

Neutrophil Pool Sizes and Granulocyte Colony-Stimulating Factor Production in Human Mid-Trimester Fetuses

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ABSTRACT

We quantified neutrophils and neutrophil progenitors, and assessed granulocyte colony-stimulating factor (G-CSF) production in the liver and bone marrow of 20 human abortuses after elective pregnancy termination between 14 and 24 wk of gestation. Mature neutrophils were not observed in any of the liver specimens, but were present in the bone marrow as early as 14 wk. The concentrations of neutrophils in the fetal marrow were extremely low, by comparison with term infants and adults, with less than 5% of the nucleated cells being segmented neutrophils, band neutrophils, or metamyelocytes compared with 31–69% in term infants. Despite the low neutrophil populations, progenitors which had the capacity for clonal maturation into neutrophils *in vitro* were abundant in the fetal liver and fetal bone marrow. In addition, such progenitors had a dose-response relationship to recombinant G-CSF similar to that of progenitors from the bone marrow of healthy adults. At each gestational age tested, stimulation of mononuclear cells from fetal liver with IL-1 α generated less G-CSF protein and fewer G-CSF mRNA transcripts than did stimulation of mononuclear cells from fetal bone marrow. Mononuclear cells from the fetal bone marrow produced less G-CSF protein and mRNA than did mononuclear cells from the blood of

adults. Thus, the liver of the mid-trimester human fetus is almost devoid of neutrophils, and the bone marrow contains a significantly lower proportion of neutrophils than does the marrow of term neonates or adults. These findings correlate with IL-1 α -induced production of G-CSF in these organs. The lack of G-CSF production might explain the small neutrophil reserves found in extremely preterm infants. (*Pediatr Res* 37: 806–811, 1995)

Abbreviations

NSP, neutrophil storage pool
G-CSF, granulocyte colony-stimulating factor
GM-CSF, granulocyte-macrophage colony-stimulating factor
CFU-MIX, colony-forming unit-mixed (multipotent hematopoietic progenitor)
CFU-GM, colony-forming unit-granulocyte-macrophage progenitor
BFU-E, burst forming unit-erythroid (primitive erythroid progenitor)
 α -MEM, α -modified minimal essential medium
r, recombinant

The NSP is the sum of all segmented neutrophils, band neutrophils, and metamyelocytes held in reserve for ready release into the circulation (1). Rat pups delivered prematurely have a much smaller NSP, per g of body weight, than do those delivered at term, and pups delivered at term have a smaller NSP than do adult rats (2–6). It is unknown, however, whether this observation applies to humans. Studies employing cells

from human umbilical cord blood suggest that preterm infants have a significant impairment in their capacity to produce the neutrophil growth factor, G-CSF (7, 8). Thus, if human preterm infants, like preterm rat pups, have a small NSP, this might be on the basis of their lack of G-CSF production. However, the significance of these umbilical cord blood studies is not clear. Specifically, it is more likely that cells within the hematopoietic organs play a much more significant role in producing hematopoietic growth factors than do cells in the umbilical cord blood; cord blood cells have been studied primarily because of the convenience of their acquisition. We maintained that additional studies were needed to determine the applicability of the animal and umbilical cord blood studies to human developmental biology. With this in mind, we obtained human fetuses ranging from 14 to 24 wk of gestation, immediately after elective termination of pregnancies. From these we as-

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sessed the relative sizes of the neutrophil reserves and the capacity of the hematopoietic organs to generate G-CSF.

METHODS

Liver and marrow studies. Between July 1993 and March 1994, liver and bone marrow were obtained from 20 human abortuses ranging from 14 to 24 wk gestation. Only fetuses that were normal by ultrasound examination and underwent elective pregnancy termination were studied. Pregnancy terminations were carried out by suction curettage (14-wk gestation) or by cervical dilatation and extraction curettage (15–24 wk gestation). In some of the abortuses only bone marrow was obtained, and in some only liver was obtained. Blood and bone marrow were also obtained from five healthy adult volunteers. The marrow was aspirated from the adult's anterior-superior iliac crest (after local Xylocaine infiltration) into a sterile syringe containing preservative-free heparin (Squibb-Marsham, Cherry Hill, NJ). All of the studies were performed in accordance with protocols approved by the University of Florida Institutional Review Board, and informed consent was obtained from the donors of blood and marrow cells.

Cell suspensions. Suspensions of cells from the fetal livers were prepared by triturating the tissues in sterile α -MEM (Hyclone Laboratories, Logan UT) and passing the pieces through serially smaller gauge needles, beginning with a #16 and ending with a #23. Fetal bone marrow cell suspensions were prepared by flushing the femurs and tibias three times with 5 mL sterile α -MEM. Nucleated cell counts were performed electronically on the cell suspensions (Coulter Electronics, Hialeah, FL). Differential cell counts (500–1000 cells) were performed on Wright-stained films.

Flow cytometric analysis. Single cell suspensions were prepared in medium 199 (Life Technologies, Inc., Bethesda, MD). Dual staining was performed using phycoerythrin-stained goat anti-mouse Ig secondary antibodies (Becton Dickinson Immunocytometry Systems, San Jose, CA) and FITC-stained primary antibodies directed against mature myeloid/monocyte cells. Samples were analyzed using primary MAB against CD14 and CD15 (kindly provided by Edward Ball, M.D., through Meadrex, West Lebanon, NH). Flow cytometry was performed with an Epics Profile Analyzer (Coulter Electronics).

Quantification of hematopoietic progenitors. Clonogenic assays were performed to assess fetal liver and marrow concentrations of multipotent hematopoietic progenitors (CFU-MIX), primitive erythroid progenitors (BFU-E), and granulocyte-macrophage progenitors (CFU-GM). Assays were performed in methyl cellulose cultures containing 5 U/mL erythropoietin (Amgen, Thousand Oaks, CA) and 10 ng/mL each of G-CSF (Amgen), GM-CSF (Immunex Corp, Seattle, WA), IL-3 (R&D Systems, Minneapolis, MN), and M-CSF (R&D Systems). Whole cell suspensions (not fractionated) from fetal liver and fetal bone marrow were cultured at a density of $1-5 \times 10^3$ /mL in α -MEM containing 5×10^{-4} mol/L β -mercaptoethanol (Eastman Kodak Co., Rochester, NY), 30% FCS (Hyclone), 1% BSA (Sigma Chemical Co., St. Louis, MO), and 1.1% methyl cellulose (Sigma Chemical Co.).

After 14 d, colonies were evaluated *in situ* using an inverted microscope. All studies were performed in quadruplicate. To determine the varieties of cells present in the clones, the dishes were scraped, and their contents were rinsed three times, after which the cells rinsed from the dishes were smeared on glass slides and stained with Wright stain for 500 cell differential counts.

Response of granulocytic progenitors to rG-CSF. The effect of rG-CSF on production of neutrophils by progenitors was tested by plating light-density (specific gravity <1.077 , Ficoll-Paque, Pharmacia Biotech Inc., Uppsala, Sweden) cell suspensions (5×10^3 cells/mL) from fetal liver or adult marrow in methylcellulose cultures with 10 ng/mL recombinant stem cell factor (Amgen) and either 0, 0.01, 0.1, 1, 0, or 100 ng/mL rG-CSF. After 10–12 d in culture, colonies were counted, and the cells were scraped from the plates and collected in 5 mL α -MEM. The nucleated cells present in each culture plate were then determined by hemocytometer counting of the cell suspensions. The neutrophils per plate were determined by multiplying the percentage of neutrophils from Wright-stained differential cell counts by the nucleated cells per plate.

Quantification of G-CSF. To assess the ability of fetal liver and bone marrow to generate G-CSF, 10^6 light-density mononuclear cells from fetal liver, fetal marrow, or for comparison, mononuclear cells from the blood of five adults were incubated for 24 h in 1 mL α -MEM in 5% CO₂ at 37°C with or without the addition of 10 ng/mL recombinant IL-1 α (R&D Systems), an effective stimulator of mononuclear cells used in previous studies (7, 8). All studies were performed in quadruplicate. The purified rIL-1 α had a specific activity of $>5 \times 10^8$ U/mg and a purity of $>95\%$ by SDS-polyacrylamide gel electrophoresis. Concentrations of G-CSF in the supernatants were measured by ELISA (R&D Systems). The minimal G-CSF concentration detectable was 11 pg/mL. The ELISA recognized both natural and recombinant G-CSF and had no cross-reactivity with the following cytokines: IL-1 α , IL-1 β , IL-2, IL-3, IL-4, IL-5, IL-6, tissue necrosis factors α and β , GM-CSF, transforming growth factor β 1, and platelet-derived growth factor.

RNA isolation and Northern blot analysis. RNA was prepared from light density mononuclear cells obtained from fetal liver, fetal bone marrow, and from adult blood, after such cells were cultured for 24 h in the presence or absence of IL-1 α (10 ng/mL). A guanidinium extraction buffer was added to the cells, and RNA was isolated using the method described by Chomczynski and Sacchi (9). The RNA was subjected to electrophoresis through a 1% agarose-formaldehyde gel, transferred to a biotrace nylon membrane (Gelman Sciences, Ann Arbor, MI), and immobilized by UV cross-linking (10). The human G-CSF probe (kindly supplied by Dr. Steve Gillis, Immunex Corp., Seattle, WA) was labeled, hybridized to the blot, washed, and subjected to autoradiography. A human β -actin probe was used as a control (University of Utah core nucleotide synthesis laboratory).

Statistical analysis. Specimens were grouped for analysis by gestational age as follows: 14–17 wk, 18–20 wk, and 21–24 wk. The Wilcoxon signed ranks test was used to assess differences in cell populations at the different gestational ages and

between liver and bone marrow. Differences in concentrations of G-CSF in cell suspensions between control and IL-1-stimulated cells were assessed by a paired *t* test. Differences in concentrations of G-CSF in cell suspensions between different groups were assessed by a nonpaired *t* test. A two-tailed *p* value of less than 0.05 was considered to indicate a significant difference.

RESULTS

Differential cell counts of fetal liver and fetal bone marrow. The differential counts of the fetal liver cell suspensions are shown in Table 1. At 14–17 wk no neutrophils were observed in the liver. At 18–20 wk a small number of promyelocytes, but no postmitotic neutrophils, were observed. At 21–24 wk the percentage of promyelocytes and myelocytes had increased, but no postmitotic neutrophils were observed, and total neutrophils were less than 2% of the total nucleated cells.

Differential counts of the fetal bone marrow cell suspensions are shown in Table 2. At all gestational ages studied the marrow and liver populations were very different from one another. The bone marrow invariably contained a much smaller proportion of normoblasts than did the liver, and segmented neutrophils were evident within the fetal bone marrow as early as 14 wk. Compared with term neonates and adults, however, the percentages of cells in the fetal marrow categorized as NSP cells were very low. Thirty one to 69% of the nucleated cells in the marrow of term neonates and 25–50% in the marrow of adults are NSP cells (11, 12) whereas less than 5 percent of the cells in the fetal bone marrow were NSP cells.

In four specimens less than 20 wk of gestation, the relative lack of neutrophils in fetal liver and marrow was validated by flow cytometry. Specifically, less than 1% of the liver cells were positive for CD14 or CD15, while 2.6% of the marrow cells were CD14 positive, and 8.5% were CD15 positive.

Granulocyte progenitors in the fetal liver and bone marrow. Fetal liver contained a higher concentration of CFU-MIX and BFU-E than did fetal bone marrow (Table 3). Although the fetal liver cell suspensions contained no postmitotic neutrophils (Table 1), CFU-GM were indeed present. In fact, the concentration of CFU-GM in fetal liver was not significantly less than in fetal bone marrow. It was not clear from these studies, however, whether the CFU-GM in fetal liver had matured *in vitro* into neutrophil/macrophage clones or, rather, into macrophage clones only. Therefore the cells were washed from the culture dishes, smeared on glass slides, and stained for differential cell counts. Of the total cells developing from fetal liver progenitors, $5.5 \pm 1.8\%$ were promyelocytes and myelocytes, and $5.6 \pm 1.1\%$ were postmitotic neutrophils.

The rG-CSF dose-response relationship of light density cell suspensions (containing progenitors and accessory cells) from fetal liver and adult bone marrow is shown in Figure 1. At every concentration of rG-CSF tested, progenitors from fetal liver produced more neutrophils per plate than did progenitors from adult marrow.

G-CSF production by fetal liver and marrow. After a 24-h incubation with IL-1 α , the supernatants of fetal liver cells contained a greater concentration of G-CSF than did liver cells after incubation without IL-1 α (Fig. 2, upper panel). At all of the three gestational ranges tested, fetal bone marrow (Fig. 2, lower panel) generated more G-CSF than did fetal liver. Liver cells of fetuses <18 wk produced more G-CSF than liver cells of fetuses ≥ 18 wk (1857 ± 1623 versus 165 ± 156 , *p* = 0.05). Neither fetal liver nor fetal marrow generated as much G-CSF as did monocytes from the blood of healthy adults. Similarly, fewer G-CSF mRNA transcripts were present in cells of fetal liver than fetal marrow, and adult blood cells contained greater G-CSF mRNA transcripts than did fetal liver (Fig. 3).

Table 1. Differential counts, in percentages (mean and 95% confidence intervals) of liver cell suspensions from elective abortions at 14–24-wk gestation

Cells	Gestation (wk)		
	14–17 (<i>n</i> = 5)	18–20 (<i>n</i> = 7)	21–24 (<i>n</i> = 8)
Normoblasts			
Pronormoblast	3.1 (1.7–4.5)	3.4 (2.8–4.0)	2.9 (1.1–4.7)
Basophilic N	18.4 (10.8–26.0)	13.6 (12.0–15.2)	13.7 (9.0–18.4)
Polychromatophilic N	57.5 (44.6–69.8)	55.3 (48.0–63.0)	51.1 (45.6–56.6)
Orthochromic N	13.9 (8.7–19.1)	15.9 (12.2–19.5)	14.2 (9.2–19.2)
Total erythroid	92.9 (86.6–99.2)	87.8 (83.0–92.6)	81.9 (76.9–86.9)
Neutrophil			
Promyelocyte	0 (0–0)	0.2 (0.0–0.4)*	1.2 (0.8–1.6)**†
Myelocyte	0 (0–0)	0 (0–0)	0.2 (0.0–0.4)**†
Metamyelocyte	0 (0–0)	0 (0–0)	0 (0–0)
Band	0 (0–0)	0 (0–0)	0 (0–0)
Segmented	0 (0–0)	0 (0–0)	0 (0–0)
Total neutrophils	0 (0–0)	0.2 (0.0–0.4)*	1.4 (0.9–1.9)**†
Undifferentiated blast	0.5 (0.0–1.1)	3.1 (1.3–4.9)*	2.2 (1.6–3.0)*
Macrophage	0.5 (0.1–1.2)	1.2 (0.8–1.6)	1.3 (0.9–1.9)
Lymphocyte	5.4 (2.6–7.8)	3.9 (0.3–7.5)	11.3 (9.1–17.5)**†
Eosinophil	0 (0–0)	0 (0–0)	0 (0–0)
Other	0.7 (0.0–1.5)	3.4 (1.0–5.8)	1.9 (0.0–4.7)

N = normoblast; other = hepatocyte, megakaryocyte, or cell of undetermined origin.

* *p* < 0.05 vs 14–17 wk.

† *p* < 0.05 vs 18–20 wk.

Table 2. Differential counts, in percentages (mean and 95% confidence intervals) of bone marrow cell suspensions from elective abortions at 14–24-wk gestation

Cells	Gestation (wk)		
	14–17 (n = 6)	18–20 (n = 6)	21–24 (n = 8)
Normoblasts			
Pronormoblast	0.3 (0.1–0.5)*	1.1 (0.7–1.5)*	0.5 (0.3–0.7)*
Basophilic N	1.4 (0.8–2.0)*	3.1 (2.3–3.9)*	1.7 (0.7–2.7)*
Polychromatophilic N	9.2 (4.2–14.2)*	12.9 (6.7–19.1)*	12.3 (7.1–17.5)*
Orthochromic N	12.1 (1.5–23.7)	20.7 (8.5–32.9)	8.1 (3.7–12.5)
Total erythroid	23.0 (9.7–36.3)*	37.8 (21.0–54.6)*	22.6 (13.6–31.6)*
Neutrophils			
Promyelocyte	5.9 (2.6–8.4)*	5.8 (3.2–8.4)*	5.0 (3.8–7.2)*
Myelocyte	2.7 (1.1–3.9)*	3.4 (1.8–5.0)*	2.1 (1.1–3.1)*
Metamyelocyte	2.1 (1.1–3.1)*	2.5 (0.7–4.3)*	1.7 (0.7–2.7)*
Band	3.1 (1.3–4.9)*	3.2 (0.6–6.0)*	1.6 (0.4–2.8)*
Segmented	1.2 (0–2.6)*	1.1 (0–2.5)*	0.4 (0.2–0.6)*
Total neutrophils	13.8 (8.0–19.6)*	16.1 (8.1–24.1)*	10.7 (5.7–15.7)*
Undifferentiated blast	8.9 (2.7–15.1)*	11.4 (6.0–16.8)*	11.6 (8.0–15.2)*
Macrophage	5.4 (1.6–9.2)*	4.2 (1.8–6.6)*	3.1 (1.5–4.7)
Lymphocyte	35.3 (19.5–51.1)*	28.8 (20.8–36.8)*	50.3 (43.5–57.1)*
Eosinophil	0.5 (0–1.1)	1.1 (0–2.3)	1.2 (0.4–2.0)*
Other	0.8 (0.4–1.2)	0.5 (0.1–0.9)	0.4 (0.2–0.6)

N = normoblast; other = megakaryocyte, or cell of undetermined origin.

* $p < 0.05$ vs fetal liver.

Table 3. Hematopoietic colonies (mean \pm SEM) per 5×10^3 plated cells, quantified 14 d after plating nucleated cell suspensions (nonfractionated) from fetal liver or fetal bone marrow

Cells	CFU-MIX	BFU-E	CFU-GM
Liver (n = 7)	12.7 \pm 2.1	20.7 \pm 3.1	22.3 \pm 4.5
Marrow (n = 6)	6.7 \pm 1.4*	9.3 \pm 2.7*	33.1 \pm 7.2

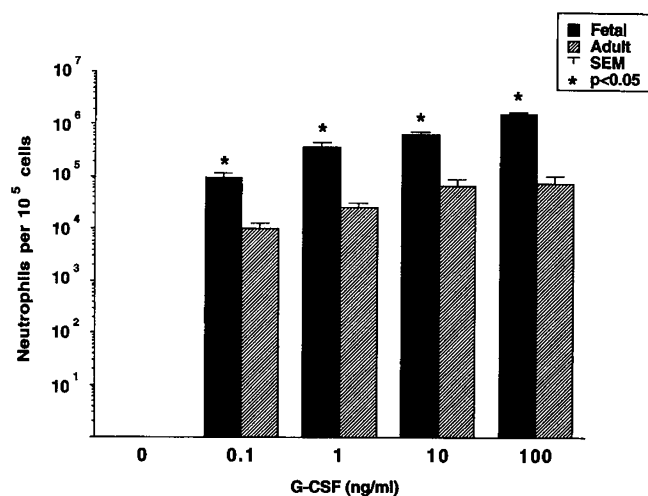
Cells were cultured in methylcellulose in the presence of recombinant erythropoietin, rG-CSF, rGM-CSF, rIL-3, and rM-CSF.

* $p < 0.05$ vs liver.

DISCUSSION

During pyogenic infections in otherwise normal adults, the daily neutrophil utilization rate increases up to 8-fold over steady-state utilization (13–15). Despite this increased demand for neutrophils, the NSP (composed of $5\text{--}7 \times 10^9$ segmented neutrophils, band neutrophils, and metamyelocytes per kg of body weight) rarely becomes completely depleted (1, 13, 15, 16). In prematurely delivered neonates, however, complete depletion of the NSP is a relatively common sequelae of bacterial infection, frequently accompanied by profound neutropenia, overwhelming sepsis, and death (2, 3, 17–21).

The propensity of preterm infants to deplete their NSP during bacterial infection has been ascribed, in part, to qualitative and quantitative developmental deficiencies of neutrophils and their progenitors (22–24). Substantial evidence exists regarding poor chemotaxis, decreased bacterial killing, and diminished oxidative metabolism of neutrophils from human preterm infants. The notion of a deficient NSP in preterm infants, however, has been inferred solely from animal models. For instance, in a previous study we observed that newborn rats had only 25% of the NSP, per g body weight, of adult rats (2). We also observed that the quantity of CFU-GM per g body weight in newborn rats was only about 10% that of adult rats (4). In this study, granulocytic progenitors were indeed present in human fetal liver and marrow, excluding the hypothesis that

**Figure 1.** Neutrophils washed from culture dishes 10–12 d after plating 5×10^3 light density cells from fetal liver or adult bone marrow. Cells were cultured in the presence of recombinant stem cell factor (10 ng/mL) plus various concentrations of rG-CSF. Neutrophil counts are listed per 10^5 cells plated.

the lack of granulocytopoiesis in the fetal liver was the result of lack of CFU-GM in that organ. Fetal liver contained a higher concentration of CFU-MIX and BFU-E than did fetal bone marrow, which likely forms the basis for the erythropoietic nature of mid-trimester fetal liver documented in this study.

Little information is available regarding granulocytopoiesis in the human fetus (25–27), and available data are not sufficient to judge whether the NSP of mid-trimester human fetuses is small, as is the case in rats. In the present study we observed no neutrophils beyond the myelocyte stage in the livers of any of 20 human fetuses at 14–24-wk gestation. Although neutrophils were observed in the bone marrow, even as early as 14 wk of gestation, the NSP invariably constituted less than 5% of the nucleated marrow cells. This is in marked contrast to the

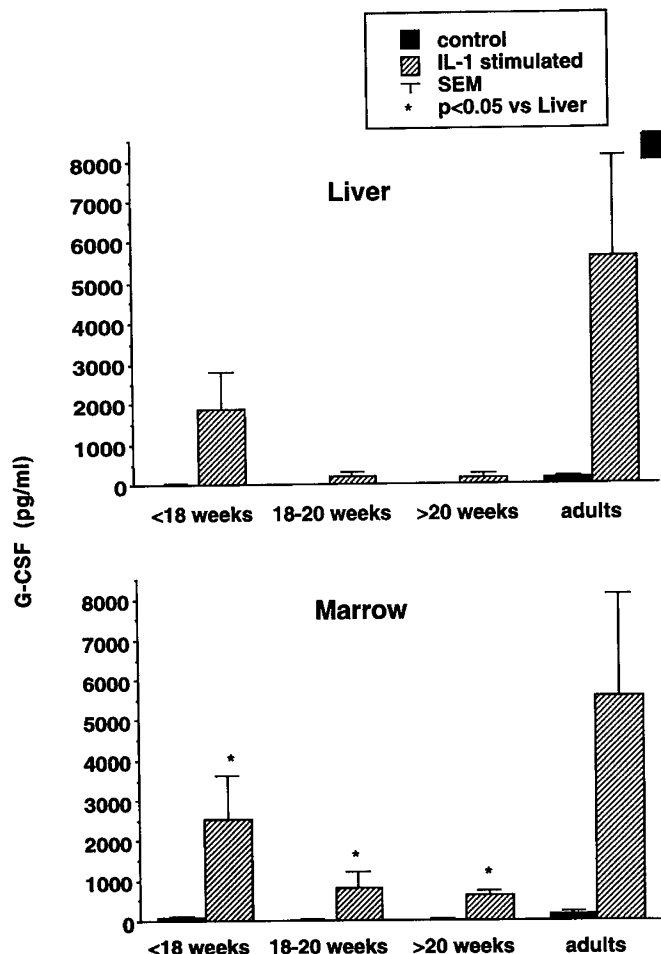


Figure 2. Accumulation of G-CSF in 1-mL culture dishes containing 10^6 mononuclear cells obtained from fetal liver vs adult blood (upper panel), and obtained from fetal bone marrow vs adult blood (lower panel). Cells were cultured for 24 h with or without IL-1 α (10 ng/mL).

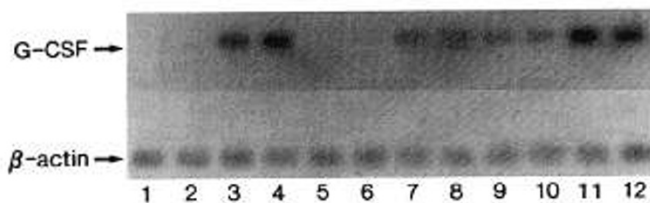


Figure 3. Northern analysis of G-CSF mRNA accumulation in cells of fetal or adult origin, incubated for 24 h with or without IL-1 α (10 ng/mL). β -actin bands for each sample are shown as a control in the lower section of the blot. Lanes 1 and 2, two separate samples of fetal marrow incubated without IL-1 α ; lanes 3 and 4, fetal marrow samples incubated with IL-1 α ; lanes 5 and 6, two separate samples of fetal liver incubated without IL-1 α ; lanes 7 and 8: fetal liver samples incubated with IL-1 α ; lanes 9 and 10, two separate adult samples incubated without IL-1 α ; lanes 11 and 12, adult samples incubated with IL-1 α .

31–69% NSP cells observed in marrow aspirates of term infants (11), and the 25–52 percent NSP cells observed in the marrow of adults (12). In the present studies, we made no attempt to sum the NSP from all bones, liver, and spleen, as would be required to calculate the absolute NSP per kg of body weight. Although the possibility that a quantitative difference in absolute numbers of granulocyte progenitors still exists, the complete absence of mature neutrophils from the liver and their

very small percentages in the femurs and tibias lead us to conclude that, indeed, human fetuses up to 24 wk of gestation have an extremely small NSP.

The size of the NSP is determined not only by the absolute quantity of granulocyte progenitors, but also by the growth factors that stimulate and regulate granulocyte development. Previous studies by Cairo *et al.* (7) illustrated that mononuclear cells obtained from the umbilical cord blood of term infants produce G-CSF poorly, compared with mononuclear cells obtained from the blood of adults. We previously observed that cells from the umbilical cord blood of preterm infants produce G-CSF less well than do those of term infants (8). Although elevated levels of G-CSF and GM-CSF have been reported in the cord blood of normal term infants (28, 29), the source of growth factor production was not evaluated.

The physiologic relevance of these observations can be questioned, however, because cells within the hematopoietic organs are probably of much greater significance in hematopoietic regulation than are blood cells. We interpret our present observations as validating the conclusions of the previous cord blood studies. It is now clear that when stimulated with the inflammatory cytokine IL-1 α , fetal bone marrow, fetal liver, and fetal blood all generate lower concentrations of G-CSF, and G-CSF mRNA, than do cells of adults.

The physiologic role of G-CSF includes up-regulation of neutrophil function and neutrophil production (30, 31). This appears to be the case for cells of the neonate as well as for adults (32). Perhaps the deficiencies in neutrophil function in preterm neonates, as well as the small NSP, can be explained, at least in part, by lack of G-CSF. Since the fetus exists *in utero* in a sterile environment, it may not be advantageous, during the first or second trimester of pregnancy, to produce G-CSF. However, when delivery occurs prematurely, perhaps the preterm neonate's limitation in producing G-CSF becomes a disadvantage, but one that could theoretically be remedied pharmacologically. In support of that potential approach, our present studies indicate that neutrophil progenitors are present in the fetal liver and bone marrow, and that these progenitors are responsive to rG-CSF.

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