Essential Requirement for c-kit and Common γ Chain in Thymocyte Development Cannot be Overruled by Enforced Expression of Bcl-2

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Abstract

The thymus in mice lacking both the receptor tyrosine kinase c-kit and the common cytokine receptor γ chain (γc) is alymphoid because these receptors provide essential signals at the earliest stages of thymocyte development. The signals transduced by these receptors potentially regulate proliferation, survival, or differentiation, but the contribution of each receptor to distinct intracellular signaling cascades is only poorly defined. Here, we have examined whether enforced expression of Bcl-2 can rescue thymocyte development in c-kit and γc single or double mutant mice. A bcl-2 transgene (Eμ-bcl-2-25; expressed in the T cell lineage) was introduced into (a) c-kit and γc wild-type (c-kit+/γc+/bcl-2+), (b) c-kit–deficient (c-kit−/γc+/bcl-2+), (c) γc–deficient (c-kit+/γc−/bcl-2+), or (d) c-kit and γc double-deficient mice (c-kit−/γc−/bcl-2−). The bcl-2 transgene was functionally active in wild-type and c-kit or γc single mutants, as it promoted survival of ex vivo isolated thymocytes, including pro-T cells. In vivo, however, transgenic Bcl-2 did not release T cell precursors from their phenotypic block and failed to increase progenitor or total thymocyte cellularity in c-kit or γc single or double mutants. These data argue strongly against a role for Bcl-2 as a key mediator in signaling pathways linked to cytokine and growth factor receptors driving early thymocyte development.

Key words: growth factors • c-kit • common cytokine receptor γ chain • bcl-2 • T cell development

Introduction

Prothymocytes colonizing the thymus are very rare cells. This small pool of thymic immigrants undergoes in the order of >100 fold (unpublished data) expansion before rearrangement of the TCR β chain. This expansion is mediated by two key growth/cytokine receptors expressed on prothymocytes, the tyrosine kinase c-kit and common cytokine receptor γ chain (γc)–IL-7Rα chain complexes. The ligands binding to these receptors, stem cell factor (SCF) and IL-7, are provided by the epithelial thymic stroma (for reviews, see references 1–3). In mutants lacking c-kit and γc together, but not in single c-kit– or γc– deficient mice, the thymus is alymphoid, and TCR-β, -γ, and -δ rearrangements are essentially undetectable (4, 5). Thus, c-kit and γc provide essential, synergistic signals in pro-T cells (for a review, see reference 2). Synergy between growth factor and cytokine signaling is a common theme in many hematopoietic cells, including stem cells (see, for example, reference 6). Due to the complete lack of thymocytes, c-kit and γc double mutant mice may provide a useful model in which individual signaling pathways can be selectively rescued in vivo. At present, the downstream signaling pathways of each receptor are, at least in primary thymocytes, only poorly defined.

Expression of the prototypic antiapoptotic protein Bcl-2 is developmentally regulated in thymocytes. Bcl-2 expression is high in immature CD3−CD4−CD8+ (triple negative [TN]) and mature CD3+CD4+CD8− or CD3+CD4−CD8+ (single positive [SP]), and is downmodulated in intermediate CD3lowCD4+CD8+ (double positive [DP]) thymocytes (7, 8). Bcl-2 expression is reduced in thymocytes from mice lacking IL-7Rα (9), γc (8), or IL-7 (10), and is upregulated after exposure of thymocytes from IL-7–deficient mice to IL-7 in vitro (1, 10). Hence, signaling via IL-7R/γc can upregulate Bcl-2 expression in thymocytes. Thymocyte development in bcl-2–deficient mice is normal until early adult life (~4 wk of age) followed by a strongly increased sensitivity to apoptotic stimuli affecting, among other cell types, all stages of thymocytes and peripheral lymphocytes (11–13). Similarly, fetal liver–derived T cell
development is more Bcl-2 independent than bone marrow–derived T cell development (13). Collectively, Bcl-2 can act as an antiapoptotic factor in thymocytes, in particular during adult life. As early T cell development is cytokine/growth factor dependent, it is of interest whether growth factor withdrawal–induced thymocyte death can be prevented by Bcl-2 expression. There are conflicting data on the role of Bcl-2 for the rescue of T cell development in the absence of IL-7Rα/γc signaling. Transgene-driven overexpression of Bcl-2 was reported to rescue T cell development in IL-7Rα chain (9, 14) or γc (8, 15) mutant mice (for a review, see reference 1). In contrast, other groups failed to observe a Bcl-2–mediated rescue of γc-deficient thymocytes (16, 17; for a review, see reference 2).

SCF stimulation can provide antiapoptotic signals in hematopoietic cells. This survival signal may be relayed via activation of phosphatidylinositol 3 (PI-3) kinase and Akt kinase (for a review, see reference 18), but c-kit–induced survival may also involve Bcl-2 expression (19). Given that c-kit−γc− mice are completely devoid of thymocytes, we reasoned that these mutants would be a very sensitive model to reveal potential, even minor, effects of Bcl-2 downstream of c-kit and γc. In this paper, we show that the bcl-2 transgene was functionally active, and efficiently promoted survival of thymocytes, including the earliest T cell progenitors ex vivo. In vivo, however, transgenic bcl-2 expression failed to release T cell progenitors, both in terms of phenotype and cell numbers, from their developmental block in all mutants tested. As our data, using postnatal day 5 old mutant mice, did not confirm other reports analyzing adult γc-deficient bcl-2 transgenic mice (1, 8, 15), we extended our studies to large numbers of adult γc-deficient mice with or without the bcl-2 transgene. Notably, thymus cell numbers in adult γc− mutants without the bcl-2 transgene showed considerable heterogeneity (ranging over two logs), an observation pointing at a possible role for modifier genes in the genetic background of the γc mutants. In line with our analysis of neonatal γc− mutants, enforced expression of Bcl-2 also failed to rescue T cell development in adult γc− mice. Our study, therefore, challenges the current view that enforced expression of Bcl-2 can rescue defects in thymocyte development caused by lack of γc, and suggests that gene products other than Bcl-2 might modulate signaling pathways linked to cytokine (IL-7) and growth factor (SCF) receptors in early thymocyte development in vivo.

Materials and Methods

Mice. WB-kit+/+ (20) (SLC), γc− (21), and Eμ-bcl-2-25 (22) (The Jackson Laboratory) were maintained under specific pathogen free-conditions. C-kit and γc single and double mutant mice were bred as described (4), and crossed to Eμ-bcl-2-25 mice. Experiments involving c-kit mutant mice were performed on postnatal day 5 old mice, as white spotting locus (W/W) mice die within 10 d after birth (23). W mutant mice were phenotyped by white skin color. Experiments on γc mutants were done either on postnatal day 5, or on adult 6–12-wk-old mice. γc mutant mice were genotyped by PCR on genomic DNA as described (4). Eμ-bcl-2-25 mice (22) were genotyped by PCR on genomic DNA using oligonucleotides amplifying the human bcl-2 gene (5′ oligo: 5′-GGTCATGTGTGTGGAGAGGTTCA-3′ linked to the SV40 polyA site (3′ oligo: 5′-GTTTCAAGGAAAGCCTA-3′) yielding a diagnostic DNA fragment of ∼1 kb. In addition, in some experiments, thymocytes were typed by Western blotting using anti–human Bcl-2–specific antibodies (Santa Cruz Biotechnology, Inc.).

Monoclonal Antibodies and Flow Cytometry. The following monoclonal antibodies were used for flow cytometric analyses: Anti–“lineage” antibodies were 2C11 (CD3ε), H129.19 (CD4), 53-6.7 (CD8α) (both from GIBCO BRL), M1/70.15 (Mac-1; Caltag), Gr-1 (granulocytes), TER119 (red blood cells), DX-5 (NK lineage), B220 (B cells; all PE-labeled), biotinylated ACK-4 (c-kit), FITC-labeled 3C7 (CD25), Cy-Chrome-labeled Pgp-1 (CD4; all from BD PharMingen, unless otherwise indicated); streptavidin-allophycocyanin (APC) (Molecular Probes) was used as second step reagent. For FACS® analysis, thymocytes were stained with monoclonal antibodies at 5–10 μg/ml in PBS/5% FCS for 15–30 min on ice and washed once in PBS/5% FCS. Cell sorting and FACS® analysis were done on FACSStar® and FASCALibur™, respectively (Becton Dickinson). Fluorescence data are displayed as dot plots using CELLQuest™ software (Becton Dickinson).

Ex Vivo Survival Assay. Total or cell sorter purified thymocyte subpopulations were placed in normal cell culture medium (DMEM plus 10% FCS plus 1 mM pyruvate plus 2 mM glutamine plus 2×10−5 M β-mercaptoethanol) and cultured 5×106 cells/ml in PBS/5% FCS for 15–30 min on ice and washed once in PBS/5% FCS. Cell sorting and FACS® analysis were done on FACStar® and FACS Calibur™, respectively (Becton Dickinson). The Eμ-bcl-2-25 transgene was functional in these growth factor receptor mutants without the bcl-2 transgene ex vivo. To analyze whether enforced expression of the bcl-2 transgene could promote survival of thymocyte populations in c-kit or γc mutant mice, the Eμ-bcl-2-25 mouse line was crossed to c-kit− and γc− single and double mutants. In the Eμ-bcl-2-25 mouse line, transgenic expression is T lineage specific (22) with an early onset of transgene expression at the CD3ε−CD4ε−CD8ε− stage of development (24). Functional transgenic Bcl-2 expression can be monitored by extended survival times of primary thymocytes isolated ex vivo (24). To test whether the Eμ-bcl-2-25 transgene was functional in these growth factor receptor mutants, total thymocytes from each strain were cultured in normal cell culture medium, and, at various times after initiation of the culture, numbers of viable cells were determined (Fig. 1, A–C). The results from γc− and c-kit wild-type mice demonstrated, as expected, that cell death of transgenic thymocytes was delayed by about 2 d when compared with nontransgenic cells cultured in parallel (Fig. 1 A). Bcl-2–transduced mice were assayed in vivo survival of thymocytes from γc− (Fig. 1 B) or c-kit− (Fig. 1 C) single mutant mice. Due to the complete absence of thymocytes (see below), the corresponding analysis could not be performed in c-kit−γc− mice.
progenitors in c-kit and stages of T cell development defined by the phenotypes analysis could not be performed on c-kit previous analysis of the E c-kit TN 1 cells (24) but c-kit expression was not used to iden-
thymocytes in the combined absence of c-kit and (15) reported that the E c-kit TN 1 subset. Kondo et al. TN 1 (Fig. 1 D) and c-kit bcl-2 transgene clearly promoted survival of both c-kit of cell sorter–purified progenitors from wild-type and E TN 2 cells by comparison of the ex vivo survival potential /H11001
bcl-2 transgene (E C), or cell sorter–purified progenitor subsets (CD3 CD25 CD44 CD44 CD44 CD44 CD25 CD44) from 5-d-old c-kit (25). Therefore, we analyzed mice were compared for their survival capacity ex vivo. Each symbol represents an individual mouse for each genotype. Thy
mice. C-kit and γc play essential roles at the most immature stages of T cell development defined by the phenotypes CD25 CD44+ (TN 1) and CD25 CD44+ (TN 2) (4). In a previous analysis of the E bcl-2 TN 1 cells (24) but c-kit expression was not used to iden-
ify pro-T cell within TN 1. The TN 1 population contains c-kit+ cells and c-kit− cells, and the pro-T cell activi
ty resides only in the c-kit+ fraction (25). Therefore, we analyzed bcl-2 transgene function in c-kit+ TN 1 as well as in c-kit+ TN 2 cells by comparison of the ex vivo survival potential of cell sorter–purified progenitors from wild-type and E bcl-2 transgenic mice. The results demonstrate that this bcl-2 transgene clearly promoted survival of both c-kit+ TN 1 (Fig. 1 D) and c-kit+ TN 2 (Fig. 1 E) cells in vitro. Thus, the E bcl-2 TN 2 cells, and extend functional bcl-2 expression to the c-kit+ (pro-T cell fraction) within the TN 1 subset. Kondo et al. (15) reported that the E bcl-2 transgene was not expressed in the earliest thymic precursors defined as CD3 CD25− c-kit+ (which would include c-kit+TN 1 cells). However, our experiments clearly show that this is not the case, as both c-kit+ TN 1 and c-kit+ TN 2 cells from E bcl-2 were markedly protected from growth factor withdrawal–induced apoptosis. Collectively, the E bcl-2 transgene is functional at exactly those stages of development which are affected by lack of SCF and IL-7.

Enforced Expression of Bcl-2 Does Not Normalize Thymus Cellularity in Postnatal c-kit and γc Single and Double Mutant Mice In Vivo. Given that the bcl-2 transgene was functional in the relevant progenitor subsets, we next analyzed the effect of the transgene on thymus cellularity of various growth factor receptor mutants. Mice of eight different genotypes were generated and analyzed: (a) wild-type mice with (c-kit+γc+bcl) or (b) without (c-kit+γc+bcl−) the transgene, (c) mice deficient for c-kit with (c-kit−γc+bcl+), or (d) without (c-kit−γc+bcl−) the transgene, (e) mice defi-
cient for γc with (c-kit+γc−bcl+), or (f) without (c-kit+γc−bcl−) the transgene, and, finally, mice deficient for both c-kit and γc with (g) (c-kit−γc−bcl+) or (h) without (c-kit−γc−bcl−) the transgene. In Fig. 2, thymocyte numbers are plotted for all individual mice of the indicated genotypes. In total, 121 mice were analyzed (numbers of mice for each group are indicated in Fig. 2). As null mutations in c-kit (c-kit/) are lethal within 10 d after birth, these analyses were done on day 5.

Thymocyte numbers are reduced ~15-fold in γc single and ~6-fold in c-kit single mutants when compared with wild-type mice (4). As shown in Fig. 2, and as reported previously (22), the bcl-2 transgene did not increase the total cell number in wild-type mice (Fig. 2 A). Notably, there was no significant (for the statistical evaluation, see the legend to Fig. 2) effect of the bcl-2 transgene on thymocyte cellularity in γc single (Fig. 2 B) and c-kit single (Fig. 2 C) mutants. Thus, enforced expression of bcl-2 does not rescue overall thymic cellularity caused by lack of γc or c-kit. The thymus anlage in c-kit− mice, lacking lymphoid cells, is composed of disorganized thymic epi-
thelium (26) and thymic dendritic cells, the latter cell type developing independently of these growth factor receptors and any detectable pro-T cell compartment (27). Given that thymocytes are undetectable in the double mutant thymus, this “null base line mutant” should be very sensi-
tive to uncover even marginal effects of the bcl-2 trans-
gene, i.e., even a partial Bcl-2–mediated rescue should be evident by the reappearance of thymocytes. However, even in bcl-2 transgenic c-kit−γc− mice, thymocytes failed to appear.

In the Adult Thymus, the Developmental Block Caused by γc Deficiency Is Not Rescued by Transgenic Bcl-2. Given that c-kit null mutations are lethal by postnatal day 10, all experiments involving c-kit single and double mutants, including the relevant control mice (Fig. 2), were performed on 5-d-
old mice. While these experiments were underway, other investigators reported that bcl-2 transgenes, when bred onto γc-deficient mice, rescued T lymphopoiesis, but not B or NK cell development (15). Similarly, bcl-2 transgenes were

Figure 1. The bcl-2 transgene (E bcl-2) is functional in T cell pro-
genitors in c-kit and γc mutant thymocytes. Nontransgenic (open symbols) or E bcl-2 transgenic (filled symbols) total thymocytes from 5-d-old c-kit−γc− (A) or c-kit+γc− (B), or c-kit−γc+ (C), or cell sorter–purified progenitor subsets (CD3 CD25 CD44−[TN]) c-kit+CD25 CD44+ (TN 1) (D), or c-kit−CD25 CD44+ (TN 2) (E) from 5-d-old c-kit+γc− mice were compared for their survival capacity ex vivo. Each symbol represents an individual mouse for each genotype. Thy
mice.
transgenic mice (Wilcoxon rank-sum test: A: transgenic vs. nontransgenic, rank-sum normal statistic with correction Z = −1.0839, P value = 0.2784, alternative hypothesis: true μ is not equal to 0; in D, no statistical comparison was done because thymocytes were undetectable in both groups of mice). Thymocytes were undetectable in c-kit− mice regardless of the bcl-2 transgene.

Figure 2. Failure of transgenic Bcl-2 to Release γ-deficient Postnatal or Adult Pre-T Cells from Their Phenotypic Block at the CD25+CD44+/− Stage. A genetic rescue of a developmental block in the thymus should be evident by increased cell numbers, and by a release from the phenotypic developmental block. For instance, introduction of a TCR- chain into SCID or recombination activating gene (RAG) mutant mice releases thymocytes from their block in differentiation at the TN 3 stage, and promotes the development of TN 4 cells and large numbers of CD4+CD8+ thymocytes (for a review, see reference 29). In mice lacking γc or IL-7Rα or IL-7, TN thymocyte development is severely affected. The transition from TN 2 (CD44+CD25+) to TN 3 (CD44−CD25+) stages is (incompletely) blocked causing increased percentages of TN 2 and decreased percentages of TN 3 cells (30, 31). As a result of this block, TN 4 cells and large numbers of CD44−CD8+ thymocytes were analyzed by flow cytometry for expression of CD44 versus CD25 (Fig. 4). This analysis included postnatal 5-d-old (Fig. 4, A–D) and adult (Fig. 4, D–F) mice which had a wild-type (Fig. 4, A and D), γ-deficient (Fig. 4, B and E), or γ-deficient bcl-2 transgenic (Fig. 4, C and F) genotype. In agreement with previous re-

also reported to rescue thymocytes cellularity in IL-7Rα−
deficient mice (9, 14). The discrepancy between lack of rescue of the γc deficiency phenotype, in our hands, and rescue by bcl-2 transgenes, reported by others, prompted us to analyze whether a Bcl-2 effect was only evident in adult mice, but not in postnatal mice. Therefore, the impact of the bcl-2 transgene on thymocyte cellularity in adult γc− mice was analyzed. In Fig. 3, thymus cellularity is compared between γc− and γc−bcl− mice. When thymus cell numbers were analyzed in large numbers (n = 25) of γc− mice, we surprisingly observed a wide range of cell numbers spreading over more than two logs. This data set suggests that γc− mice fall into three groups: (a) mice with a relatively mild phenotype (>107 thymocytes), (b) mice with an intermediate phenotype (between 106 and 107 thymocytes), and (c) mice with the most severe phenotype (between 105 and 106 thymocytes). Of note, a similar distribution was found in γc−bcl− (n = 34) mice. As these mice, but also mice examined by Kondo et al. (15), were on mixed genetic backgrounds (here WB, C57Bl/6, Balb/c, and 129/ola), it is possible that the penetrance of the γc mutation is subject to modifier genes, reminiscent of phenotypic heterogeneity in IL-7R mutants (28). Nevertheless, comparing γc− versus γc−bcl− mice, both range and mean values were overlapping and not significantly different. Thus, in our hands, a functionally active bcl-2 transgene does not rescue thymus cellularity in adult γc− mice.
portions (30, 31), data shown in Fig. 4 confirmed that the transition from TN 2 to TN 3 was inhibited, and that TN 4 cells were almost absent in postnatal (γ−: 16% TN 2, 63% TN 3, and 14% TN 4, versus γ−: 42% TN 2, 36% TN 3, and 1% TN 4), and in adult (γ−: 11% TN 2, 63% TN 3, and 21% TN 4, versus γ−: 40% TN 2, 57% TN 3, and 1% TN 4) γ− mice. A release from the "γ block" should be detectable by decreased percentages of TN 2, increased percentages of TN 3, and, most visibly, by rescue of the lacking TN 4 compartment. Notably, the pattern of TN subsets was unaltered comparing γ− and γ−bcl+ mice. This holds true for both young (Fig. 4, compare B versus C) and adult (Fig. 4, compare E versus F) mice.

In addition to the overall reduction in thymocyte cellularity, cell numbers of lineage marker-negative thymocytes are also reduced in γ− mice (for a review, see reference 2). To analyze lin− cell numbers in these mutant mice, CD4+CD8− (DN) thymocytes were prepared (by αCD4 and αCD8 antibody binding and C′ lysis), and the resulting populations were stained with the lineage-mix antibodies (see above). In postnatal γ− mice, the DN population contained 43% lin− cells (which split into TN 1 to 4 subsets as shown in Fig. 4). In contrast, in postnatal γ− mice, the DN population contained only 27% lin− cells, demonstrating an overall reduction in thymic lin− cellularity of ∼30-fold (15-fold reduction in overall thymocyte cell numbers and 2-fold reduction of lin− cells). The proportion of thymic lin− cells among DN thymocytes was strongly reduced in adult mice compared with young mice (young γ− mice: 43% lin− cells; adult γ− mice: 7.8% lin− cells). Of note, the proportion of thymic lin− cells among DN thymocytes was not increased by Bcl-2 (postnatal mice: γ− = 43%; γ− = 27%; γ−bcl+ 25% lin− cells; adult mice: γ− = 7.8%; γ− = 4.4%; γ−bcl+ 3.1% lin− cells). Collectively, enforced expression of Bcl-2 failed to rescue the phenotypic block in γ− mice, and did not increase the lin− thymic progenitor pool. This finding is consistent with the observed inability of Bcl-2 to rescue overall thymus cell numbers.

In two previous reports suggesting that transgenic bcl-2 can rescue T cell development in γ− mice (8, 15), the phenotype of TN subsets, according to CD44 versus CD25 staining on lineage-negative thymocytes, had not been investigated. Kondo et al. analyzed progenitor populations
for expression of c-kit versus CD25 gated on CD3− cells, an analysis which may have obscured phenotype and distribution within TN subsets because many other nonprogenitor cells types such as NK cell or myeloid cells are included in the CD3− cells. Clearly, failure of a Bcl-2-mediated rescue of the strongest γc− phenotype, i.e., restoration of TN 4 cells, cannot be detected by this type of analysis. Phenotypic and quantitative analysis of the TN 1 to 4 subsets in γc− mice now demonstrates that enforced expression of Bcl-2 does not rescue the γc− mutant phenotype.

**Growth Factor- and Cytokine Receptor-mediated Signals Are Essential at the First Stages of Thymocyte Development: A Role for Bcl-2?** Thymocyte development can be separated into three major stages: (a) growth factor- and cytokine-driven expansion of rare colonizing T cell progenitors; (b) selection of thymocytes for further development on the basis of productive TCR β chain/pre-TCR complexes (β-selection); and (c) α/β TCR-based repertoire selection. In this first expansion, c-kit and γc act in synergy, i.e., each receptor mutant alone is permissive for T cell development but mice lacking both receptors lack all thymocytes. The molecular mechanisms underlying this signaling synergy in thymocytes are unknown. Clearly, one or both of these receptors may induce an intracellular antiapoptotic signal.

In this paper, we have analyzed bcl-2 transgenic young c-kit and γc single and double mutants, as well as adult γc mutants. Enforced expression of Bcl-2 (a) failed to increase thymocyte cell numbers, and (b) TN 2/TN 3 cells were not released from their block in differentiation. To account for possible variability within each group of mice, large numbers of animals were analyzed, both for young and adult mice. Consistent with our finding that Bcl-2 expression cannot overcome defects in c-kit− thymocytes, Domen and Weissman showed that development of mast cells, a c-kit−dependent lineage, was not rescued by expression of transgenic bcl-2 in viable c-kit− mutants (W641/W641) (6). In stem cells, Bcl-2 has been suggested to synergize with c-kit signaling in preventing cell death but here, as in early thymocytes (this study), Bcl-2 does not act downstream of c-kit (6).

The effect of bcl-2 transgenes on the thymus in γc− deficient mice, has been studied by us (this paper) and others (8, 15), and the results are clearly conflicting. Kondo et al., crossed the very same bcl-2 transgenic line (Ep−bcl-2−25) into γc− deficient mice, and reported partial rescue of thymocyte cell numbers (6.8 × 10^6 [γc−] vs. 2.8 × 10^7 [γc− Ep−bcl-2−25]). Although these authors observed almost no effect of the bcl-2 transgene on the precursor pool (defined as CD3− c-kit+), the increase in cell numbers was somewhat stronger in more mature thymocytes (defined as CD69+ cells). We cannot reproduce the findings by Kondo et al. on overall cellularity, and have no evidence that bcl-2 expression can release γc− deficient thymocytes from their phenotypic block (Fig. 4). This discrepancy is not easily explained. One caveat in this type of genetic rescue experiment by crossing mouse lines is based on the fact that γc deficiency appears to be sensitive to modifier genes, and mice used in our study, but also mice used by others, were on mixed genetic backgrounds (only three backcrosses to B6 [15], or two backcrosses to B6 or Balb/c [8]). As we observed a massive heterogeneity of cell numbers ranging over two logs which became evident only after analysis of large numbers of mice (Fig. 3), such heterogeneity might account for the observed discrepancies. A detailed analysis of the developmental block in γc− mutants with “mild” versus “intermediate” versus “severe” reduction in thymus cell numbers could give insights into the nature of any modifier gene(s). Enforced Bcl-2 expression may “tilt” death versus differentiation choices constantly towards survival along various intra- and extrathymic T cell differentiation stages, causing “nonspecific” accumulation of thymocytes. Nevertheless, we conclude that a Bcl-2–mediated survival signal is insufficient to substitute for the stage-specific signals provided by c-kit and γc in prothymocyte development.

We thank Drs. H.J. Fehling, U. Gravunder (Ulm), J.P. DiSanto (Paris), and H. Spits (Amsterdam) for discussions, and Dr. Louise Ryan (Dana-Farber Cancer Institute, Boston, MA) for statistical evaluations.

The Basel Institute for Immunology was founded and supported by F. Hoffmann-La Roche (Basel, Switzerland). H.R. Rodewald was supported by the Deutsche Forschungsgemeinschaft (SFB-497-BS), and by Landesforschungsschwerpunkt Baden-Württemberg.

**Submitted:** 26 February 2001  
**Revised:** 25 April 2001  
**Accepted:** 27 April 2001

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