

The Rat *c-kit* Ligand, Stem Cell Factor, Induces the Development of Connective Tissue-type and Mucosal Mast Cells In Vivo. Analysis by Anatomical Distribution, Histochemistry, and Protease Phenotype

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Summary

Mast cell development is a complex process that results in the appearance of phenotypically distinct populations of mast cells in different anatomical sites. Mice homozygous for mutations at the *W* or *Sl* locus exhibit several phenotypic abnormalities, including a virtual absence of mast cells in all organs and tissues. Recent work indicates that *W* encodes the *c-kit* tyrosine kinase receptor, whereas *Sl* encodes a *c-kit* ligand that we have designated stem cell factor (SCF). Recombinant or purified natural forms of the *c-kit* ligand induce proliferation of certain mast cell populations in vitro, and injection of recombinant SCF permits mast cells to develop in mast cell-deficient WCB6F₁-*Sl*/*Sl*^d mice. However, the effects of SCF on mast cell proliferation, maturation, and phenotype in normal mice in vivo were not investigated. We now report that local administration of SCF in vivo promotes the development of connective tissue-type mast cells (CTMC) in the skin of mice and that systemic administration of SCF induces the development of both CTMC and mucosal mast cells (MMC) in rats. Rats treated with SCF also develop significantly increased tissue levels of specific rat mast cell proteases (RMCP) characteristic of either CTMC (RMCP I) or MMC (RMCP II). These findings demonstrate that SCF can induce the expansion of both CTMC and MMC populations in vivo and show that SCF can regulate at least one cellular lineage that expresses *c-kit*, the mast cell, through complex effects on proliferation and maturation.

Studies in mice and rats indicate that mast cells are derived from hematopoietic precursors that arise in the bone marrow and circulate in the blood, but complete their program of differentiation and maturation within interstitial tissues, epithelia, or serosal cavities (reviewed in references 1–6). In murine rodents, this process results in the generation of at least two distinct mast cell populations that vary in many aspects of phenotype, the connective tissue-type mast cells (CTMC)¹, which occur in such sites as the skin and peritoneal cavity, and the mucosal mast cells (MMC), which occur in the mucosa of the gastrointestinal tract (reviewed in references 1–4). Evidence derived from both in vitro and in vivo studies indicates that IL-3 represents one major growth

factor for MMC, whereas the regulation of CTMC proliferation requires additional and/or alternative factors (reviewed in references 1–6). For example, in vitro work indicates that one representative CTMC, peritoneal mast cells (PMC), do not proliferate in response to IL-3 alone but divide when IL-3 is provided with a second stimulus such as IL-4 (7) or PMA (8).

Several important insights into mast cell development have been derived from analyses of mice that virtually lack mast cells as a result of a double dose of mutant genes at either the *W* locus on chromosome 5 or the *Sl* locus on chromosome 10 (reviewed in references 1, 2, and 5). For example, *W* or *Sl* mutant mice lack both CTMC and MMC, indicating that the *W* or *Sl* gene products are important in the development of both of these mast cell populations in vivo (reviewed in references 1, 2, 5, and 6). Moreover, both in vitro and in vivo studies suggested that the mast cell deficiencies of these mutant mice reflected, in the case of *W* mutants, defects in the responsiveness of cells in the mast cell

¹ Abbreviations used in this paper: BrdU, 5-bromo-2'-deoxyuridine; CTMC, connective tissue-type mast cell; Hct, hematocrit; MMC, mucosal mast cell; PMC, peritoneal mast cell; RMCP, rat mast cell protease; rrSCF¹⁶⁴, recombinant rat stem cell factor¹⁶⁴.

lineage to a stromal cell-derived growth factor and, in the case of *Sl* mutants, inadequate production of this growth factor by stromal cells (reviewed in references 1, 2, and 5).

The products encoded at *W* and *Sl* recently have been defined. *W* encodes the *c-kit* tyrosine kinase growth factor receptor (9, 10), whereas *Sl* encodes a newly recognized multifunctional growth factor that represents a ligand for *c-kit* (11–18). Several lines of evidence in addition to the lack of mast cells in *W* or *Sl* mutant mice indicate that one of the important biological activities of the *c-kit* ligand is to regulate mast cell development. The recombinant *c-kit* ligand can promote the proliferation of certain populations of immature, IL-3-dependent mast cells in vitro (11, 14, 15, 17, 18), and the *c-kit* ligand purified from the supernatants of BALB/3T3 fibroblasts (19), or the recombinant ligand (20), can induce proliferation of IL-3-independent mouse PMC in vitro. In addition, we demonstrated that daily subcutaneous injection of a recombinant *c-kit* ligand, recombinant rat stem cell factor¹⁶⁴ (rrSCF¹⁶⁴), for 3 wk permitted mast cells to develop in the skin of genetically mast cell-deficient WCB6F₁-*Sl*/*Sl*^h mice (16). However, the mast cells that developed in WCB6F₁-*Sl*/*Sl*^h mice injected with rrSCF¹⁶⁴ were not characterized according to phenotype. Nor was it determined whether rrSCF¹⁶⁴ could influence mast cell development or phenotype in normal mice. In the present study, we therefore examined the effects of rrSCF¹⁶⁴ on mast cell populations in normal mice and rats in vivo, using approaches that permitted the anatomical and phenotypic characterization of the responding cells as CTMC or MMC. Some of these results have been reported in abstract form (20).

Materials and Methods

Studies in Mice. Groups of five to seven female 8–12-wk-old WCB6F₁-+/+, WCB6F₁-*Sl*/*Sl*^h, WBB6F₁-+/+, and WBB6F₁-*W*/*W*^v mice (The Jackson Laboratory, Bar Harbor, ME) received for 3 wk a daily subcutaneous injection of rrSCF¹⁶⁴ purified from *Escherichia coli* and modified by the covalent attachment of polyethylene glycol (16) or vehicle alone (0, 30, or 100 µg rrSCF¹⁶⁴/kg in 150–200 µl of sterile saline containing 0.1% BSA) (fraction V, fatty acid free; ICN Immunobiologicals, Lisle, IL). Injections were performed with the mice under light ether anesthesia and were delivered to approximately the same site on the dorsal back skin. Blood for determination of hematocrit (Hct) was obtained from the mice by retroorbital puncture under light ether anesthesia on the day before initiation of treatment and on the day of death. For labeling of tissue mast cells proliferating in vivo (21), 5-bromo-2'-deoxyuridine (BrdU) (Sigma Chemical Co., St. Louis, MO) was injected intraperitoneally (100 mg/kg in sterile saline) 1 h before death. After death by cervical dislocation, the cutaneous injection site was excised and fixed in Carnoy's fixative, and embedded in paraffin. 4-µm sections were cut, placed onto polylysine-coated slides, and processed as in reference 21 for staining of mast cells with 1.0% alcian blue, pH 1.0, and for immunohistochemical staining of BrdU-labeled nuclei, using an anti-BrdU mAb (Becton Dickinson & Co., Mountain View, CA). The slides were examined at 400× to quantify mast cells/mm² of dermis (22) and to determine the percent of mast cells positive for BrdU incorporation (21). Other slides of the Carnoy's-fixed specimens were stained with the heparin-binding fluorescent dye berberine sulfate (23) or

with alcian blue/safranin, and were examined in an epifluorescent or light microscope, respectively, as previously described (1, 22).

Studies in Rats. Groups of five to eight female 16-wk-old Wistar rats received 14-d daily intravenous injections of either rrSCF¹⁶⁴ (25 µg/kg/d in 1.0 ml of sterile, pyrogen-free 0.9% NaCl "saline") or 1.0 ml of saline alone. 24 h after the last injection, the rats were killed by exsanguination under deep ether anesthesia, and samples of tissues were fixed in 4.0% paraformaldehyde, processed into paraffin, and assessed in sections stained with 0.5% toluidine blue at pH 0.5 for 2 h (24). Mast cells were quantified at 400×, and the results were expressed as no. (mean ± SEM)/mm² of tissue (22). Other samples of the same tissues were processed as described (24) for determination of content of rat mast cell protease I (RMCP I) and RMCP II by ELISA, and the results were expressed as micrograms protease (mean ± SEM)/gram of tissue. The bone marrow cells were harvested from a single femur of each rat by flushing the marrow cavity with 0.9% NaCl and then washing the cells three times with HBSS/5% FCS (HBSS/FCS). The peritoneal lavage cells of each rat were obtained by lavaging each peritoneal cavity with 20 ml of HBSS/FCS and then washing the cells three times with HBSS/FCS. Aliquots of the total femoral bone marrow cells and peritoneal lavage cells were diluted 1:10 in methylene blue/propylene glycol for counting the numbers of total cells and mast cells in each preparation, the remainder of the femoral blue marrow or peritoneal cells were used for determination of RMCP I and II content by ELISA (24).

Results and Discussion

Local Administration of SCF Induces the *c-kit*-dependent Development of CTMC in the Skin of Mice. Recombinant rat SCF¹⁶⁴ administered in daily subcutaneous injections for 3 wk induced a striking expansion of dermal mast cell populations in normal WBB6F₁-+/+ or WCB6F₁-+/+ mice, and in genetically mast cell-deficient WCB6F₁-*Sl*/*Sl*^h mice (Table 1). In WBB6F₁-+/+ mice, mast cell numbers at sites injected with 100 or 30 µg of rrSCF¹⁶⁴/kg were 166 or 15 times that in control sites injected with vehicle alone. In striking contrast to its effects on populations of dermal mast cells, rrSCF¹⁶⁴ had little or no effect on the Hct of normal mice of either genotype tested (Table 1).

The majority of the mast cells at rrSCF¹⁶⁴ injection sites, like dermal CTMC in the skin of untreated normal mice (1, 22), exhibited cytoplasmic reactivity with the heparin-binding fluorescent dye berberine sulfate, and many of them stained with safranin (Fig. 1, A, B, D, and E). Thus, injection of rrSCF¹⁶⁴ as the sole exogenous cytokine resulted in the development of dermal mast cells with characteristics of CTMC. Colocalization of immunohistochemical staining for BrdU incorporated into nuclear DNA and mast cell cytoplasmic granule staining with alcian blue (21) indicated that morphologically identifiable mast cells were proliferating at sites of rrSCF¹⁶⁴ injection (Table 1 and Fig. 1 F). This finding indicated that the increased numbers of mast cells at sites of rrSCF¹⁶⁴ injection reflected at least in part the proliferation of differentiated dermal mast cells, not merely the proliferation and/or maturation of mast cell precursors.

However, the skin of *Sl*/*Sl*^h mice ordinarily contains virtually no dermal mast cells (Table 1 and reference 25). Thus, even though many differentiated mast cells were proliferating

Table 1. *c-kit*-dependent Stimulation of Proliferation of Mouse Dermal Mast Cells *in vivo* by *rrSCF*¹⁶⁴

| Mouse genotype | <i>rrSCF</i> ¹⁶⁴ $\mu\text{g}/\text{kg}/\text{d}$ | Hct | | Mast cells <i>no./mm</i> ² of dermis | BrdU ⁺ (percent of mast cells) |
|--|---|------------------|-----------------|--|--|
| | | Before treatment | After treatment | | |
| WBB6F ₁ ^{-/+} | 100 | 48 ± 1.3 | 50 ± 0.7 | 4,710 ± 1,190* | 16 ± 2.2* |
| | 30 | 50 ± 0.8 | 51 ± 0.3 | 454 ± 155* | 13 ± 2.7* |
| | 0 | 51 ± 0.6 | 51 ± 0.5 | 31 ± 2 | 3.3 ± 1.2 |
| WBB6F ₁ ^{-W/W^v} | 100 | 39 ± 0.8 | 36 ± 3.4 | 0 | - |
| | 30 | 40 ± 0.7 | 40 ± 2.5 | 0 | - |
| | 0 | 40 ± 0.6 | 39 ± 0.2 | 0 | - |
| WCB6F ₁ ^{-/+} | 30 | 46 ± 0.8 | 44 ± 2.7 | 122 ± 17* | 36 ± 3.5* |
| | 0 | 47 ± 0.3 | 44 ± 0.4 | 29 ± 5 | 4.9 ± 0.7 |
| WCB6F ₁ ^{-Sl/Sl^d} | 30 | 30 ± 1.1 | 38 ± 1.7* | 61 ± 24* | 29 ± 5.7 |
| | 0 | 30 ± 0.3 | 33 ± 0.6 | 0 | - |

Mice were killed 24 h after the last of 21 daily subcutaneous injections of *rrSCF*¹⁶⁴ or vehicle for assessment of the number of mast cells/mm² of dermis at the injection site and for quantification of the percent of these mast cells that were proliferating, based on nuclear incorporation of BrdU (See Materials and Methods). All results are expressed as mean ± SEM (*n* = 5–7).

* *p* < 0.001 vs. value for mice treated with 0 $\mu\text{g}/\text{kg}/\text{d}$ by Student's *t* test (two tailed).

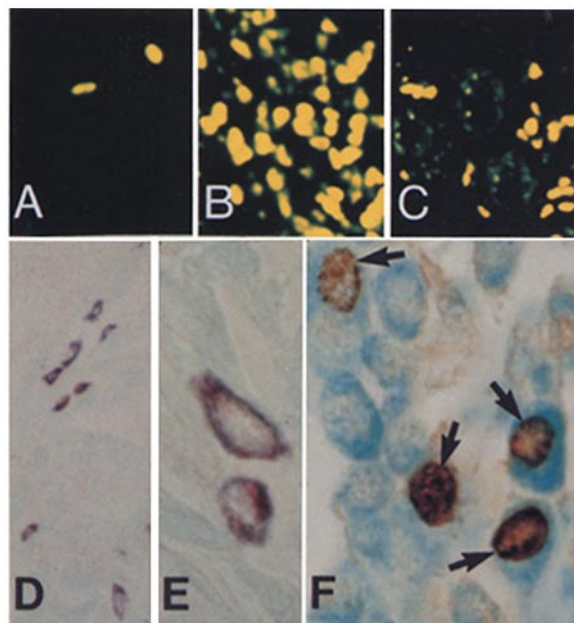


Figure 1. (A–C) Skin at sites injected daily for 3 wk with either *rrSCF*¹⁶⁴ or vehicle and then stained with berberine sulfate. (A) Vehicle injection site in a WBB6F₁^{-/+} mouse exhibits only occasional brightly fluorescent mast cells. (B) Site in a WBB6F₁^{-/+} mouse injected with *rrSCF*¹⁶⁴ (100 $\mu\text{g}/\text{kg}$) exhibits many mast cells. (C) Many berberine sulfate-positive mast cells in the skin of a WCB6F₁^{-Sl/Sl^d} mouse injected with *rrSCF*¹⁶⁴ (30 $\mu\text{g}/\text{kg}$). (D and E) Low (D) and high (E) power views of safranin-positive dermal mast cells at a *rrSCF*¹⁶⁴ injection site (30 $\mu\text{g}/\text{kg}$) in a WCB6F₁^{-/+} mouse. (F) A *rrSCF*¹⁶⁴ (100 $\mu\text{g}/\text{kg}$) injection site in a WBB6F₁^{-/+} mouse that had received BrdU to label proliferating cells. There are many mast cells (stained with alcian blue in this preparation); four of these mast cells have nuclei that are positive (brown immunoperoxidase reaction product) for BrdU (arrows) (A–C, E, and F; $\times 250$; D: $\times 1,000$).

at *rrSCF*¹⁶⁴ injection sites in *Sl/Sl^d* mice 3 wk after initiation of treatment (Table 1), some of the effect of *rrSCF*¹⁶⁴ in these mice must have reflected the recruitment, proliferation, and/or induction of maturation of the mast cell precursors present in these animals (25). An effect of *rrSCF*¹⁶⁴ on mast cell maturation is also supported by the finding that many of the mast cells at *rrSCF*¹⁶⁴ injection sites in *Sl/Sl^d* mice stained with berberine sulfate (Fig. 1 C) or safranin (data not shown), and by data indicating that *rrSCF*¹⁶⁴ can induce IL-3-dependent bone marrow-derived cultured mast cells to mature and acquire phenotypic characteristics of CTMC *in vitro* (20, 26).

In contrast to the effects of *rrSCF*¹⁶⁴ in *Sl/Sl^d* mice, injection of *rrSCF*¹⁶⁴ into genetically mast cell-deficient WBB6F₁^{-W/W^v} mice influenced neither the profound mast cell deficiency, nor the anemia, of these animals (Table 1). This result, together with work indicating that the *W* and *W^v* alleles encode *c-kit* products that express no (*W*) or markedly diminished (*W^v*) tyrosine kinase activity (27), indicates that the ability of *rrSCF*¹⁶⁴ to induce the development of dermal CTMC *in vivo* requires that the *rrSCF*¹⁶⁴ interacts with a functionally active *c-kit* receptor. The complete failure of even very high doses of *rrSCF*¹⁶⁴ to induce cutaneous mast cell development in *W/W^v* mice is also noteworthy in light of reports that this mutant can develop mature dermal mast cells in association with a chronic idiopathic dermatitis (22), in response to repeated epicutaneous applications of PMA (28), or as a result of treatment with IL-3 (29). The results reported here indicate that the appearance of mast cells in the skin of *W/W^v* mice in these settings probably does not reflect increased local production of endogenous *c-kit* ligand, but instead may be due to other

effects such as the activation of either alternative signaling mechanisms or processes distal to the interaction between *c-kit* and its ligand.

Systemic Administration of SCF Expands Populations of CTMC and MMC in the Rat. Even though subcutaneous administration of rrSCF¹⁶⁴ increased the Hct of *Sl/Sl^h* mice (Table 1), the effects of rrSCF¹⁶⁴ on mast cell populations were most striking in the vicinity of the subcutaneous injection sites. For example, no mast cells appeared in the skin contralateral to rrSCF¹⁶⁴ injection sites in *Sl/Sl^h* mice or in the gastric tissues of these animals (data not shown). To assess the systemic effects of rrSCF¹⁶⁴ on mast cell development and phenotype, we turned to the rat. Daily intravenous injections are more readily performed in rats than in mice. More importantly, Huntley et al. (24) have reported highly sensitive and specific ELISA methods for quantifying tissue content of mast cell-specific proteases associated with CTMC

and MMC (RMCP I and RMCP II, respectively) in this species.

We found that daily intravenous administration of rrSCF¹⁶⁴ to rats at a dose of 25 $\mu\text{g}/\text{kg}/\text{d}$ for 14 d increased mast cell levels systemically and also produced striking increases in levels of both types of rat mast cell-associated proteases (Tables 2 and 3). In the skin, lung, and liver (Table 2), rrSCF¹⁶⁴ treatment resulted in marked increases in numbers of mast cells and in tissue content of the CTMC-associated protease, RMCP I, but no significant changes in levels of the MMC-associated protease, RMCP II. In the spleen, bone marrow, and peritoneal cavity (Tables 2 and 3), rrSCF¹⁶⁴ treatment resulted in striking expansion of mast cell populations, as well as marked increases in levels of both RMCP I and RMCP II. Note, however, that the total amounts of RMCP II in these sites in the rrSCF¹⁶⁴-treated animals were much less than the corresponding amounts of RMCP

Table 2. Systemic Effects of rrSCF¹⁶⁴ on Mast Cell Populations and Tissue Mast Cell Protease Content in the Rat

| Tissue | rrSCF ¹⁶⁴ | Mast cells | RMCP I | RMCP II |
|--------------------|----------------------------------|-------------------------------------|-----------------------------------|-----------------------------------|
| | $\mu\text{g}/\text{kg}/\text{d}$ | <i>no./mm² of tissue</i> | $\mu\text{g}/\text{gm of tissue}$ | $\mu\text{g}/\text{gm of tissue}$ |
| Skin | 25 | 87 \pm 7* | 69 \pm 7* | ND |
| | 0 | 26 \pm 3 | 27 \pm 3 | ND |
| Lung | 25 | 48 \pm 6* | 13 \pm 1.2* | 10 \pm 1.6 |
| | 0 | 7 \pm 0.4 | 3.8 \pm 0.8 | 11 \pm 1.2 |
| Liver | 25 | 62 \pm 10* | 35 \pm 8 [†] | 1.2 \pm 0.3 |
| | 0 | 3.6 \pm 0.4 | 0.4 \pm 0.06 | 0.8 \pm 0.1 |
| Spleen | 25 | 204 \pm 40* | 92 \pm 24 [†] | 14.2 \pm 5.0 [§] |
| | 0 | 0.07 \pm 0.02 | 0.7 \pm 0.08 | 0.10 \pm 0.004 |
| Glandular stomach: | | | | |
| Mucosa | 25 | 50 \pm 9 [†] | 5.9 \pm 1.2 | 19 \pm 1.3 [§] |
| Submucosa | 25 | 144 \pm 36 [§] | | |
| Muscularis propria | 25 | 32 \pm 5 [†] | | |
| Mucosa | 0 | 12 \pm 3 | 3.3 \pm 0.4 | 13 \pm 1.7 |
| Submucosa | 0 | 45 \pm 8 | | |
| Muscularis propria | 0 | 12 \pm 2 | | |
| Ileum: | | | | |
| Mucosa | 25 | 80 \pm 14 [§] | 1.4 \pm 0.5 [§] | 508 \pm 83 [§] |
| Submucosa | 25 | 146 \pm 51 [§] | | |
| Muscularis propria | 25 | ND | | |
| Mucosa | 0 | 38 \pm 12 [†] | 0.2 \pm 0.1 | 272 \pm 5 |
| Submucosa | 0 | 19 \pm 4 [§] | | |
| Muscularis propria | 0 | ND | | |

Rats were killed \sim 24 h after the last injection of 14 daily intravenous injections of rrSCF¹⁶⁴ or vehicle for assessment of mast cell numbers and content of RMCP I and RMCP II in various tissues (See Materials and Methods). All results are expressed as mean \pm SEM ($n = 5-8$).

*, †, or § = $p < 0.001$, 0.01, or 0.05, respectively, vs. value for rats treated with 0 $\mu\text{g}/\text{kg}/\text{d}$ by Student's *t* test (two tailed).

Table 3. Systemic Effects of rrSCF¹⁶⁴ on Peritoneal and Bone Marrow Mast Cells and Mast Cell Protease Content in the Rat

| Site | rrSCF ¹⁶⁴ $\mu\text{g}/\text{kg}/\text{d}$ | Cells/peritoneal cavity or femur | | Protease content of total peritoneal cells or femoral bone marrow cells | |
|---------------------|--|----------------------------------|-----------------------|---|------------------------|
| | | Total cells (10^7) | Mast cells (10^6) | RMCP I μg | RMCP II ng |
| Peritoneal cavity | 25 | 2.3 \pm 0.1 | 7.6 \pm 0.5* | 134 \pm 7* | 17 \pm 5* |
| | 0 | 2.1 \pm 0.2 | 1.7 \pm 0.3 | 49 \pm 3 | 0.2 \pm 0.2* |
| Femoral bone marrow | 25 | 9.0 \pm 0.3 | 3.9 \pm 0.4* | 34 \pm 4* | 42 \pm 6* |
| | 0 | 8.7 \pm 0.6 | 0.3 \pm 0.1 | 7 \pm 1 | 8 \pm 3 |

These data (mean \pm SEM) are from the same rats shown in Table 2 (see Materials and Methods).

* $p < 0.001$ vs. value for rats treated with 0 $\mu\text{g}/\text{kg}/\text{d}$ by Student's t test (two tailed).

I. rrSCF¹⁶⁴ treatment significantly expanded populations of mast cells in the mucosa, submucosa, and muscularis propria of the glandular stomach, and in the mucosa and submucosa of the ileum (Table 2). MMC predominate in the mucosa of these organs, whereas CTMC predominate in the submucosa and muscularis propria (3, 4, 24). rrSCF¹⁶⁴ treatment increased the content of both RMCP II ($p = 0.018$ by the two-tailed Student's t test) and RMCP I ($p = 0.045$ by the one-tailed Student's t test) in the glandular stomach, and also significantly increased the content of both proteases in the ileum (Table 2). Thus, according to anatomical distribution and mast cell protease phenotype (reviewed in references 3-5 and 24), intravenous administration of rrSCF¹⁶⁴ to rats markedly expanded populations of CTMC and also increased levels of gastrointestinal MMC.

By immunohistochemistry (24), mast cells in rrSCF¹⁶⁴-treated rats exhibited a pattern of staining for RMCP I and II that was similar to that of mast cells in control rats (24). For example, dermal mast cells were exclusively RMCP I⁺ and RMCP II⁻, whereas mast cells in the mucosa of the ileum were predominantly RMCP II⁺ and RMCP I⁻. However, the effect of rrSCF¹⁶⁴ treatment on the numbers of mast cells in various sites generally was more marked than the corresponding effect on RMCP content. For example, in the spleen, femoral bone marrow, and glandular stomach, mast cell numbers were $\sim 3,000$, 13, and 3 times those in control rats, whereas the corresponding ratios for total content of RMCP I and II in rrSCF¹⁶⁴-treated vs. control rats were ~ 130 , 5, and 1.5. This finding may reflect any of a

number of factors, including the recent expansion of mast cell populations in the rrSCF¹⁶⁴-treated animals. It is well established that the cytoplasmic granule-associated mediator content of rat mast cells increases progressively with the age of the cells (reviewed in references 30 and 31).

These data show that the local injection of SCF promoted the expansion of dermal CTMC populations in mice in vivo, and indicate that this effect required a functionally active *c-kit* receptor. In rats, intravenous administration of rrSCF¹⁶⁴ resulted in striking expansions of mast cell populations in all of the anatomical sites examined. These included sites that usually contain large populations of CTMC, such as the skin and peritoneal cavity, and sites that ordinarily contain substantial numbers of MMC, such as the mucosa of the glandular stomach and ileum. These findings, taken together with the observation that *W/W^v* and *Sl/Sl^h* mice virtually lack both CTMC and MMC (1, 2, 6), indicate that interactions between *c-kit* and its ligand importantly regulate the normal development of both CTMC and MMC. However, administration of exogenous SCF not only produced significant expansions of populations of CTMC and MMC in sites ordinarily containing large numbers of mast cells, such as the skin and stomach, but also resulted in the appearance of many mast cells at sites that typically contain very few mast cells, such as the liver, spleen, and bone marrow. These findings raise the possibility that the mast cell proliferation associated with certain immunological or pathological responses may reflect, in part, excessive local or even systemic production of *c-kit* ligand.

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