The Caspase 8 Inhibitor c-FLIP_L Modulates T-Cell Receptor-Induced Proliferation but Not Activation-Induced Cell Death of Lymphocytes

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The caspase 8 inhibitor c-FLIP_L can act in vitro as a molecular switch between cell death and growth signals transmitted by the death receptor Fas (CD95). To elucidate its function in vivo, transgenic mice were generated that overexpress c-FLIP_L in the T-cell compartment (c-FLIP_L Tg mice). As anticipated, FasL-induced apoptosis was inhibited in T cells from the c-FLIP_L Tg mice. In contrast, activation-induced cell death of T cells in c-FLIP_L Tg mice was unaffected, suggesting that this deletion process can proceed in the absence of active caspase 8. Accordingly, c-FLIP_L Tg mice differed from Fas-deficient mice by showing no accumulation of B220⁺ CD4[−] CD8[−] T cells. However, stimulation of T lymphocytes with suboptimal doses of anti-CD3 or antigen revealed increased proliferative responses in T cells from c-FLIP_L Tg mice. Thus, a major role of c-FLIP_L in vivo is the modulation of T-cell proliferation by decreasing the T-cell receptor signaling threshold.

Death receptors belonging to the tumor necrosis factor receptor family (e.g., Fas/CD95, TNFR1, TRAIL-R1, TRAIL-R2, TRAMP/DR3, and DR6) play an important role in the regulation of lymphocyte homeostasis (35, 42, 44). Upon encounter of specific antigenic peptides presented by antigen-presenting cells, T cells become activated and enter the cell cycle. Concomitantly with proliferation, T cells differentiate into effector cells that are either cytolytic or able to provide help to B lymphocytes. Once the effector cells have successfully performed their prescribed function, the expanded pool of antigen-specific T-cell clones needs to be reduced to its original size. Both for the effector function of cytolytic T cells and in the elimination of expanded T cells, cell death signals transmitted by Fas/CD95 are of key importance (30). This is underscored by the observation that mice and humans lacking a functional Fas receptor or Fas ligand develop a lymphoproliferative disease and systemic autoimmunity accompanied by the production of autoantibodies (12, 40, 52, 59).

As with all death receptors, the prototypic death receptor Fas/CD95 contains within its cytoplasmic tail a 60-amino-acid death domain (DD) motif (35, 44). Upon activation of Fas by its ligand, the DD undergoes homotypic interaction with a DD in the adaptor protein FADD, which then recruits the initiator caspase 8 via their mutual N-terminal death effector domains (DED) (3). A high local concentration of caspase 8 zymogens is thought to facilitate self-processing and cleavage to the active enzyme (34). Activated caspase 8 then initiates apoptosis by cleavage of the downstream effector caspases 3, 6, and 7 (10).

A number of gammaherpesviruses and molluscipoxviruses encode a molecule termed FLIP (FLICE inhibitory protein) that can inhibit FasL-induced cell death (5, 53). v-FLIP resembles caspase 8 in containing two DED but lacks the enzymatic C-terminal portion. As such, v-FLIP can be recruited into the death-inducing signaling complex (DISC) of Fas, thereby competing with recruitment of caspase 8 to FADD. In this manner, v-FLIP may function to promote viral persistence and dissemination by inhibiting death receptor-mediated elimination of infected cells (56). A mammalian cellular homologue (c-FLIP) has been described that exists in at least two splice variants, c-FLIP_S and c-FLIP_L (15, 16, 20–22, 38, 47, 49). Like v-FLIP, the 26-kDa c-FLIP_S has two DED and functions in a similar manner to inhibit death receptor-induced apoptosis (22). The full-length 55-kDa form of c-FLIP (c-FLIP_L) shows overall structural homology to caspase 8. It contains two DED that interact with FADD but bears a mutation in the caspase-like domain that renders it enzymatically inactive. Following Fas ligation, both c-FLIP_L and caspase 8 are recruited into the DISC and are subsequently partially cleaved. The affinity for FADD of the c-FLIP_L/caspase 8 heterodimer appears to be considerably greater than that of the caspase 8 homodimer; therefore, the ratio of c-FLIP_L to caspase 8 is critical in determining cell fate after Fas ligation (24, 43).

c-FLIP_L is expressed in resting T cells, but after activation its expression is gradually decreased in an interleukin-2 (IL-2)-dependent manner (2, 39). Paralleling this, resting T cells are resistant to Fas-induced death, whereas cycling T cells, especially in the presence of exogenous IL-2, become highly sensitive to Fas-induced death, as occurs during activation-induced cell death (AICD) following restimulation of cycling T cells (39). This agrees with the view that AICD is mediated in part by functional interactions between Fas and FasL (1, 6, 9, 23, 37, 63). In view of its potent capacity to inhibit FasL-induced

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apoptosis in vitro (22, 61), c-FLIP_L is suggested to play a central role in the regulation of T-lymphocyte homeostasis in vivo.

In addition to inhibition of cell death, we recently observed that c-FLIP_L is also capable of binding Raf-1, which leads to the activation of the mitogen-activated protein kinase, ERK, and of binding TRAF-1 and -2, which activate NF- κ B. Overexpression of c-FLIP_L in Jurkat T cells increased ERK and NF- κ B activities upon T-cell receptor (TCR) ligation. Hence, c-FLIP_L not only has the capacity to block Fas-mediated cell death but can divert Fas signals toward pathways leading to cell growth and differentiation. To assess how these two aspects of c-FLIP_L function in primary T cells, murine c-FLIP_L was overexpressed transgenically. As expected, overexpression of c-FLIP_L in T cells decreased Fas-mediated cell death but surprisingly had no influence on AICD. Rather, c-FLIP_L overexpression appears to render primary T cells more sensitive to TCR-triggered proliferation.

MATERIALS AND METHODS

Antibodies and reagents. The following antibodies (Abs) were used for fluorescence-activated cell sorting (FACS) staining: fluorescein isothiocyanate (FITC)-anti-CD3, phycoerythrin (PE)- and CyChrome (CyChr)-anti-CD8, CyChr and antigen-presenting cell anti-CD4, PE-anti-V α 2, FITC-anti-V β 8, FITC-anti-V β 14, PE-anti- α TCR, PE-anti-heat-stable antigen, and biotinylated anti-Fas Jo2 (all purchased from Pharmingen, San Diego, Calif.) and PE-anti-Thy1 (Caltag Laboratories, South San Francisco, Calif.). Anti-FLAG polyclonal Ab (Zymed, South San Francisco, Calif.), anti-FLIP monoclonal Ab (MAb) Dave-2 (Alexis, Lausanne, Switzerland), and anti-FLIP polyclonal Ab (R & D Systems, Abingdon, England) were used for Western blotting. Anti-CD3 (145-2C11), anti-CD28 (37.51), and anti-IL-2R MAb (PC66) were purified from hybridoma supernatants and used in T-cell proliferation assays. Recombinant human FasL, staurosporine, and z-VAD-fmk were obtained from Alexis. Bromodeoxyuridine (BrdU) was purchased from Sigma (St. Louis, Mo.), and *Staphylococcus enterotoxin B* (SEB) was from Toxin Technology (Sarasota, Fla.).

Generation of c-FLIP_L transgenic mice. FLAG-tagged mouse FLIP_L cDNA was inserted into a target vector containing the β -globin promoter and a downstream human CD2 locus enhancer element (11). The resulting construct was injected into BALB/c $\times$ C57BL/6 single-cell-stage embryos, and transgenic founders were screened by PCR of tail DNA with the following primers: JT766 (5'-GGAGCCAGGGCTGGGCATAAAA-3') and JT767 (5'-GACTCACCTGAAGTCTCAGGATCC-3'). Western blotting using anti-FLIP MAb (Dave-2) and anti-FLAG Ab (Zymed) further confirmed expression of the transgene. The c-FLIP_L Tg mouse strain was maintained by serially mating heterozygous transgenic animals with wild-type C57BL/6 mice, and the Tg mice used for the indicated experiments were backcrossed at least six times. For certain experiments c-FLIP_L Tg mice were crossed with DBA/2 mice, expressing the endogenous superantigens *Mtv* 1, 6, 7, 8, 11, 13, 14, and 17 together with I-E major histocompatibility complex (MHC) class II, or with OT-1 Tg mice (C57BL/6 background) expressing a transgenic TCR specific for ovalbumin (Ova) peptide 257-264 (SIINFEKL) in the context of H-2K^b (18). Expression of the transgenic TCR was determined by FACS analysis of peripheral blood with CyChr anti-CD8 and PE anti-V α 2 Abs. For most experiments, mice were used between 6 and 12 weeks of age.

Purification and activation of T cells. Single-cell suspensions of pooled mesenteric, inguinal, and axillary lymph nodes were depleted of B cells and monocytes by incubation with anti-B220 and anti-MHC class II immunoglobulin M Abs and rabbit complement (Saxon Europe, Cambridge, United Kingdom) for 45 min at 37°C. The remaining viable cells were collected by Ficoll-Hypaque density centrifugation. T-cell purity was 95% as confirmed by FACS analysis with FITC-labeled anti-CD3. Lymph node T cells (1.5×10^5 /well) were stimulated with graded amounts of plate-bound anti-CD3 (145-2C11) in the absence or presence of recombinant mouse IL-2 (50 U/ml; Roche Diagnostics, Rotkreutz, Switzerland). Cells were cultured in RPMI 1640 supplemented with 10% heat-inactivated fetal calf serum, 50 μ M β -mercaptoethanol, and antibiotics (Life Technologies, Paisley, Scotland) for 4 or 6 days. To measure proliferation, 0.5 μ Ci of [³H]thymidine (Amersham, Little Chalfont, Buckinghamshire, England) was added during the last 16 h of culture. IL-2 production was measured by

enzyme-linked immunosorbent assay in culture supernatants 24 h after activation with anti-CD3. To assess AICD, purified T cells were stimulated with 0.1 μ g of anti-CD3/ml and IL-2 (20 U/ml). After 3 days cells were collected, extensively washed, and put back into culture in the presence of IL-2 (50 U/ml). After 24 h viable cells were recovered by Ficoll-Hypaque density centrifugation and subsequently restimulated with 5 μ g of plate-bound anti-CD3/ml for 24 h. The percentage of viable cells was determined by annexin V staining and FACS analysis.

In case of peptide stimulation, total splenocytes (1.5×10^6 /well) of single OT-1 Tg mice or double (c-FLIP_L Tg $\times$ OT-1) Tg mice were stimulated with various concentrations of Ova peptide 257-264 (SIINFEKL) for 4 days. During the final 4 h of culture 0.5 μ Ci of [³H]thymidine was added.

Western blot analysis. Cells were washed once in ice-cold phosphate-buffered saline (PBS) and solubilized in lysis buffer (1% NP-40, 50 mM Tris-HCl [pH 7.6], 150 mM NaCl, 2 mM dithiothreitol, protease inhibitor cocktail [Complete; Boehringer Mannheim, Indianapolis, Ind.]). Postnuclear lysates were collected after centrifugation (15,000 $\times$ g), and proteins (60 μ g) were separated in sodium dodecyl sulfate-12% polyacrylamide gels. Proteins were transferred to nitrocellulose (Hybond-ECL; Amersham), and blots were blocked and probed with the indicated Abs in 5% nonfat milk in PBS-Tween 20 (0.1%). Immunoreactive proteins were visualized with horseradish peroxidase-labeled conjugates (Jackson Laboratories, West Grove, Pa.) and ECL blotting substrate (Amersham).

Flow cytometry. Cells (5×10^5) were washed with ice-cold FACS buffer (PBS, 3% fetal calf serum, 0.02% sodium azide) and incubated with saturating amounts of conjugated Abs or isotype-matched controls for 30 min at 4°C. After two washes with FACS buffer, PE-labeled streptavidin was added to samples that had been incubated with biotinylated Abs. After washing, 10^5 viable cells were analyzed on a FACScan apparatus (Becton Dickinson, San Jose, Calif.) with the CellQuest program. Phosphatidylserine exposure on apoptotic cells was measured by washing cells once in ice-cold HEPES buffer (10 mM HEPES, 150 mM NaCl, 5 mM KCl, 1 mM MgCl₂, and 1.8 mM CaCl₂, pH 7.4) supplemented with 1 mg of glucose/ml and 0.5% bovine serum albumin (BSA) and subsequent incubation with FITC-labeled annexin V (2.5 μ g/ml). Cells were incubated for 20 min at 4°C and washed twice with HEPES buffer. Before analysis on a FACScan apparatus, propidium iodide (PI) was added (final concentration, 5 μ g/ml) to the samples to discriminate necrotic cells (annexin V⁻, PI⁺) from apoptotic cells (annexin V⁺, PI⁻, annexin V⁺, PI⁺).

Detection of apoptosis by TUNEL and cell cycling by BrdU. To analyze apoptotic cells by flow cytometry, the terminal deoxynucleotidyltransferase-mediated dUTP-biotin nick end labeling (TUNEL) assay was used as described previously (8, 41, 46). Thymocytes were cultured at 37°C for 4 h to reveal apoptotic cells. Cells (0.75×10^6) were stained for TCR β , CD4, and CD8 expression by using a MAb in PBS containing 1% BSA for 30 min at 4°C and then washed with cold PBS. The cells were fixed with 1% methanol-free formaldehyde for 15 min on ice, washed with cold PBS, and pelleted by spinning at 10,000 $\times$ g. The thymocytes were resuspended in cold 70% ethanol for 15 min and then washed twice with cold PBS. For the TUNEL reaction, the cells were incubated in TUNEL reaction mix containing 10 U of terminal deoxynucleosyltransferase, 10 mM dUTP-biotin, and 2.5 mM cobalt chloride in 1 $\times$ terminal transferase reaction buffer (Boehringer Mannheim) for 1 h at 37°C. The samples were washed twice with PBS-1% BSA and incubated with streptavidin tricolor (Caltag Laboratories) for 20 min on ice. Cells were washed twice, fixed in 1% methanol-free formaldehyde in PBS-1% BSA, and stored at 4°C until analysis. Cells stained with the same protocol in the absence of dUTP-biotin were used as negative controls.

To determine proliferation of thymocyte subsets, mice received 4 intraperitoneal (i.p.) injections of 1 mg of BrdU (Sigma) over a 24-h period with the last injection given 1 h prior to tissue harvest. Single-cell suspensions were stained for TCR β , CD4, and CD8 expression with a MAb in PBS containing 1% BSA for 30 min at 4°C, washed with cold PBS, and fixed with 70% ethanol for 30 min on ice. By a modification of published methods (54), the thymocytes were washed, fixed with 1% methanol-free formaldehyde for 15 min on ice, washed, and fixed overnight in PBS containing 1% methanol-free formaldehyde and 0.01% Tween 20. The following day, the samples were washed and incubated with 50 U of DNase I (Sigma)/ml in 0.15 M NaCl-4.2 mM MgCl₂ (pH 5) for 15 min at 37°C. After two washes with PBS-1% BSA, the cells were incubated with anti-BrdU-FITC (Becton Dickinson) for 30 min on ice. The samples were washed twice with PBS-1% BSA, fixed in 1% methanol-free formaldehyde in PBS-1% BSA, and stored at 4°C until analysis. Samples were analyzed on a FACScalibur apparatus with the CellQuest program.

In vivo experiments. SEB (100 μ g) was injected into each of eight c-FLIP_L Tg mice and eight C57BL/6 control littermates. Blood was collected from tail vein on the indicated days, and peripheral T-cell subsets were analyzed. Cells were stained for V β 8, V β 14, CD4, and CD8, and the percentages of the responsive

V β 8⁺ T cells and nonresponsive V β 14⁺ T cells were determined within the CD4⁺ and CD8⁺ subsets. The mean percentage $\pm$ standard deviation of eight different animals per group was calculated.

To determine the in vivo proliferative responses of T cells in c-FLIP_L Tg mice, c-FLIP_L Tg $\times$ OT-1 double Tg mice (four mice per group) and OT-1 Tg control littermates (four mice per group) received i.p. injections of 2 mg of BrdU and 250 μ l of 100 μ M Ova peptide or the same volume of PBS and BrdU. BrdU (1 mg/ml) and 5% glucose were added to the drinking water, and 36 h after injection mice were sacrificed and spleens were removed. Single-cell suspensions were surface stained with PE-labeled anti-V α 2 and CyChr-labeled anti-CD8 followed by nuclear staining of BrdU-labeled DNA with FITC-labeled anti-BrdU (Pharmingen). Samples were analyzed on a FACScan apparatus, and the percentage of cells incorporating BrdU was determined within the CD8⁺ V α 2⁺ T-cell subset.

RESULTS

Generation of transgenic mice that constitutively express murine c-FLIP_L in T cells. To investigate the role of c-FLIP_L in T-cell development and activation in vivo, we generated transgenic mice expressing mouse c-FLIP_L under the control of a β -globin promoter and the downstream human CD2 enhancer element (11) (Fig. 1A). Two independent transgenic lines were obtained, both expressing high levels of transgenic protein (Fig. 1A). Transgenic mice expressed the transgene at levels approximately 5 (Tg 2)- to 10 (Tg 1)-fold higher than endogenous c-FLIP_L (Fig. 1A and B). One line (Tg 1), expressing slightly higher concentrations of transgenic FLIP, was chosen for detailed analysis, but similar results were also obtained with the second line. Whereas endogenous lymphoid expression of c-FLIP_L is mainly found in lymph node, with lowest levels in spleen and thymus (Fig. 1B), transgenic FLAG-tagged c-FLIP_L expression was highest in the thymus and lymph node, low in the spleen, and absent in bone marrow, consistent with the expected T-cell-directed expression of the human CD2 enhancer element (11) (Fig. 1B). Sorting of T and B cells from spleen confirmed the T-cell-specific expression of the transgene. Within T-cell compartments, the transgene was equally expressed in CD4⁺ and CD8⁺ T cells (Fig. 1C). Interestingly, in addition to its full-length 55-kDa form, an additional 43-kDa cleavage product of transgenic c-FLIP_L was observed. This fragment is usually generated by caspase 8 when the caspase 8/FLIP_L heterodimeric complex is recruited to death receptors (24, 43). Finally, whereas the expression of endogenous c-FLIP_L was only transiently induced after T-cell activation (Fig. 1D) (2, 39), transgenic c-FLIP_L was increased after T-cell stimulation and remained highly expressed in activated T cells (Fig. 1D).

Overexpression of c-FLIP_L protects T cells against FasL-induced apoptosis. In vitro, Fas-sensitive cell lines are protected against FasL-induced cell death when c-FLIP_L is overexpressed (22, 25). Primary T cells become sensitive to FasL-induced apoptosis 3 to 5 days after activation, which coincides with downregulation of c-FLIP_L protein levels (22, 28, 39) (Fig. 1D). Consistent with this, we found that activated T cells from control mice were rapidly killed after incubation with recombinant FasL, whereas the majority of day 3 T-cell blasts derived from c-FLIP_L Tg mice still expressed high levels of transgenic c-FLIP_L (Fig. 1D) and were protected against FasL-induced apoptosis (Fig. 2A). Similar to peripheral T cells, the high expression levels of c-FLIP_L in the thymi of Tg mice derived from both lines (Fig. 1A and B) were sufficient to inhibit

FasL-induced apoptosis of thymocytes (Fig. 2C). As opposed to T-cell blasts, resting T cells from either c-FLIP_L Tg or control mice were insensitive to killing by recombinant FasL (Fig. 2B). Resting and activated T cells from both strains of mice expressed similar levels of Fas on the cell surface (data not shown). In contrast to FasL-induced cell death, death receptor-independent apoptosis, as induced by staurosporine, was unaffected by c-FLIP_L transgene expression (Fig. 2D). Likewise, apoptosis induced by an established Bcl-2 family-regulated pathway (e.g., IL-2 withdrawal) (51) was also not inhibited by c-FLIP_L but appeared to be somewhat enhanced (Fig. 2E). These data show that the transgene-encoded c-FLIP_L is functional. However, as shown in Fig. 2A and F, protection against FasL-induced cell death in the c-FLIP_L Tg mice was not complete, whereas it was completely inhibited in Fas-deficient *lpr* mice (Fig. 2F). To investigate whether this difference was due to insufficient expression and caspase-inhibitory function of the c-FLIP_L transgene, z-VAD was added to cultures of activated T cells of c-FLIP_L Tg mice and normal littermate controls (NLC). z-VAD was found to protect T cells of NLC from recombinant FasL-induced death to a level similar to, but no better than, c-FLIP_L Tg T cells (Fig. 2A). Moreover, z-VAD did not significantly affect the remaining cell death in T cells derived from the c-FLIP_L Tg mice. These experiments indicate that the transgene-encoded c-FLIP_L levels are sufficient to block completely the Fas-triggered, caspase-mediated cell death pathway and that the remaining FasL-induced cell death in T cells is most likely caspase independent, in agreement with previous observations (17, 19). However, it is also possible that the lack of full inhibition of FasL-mediated death with z-VAD reflects incomplete permeability of the blocker or caspase-dependent cell death mediated by a z-VAD-insensitive caspase.

c-FLIP_L Tg mice have a decreased proportion of CD8⁺ T cells. Although T cells from c-FLIP_L Tg mice were significantly protected against Fas-induced cell death, c-FLIP_L Tg mice did not develop profound lymphadenopathy and age-dependent autoimmune disease characteristic of Fas-deficient *lpr* and FasL-deficient *gld* mice (7). The c-FLIP_L Tg mice appear healthy with normal cell counts for spleen and lymph nodes even at 12 to 18 months of age (Table 1 and data not shown). Consistent with these data, FACS analysis of mature lymphocyte populations from lymph node and spleen did not reveal an accumulation of B220⁺ CD4⁺ CD8⁺ T cells, characteristic of *lpr* and *gld* mice (Fig. 3). However, the c-FLIP_L Tg mice demonstrated a small reduction in the number of T cells in the spleen and lymph node compared to control mice, which was primarily confined to the CD8⁺ T-cell subset (Fig. 3B and Table 1).

Analysis of thymi from c-FLIP_L Tg mice did not reveal overt deviations in thymus size or thymocyte differentiation (Fig. 4A and data not shown). However, similar to peripheral T cells, we found a small reduction in the percentage of mature single-positive CD8 thymocytes. These are discriminated from immature single-positive CD8⁺ thymocytes by the expression of high surface levels of CD3 and TCR and decreased expression of Thy1 and heat-stable-antigen (Fig. 4A and data not shown). Since the upregulation of CD5, CD69, and Bcl-2 expression coincident with increasing surface TCR expression on c-FLIP_L Tg CD4⁺ CD8⁺ thymocytes was comparable to that for control

FIG. 1. Expression of the c-FLIP_L transgene. (A) FLAG-tagged mouse c-FLIP_L was cloned into a targeting vector, which is driven by the β -globin promoter and controlled by the human CD2 enhancer element. Protein levels of transgenic c-FLIP_L in thymocytes derived from two independent transgenic mice were compared. Numbers indicate the amount of total protein loaded per lane. (B) Western blot analysis of transgene protein expression in different immunological compartments. Cell lysates were prepared from single-cell suspensions of thymus, bone marrow, lymph nodes, and spleen of a transgenic mouse (Tg 1) and an NLC. The blots were probed with an anti-FLIP MAb that recognizes both endogenous and transgenic c-FLIP_L (Dave-2, top panel) and with an anti-FLAG Ab detecting only the transgenic form of c-FLIP_L (middle panel). Ponceau staining was included to show the presence of similar amounts of protein in each lane (bottom panel). (C) T and B cells were FACS sorted from single-cell suspensions of spleens, and purified CD4⁺ and CD8⁺ T cells were sorted from lymph nodes of c-FLIP_L Tg mice (line 1). Lysates were subjected to Western blotting with a polyclonal anti-FLIP Ab. (D) Lymph node cells of transgenic (line 1) and control littermate mice were stimulated with plastic-coated anti-CD3 MAb (1 μ g/ml). At the indicated time points cell lysates were prepared and subjected to Western blotting. The blot was probed with an anti-FLIP MAb. In order to view differences in expression, less protein was loaded than in panel B.

FIG. 2. T cells from c-FLIP_L Tg mice are resistant to FasL-induced cell death. T cells purified from lymph nodes (A, B, and D to F) or thymocytes isolated from thymi (C) of c-FLIP_L Tg mice (Tg 1 and Tg 2), NLC, or *lpr* mice (F) were either directly incubated with recombinant FasL for 3 h (B and C) or stimulated with plastic-coated anti-CD3 (1 $\mu\text{g/ml}$) and recombinant mouse IL-2 (50 U/ml) (A and D to F). After 2 to 3 days T-cell blasts were harvested and depleted of dead cells by Ficoll gradient centrifugation. Viable cells were incubated with titrated amounts of recombinant FasL, either in the absence or in the presence of 50 μM z-VAD (A), with staurosporine (D), or with or without IL-2 (50 U/ml) (E). After 6 h of incubation the cells were stained with FITC-labeled annexin V and PI, and the percentage of apoptotic cells was determined by FACS analysis. In panel F activated T cells were exposed to 0.5 μg of FasL/ml for 3 h. The percentage of spontaneous cell death 6 h after incubation with medium only was 11% for controls and 23% for c-FLIP_L Tg T cells (in panels A and D). wt, wild type.

TABLE 1. Absolute cell numbers of CD3⁺ (T) and B220⁺ (B) cells and CD4⁺ and CD8⁺ T cells in lymph nodes and spleens of c-FLIP_L Tg mice (line 1) and NLC^a

Cell type	Value for:			
	Lymph node (10 ⁶)		Spleen (10 ⁶)	
	NLC (n = 7)	c-FLIP _L Tg (n = 7)	NLC (n = 7)	c-FLIP _L Tg (n = 7)
CD3 ⁺ cells	1.53 ± 0.43	1.45 ± 0.33	28.5 ± 1.15	18.6 ± 0.61
B220 ⁺ cells	0.67 ± 0.42	1.37 ± 0.43	58.1 ± 7.76	50.2 ± 0.52
Ratio T/B	2.30 ± 0.6	1.08 ± 0.2	0.50 ± 0.1	0.37 ± 0.1
CD3 ⁺ gate				
CD4 ⁺ T cells	0.86 ± 0.05	0.94 ± 0.06	19.3 ± 0.54	14.3 ± 0.9
CD8 ⁺ T cells	0.63 ± 0.04	0.44 ± 0.05	7.81 ± 0.63	3.16 ± 0.8
Ratio CD4/CD8	1.37 ± 0.3	2.14 ± 0.6	2.47 ± 0.35	4.52 ± 0.2

^a Spleens and inguinal lymph nodes of 8- to 12-week-old mice were removed, and lymphocytes were counted. The number of cells in each subset was calculated from flow cytometric analysis of two- and three-color immunofluorescence on 10,000 cells per sample. Values are the means ± standard deviations.

mice, defects in positive selection do not appear to explain the reduced percentage of single-positive CD8 thymocytes in c-FLIP_L Tg mice (data not shown). However, TUNEL analysis revealed an increased percentage of apoptotic cells within the CD4⁺ CD8⁺ double-positive and single-positive CD8 thymocyte subsets of c-FLIP_L Tg mice compared to the same subsets in control mice (Fig. 4B). This suggests that the reduced percentages of mature CD8⁺ thymocytes and peripheral CD8⁺ T cells in c-FLIP_L Tg mice are likely caused by an increased cell death turnover of single-positive CD8 thymocytes. The increased spontaneous thymocyte death was balanced by augmented proliferation based on in vivo BrdU labeling (Fig. 4C). This balance most likely accounts for the overall similar thymus size of c-FLIP_L Tg and control mice. Finally, when c-FLIP_L Tg mice were backcrossed onto the DBA/2 background, which expresses endogenous mammary tumor virus (*mtv*) superantigen as well as I-E MHC class II (31), deletion of the *mtv*-reactive Vβ3 and Vβ17 TCR-expressing thymocytes and T cells was unaffected (Fig. 4D and data not shown), demonstrating that c-FLIP_L does not affect negative selection.

Increased proliferation of T cells from c-FLIP_L Tg mice upon suboptimal triggering of the TCR/CD3 complex. In addition to its antiapoptotic function, we recently showed that c-FLIP_L can also promote IL-2 production in T-cell lines by connecting with the ERK and NF-κB pathways (24). To investigate this alternative function of c-FLIP_L in nontransformed primary T cells, proliferation and IL-2 production were measured in purified lymph node T cells stimulated with plate-bound anti-CD3. Interestingly, the dose-response curve for CD3 stimulation of c-FLIP_L Tg T cells was shifted toward a lower dose of anti-CD3. At high doses of anti-CD3, T cells from c-FLIP_L Tg mice actually proliferated less well than T cells derived from control mice (Fig. 5A). Since increased percentages of annexin V-positive cells (47% in NLC versus 65% in Tg 1 with 0.2 μg of anti-CD3/ml and 50% in NLC versus 74% in Tg 1 with 5 μg of anti-CD3/ml) were observed in the cultures of c-FLIP_L Tg T cells, enhanced cell death is the likely reason for the reduced proliferation rates at higher doses of anti-CD3.

Kinetics of the proliferative response did not appear to be

faster in the c-FLIP_L Tg mice at high doses of anti-CD3. Addition of [³H]thymidine 1, 2, and 3 days after stimulation revealed no early increase in proliferation in c-FLIP_L Tg mice (Fig. 5B). However, with the use of doses of anti-CD3 (0.02 and 0.04 μg/ml) at which T cells from control mice no longer proliferated, the c-FLIP_L Tg T cells still proliferated vigorously and produced significant levels of IL-2 (Fig. 5A, B, C, and E). The increased proliferative potential at low doses of anti-CD3 was observed in both CD4⁺ and CD8⁺ T cells of c-FLIP_L Tg mice, but the amplitude of the response was higher in CD4⁺ Tg T cells than in CD8⁺ Tg T cells (Fig. 5D). When TCR signaling was circumvented by direct activation of protein kinase C with phorbol myristate acetate and addition of the calcium ionophore ionomycin, T cells from the two types of mice showed similar proliferative responses (Fig. 5A). To examine whether the enhanced IL-2 production was responsible for the increased proliferation of c-FLIP_L Tg T cells, IL-2 was added to the cultures. In the presence of exogenous IL-2, T cells of control mice were able to proliferate at low doses of anti-CD3, and, more importantly, their proliferative response was similar to the response of FLIP_L Tg T cells (Fig. 5F). In line with this, addition of a blocking anti-IL-2R MAb to suboptimally triggered c-FLIP_L Tg T cells completely eliminated proliferation, showing that the increased proliferation rate of c-FLIP_L Tg T cells at low doses of anti-CD3 is IL-2 mediated (Fig. 5G).

Finally, to assess whether these in vitro observations could also be made in vivo, c-FLIP_L Tg mice were crossed with OT-1 mice expressing a transgenic TCR specific for the Ova peptide 257-264 restricted to MHC class I K^b. Cell cycling following injection of Ova peptide was monitored by BrdU incorporation. As shown in Fig. 6A, c-FLIP_L × OT-1 mice manifested enhanced BrdU incorporation in the responsive CD8⁺ Vα2⁺ T cells compared to the same subset in OT-1 control littermates. Furthermore, when splenocytes of OT-1 Tg mice and double (c-FLIP_L Tg × OT-1) Tg mice were stimulated in vitro with Ova peptide, we observed a similar effect as that with anti-CD3 stimulation: higher proliferative responses in the c-FLIP_L Tg mice at low doses of peptide and lower proliferative responses at high doses of peptide (Fig. 6B). Thus, T cells from c-FLIP_L Tg mice have increased proliferative potential both in vitro and in vivo.

AICD of T cells is not inhibited by overexpression of c-FLIP_L in vivo. Despite the increased proliferative potential of c-FLIP_L Tg T cells and their increased resistance to FasL-induced cell death, c-FLIP_L Tg mice did not accumulate vast numbers of T cells in secondary lymphoid organs. T-cell homeostasis after immune activation is complex and maintained both by deprivation of cytokines, which become limiting at the end of an immune response (30), and by AICD that is, at least in part, mediated by Fas-FasL interactions (6, 9, 23). In vitro, AICD is mimicked by restimulation of activated T cells via their antigen receptor (37). Surprisingly, T cells from c-FLIP_L Tg mice were not protected against this type of cell death, whereas FasL-deficient *gld* mice were partially protected (Fig. 7A), in agreement with previous results (14). Injection of the superantigen SEB in mice induces T-cell activation and expansion followed by deletion of the reactive Vβ8⁺ T cells (27). Injection of SEB resulted in increased percentages of SEB-reactive Vβ8⁺ T cells at day 2 followed by markedly decreased

FIG. 3. Flow cytometric analysis of peripheral lymphocytes in c-FLIP_L Tg mice. Lymph node cells from 8- to 12-week-old c-FLIP_L Tg mice and control littermates were stained with PE-B220 and FITC-CD3 MAb (A) or with FITC-CD3, PE-CD8, and CyChr-CD4 MAb (B). The percentages of B (B220⁺) and T (CD3⁺) cells and CD4⁺ and CD8⁺ T cells within the CD3⁺ subset are shown and are representative of seven independent experiments.

percentages at day 7 in both strains of mice (Fig. 7B). No effect was found in the nonresponsive V β 14 TCR-expressing T cells. Thus, both the in vitro and in vivo experiments imply that AICD is not affected by overexpression of c-FLIP_L.

DISCUSSION

The present findings show that constitutive expression of the caspase 8 inhibitor c-FLIP_L in primary mouse T cells not only

FIG. 4. Thymic development and negative selection in c-FLIP_L Tg mice. (A) Single-cell suspensions of thymi of 4-week-old c-FLIP_L Tg mice and control littermates were prepared and stained with FITC-CD3, CyChr-CD4, or PE-CD8 MAb. The percentages of double-negative, double-positive, and single-positive thymocytes of the total thymocyte population are shown and are representative of four independent experiments. (B) Thymi of 8- to 12-week-old Tg mice or NLC were isolated, and single-cell suspensions were surface stained for TCR β , CD4, and CD8. Spontaneous apoptosis in the different thymocyte subsets was determined by TUNEL assay. The experiment shown is one representative of three. (C) c-FLIP_L Tg mice and NLC received four i.p. injections of BrdU (1 mg/ml) over a 24-h period. Thymi were harvested, and single-cell suspensions were surface stained prior to staining with anti-BrdU. One representative experiment of three is shown. (D) c-FLIP_L Tg mice were backcrossed into a DBA/2 background that expresses the endogenous superantigens *mtv* 1, 6, 7, 8, 11, 13, 14, and 17 as well as I-E MHC class II. Deletion of V β 3 TCR-expressing T cells was monitored by staining total splenocytes of DBA/2 mice with V β 3-specific Abs followed by FACS analysis. Since C57BL/6 mice do not express I-E MHC class II molecules, V β 3-expressing T cells are present in the spleen. V β 8.2 T cells are not deleted by the above-mentioned *mtv* subtypes and are thus present both in C57BL/6 and in DBA/2 mice.

confers protection from FasL-induced cell death but also lowers the threshold for TCR-triggered proliferation and IL-2 production. The former is likely due to the ability of c-FLIP_L to bind caspase 8 and block its proapoptotic activity (55). The augmented T-cell activation may reflect the more recently discovered ability of c-FLIP to augment signaling via ERK and NF- κ B (24).

Despite the resistance of their T cells to FasL-induced death, the c-FLIP_L Tg mice did not accumulate B220⁺ CD4⁺ CD8⁺ T cells, typical of Fas-deficient *lpr* mice. The fact that AICD was not impaired in c-FLIP_L Tg mice probably explains why progression to T-cell hyperplasia does not occur in c-FLIP_L Tg mice. The unaffected AICD stands in contrast to findings of Van Parijs et al. (57) for mice reconstituted with bone marrow cells that had been retrovirally transduced with c-FLIP_L. This approach allowed the expression of c-FLIP_L in both T and B cells. In this study, both restimulation of activated T cells with anti-CD3 and injection of SEB resulted in impaired cell death responses in the FLIP-expressing mice. Similar to the c-FLIP_L Tg mice, however, the retrovirally transduced c-FLIP mice did not accumulate aberrant T cells but rather developed an autoimmune disease due to defects in the B-cell compartment.

Several reasons may explain why AICD was not impaired in c-FLIP_L Tg mice. First, T-cell deletion in response to superantigen stimulation is only mildly affected in *lpr* mice (45, 51). Therefore, this type of cell death most likely has a major death

receptor-independent component, which cannot be inhibited by c-FLIP_L. Second, we cannot formally exclude the possibility that the levels of c-FLIP_L in T cells from c-FLIP_L Tg mice are insufficient to block caspase 8 activity required for AICD in vivo, although it efficiently decreases FasL-mediated death in vitro. It is, however, more likely that AICD can occur in the absence of active caspase 8. Recently, Holler et al. (19) and Hildeman et al. (17) have demonstrated that AICD of human and murine T cells occurs in the presence of caspase inhibitors. Since some caspases are not readily blocked by the concentrations that can be achieved in cultured cells, these results do not completely rule out the involvement of caspases. However, FasL, TRAIL, and TNF- α were found to kill both primary and transformed T cells in the absence of caspase 8, inducing morphological changes that are reminiscent of necrosis. This type of cell death is transmitted via the DD-containing kinase RIP (19). Together with the classical caspase-dependent route, this caspase-independent cell death pathway may regulate T-cell homeostasis. The differences in the phenotypes of *lpr* mice and c-FLIP_L Tg mice may therefore be explained as follows. Whereas the loss-of-function mutation *lpr* affects both the caspase-dependent and -independent cell death pathways due to impaired recruitment of both FADD-caspase 8 and RIP to the DD of Fas, overexpression of c-FLIP_L inhibits only the caspase 8-dependent route and leaves the RIP-dependent cell death pathway intact. In line with this, we observed that the pancaspase inhibitor z-VAD inhibited FasL-induced cell death

FIG. 5. Enhanced in vitro proliferative responses in c-FLIP_L Tg mice. (A) Purified lymph node T cells from c-FLIP_L Tg mice and NLC were activated with the indicated concentrations of plate-bound anti-CD3 or phorbol myristate acetate (1 ng/ml) and ionomycin (0.5 μ M) for 6 days. [³H]thymidine (0.5 μ Ci/well) was added during the final 16 h of culture. (B) Purified lymph node T cells from c-FLIP_L Tg mice and NLC were activated with a high (0.2- μ g/ml) or suboptimal (0.04- μ g/ml) dose of plate-bound anti-CD3 for the indicated number of days, and [³H]thymidine (0.5 μ Ci/well) was added during the final 16 h of culture. (C) Purified lymph node T cells of NLC and Tg mice derived from the second c-FLIP_L Tg line (Tg 2) were stimulated with suboptimal doses of anti-CD3 for 4 days, and [³H]thymidine was added during the final 16 h of culture. (D) Total lymph node cells and FACS-sorted CD4⁺ and CD8⁺ T cells were stimulated with a suboptimal dose of plate-bound anti-CD3 (0.04 μ g/ml) for 4 days, and [³H]thymidine was added during the final 16 h of culture. (E) Purified lymph node T cells were stimulated with 0.04 μ g of plate-bound anti-CD3/ml, and IL-2 production was measured in 24-h culture supernatants. (F) Purified lymph node T cells were stimulated with 0.04 μ g of plate-bound anti-CD3/ml in the presence or absence of IL-2 for 4 days, and [³H]thymidine was added during the final 16 h of culture. (G) Purified lymph node T cells were stimulated for 4 days with a low dose of coated anti-CD3 (0.04 μ g/ml) in the presence or absence of a blocking anti-IL-2R MAb (10 μ g/ml). [³H]thymidine was added during the final 16 h of culture.

A **B**

FIG. 6. c-FLIP_L Tg mice show enhanced proliferative responses in vivo. (A) c-FLIP Tg mice were crossed with OT-1 mice that express an MHC class I-restricted transgenic TCR specific for Ova peptide 257-264. Four OT-1 c-FLIP Tg mice or four single OT-1 Tg littermates were injected with 2 mg of BrdU together with 100 μM Ova peptide or the carrier PBS. After 36 h, single-cell suspensions of spleens were prepared and stained with anti-CD8, anti- $V\alpha 2$, and anti-BrdU. BrdU expression within the responsive CD8⁺ $V\alpha 2$ ⁺ T cells is shown and is representative of four experiments. (B) Splenocytes derived from single OT-1 transgenic mice or double OT-1 × c-FLIP_L Tg mice were stimulated in vitro with the indicated amount of Ova peptide for 4 days. [³H]thymidine was added during the final 4 h of culture.

A **B**

FIG. 7. AICD is not inhibited in c-FLIP_L Tg mice. (A) Purified lymph node T cells of c-FLIP_L Tg and NLC mice (left panel) and of wild-type C57BL/6 (wt) and *gld* C57BL/6 mice (right panel) were stimulated with CD3 MAb (0.1 μg/ml) and IL-2 (20 U/ml) for 3 days. Cells were washed and put into culture in the presence of IL-2 (50 U/ml) only. After 24 h dead cells were removed by Ficoll gradient centrifugation and the preactivated T cells were restimulated with plate-bound CD3 MAb (5 μg/ml) for 24 h. Cell viability was determined by annexin V staining and FACS analysis. Means ± standard deviations of triplicate determinations are shown. (B) SEB (100 μg/animal) was injected into eight c-FLIP_L Tg mice and eight NLC, and peripheral blood T cells were analyzed on the indicated days. Cells were stained for Vβ8, Vβ14, CD4, and CD8, and the percentages of Vβ8⁺ and Vβ14⁺ within CD4⁺ (left) and CD8⁺ (right) T cells are shown.

in normal mice nearly to the same extent as in c-FLIP_L Tg mice. In addition, the remaining Fas-induced cell death in the c-FLIP_L Tg mice was not further blocked by addition of z-VAD. T-cell accumulation leading to hyperplasia is therefore expected to take place only in *lpr* and *gld* mice, not in c-FLIP_L Tg mice. Similarly, T cells from transgenic mice expressing the

caspase 8 inhibitor CrmA were protected from FasL-induced cell death in vitro, but these mice also did not develop a lymphoproliferative disease (48, 64). Finally, whereas in the CD2 enhancer-driven c-FLIP_L Tg mice Fas-mediated responses are affected only in T cells, in *lpr* and *gld* mice Fas and FasL are absent on other cell types, such as B cells, NK cells,

and macrophages. Impairment of Fas-induced death in these cell types may also contribute to the development of hyperproliferative disease in these mice.

Rather than the anticipated T-cell accumulation, we observed a reduction in the percentage of CD8⁺ T cells in the periphery of both young and old c-FLIP_L Tg mice. This reduction became apparent developmentally as early as the mature CD8 single-positive thymocyte stage. Despite the fact that thymocytes were resistant to FasL-induced death, studies of c-FLIP_L Tg thymi showed that neither positive nor negative selection was affected. However, both the double-positive and the CD8 single-positive thymocytes manifested increased cell cycling and, at the same time, were more prone to undergo spontaneous cell death, suggesting that increased (or noncontrolled) proliferation is kept in check by subsequent cell death. Therefore, the decrease in the number of peripheral CD8⁺ T cells is at least in part due to decreased survival of its thymocyte precursor. Furthermore, the observation that predominantly the CD8⁺ T cells from c-FLIP_L Tg mice are more prone to undergo apoptosis upon IL-2 withdrawal (unpublished observations) suggests also that the mature CD8⁺ T cells have a decreased survival potential. Why and how overexpression of c-FLIP_L decreases the survival of CD8⁺ T cells are unclear at this point.

A similar duality of proliferation and cell death was observed in activated peripheral T cells. Whereas the proliferative response of c-FLIP_L Tg T cells was markedly enhanced at suboptimal doses of anti-CD3, at optimal doses the c-FLIP_L Tg T cells had decreased proliferative rates due to increased cell death. This finding may be linked to our recent observation that c-FLIP_L can act as an adaptor molecule, associating with Raf-1 and TRAF-1/TRAFF-2, which activate ERK and NF-κB, respectively, important mediators of cell proliferation and survival (24). Interestingly, T cells derived from c-FLIP_L Tg mice were observed to have increased ERK phosphorylation and NF-κB activation after triggering of the TCR/CD3 complex, resulting in enhanced IL-2 production (24). It can thus be envisaged that at low doses of anti-CD3 the increased ERK and NF-κB activation may help to overcome the signaling threshold necessary for IL-2 production and T-cell proliferation. However, at high doses of anti-CD3, these same TCR signal pathways may provoke excessive proliferation and subsequent cell death. Consistent with this view, enhanced cell death has been observed with both augmented activation of ERK and sustained activation of NF-κB (4, 29, 32, 50).

Cross-linked recombinant FasL can costimulate proliferation and IL-2 production of primary human T cells, triggered with suboptimal doses of anti-CD3. Furthermore, addition of Fas-Fc as well as caspase inhibitors to cultures of anti-CD3-activated T cells was shown to inhibit their proliferation, implying that Fas-induced caspase activity can result in augmentation of T-cell proliferation (26). Since c-FLIP_L is expressed in resting T cells, activation of these cells might first result in FasL expression, and subsequent Fas-FasL interactions may lead to the recruitment of both caspase 8 and c-FLIP_L into the DISC. Caspase 8 is known to process c-FLIP_L between its p10 and p18 caspase-like subunits, resulting in a cleaved c-FLIP_L protein of 43 kDa (22, 43). Cleaved c-FLIP_L may now be able to activate ERK and NF-κB and thus contribute to proliferation. In this context, it is noteworthy that thymocytes or T cells

deficient in FADD or overexpressing a dominant negative form of FADD show highly impaired proliferation capacities (36, 58, 62). Since FADD is essential for c-FLIP_L recruitment to death receptors, it is tempting to speculate that c-FLIP_L-dependent ERK and NF-κB activation is required for T-cell growth. The costimulatory effect observed in the c-FLIP_L Tg mice, however, was already apparent without further triggering of the Fas receptor by exogenously added FasL. Interestingly, in both resting and activated T cells of these mice, a large proportion of transgenic c-FLIP_L is constitutively cleaved at its C terminus, giving rise to a product of 43 kDa. This cleavage is normally induced by death receptor-activated caspase 8 (43), but the high expression levels of c-FLIP_L in the transgenic mice may lead to spontaneous processing of c-FLIP_L that is sufficient to support CD3-triggered IL-2 production and proliferation in the absence of exogenous death receptor ligation. Alternatively, death ligands may constitutively activate death receptor-dependent signaling pathways in thymocytes and T cells. In line with this, high FasL expression has been detected in stromal cells of the thymus (13). The function of such signals, however, remains to be determined.

Collectively, our observations demonstrate that c-FLIP_L possesses the potential of modulating signals leading to both cell death and cell growth. These crucial functions of c-FLIP_L require tight control of c-FLIP_L expression levels. The c-FLIP_L protein is short lived (60), and its transcription is regulated by IL-2 in T cells (39). Since c-FLIP_L is also upregulated in various types of tumors (22), rendering tumor cells resistant to immune surveillance (33), the understanding of the regulation of c-FLIP_L will be of utmost importance.

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