Stat5a (Fig. 5 C). Wild-type and mutants of Flt3 were expressed at similar levels as measured by flow cytometry (data not shown). These results demonstrate that phosphorylation of Stat5a requires tyrosine kinase activity of Flt3. Thus, Stat5a is either a direct substrate of Flt3, or Flt3 activates an endogenous tyrosine kinase that in turn phosphorylates Stat5a, or both. Myeloid Progenitors from $Stat5a^{-/-}$ Mice Do Not Respond to the Stimulating/Costimulating Effects of FL. Use of cells from mice functionally deleted in genes for certain intracellular molecules has allowed the elucidation of the roles of these gene-encoded proteins in blood cell regulation (39). To this end, we evaluated the responsiveness in vitro of myeloid progenitor cells from femurs of $Stat5a^{-/-}$ and $Stat5a^{+/+}$ mice to the proliferation effects of FL. By itself, FL is a weak stimulator of the proliferation of myeloid progenitor cells. However, in combination with GM-CSF, M-CSF, or G-CSF, it is a potent costimulating molecule that acts synergistically with these CSFs to enhance the number and size of colonies (4).



with anti-Stat5 antibody that recognizes both Stat5a and Stat5b. The expression of Flt3 was confirmed by immunoblot with anti-Flt3 antibody on total cell lysates. (B) Nuclear extracts from COS7 cells transfected with Stat5a, Stat5b, Flt3, or vector control in the combination shown were incubated with 32 P-labeled probe and subjected to EMSA. Oligonucleotide competition was performed by preincubating nuclear extracts with the cold probe. For supershifts, nuclear extracts were preincubated with the anti-Stat5a or -Stat5b antibodies. The arrow indicates the DNA–Stat5 complex. (C) COS-7 cells were transfected with Stat5a plus control vector, Flt3, Flt3– Δ CT, or Flt3– Δ CT/TKII, as shown. The tyrosine phosphorylation of Stat5a was examined by immunoprecipitation and immunoblotting with antiphosphotyrosine antibody. The same membrane was stripped and reblotted with anti-Stat5a antibody. pY, antiphosphotyrosine antibody; SS, supershift.

723 Zhang et al.

Bone marrow cells from Stat5a^{-/-} and littermate control Stat5a^{+/+} mice were plated in the absence and presence of varying concentrations of FL, SLF, GM-CSF, M-CSF, or G-CSF, or varying concentrations of SLF or FL with each other, or varying concentrations of each of the CSFs (Fig. 6). FL by itself had minimal activity on stimulation of colony formation of Stat5a^{+/+} progenitors and little or no stimulation of Stat5a^{-/-} progenitors. No significant differences (P > 0.05) were noted in response of Stat5a^{-/-} and Stat5a^{+/+} progenitors to stimulation by SLF, GM-CSF, M-CSF, or G-CSF, or to the synergistic effects of 50 ng/ml SLF plus GM-CSF, M-CSF, or G-CSF. However, the costimulating synergistic effects of 100 ng/ml FL with SLF, GM-CSF, M-CSF, or G-CSF seen with cells from Stat5a^{+/+} mice (P < 0.001) were not apparent with cells from Stat5a^{-/-} mice. The effects of FL in combination with the different concentrations of SLF or CSFs on Stat5a^{-/-} cells were equivalent to the single actions of these concentrations of SLF or CSFs. Similar differences were noted using 10



ng/ml FL or SLF with varying concentrations of CSFs (data not shown). Greater than 90% of colonies stimulated with M-CSF with or without FL or SLF were composed purely of macrophages. Greater than 90% of colonies stimulated with G-CSF with or without FL or SLF were composed of neutrophilic granulocytes. Colonies stimulated with GM-CSF with or without FL or SLF were >50% composed of granulocytes and macrophages with the remainder being pure granulocyte or pure macrophage colonies.

To evaluate the specificity of these effects, similar comparative experiments were done with myeloid progenitor cells from Stat5b^{-/-} and their littermate control Stat5b^{+/+} mice (Fig. 7). No significant differences (P > 0.05) were noted in the responses of Stat5b^{-/-} and Stat5b^{+/+} cells to the individual effects of FL, SLF, GM-CSF, M-CSF, G-CSF, or to the combined effects of SLF or FL with each other or with any of the CSFs. At the same time these experiments were done, experiments were also done with Stat5a^{-/-} and Stat5a^{+/+} cells with results comparable to those seen in Fig. 6.

These results demonstrate the critical role played by Stat5a, but not Stat5b, in myeloid progenitor cell respon-



Figure 6. FL stimulates/costimulates proliferation of myeloid progenitor cells from marrows of Stat5a^{+/+} but not Stat5a^{-/-} mice. Results are shown as the average percentage of control of cells incubated with 50 ng/ml SLF plus 10 ng/ml GM-CSF for a total of six experiments each. The control colony numbers upon which the percent of control are based were 52 ± 2 (mean ± 1 SEM), 70 ± 3, 89 ± 4, 60 ± 4, 152 ± 3, and 132 ± 2 for Stat5a^{-/-} cells and 62 ± 3, 87 ± 3, 101 ± 8, 71 ± 2, 70 ± 2, and 55 ± 3 for Stat5a^{+/+} cells.

Figure 7. FL stimulates/costimulates proliferation of myeloid progenitor cells from marrows of Stat5b^{+/+} and Stat5b^{-/-} mice to the same extent. Results are shown as the average percentage of control of cells incubated with 50 ng/ml SLF plus 10 ng/ml GM-CSF for a total of two experiments each. The control colony numbers upon which the percentage of control are based were 66 ± 2 and 93 ± 1 for Stat5b^{-/-} cells, and 93 ± 2 and 74 ± 3 for Stat5b^{+/+} cells.



Figure 8. Flt3 receptor expression on the surface of bone marrow cells of $5tat5a^{+/+}$, $5tat5a^{-/-}$, $5tat5b^{+/+}$, and $5tat5b^{-/-}$ mice as analyzed by flow cytometry. Mouse bone marrow cells were depleted of lineage-positive cells and stained with anti–Flt3–PE and anti–c-kit–FITC antibodies. Large granulocytes and debris were gated out on the basis of their light scatter pattern. The quadrants were set up based on isotype control antibodies. The numbers shown on the two-dimensional dot plots are percentages of single positive cells (Flt3⁺ or c-kit⁺) and double positive cells (Flt3⁺c-kit⁺).

siveness to the stimulating/costimulating effects of FL. To make sure that the differences noted between Stat5a^{-/-} cells compared with those of Stat5a^{+/+}, Stat5b^{-/-}, and Stat5b^{+/+} cells were not due to differential expression of Flt3 on the cells, protein levels of Flt3 were assessed by flow cytometry on the surface of bone marrow cells that were depleted of lineage-positive cells. (Fig. 8). Expression of Flt3 on Stat5a^{-/-} c-kit⁺Lin⁻ cells, a population highly enriched for myeloid progenitor cells (40), was at least as high as on the Stat5a^{+/+}, Stat5b^{+/+}, and Stat5b^{-/-} cells. Thus, the expression levels of Flt3 on the cells from these mice could not explain the nonresponsiveness of Stat5a^{-/-} progenitors to stimulation/costimulation by FL.

Discussion

In this study we have found that FL preferentially activates Stat5a over Stat5b, and that myeloid progenitors from the bone marrow of Stat5a^{+/+}, Stat5b^{+/+}, and Stat5b^{-/-}, but not from Stat5a^{-/-}, mice respond to the stimulating/ costimulating effects of FL. These results demonstrate that Stat5a plays a critical role in mediating FL effects. Moreover, this adds additional evidence to the literature showing the nonoverlapping effects of Stat5a and Stat5b.

Among the different Stat proteins examined, only Stat5a was potently tyrosine phosphorylated by FL stimulation. FL did not seem to activate any Janus kinase as measured by phosphotyrosine immunoblotting (Fig. 3). Longer stimulation with FL (30 min) did not result in any phosphorylation of Jak1 and Jak2 (data not shown). This is in contrast to most growth factors whose receptors are tyrosine kinases, such as PDGF, SLF, and M-CSF (or CSF-1). PDGF, SLF,

and M-CSF have been shown to activate multiple Jak and Stat proteins. M-CSF induces tyrosine phosphorylation of Jak1, Tyk2, and Stat1, 3, and 5; PDGF activates Jak1, Jak2, Jak3, and Stat1, 3, 5, and 6 (41–44); and SLF activates Jak2 and Stat1, 3, and 5 (45, 46). These growth factors activate both Stat5a and Stat5b. Our results suggest that FL differs from these RTKs in that it activates only Stat5a without an apparent activation of Jaks.

Recently, a group reported that a tandem-duplicated form of Flt3 constitutively activates Stat5 and mitogen-activated protein (MAP) kinase, whereas FL stimulation leads to activation of MAP kinase but not Stat5 activation in Baf3 and 32D cells expressing wild-type Flt3 (47). In that study (47), it was not specified whether Hayakawa et al. were dealing with Stat5a or Stat5b or both Stat5 proteins. Our data show that FL stimulation of wild-type Flt3 preferentially activates Stat5a, and our in vivo data strongly suggest that Stat5a, but not Stat5b, plays a critical and potentially physiological role in mediating the synergistic effects of FL. Although the same cell line, Baf3, was used, the discrepancy between the study by Hayakawa et al. and ours may be due to different levels of Flt3 expression and the assays used.

Stat5a and Stat5b are two highly related transcription factors encoded by two different genes. Stat5a and Stat5b are activated by a broad range of cytokines and growth factors. Although most cytokines and growth factors activate both Stat5a and Stat5b, there are some studies showing that Stat5a and Stat5b can be differentially activated and regulate different sets of genes. For example, GM-CSF preferentially activates Stat5b, but not Stat5a, in human neutrophils, whereas IFN- α and - γ predominantly activate Stat5a in promonocytic U937 cells (48, 49). Insulin preferentially activates Stat5b, but not Stat5a, via a Jak2-independent signaling pathway in kym-1 rhabdomyosarcoma cells (50). Not only can Stat5a and Stat5b be differentially activated, they also have distinctive functions. In support of this, Stat5a^{-/-} and Stat5b^{-/-} mice have different phenotypes (22-25, 51). Our findings that FL preferentially activates Stat5a in both Baf3/Flt3 and COS7 cells and that myeloid progenitors from the marrow of Stat5a^{+/+}, Stat5b^{+/+}, and Stat5b^{-/-}, but not from Stat5a^{-/-}, mice respond to the stimulating/costimulating effects of FL provide another example of differential activation of Stat5 protein. Moreover, our results strongly suggest that Stat5a, but not Stat5b, plays a critical role in myeloid progenitor cell responsiveness to the stimulating/costimulating effects of FL.

The Jak/Stat pathway has been well characterized in cytokine signaling. In the type I cytokine receptor family, ligand stimulation induces rapid activation of Jaks which then phosphorylate Stat proteins. In contrast, in tyrosine kinase receptors, the role of Jak proteins in Stat activation is not clearly defined. Some RTKs like PDGFR and epidermal growth factor receptor (EGFR), although they can activate Jaks, activate Stat proteins in a Jak-independent fashion (43, 52). In contrast, the activity of the intrinsic RTK is absolutely required for Stat activation by EGF or PDGF. A mutant PDGFR, which lacks kinase activity, is unable to stimulate Stat1 and Stat3 activation in response to PDGF stimulation (43). Stat1 can directly interact with EGF, PDGF, and SLF receptors and these receptors can phosphorylate Stat1 in vitro (46, 53, 54). EGFR can also directly phosphorylate Stat3 in vitro (55). These results suggest that Stat proteins may be direct substrates of receptor tyrosine kinases. Our findings that the kinase activity of Flt3 is required for Stat5a activation supports this hypothesis. However, our results do not rule out the possibility that Flt3 may activate an endogenous tyrosine kinase that in turn phosphorylates Stat5a. Regardless of the mechanism, our data confirm that Stat5a and Stat5b are not redundant, but rather are at least partially distinctive in their functions.

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