In Vitro Suppression of Programmed Cell Death of B Cells by Tissue Inhibitor of Metalloproteinases-1

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Abstract

Cellular pathways for induction of programmed cell death (PCD) have been identified, but little is known about specific extracellular matrix processes that may affect apoptosis along those pathways. In this study, a series of Burkitt’s lymphoma (BL) cell lines were assayed for their expression of tissue inhibitor of metalloproteinases (TIMP)-1. Results indicate that TIMP-1–positive BL lines show resistance to cold-shock–induced apoptosis. Furthermore, recombinant TIMP-1, but not TIMP-2 or a synthetic metalloproteinase inhibitor (BB-94), confers resistance to apoptosis induced by both CD95-dependent and -independent (cold shock, serum deprivation, and γ-radiation) pathways in TIMP-1–negative BL lines. TIMP-1 suppression of PCD is not due to metalloproteinase inhibition, as reduction and alkylation of the TIMP-1 did not abolish this activity. Retroviral induction of TIMP-1 not only resulted in cell survival but also in continued DNA synthesis for up to 5 d in the absence of serum, while controls underwent apoptosis. This resistance to apoptosis is reversed by anti–TIMP-1 antibodies, demonstrating that secreted TIMP-1 is active in blocking apoptosis. Furthermore, TIMP-1 upregulation induced expression of Bcl-XL but not Bcl-2 as well as decreased NF-κB activity as compared with controls. These results demonstrate that TIMP-1 suppresses apoptosis in B cells and suggests a novel activity for TIMP-1 in tissue homeostasis. (J. Clin. Invest. 1998. 102:2002–2010.) Key words: Bcl-X • Bcl-2 • extracellular matrix • Burkitt’s lymphoma • metalloproteinases

Introduction

Programmed cell death (PCD),1 or apoptosis, is now recognized as playing a central role in development and tissue-specific functions in such diverse processes as regulation of immune responses, angiogenesis, and organogenesis (1–3).

In these processes, components of the extracellular matrix (ECM) (4) and in particular basement membrane act as survival factors and suppress apoptosis. Proteolytic modification of matrix organization or disruption of cell–matrix contacts can result in initiation of PCD in epithelial cells, with induction of specific molecular effectors of apoptosis such as caspases (3, 4). Recently, it was shown that proteolytic processing of TNF-α and FAS ligand (FasL) from the surface of lymphoid cells by matrix metalloproteinase-like activity can also alter PCD (5, 6). We have recently examined the expression of matrix metalloproteinases (MMPs) and their endogenous inhibitors, the tissue inhibitors of metalloproteinases (TIMPs), in human normal and neoplastic lymphoid cell lines (7). In this study, TIMP-1 secretion was shown only by normal activated tonsillar B cells and by a subset of high-grade Burkitt’s lymphomas but not by quiescent peripheral blood cells or follicular lymphoma cell lines. MMP expression was variable and did not correlate with TIMP-1 production. These results are in agreement with previous reports from our laboratory and others that TIMP-1 expression correlates with clinical grade in non–Hodgkin’s lymphomas as well as in other malignancies but did not correlate with MMP expression (8–10). Therefore, these effects of TIMPs may be independent of their ability to inhibit MMPs (11, 12). In addition, TIMP-1 promotes survival of serum-dependent cell lines, including Burkitt’s lymphomas, in serum-free conditions (13). Together, these findings suggest that extracellular matrix proteases or their inhibitors (TIMPs) may directly influence malignancy and PCD in B cells as well as in other cell types. The present study was therefore undertaken to determine whether TIMP-1 regulates apoptosis in B cells. The results of our study demonstrate that TIMP-1 directly controls apoptosis through a novel, non-MMP inhibitory pathway and suggest that this protein plays a pivotal role in maintenance of B cell homeostasis.

Methods

Cells. Burkitt’s lymphoma cell lines used in this study (see Table I) were kindly provided by Dr. Ian Magrath, Chief of Lymphoma Biology Section, National Cancer Institute (NCI) (14–17) or purchased from ATCC (Rockville, MD). Hyperplastic tonsils were obtained fresh in saline. A cell suspension was prepared. The tonsillar cells were cultured in serum-free conditions overnight in the absence or presence of 400 ng/ml of rTIMP-1. Cell lines were cultured in RPMI 1640 obtained from Gibco BRL (Rockville, MD), supplemented with 10% FBS (without serum as indicated), and antibiotics, 5% CO2, at 37°C.

TIMP-1 specificity. JD38 cells were preincubated with both native recombinant TIMP-1 and reduced and alkylated TIMP-1 (50–250 ng/ml) recombinant TIMP-2 (250–500 ng/ml), and the synthetic MMP inhibitor BB-94 (5 μM) obtained from British Biotech (Oxford, England) for 24 h before induction of apoptosis.

TIMP-1 dose response. JD38 cells (2.5 × 105) were cultured in a 96-well plate with serum-free media and in the presence or absence of


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rTIMP-1 (1.43–143 nM) prepared as previously described (18). After 48 h, the number of viable cells was determined.

Apoptosis-inducing treatments. In all methods, cells were monitored for apoptosis over extended periods of time (4–72 h), and optimal time points for detection of apoptosis were determined. Cold-shock induction of apoptosis was as described previously (19). In brief, 1 × 10^6 cells/ml were incubated at 4°C for 4 h and returned to 37°C for an additional 4–8 h. Anti-Fas apoptosis was induced as follows: 2 × 10^5 cells/ml were treated for 6–24 h with 200 ng/ml anti-CD95 monoclonal antibody clone CH-11 purchased from Immunotech (Miami, FL) at 37°C. For serum deprivation, cells were incubated for 48 h in serum-free RPMI at 37°C. Cells were also treated with 800 rads using a Gamma Cell 40 irradiator, followed by incubation in fresh RPMI at 37°C. Apoptosis was also induced in TIMP-1 positive cells grown in serum-free media and with anti–TIMP-1 monoclonal antibody (5 μg/ml). Anti–TIMP-2 and irrelevant isotype were used as controls. The TIMP-1 and TIMP-2 antibodies were purchased from Oncogene Research Products (Cambridge, MA).

Detection of apoptosis. Apoptosis was measured by at least three methods for each condition. All cells were reviewed microscopically for morphological evidence of apoptosis such as blebbing and nuclear condensation. Apoptosis was detected using flow cytometry by the following methods: light scatter as previously described (20) and decreased DNA content by using propidium iodide (PI) staining (sub-diploid peak characteristic of DNA fragmentation), Annexin V staining, assay of Caspase-3 activity, and TUNEL method (detects TdT incorporation of labeled nucleotides into DNA strand breaks). DNA content analysis by PI staining and the TUNEL method were combined using the Fluorescein Cell Death Detection Kit in a modification of the manufacturer’s protocol (Boehringer Mannheim, Indianapolis, IN). In brief, 2 × 10^5 cells were washed twice in PBS with 1% BSA and fixed in 2% paraformaldehyde for 30 min at room temperature. Cells were then washed twice with PBS, permeabilized with 0.1% Triton-X 100 and 2× SSC for 2 min on ice and resuspended in 50 l of TUNEL reaction media from the Boehringer’s kit. After 1 h incubation at 37°C, cells were washed with PBS and resuspended in 1 ml (5 mg/ml) PI from Calbiochem (La Jolla, CA) and 1 μl of 5 mg/ml Ribonuclease A from Worthington Biochemical (Freehold, NJ). Apoptosis was detected in tonsillar cells by using Annexin V–FITC from Caltag (Burlingame, CA) in conjunction with PE–anti-CD3 antibody (T cells), anti-CD10 antibody (germinal center B cells) obtained from Dako (Carpinteria, CA), and PerCP–anti-CD20 antibody from Becton Dickinson (San Jose, CA). Activation of caspase-3 was detected by using fluorogenic substrate PhiPhlux obtained from Oncolimmunin (College Park, MD) according to the manufacturer’s protocol.

Flow cytometric analysis was done on a FACSscan® from Becton Dickinson (San Jose, CA).

TIMP-1 retroviral construct. The TIMP-1–negative Burkitt’s lymphoma cell line JD-38 was selected, and LXSX retroviral transduction was used to induce TIMP-1 expression. JD-38 cells (see Table I) were obtained from Dr. Ian Magrath (Lymphoma Biology Section, NCI). TIMP-1 cDNA was obtained by PCR and subcloned into LXSX by using DNA recombinant techniques. Empty LXSX or TIMP-1–LXSX constructs were transfected into packaging lines. Nonadherent JD38 cells were then cocultured with adherent LXSX–TIMP-1 packaging cells (GP+envAM12) for 48 h (21). LXSX TIMP-1–infected JD38 cells were then selected using 2,500 μg/ml G418 for 10 d. Individual clones were isolated by repeated limiting dilution. Clonality was confirmed by restriction enzyme digestion with Sma I and Southern blot analysis, with unique sites of integration detected in the JD38 cell clones used in this study.

Northern blot analysis. Total RNA (7.5 μg), extracted from cell suspensions by using RNAzol purchased from Cinna/Biotec Laboratories (Houston, TX) was electrophoresed on 1% wt/vol agarose–formaldehyde gel before transfer onto nylon filters GeneScreen Plus from Dupont, NEN products (Boston, MA) and ultraviolet (UV) cross-linking to the filter. Blots were hybridized using standard conditions with TIMP-1 and GAPDH probes labeled with a [32P]dCTP by using a random primer labeling kit from Bethesda Research Laboratories (Gaithersburg, MD).

Western blot analysis. Conditioned media was collected from Burkitt’s lymphoma cell lines after 48 h incubation of 3 × 10^5 cells/ml at 37°C and 5% CO2 in AIMV or serum-free RPMI media both purchased from GIBCO BRL (Gaithersburg, MD). Supernatants were cleared of cells by centrifugation and protein concentration determined by BCA protein assay from Pierce (Rockford, IL). Equal amounts of proteins in 20–40 μl of conditioned media were electrophoresed in a 4–20% (wt/vol), polyacrylamide/SDS gel at 100 V for 60 min at room temperature and electroblotted onto polyvinylidene difluoride membranes from Novex (San Diego, CA) and at 10 V for 25 min at room temperature by using a semidry BIO-RAD electroblotter (Indianapolis, IN). Rabbit polyclonal antibodies against TIMP-1 kindly provided by Dr. Bente Birkedahl-Hansen (Extracellular Matrix Pathology Section, Laboratory of Pathology, NCI) and murine monoclonal TIMP-1 antibody purchased from Oncogene Sciences (La Jolla, CA); dilutions of 1:1,000 were used. Also, cytoplasmic proteins were extracted using RIPA buffer (150 mM NaCl, 1% NP-40, 0.1% SDS, 50 mM Tris-HCL, pH 8.0) containing protease inhibitors from JD38 cells, LXSX–JD38 cells and TIMP-1–JD38 cells, and 50 μg of proteins fractionated by electrophoresis as explained previously. Antibodies against Bcl-2, Bcl–X, p–65 NFκB, and IκBα were purchased from Santa Cruz Biotechnology (San Diego, CA) and used. These blots were developed with a horseradish peroxidase–conjugated goat anti–rabbit immunoglobulin or goat anti–mouse both from Pierce and a chemiluminescence kit from DuPont, NEN.

Cell cycle and DNA synthesis. TIMP-1–JD38 cell clones and LXSX–JD38 cells (5 × 10^5/ml) were cultured in the presence or absence of serum for 1–5 d. 10 μM bromodeoxyuridine (BRDU) from Calbiochem (La Jolla, CA) was added 30 min before harvesting cells. Cells were rinsed in PBS and fixed with 1% paraformaldehyde, followed by treatment with cold ethanol, and 2 N HCL. Cell suspension pH was normalized with borate buffer (pH 8.9). After rinsing with PBS, cells were incubated with FITC mouse anti-BRDU antibody from Becton-Dickinson, 30 min on ice, in the dark. Total DNA was counterstained with solution containing 50 μg/ml propidium iodine from Calbiochem and 1 μl Ribonuclease A (5 mg/ml) from Worthington Biochemical. Cell cycle distribution determined by PI staining and DNA synthesis detected by BRDU uptake were analyzed by flow cytometry using a Becton-Dickinson FACSscan® and CellQuest software.

Results

TIMP-1 expression and apoptosis. To determine whether soluble components of the ECM such as MMPs and their inhibitors regulate apoptosis in nonadhesion-dependent fashion, we correlated the expression of these components with the response of a series of human, Burkitt’s lymphoma cell lines to cold-shock–induced apoptosis (Table I). Apoptosis was determined by morphology and flow cytometry as explained in Methods. TIMP-1 expression in these Burkitt’s cell lines, determined by Northern and Western blot analysis (Fig. 1), showed 100% correlation with resistance to cold-shock–induced apoptosis. Four of the nine cell lines tested are TIMP-1 positive and did not undergo induction of apoptosis after cold shock while TIMP-1–negative cell lines readily underwent apoptosis. This response was not associated with other markers of biologic behavior that have been well studied in these and other Burkitt’s lymphoma cell lines such as patient origin, chromosomal translocation (t(8;14) vs. t(8;22)), presence of p53 mutation, or level of c-myc expression (14–17). Although all TIMP-1–positive lines were infected with EBV, both EBV–positive and negative lines were TIMP-1 negative. Some of the Burkitt’s cell lines studied expressed MMP-9 as assayed by zy-
TIMP-1, while there is no decrease in apoptosis in TIMP-2–DNA content and TUNEL method in cells pretreated with reduction in percentage of apoptotic cells as determined by BB-94 (5) demonstrated by incubating JD38 cells with TIMP-1 (50–250 ng/ml), (22). Specificity of TIMP-1 antiapoptotic effect is demonstrated by incubation of JD38 cells with TIMP-1 and GAPDH. Lane 1, ST46; lane 2, AG876; lane 3, EW36; lane 4, PA682; lane 5, Daudi; lane 6, DW6; lane 7, JD38; lane 8, Raji; lane 9, Jijyoe. (bottom) Western blot analysis of TIMP-1 in conditioned media from Burkitt’s lymphoma cell lines. Lane 1, ST46; lane 2, AG876; lane 3, EW36; lane 4, PA682; lane 5, Daudi; lane 6, DW6; lane 7, JD38; lane 8, Raji; lane 9, Jijyoe.

Gelatinase B expression was determined by zymography as previously reported. Apoptosis was induced by cold shock and detected by examination of morphology in culture for typical apoptotic changes and by flow cytometry using changes in light scatter characteristics as described in Methods. Gelatinase B expression was determined by zymography as previously reported. Apoptosis was induced by cold shock and detected by examination of morphology in culture for typical apoptotic changes and by flow cytometry using changes in light scatter characteristics as described in Methods.

Table I. Correlation of Burkitt’s Lymphoma Cell Line Phenotype with Susceptibility to Programmed Cell Death

<table>
<thead>
<tr>
<th>Cell line</th>
<th>Origin/translocation</th>
<th>EBV</th>
<th>p53</th>
<th>TIMP-1*</th>
<th>Gel B</th>
<th>Apoptosis</th>
</tr>
</thead>
<tbody>
<tr>
<td>PA682</td>
<td>American [t(8;22)]</td>
<td>+</td>
<td>+</td>
<td>+</td>
<td>+</td>
<td>-</td>
</tr>
<tr>
<td>AG876</td>
<td>African [t(8;14)]</td>
<td>+</td>
<td>-</td>
<td>+</td>
<td>+</td>
<td>-</td>
</tr>
<tr>
<td>DW6</td>
<td>American [t(8;14)]</td>
<td>+</td>
<td>+</td>
<td>-</td>
<td>+</td>
<td>-</td>
</tr>
<tr>
<td>Jijyoe</td>
<td>African [t(8;14)]</td>
<td>+</td>
<td>+</td>
<td>+</td>
<td>+</td>
<td>-</td>
</tr>
<tr>
<td>Raji</td>
<td>African [t(8;14)]</td>
<td>+</td>
<td>+</td>
<td>+</td>
<td>+</td>
<td>-</td>
</tr>
<tr>
<td>JD38</td>
<td>American [t(8;14)]</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>+</td>
<td>+</td>
</tr>
<tr>
<td>Daudi</td>
<td>African [t(8;14)]</td>
<td>+</td>
<td>+</td>
<td>-</td>
<td>+</td>
<td>-</td>
</tr>
<tr>
<td>EW36</td>
<td>American [t(8;14)]</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>+</td>
<td>-</td>
</tr>
<tr>
<td>ST46</td>
<td>American [t(8;14)]</td>
<td>-</td>
<td>+</td>
<td>-</td>
<td>-</td>
<td>-</td>
</tr>
</tbody>
</table>

Cell lines were established from patients with malignant lymphoma and small nonecleaved, Burkitt’s subtype, and include African and American Burkitt’s as well as cases with the t(8;14) and (8;22) translocations. Origin, translocation, EBV, and p53 mutation data provided in Methods. TIMP-1 expression determined by Northern and Western blot (Fig. 1). Gelatinase B expression was determined by zymography as previously reported. Apoptosis was induced by cold shock and detected by examination of morphology in culture for typical apoptotic changes and by flow cytometry using changes in light scatter characteristics as described in Methods.

In most lymphoid cell lines, PCD is induced by withdrawal of serum growth factors. The fetal bovine serum commonly used in tissue culture is a source of TIMP-1, and as reported by Hayakawa et al. (13), depletion of TIMP-1 from the serum is necessary to demonstrate the growth effects of TIMP-1 in several cell lines. The Burkitt’s lymphoma cell line AG876 expresses high levels of TIMP-1 and is resistant to induction of PCD by incubation in serum-free media. Neutralization of secreted TIMP-1 by incubation of AG876 cells with anti–TIMP-1 antibody results in a fourfold increased induction of apoptosis under serum-free conditions (24.03 ± 6.8% [SD] apoptotic AG876 cells in anti–TIMP-1–treated compared with 6.9 ± 1.0% [SD] apoptotic AG876 cells in controls, determined by TUNEL/PI staining) was demonstrated in the presence of native as well as reduced and alkylated rTIMP-1 (29 ± 8.3% [SD]) and 20 ± 6.2% [SD] apoptotic cells, respectively, as compared with cells cold-shocked in the absence of rTIMP-1 (61 ± 10.2% [SD] apoptotic cells). A dose of 1.4 nM of reduced and alkylated rTIMP-1 was as effective as native rTIMP-1 in suppressing cold-shock–induced apoptosis. Similar effects are observed when specific protease activity associated with PCD induction is determined. Caspase-3 activity is a critical event in the PCD pathway. Activation of this protease is indicative of initiation of the apoptotic cascade (4). Fig. 2C shows decreased percentage of cells with caspase-3 activity in JD38 cells pretreated with both native rTIMP-1 (13 ± 4.0% [SD] of positive cells) and reduced-alkylated rTIMP-1 (9.0 ± 2.5% [SD] of cells with caspase-3 activity), compared with caspase-3 activity of cells cold-shocked in the absence of exogenous TIMP-1 (29 ± 5.0% [SD] of positive cells). JD38 cells have a demonstrable caspase-3 activity of 7.8 ± 1.6% [SD] in the absence of cold shock. This further indicates that TIMP-1 protective effect is specific and is not due to inhibition of MMP activity. Also, this result demonstrates that TIMP-1 modulation of the apoptotic pathway results in downregulation of caspase-3 activity. The proliferation of TIMP-1 expression in Burkitt’s lymphoma cell lines. No other biological parameter studied in these tumors shows correlation with apoptosis. TIMP-1 protects against PCD. Recombinant TIMP-1 inhibited PCD induced by serum starvation (Fig. 2A), cold shock (Fig. 2B), and Fas activation (data not shown) in the TIMP-1 nonexpressing, EBV-negative JD38 cell line. Cell viability in serum deprivation was dose dependent in the JD38 cells incubated with 1.4–143 nM rTIMP-1 for 48 h (Fig. 2A) with 50% survival observed at < 10 nM TIMP-1. Saturable cell surface binding of TIMP-1 by JD38 cells was detected (data not shown) at levels similar to that observed in other cell lines (22). Specificity of TIMP-1 antiproteoic effect is demonstrated by incubating JD38 cells with TIMP-1 (50–250 ng/ml), TIMP-2 (250–500 ng/ml), and the synthetic MMP inhibitor BB-94 (5 μM) before cold-shock treatment. Fig. 2B shows a reduction in percentage of apoptotic cells as determined by DNA content and TUNEL method in cells pretreated with TIMP-1, while there is no decrease in apoptosis in TIMP-2–BB-94–treated cells. Recombinant TIMP-1 inhibited PCD (33% decrease of apoptosis from controls) by serum starvation in germinal center B cells (CD20+ and CD10+) from tonsils but did not affect apoptosis in T cells (1.2% decrease of apoptosis from controls).
activity and suggests that TIMP-1 effects are mediated upstream of activation of this cell-death protease. TIMP-1 does not inhibit caspase-3 activity (data not shown).

**TIMP-1 upregulation in JD38 cells inhibits PCD.** A direct TIMP-1 effect was demonstrated by inducing TIMP-1 expression in the EBV-negative JD38 cells and selection of independent cell clones as explained in Methods. Fig. 3A shows expression of TIMP-1 in parent cells, three clones carrying empty LXSN vector, and four TIMP-1–JD38 cell clones as detected by Northern blot analysis. Western blot analysis of conditioned media demonstrated TIMP-1 expression and secretion by the TIMP-1–expressing clones T8, T20, and T24 but not detectable TIMP-1 in the parental line JD38 or LXSN controls (Fig. 3B). The TIMP-1 protein was functional as demonstrated by reverse zymography (data not shown). The JD38 parental cells, LXSN–JD38 cell clones, as well as TIMP-1–JD38 cell clones, were examined for susceptibility to induction of PCD by various stimuli. Cells were observed periodically from 4–72 h after any of the following treatments: cold shock at 4°C, anti-CD95 (anti-Fas/APO) activating antibody, serum deprivation, or γ-radiation. PCD was assessed by morphology (not shown), TUNEL method (TdT incorporation of labeled nucleotides into DNA strand breaks), PI staining (decreased DNA content due to DNA fragmentation), and light scatter changes (decreasing size and increasing granularity of cells, not shown). Table II shows a summary of percentage of apoptotic cells as determined by TUNEL/PI after treatment for each of the JD38 cell lines. Induction of TIMP-1 expression in three independent JD38 cell clones, each with a unique integration site, is associated with inhibition of apoptosis initiated by all methods studied as shown in Table II. Moreover, neutralization of secreted TIMP-1 in clones T20 and T24 with monoclonal anti–TIMP-1 antibody restored sensitivity to serum starvation while treatment with anti–TIMP-2 or isotype control did not (Fig. 4), indicating that it is secreted TIMP-1 that prevents apoptosis. These results in conjunction with our preliminary observations of cell surface binding of TIMP-1 by JD38 cells suggest that TIMP-1 effect may be receptor mediated.

To determine whether TIMP-1 protective effect was dependent on withdrawal from cell cycle, parent, LXSN–JD38, and TIMP-1–JD38 cell clones 20 and 24 were cultured in the absence or presence of serum for 1–5 d, and changes in cell cycle distribution were detected by PI staining and DNA synthesis monitored by BRDU uptake. Fig. 5 shows flow cytometric analysis of these parameters for LXSN–JD38 and TIMP-1–JD38 clone 24. A decreasing percentage of cells incorporating BRDU is demonstrated by control LXSN–JD38 cells, with 44.3% of cells synthesizing DNA in the presence of serum, decreasing to 9.6% by 48 h in the absence of serum. TIMP-1–JD38 cells demonstrate a higher percentage of cells actively incorporating nucleotides throughout the same treatment times. These results suggest both that the TIMP-1–protective effect does not involve cell cycle perturbations and that control cells die while progressing through the cycle.

**Effects of TIMP-1 on apoptotic proteins.** Bcl-2 has been shown to inhibit apoptosis in B cells as well as in numerous other cell types as a consequence of its ability to inhibit apoptosis (Fig. 2A). Exogenous rTIMP-1 suppresses PCD and caspase-3 in the Burkitt’s lymphoma cell line JD38. (A) TIMP-1 dose response in serum-deprived conditions. JD38 cells were incubated for 48 h in serum-free media with 1.43–143 nM rTIMP-1. Results are reported as the number of viable cells as determined by Trypan blue exclusion viability assay. Results represent mean ± SD of duplicates of two analyses. (B) Susceptibility to cold-shock–induced apoptosis of JD38 cells as determined by TUNEL (y-axis) and a shift in total DNA as result of DNA fragmentation (x-axis). Boxed area indicates viable cells. Increased viability (85%) is observed in cells treated with 250 ng/ml TIMP-1 as compared with untreated cells (54%) while treatment with 500 ng/ml TIMP-2 and 5 μM BB-94 shows viability of 56% and 52%, respectively. (C) Cold-shocked cells pretreated with 1.4 nM reduced and alkylated (R&A) TIMP-1 also show a decreased number of cells with caspase-3 activity (10.8%) similar to the effect of native protein (TIMP-1) while an increased caspase activity (32%) is observed in cells cold-shocked in the absence of TIMP-1 (untreated). Endogenous caspase-3 activity at 37°C is also shown. Similar results obtained with TUNEL/PI are discussed in the text.

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**Figure 2.** Exogenous rTIMP-1 suppresses PCD and caspase-3 in the Burkitt’s lymphoma cell line JD38. (A) TIMP-1 dose response in serum-deprived conditions. JD38 cells were incubated for 48 h in serum-free media with 1.43–143 nM rTIMP-1. Results are reported as the number of viable cells as determined by Trypan blue exclusion viability assay. Results represent mean ± SD of duplicates of two analyses. (B) Susceptibility to cold-shock–induced apoptosis of JD38 cells as determined by TUNEL (y-axis) and a shift in total DNA as result of DNA fragmentation (x-axis). Boxed area indicates viable cells. Increased viability (85%) is observed in cells treated with 250 ng/ml TIMP-1 as compared with untreated cells (54%) while treatment with 500 ng/ml TIMP-2 and 5 μM BB-94 shows viability of 56% and 52%, respectively. (C) Cold-shocked cells pretreated with 1.4 nM reduced and alkylated (R&A) TIMP-1 also show a decreased number of cells with caspase-3 activity (10.8%) similar to the effect of native protein (TIMP-1) while an increased caspase activity (32%) is observed in cells cold-shocked in the absence of TIMP-1 (untreated). Endogenous caspase-3 activity at 37°C is also shown. Similar results obtained with TUNEL/PI are discussed in the text.
types (1, 2, 24–26). To determine whether Bcl-2 or related proteins are implicated in the TIMP-1 protective effect, TIMP-1–JD38 cell clones and control LXSN and parent JD38 cells were assayed for their expression of these antiapoptotic proteins by Western analysis. Although both control and TIMP-1–JD38 express Bcl-2, upregulation of TIMP-1 does not further enhance expression of this protein. However, induction of TIMP-1 clearly upregulates Bcl-X<sub>L</sub> in TIMP-1–JD38 cell clones as compared with LXSN–JD38 cells (Fig. 6). Also, cells were all negative for the Bcl-2 homologue Mcl-1 (27) (data not shown). In addition to controlling several genes in B lymphocytes, including immunoglobulin light chain gene, the transcription factor NF-κB has been reported to regulate expression of genes involved in apoptosis (28–30). After immunodetection of Bcl-X<sub>L</sub>, the same blot in Fig. 6 was reprobed with an antibody against NF-κB p65 (Rel A). No change in the cytoplasmic expression of NF-κB was observed for any of JD38 cell lines (Fig. 6). However, higher expression of the NF-κB inhibitor IκBα is observed in TIMP-1–expressing cell clones (20, 24), indicating a decreased in NF-κB activity (Fig. 6). Furthermore, TIMP-1 suppression of PCD did not correlate with levels of cell surface CD95 expression (data not shown). These results suggest that TIMP-1 antiapoptotic mechanism is not mediated by Bcl-2 or suppression of CD95 expression, but likely by inducing Bcl-X<sub>L</sub>.

### Table II. Summary of Induction of Programmed Cell Death in JD38 Cells with Retroviral Transduced TIMP-1 Expression as Determined by TUNEL/PI

<table>
<thead>
<tr>
<th>Cell line</th>
<th>Untreated</th>
<th>Cold shock</th>
<th>anti-FAS</th>
<th>γ-Radiation</th>
<th>Serum free</th>
</tr>
</thead>
<tbody>
<tr>
<td>JD38</td>
<td>8.4±4.3</td>
<td>56.3±7.5</td>
<td>29.0±8.0</td>
<td>32.0±1.4</td>
<td>87.0±3.0</td>
</tr>
<tr>
<td>LXSN (Control)</td>
<td>5.8±2.7</td>
<td>43.6±9.0</td>
<td>40.0±13.0</td>
<td>34.1±5.7</td>
<td>78.0±13.0</td>
</tr>
<tr>
<td>TIMP-1 (Clone 8)</td>
<td>2.4±0.6</td>
<td>1.3±0.9</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>TIMP-1 (Clone 20)</td>
<td>5.5±3.5</td>
<td>8.1±0.6</td>
<td>5.4±3.4</td>
<td>3.2±1.2</td>
<td>6.1±2.5</td>
</tr>
<tr>
<td>TIMP-1 (Clone 24)</td>
<td>3.4±1.1</td>
<td>13.2±9.0</td>
<td>2.1±0.1</td>
<td>4.7±1.2</td>
<td>5.7±0.8</td>
</tr>
</tbody>
</table>

Apoptosis was detected by TUNEL in conjunction with PI staining for the detection of sub G0/G1 DNA peak as explained in Methods. These results were confirmed by at least three methods: light-microscopy morphology, forward and side scatter, and caspase-3 activity (not shown). Results are expressed as percentage of apoptotic cells, mean±SD of duplicate determinations of three flow-cytometric analyses.
Also, upregulation of Bcl-X\textsubscript{L} by TIMP-1 appears to be independent from NF-\kappaB activation in TIMP-1-expressing JD38 cells.

**Discussion**

The extracellular matrix has been implicated in the regulation of apoptosis of epithelial cells in an attachment-dependent fashion. In these studies, ECM was demonstrated to modulate PCD and effectors such as caspases through integrin receptors (4). More recently, extracellular matrix factors were shown to control PCD of hematolymphoid cells (5, 6, 31). Using a human melanoma cell line, Valente et al. have shown that TIMP-2 inhibits tumor cell invasion and that TIMP-2 overexpressing tumor clones were found to be more resistant to apoptosis than parental and control melanoma cells (32). A recent report in this journal demonstrates that TIMP-3 promotes apoptosis in rat vascular smooth muscle cells (33). These studies support the idea that TIMPs can exert different activities in addition to inhibition of MMP, including modulation of PCD. However, in both of these studies, the mechanism for the effect of TIMP-2 and TIMP-3 on PCD is not addressed. Our laboratory and others have previously reported on the expression of MMPs and their endogenous inhibitors (TIMPs) by normal and neoplastic lymphocytes (7, 34). We have found high-level expression of TIMP-1 in normal activated tonsillar B cells as well as in high-grade B cell lymphoma cell lines, frequently in the absence of MMP expression (7). In the present study, we have found TIMP-1 expression is associated with resistance to apoptosis in Burkitt’s lymphoma cell lines and that TIMP-1 can inhibit apoptosis in normal tonsillar B cells.

The suppression of PCD is a novel action for TIMP-1. Our data further demonstrates that this effect is TIMP-1 specific and is not secondary to MMP inhibition. The ability of TIMP-1 to inhibit apoptosis in an MMP-independent manner is consistent with observations of growth promotion by TIMPs in a variety of cell types (13). Hayakawa et al. have shown that TIMP-1 promotes cell survival in serum-free conditions for a wide variety of cell lines sensitive to serum deprivation including lymphoid, myeloid, endothelial, fibroblasts, hepatoma, breast carcinoma, and chondrocyte cell lines (13). Although these authors did not measure apoptosis in these cell lines, it is known that they do undergo apoptosis under conditions of serum starvation. Addition of TIMP-1 was sufficient to maintain cell growth and viability, suggesting that TIMP-1 may have inhibited serum-starvation induced PCD in these lines representing multiple lineages. TIMP-1 inhibits apoptosis in mouse mammary epithelial cells (3, 4). This effect was attributed to inhibition of MMP activity and preservation of cell–matrix contacts. The possible direct antiapoptotic effect of TIMP-1 was not examined in this system. Preservation of cell–matrix is clearly not the mechanism of TIMP-1 suppression of PCD in Burkitt's lymphoma cell lines. First, reduced-alkylated TIMP-1, completely devoid of MMP inhibitory activity, results in PCD suppression. In addition, TIMP-2 or the synthetic inhibitor BB-94 failed to protect from PCD, even though the levels of BB-94 used should completely inhibit a wide spectrum of metalloproteinases, including MMPs. The ability of TIMP-1 to exert antiapoptotic effects independently of its MMP inhibitory effect is consistent with other studies that demonstrated TIMP-1 cell growth modulatory effects unrelated to its ability to act as MMP inhibitor (12).
The growth promoting effects of TIMP-1 do not appear to be due to withdrawal from cell cycle, as has been observed for the survival activity of the ECM (42). In this study, TIMP-1–JD38 cells cultured in serum-deprived conditions continued cell cycle for extended periods of time. These results support TIMP-1 as an antiapoptotic factor that inhibits PCD while allowing cell growth in stress. The observation that TIMP-1 is secreted by these cell lines and binds to the cell surface in combination with the demonstration that neutralization of secreted TIMP-1 reverses suppression of apoptosis, suggests that TIMP-1 acts through a receptor-mediated autocrine loop. Other investigators (12, 13, 22) have postulated the presence of such a receptor. Studies in our laboratory are underway to isolate and characterize this receptor.

The central role of apoptosis in the homeostasis of lymphoid tissues is now widely acknowledged. In lymphocytes there are at least two independent pathways for initiation of PCD: the tumor necrosis (TNF)/nerve growth factor (NGF) receptor family pathway (involved in numerous functions including immunoresponses and T cell cytotoxicity), and the Bcl-2 family–dependent pathway, initiated by factors such as irradiation, cytotoxic drugs, and serum deprivation (43–46). Except for Bcl-X$_L$, Bcl-2 family members have not been shown to directly alter CD95/Fas pathway in lymphocytes (45, 46). Results of the present study demonstrate that TIMP-1 protects B cells from apoptosis induced by both Bcl-2–dependent pathways (serum starvation and $\gamma$-radiation) and Bcl-2 independent ones such as Fas (43–45). Most Burkitt’s lymphoma lines are prone to apoptosis and have been reported to be highly sensitive to cold shock (19). Although factors implicated in the
control of this pathway are yet to be described, some reports suggest that Bcl-2 is not one of them (19, 47), and this may be a third pathway for induction of PCD that TIMP-1 can also protect against. In the present study, TIMP-1 did not affect Bcl-2 or CD95 expression. However Bcl-XL is upregulated in the TIMP-1–JD38 cell clones, indicating that the TIMP-1 protective effect might be in part mediated through induction of Bcl-XL. Bcl-XL is known to not only overlap with Bcl-2 in protection from apoptosis but also exerts its protective effect in pathways not controlled by Bcl-2, such as the Fas/TNF pathway (45, 46, 48). Also, in contrast to B cell studies showing induction of antiapoptotic proteins by NF-κB (28, 29), the present study demonstrates upregulation of the inhibitor IκB (48) in TIMP-1–JD38 clones. Therefore, NF-κB activation may not be involved in the induction of antiapoptotic proteins by TIMP-1. Interestingly, this decreased NF-κB activity is also supported by our previous results showing downregulation of immunoglobulin expression in TIMP-1–positive cells, since immunoglobulin expression is known to be controlled by NF-κB (49). Future studies are needed to conclusively determine the role of NF-κB or to identify other transcription factors involved in TIMP-1’s antiapoptotic effect.

Overall, this study supports a novel role for TIMP-1 in the control of apoptosis in B cells. Results discussed here provide a biological basis for the observed negative prognosis associated with increased TIMP-1 expression in non–Hodgkin’s lymphomas, as well as many solid tumors (8–10, 36). Further studies are indicated to determine if TIMP-1 inhibition of apoptosis in the tumor cells alters the prognosis in these patients.

Acknowledgments

The authors thank Dr. Megan Lim for performing the reverse zymography analysis, Dr. Peter Brown, British Biotechnology, Ltd., for supplying BB-94, Dr. Anita Yu for reduced and alkylated TIMP-1, and Dr. Kevin Gardner for his valuable advice and NF-κB antibodies.

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