

Monocyte-mediated Tumoricidal Activity via the Tumor Necrosis Factor-related Cytokine, TRAIL

By Thomas S. Griffith,* Steven R. Wiley,[†] Marek Z. Kubin,*
Lisa M. Sedger,[§] Charles R. Maliszewski,[§] and Neil A. Fanger[§]

From the *Department of Immunobiology, the [†]Department of Bioinformatics, and the [§]Department of Discovery Research, Immunex Corporation, Seattle, Washington 98101

Summary

TRAIL (tumor necrosis factor [TNF]-related apoptosis-inducing ligand) is a molecule that displays potent antitumor activity against selected targets. The results presented here demonstrate that human monocytes rapidly express TRAIL, but not Fas ligand or TNF, after activation with interferon (IFN)- γ or - α and acquire the ability to kill tumor cells. Monocyte-mediated tumor cell apoptosis was TRAIL specific, as it could be inhibited with soluble TRAIL receptor. Moreover, IFN stimulation caused a concomitant loss of TRAIL receptor 2 expression, which coincides with monocyte acquisition of resistance to TRAIL-mediated apoptosis. These results define a novel mechanism of monocyte-induced cell cytotoxicity that requires TRAIL, and suggest that TRAIL is a key effector molecule in antitumor activity *in vivo*.

Key words: TRAIL • apoptosis • tumor • monocyte • human

Mononuclear phagocytes (M ϕ)¹ circulate in the peripheral blood before differentiation into non-lymphoid and lymphoid tissue-associated macrophages (1). M ϕ mediate several host defense mechanisms through activation of MHC class I- and class II-restricted T lymphocytes (2, 3), release of inflammatory mediators (4), and killing of virus-infected cells (5). In addition, M ϕ effectively kill tumor cells through both antibody-dependent and antibody-independent mechanisms (6, 7). A critical prerequisite for these M ϕ functions is cellular activation. Two potent mediators of M ϕ activation, IFN- γ and IFN- α , dramatically enhance the cytolytic potential of human M ϕ (8, 9). The mechanisms responsible for this IFN-induced tumoricidal activity remain poorly defined. Nitric oxide (NO) is known to be produced by M ϕ and is a potent mediator of tumor cell death (10, 11). However, several reports have demonstrated that IFN- γ and IFN- α stimulation of human M ϕ does not increase the level of NO release, suggesting that NO does not play a significant role in the cytotoxic activity of IFN-stimulated human M ϕ (12–14). Although TNF possesses cytolytic activity and is produced by M ϕ after certain types of activation (e.g., LPS), several studies have shown that IFN does not induce TNF production in human M ϕ (15–17). These data indicate that other cell death-inducing

molecules may be involved with the M ϕ -mediated tumoricidal activity after IFN stimulation.

Two TNF family members capable of inducing cell death are Fas ligand (FasL) and TNF-related apoptosis-inducing ligand (TRAIL). FasL participates in many physiological events, including autoimmunity, activation-induced cell death (AICD) of lymphocytes, immune privilege, and tumor evasion from the immune system (18–21). Dysregulation of physiological Fas/FasL interactions results in immune disease states characterized by enhanced levels of Fas-mediated apoptosis and a variety of forms of hepatitis (22–24) or diseases with decreased levels of lymphocyte death (e.g., human autoimmune lymphoproliferative syndrome [18]). Human M ϕ contain high levels of intracellular FasL that can be released after cellular activation (25). Recent evidence indicates that FasL-expressing M ϕ are essential in the elimination of activated effector T cells in homeostasis and in disease, suggesting that they may also play an important physiological role in a variety of other immunological settings (23, 26–29).

In contrast to FasL, the role of TRAIL in immune regulation remains enigmatic (30). Recent studies have identified four distinct cell surface TRAIL receptors, with two (DR4 and DR5/TRAIL-R2; hereafter referred to as TRAIL-R1 and -R2, respectively) that contain a cytoplasmic death domain and signal for apoptotic cell death upon receptor cross-linking, and two (TRID/DcR1/TRAIL-R3 and TRAIL-R4/DcR2; hereafter referred to as TRAIL-R3 and -R4, respectively) that lack a death domain, making

¹Abbreviations used in this paper: AICD, activation-induced cell death; L-NMMA, N^G-monomethyl-L-arginine; LZ, leucine zipper; M ϕ , mononuclear phagocyte(s); NO, nitric oxide; RT, reverse transcription; TRAIL, TNF-related apoptosis-inducing ligand.

them unable to signal for cell death (31–37). Recombinant, soluble forms of TRAIL are potent mediators of tumor cell apoptosis, while demonstrating minimal cytotoxicity toward normal tissues *in vitro* and *in vivo* (38–40). The cell death induced by TRAIL displays many of the same characteristics observed with other apoptotic molecules (i.e., FasL and TNF), such as caspase activation, DNA fragmentation into oligonucleosomal “ladders,” annexin V binding, and the morphological membrane blebbing and apoptotic body release (30, 38, 41). This tumoricidal activity occurs on ~2/3 of the >30 hematopoietic and nonhematopoietic tumor cell lines tested (30, 36, 38, 42). Such studies not only support the potential use for TRAIL as an antitumor therapeutic agent (30, 38, 40), but suggest that TRAIL may be an innate effector molecule involved in the elimination of a broad range of spontaneously arising tumor cells. The results presented here identify a novel mechanism by which M ϕ induce tumor cell apoptosis via TRAIL.

Materials and Methods

Reagents and mAbs. Reagents and sources were as follows: IL-1, IL-2, IL-3, IL-4, IL-6, IL-7, IL-15, GM-CSF (100 ng/ml; Immunex); IFN- α , IFN- γ , IL-10, IL-12 (100 ng/ml; Genzyme Corp.); LPS (5 ng/ml; DIFCO); MOPC-21, nonspecific IgG1 isotype control; MOPC-173, nonspecific IgG2a isotype control; M181, IgG1 anti-TRAIL (Immunex); mAb11, IgG1 anti-TNF; NOK-1, IgG1 anti-FasL (PharMingen). The mAbs against the four TRAIL receptors (M271, IgG2a anti-TRAIL-R1; M413, IgG1 anti-TRAIL-R2; M430, IgG1 anti-TRAIL-R3; and M444, IgG1 anti-TRAIL-R4) were produced at Immunex by immunizing BALB/c mice (The Jackson Laboratory) with a purified fusion protein consisting of the extracellular domain of human TRAIL-R1, -R2, -R3, or -R4 coupled to the constant region of human IgG1 (huTRAIL-R:Fc) in Titermax (CytRx Corporation). Mice were boosted three times, and spleen cells were fused with the murine myeloma NS1 in the presence of 50% polyethylene glycol in PBS followed by culture in DMEM/HAT and DMEM/HT selective media. Supernatants from positive wells were tested for the ability to bind the appropriate TRAIL receptor in an ELISA (cell-based ELISA using CV1 cells transfected with TRAIL receptor cDNA) and reactivity to huTRAIL-R:Fc in Western blots. Hybridomas that produced antibodies that bound to huTRAIL-R:Fc, but not human IgG1, were cloned by three rounds of limiting dilution. All mAbs were purified by protein A affinity chromatography. The matrix metalloproteinase inhibitors, TAPI (TNF protease inhibitor [43]) and KB8301, were obtained from Immunex and PharMingen, respectively. The soluble fusion proteins TRAIL-R2:Fc, Fas:Fc, and TNFR:Fc were produced at Immunex. The leucine zipper (LZ)-TRAIL expression plasmid (34) and the production and purification of LZ-huTRAIL (40) have been described previously.

Cell Lines. The ovarian carcinoma cell line (OVCAR3) was obtained from Dr. Richard F. Camalier, Developmental Therapeutics Program, Division of Cancer Treatment and Diagnosis, National Cancer Institute (Bethesda, MD). The human melanoma cell lines (WM 793 and 164) were obtained from Dr. M. Herlyn, Wistar Institute (Philadelphia, PA). The human prostate carcinoma cell line (PC-3) was obtained from Dr. Michael Cohen, University of Iowa (Iowa City, IA). The human colon carcinoma cell line (Colo205) and the human lung adenocarcinoma

cell line (H2126) were provided by Dr. Brian Gliniak and Tim Lofton, respectively (Immunex). L929 cells and normal human foreskin fibroblasts were obtained from American Type Culture Collection. All tumor cell lines were cultured as directed. The normal human lung fibroblasts were purchased from Clonetics Corporation and cultured as directed.

Isolation of Human M ϕ . Peripheral blood M ϕ were enriched using countercurrent elutriation. Cells from leukopheresis packs obtained from healthy volunteers were loaded onto a JE-5 elutriator (Beckman), and 50-ml fractions were collected while increasing the flow rate from 65 to 85 ml/min at 2,000 rpm. M ϕ -enriched fractions generated from a flow rate >75 ml/min were >90% CD14⁺ as assessed by flow cytometric analysis using TUK-4, an IgG2a anti-CD14 (Caltag Laboratories, Inc.).

Flow Cytometry. Untreated or cytokine-stimulated M ϕ were incubated with the following unlabeled primary mAbs for 1 h at 4°C: MOPC-21, MOPC-173, M181, M271, M413, M430, M444, mAb11, and NOK-1. After three washes, primary antibody binding was detected with a PE-conjugated, Fc-specific, mouse anti-human F(ab')₂ (Jackson ImmunoResearch Laboratories). Staining for TNF was done in the presence of the matrix metalloproteinase inhibitor, TAPI (50 μ M). Staining for FasL was done in the presence of the matrix metalloproteinase inhibitor, KB8301 (10 μ M). Cells were analyzed immediately after staining or fixed in 1% paraformaldehyde until analysis on a FACSCalibur™ (Becton Dickinson).

TRAIL-mediated Killing of Human Tumor Cells. M ϕ were cultured for 12 h in medium alone, GM-CSF, IFN- γ , or IFN- α , washed, and resuspended in complete medium. Tumor cells were labeled with 100 μ Ci of ⁵¹Cr for 1 h at 37°C, washed three times, and resuspended in complete medium. To determine TRAIL-induced death, ⁵¹Cr-labeled tumor cells (10⁴/well) were incubated with varying numbers of M ϕ effector cells for 8 h. As a positive control, soluble LZ-TRAIL was added to the target cells at the indicated concentrations. In some cultures, TRAIL-R2:Fc, Fas:Fc, or TNFR:Fc (20 μ g/ml) was added to the M ϕ effector cells 15 min before adding tumor cell targets. All cytotoxicity assays were performed in round-bottomed 96-well plates, and the percent specific lysis was calculated as: 100 \times (experimental cpm – spontaneous cpm)/(total cpm – spontaneous cpm). Spontaneous and total release were determined in the presence of either medium alone or 1% NP-40, respectively. The presence of TRAIL-R2:Fc, Fas:Fc, or TNFR:Fc during the assay had no effect on the level of spontaneous release of ⁵¹Cr by the target cells. For analysis of tumor cell apoptosis, tumor cell targets were incubated with unstimulated or cytokine-stimulated M ϕ as described above. Apoptotic cell death of the tumor cells was measured by flow cytometry using FITC-conjugated annexin V and propidium iodide (Apoptosis detection kit; R&D Systems) as per the manufacturer's protocol. Light scatter characteristics were used to identify the tumor cells.

Inhibition of NO Synthesis. The specific inhibitor of NO synthase, N^G-monomethyl-L-arginine (L-NMMA; AerBio, Ltd.), was used to block M ϕ NO production (44). M ϕ were stimulated with the appropriate cytokine and then incubated with the tumor cell targets as above. M ϕ were cultured in the presence of 300 μ M L-NMMA at all steps throughout the assay. The presence of L-NMMA had no effect on the level of spontaneous release of ⁵¹Cr by the target cells during the assay.

TNF-mediated Killing of L929 Cells. M ϕ were cultured for 2 or 12 h in medium alone or LPS (5 ng/ml), washed, and resuspended in complete medium. L929 cells were labeled with 100 μ Ci of ⁵¹Cr for 1 h at 37°C, washed three times, and resuspended in com-

plete medium. To determine TNF-induced death, ^{51}Cr -labeled L929 cells (10^4 /well) were incubated with varying numbers of M ϕ effector cells for 8 h. As a positive control, soluble TNF was added to the target cells at the indicated concentrations. In some cultures, TNFR:Fc or TRAIL-R2:Fc (20 $\mu\text{g}/\text{ml}$) was added to the M ϕ effector cells 15 min before adding tumor cell targets. L929 cytotoxicity assays were performed in an identical manner as for the human tumor cell lines. The presence of TNFR:Fc or TRAIL-R2:Fc during the assay had no effect on the level of spontaneous release of ^{51}Cr by the L929 target cells.

TRAIL-mediated Killing of M ϕ . M ϕ were cultured for 12 h in medium alone or with cytokine (IL-1, IL-2, IL-3, IL-4, IL-7, IL-10, IL-12, IL-15, GM-CSF, or IFN- γ); all cytokines were used at 100 ng/ml, and then labeled with 100 μCi of ^{51}Cr for 1 h at 37°C, washed three times, and resuspended in complete medium. To determine sensitivity to TRAIL-induced death, ^{51}Cr -labeled M ϕ (10^6 /well) were incubated with LZ-TRAIL for 8 h. Assays were performed as described above.

Reverse Transcription PCR. Total RNA was isolated from M ϕ with TRIzol reagent (Life Technologies) as per the manufacturer's instructions. RNA samples (1 μg each) were tested for DNA contamination by 30 cycles of PCR with human β -actin primers. After it was shown that there was no DNA contamination, cDNA synthesis was performed using an RNA PCR kit (Perkin-Elmer) with the supplied oligo d(T) $_{16}$ primer. Reverse transcription was performed using a thermal program of 25°C for 10 min, 42°C for 30 min, and 95°C for 5 min. PCR reactions were performed using the following primers: human β -actin (forward: 5'-GAACTACCTTCAACTCCATC-3'; reverse: 5'-CGAGGCCAGGATGGAGCCGCC-3'); human TRAIL-R1 (forward: 5'-CTGAGCAACGCAGATCGCTGTCCAC-3'; reverse: 5'-TCCAAGGACACGGCAGAGCCTGTGCCAT-3'); human TRAIL-R2 (forward: 5'-GCCTCATGGACAATGAGATAAAGGTGGCT-3'; reverse: 5'-CCAAATCTCAAAGTACGCACAAACGG-3'); human TRAIL-R3 (forward: 5'-GAAGAATTTGGTGCCAATGCCACTG-3'; reverse: 5'-CTCTTGGACTTGGCTGGGAGATGTG-3'); human TRAIL-R4 (forward: 5'-CTTTTCCGGCGGCGTTCATGTCTTC-3'; reverse: 5'-GTTTCTTCCAGGCTGCTTCCCTTTGTAG-3'); and human TRAIL (forward: 5'-CAACTCCGTCAGTCGTTAGAAAG-3'; reverse: 5'-TTAGACCAACAACCTATTCTAGCACT-3'), giving prod-

ucts of 219, 506, 502, 612, 453, and 443 bp, respectively. β -actin PCR cycle conditions were 95°C for 45 s, 55°C for 1 min, and 72°C for 45 s for 30 cycles. TRAIL-R1, -R2, and -R3 cycle conditions were 94°C for 1 min, 55°C for 1 min, and 72°C for 1 min for 30 cycles. TRAIL-R4 cycle conditions were 95°C for 4 min 15 s, followed by 30 cycles of 95°C for 45 s, 60°C for 45 s, and 72°C for 45 s. TRAIL cycle conditions were 95°C for 45 s, 55°C for 45 s, and 72°C for 45 s for 30 cycles. Samples were resolved on a 2% agarose gel and visualized with ethidium bromide.

Results

Human M ϕ Stimulated with IFN Upregulate TRAIL. To compare the difference in TRAIL, FasL, and TNF surface expression, peripheral blood human M ϕ were isolated and cultured with several molecules known to induce M ϕ activation and differentiation (IFN- γ , IFN- α , GM-CSF, and LPS), and then examined by flow cytometry. Significant TRAIL expression was detected on the M ϕ cultured for 12 h in the presence of either IFN- γ or IFN- α , but not with either GM-CSF or LPS (Fig. 1 A). In contrast, FasL expression on M ϕ was undetectable after either 2- or 12-h stimulation with any of the cytokines or LPS in the presence of the metalloproteinase inhibitor, KB8301 (Fig. 1 A). Analysis of the surface levels of TNF demonstrated no measurable increase after stimulation with IFN- γ or IFN- α ; however, stimulation with LPS for 2 h led to increased surface levels of TNF that disappeared by 12 h (Fig. 1 B). To inhibit the cleavage of membrane TNF, M ϕ were cultured in the presence of the metalloproteinase inhibitor, TAPI (TNF protease inhibitor [43]). These data demonstrate that TRAIL, but not FasL and TNF, is induced after IFN stimulation, and that the expression of TRAIL and TNF on M ϕ is regulated by distinct activation stimuli and with different kinetics.

IFN-stimulated M ϕ Kill Tumor Cells via a TRAIL-dependent Mechanism. The results from Fig. 1 demonstrate that M ϕ stimulated with IFN upregulate the expression of TRAIL

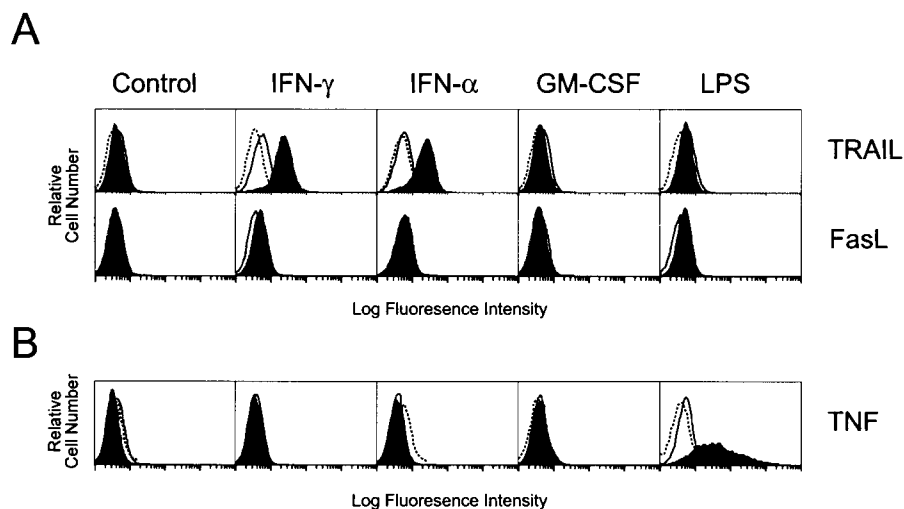


Figure 1. TRAIL, FasL, and TNF expression on human M ϕ . (A) M ϕ were incubated for 2 or 12 h in the absence or presence of IFN- γ , IFN- α , GM-CSF, or LPS and then analyzed for TRAIL or FasL surface expression. Filled histograms represent staining at 12 h by either M181 (anti-TRAIL mAb) or NOK-1 (anti-FasL mAb). Open histograms represent staining at 2 h with the same mAb, whereas dotted histograms represent staining with isotype control mAb. (B) M ϕ were incubated as in A and then analyzed for TNF surface expression. Filled histograms represent staining at 2 h by mAb11 (anti-TNF mAb). Open histograms represent staining at 12 h with the same mAb, whereas dotted histograms represent staining with isotype control mAb. Histograms represent 10^4 gated M ϕ in all conditions, and viability was >95% as assessed by propidium iodide exclusion. These observations were reproduced using M ϕ from at least five different donors.

on the cell surface. Thus, to examine the functional activity of TRAIL in this setting, M ϕ were stimulated with either GM-CSF, IFN- γ , or IFN- α for 12 h and then cultured in the presence of OVCAR3, a TRAIL-sensitive human ovarian carcinoma cell line. Although the unstimulated or GM-CSF-treated M ϕ demonstrated minimal tumoricidal activity toward OVCAR3, the M ϕ stimulated with IFN- γ or IFN- α were potent killers of these TRAIL-sensitive tumor cells over a broad range of E/T ratios (Fig. 2 A). A titration of both IFN- γ and IFN- α concentrations revealed that as little as 10 pg/ml led to enhanced M ϕ cytotoxicity against the tumor cells; however, GM-CSF did not induce any antitumor activity at any concentration tested (100 ng/ml

to 10 pg/ml; data not shown). Moreover, the IFN-stimulated M ϕ were as effective in killing the tumor target cells as recombinant, soluble TRAIL (LZ-TRAIL [34]). The tumoricidal activity of IFN-stimulated M ϕ was also examined on a TRAIL-resistant human melanoma cell line, WM 164, and a TRAIL-sensitive human melanoma cell line, WM 793 (38). The TRAIL-resistant melanoma (WM 164) was also resistant to the M ϕ -mediated cytotoxicity, whereas the TRAIL-sensitive melanoma (WM 793) was quite sensitive to the cytotoxic activity of either IFN- γ - or IFN- α -stimulated M ϕ (Fig. 2, B and C). The tumoricidal activity of both IFN- γ - or IFN- α -stimulated M ϕ was seen from multiple donors and with other TRAIL-sensitive tumor cells (Table

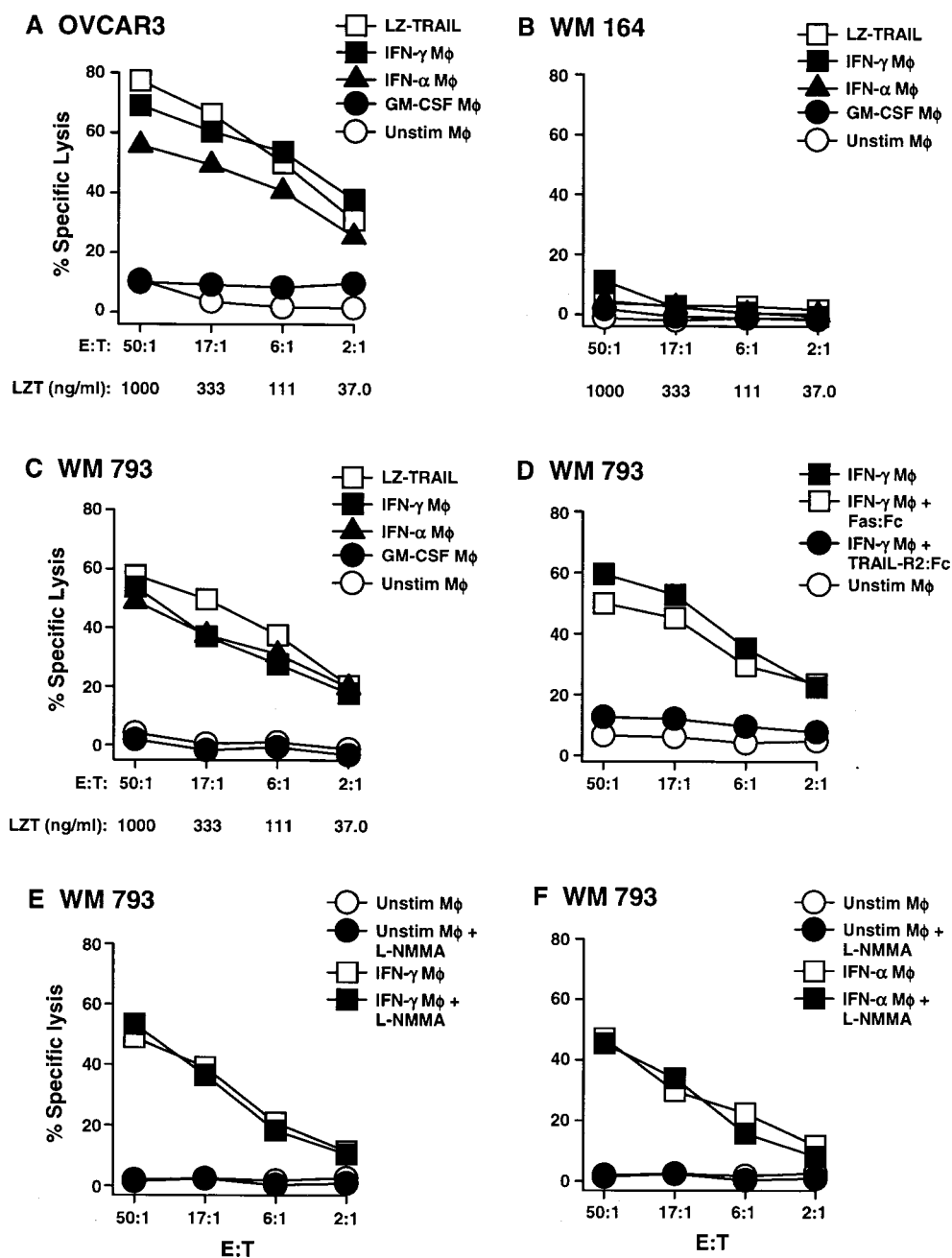


Figure 2. TRAIL-mediated tumoricidal activity by human M ϕ occurs after stimulation with IFN. (A–D) M ϕ were incubated for 12 h in the absence or presence of either GM-CSF, IFN- γ , or IFN- α and then cultured for 8 h with ^{51}Cr -labeled (A) OVCAR3, (B) WM 164, or (C) WM 793 target cells at the indicated E/T ratios. As a positive control, soluble LZ-TRAIL (LZT) was added to target cells at the indicated concentrations. (D) Inclusion of the fusion protein TRAIL-R2:Fc (20 $\mu\text{g}/\text{ml}$) to 12-h IFN- γ -stimulated M ϕ inhibited killing of WM 793 target cells, whereas addition of Fas:Fc (20 $\mu\text{g}/\text{ml}$) did not. (E and F) Addition of the NO inhibitor L-NMMA (300 μM) did not alter the antitumor activity of (E) IFN- γ - or (F) IFN- α -stimulated M ϕ against WM 793 target cells. Data points represent the mean of triplicate wells, and experiments were repeated at least three times with similar results. For clarity, SD bars were omitted from the graphs, but were <10% of the value of all points.

Table I. *Tumoricidal Activity of Cytokine-stimulated M ϕ*

Tumor cell target	Donors tested	M ϕ *				
		Unstimulated	GM-CSF	IFN- γ	IFN- α	LZ-TRAIL ‡
H2126 (lung adenocarcinoma)	3	1.3 \pm 0.9	1.3 \pm 1.3	47.5 \pm 4.0	36.1 \pm 11.9	59.0 \pm 5.9
PC-3 (prostate carcinoma)	4	3.3 \pm 2.0	2.0 \pm 1.7	57.2 \pm 7.6	41.5 \pm 5.2	67.5 \pm 4.2
Colo205 (colon carcinoma)	3	2.3 \pm 2.1	1.8 \pm 4.2	46.1 \pm 3.0	38.4 \pm 7.1	58.3 \pm 3.1
OVCAR3 (ovarian carcinoma)	5	5.8 \pm 4.3	6.3 \pm 3.8	59.4 \pm 13.2	47.5 \pm 7.5	65.4 \pm 11.0
MDA 231 (breast adenocarcinoma)	3	2.0 \pm 0.7	2.4 \pm 0.3	39.6 \pm 6.8	31.2 \pm 8.8	57.7 \pm 10.7
WM 164 (melanoma)	3	0.9 \pm 0.4	2.4 \pm 0.8	9.3 \pm 2.7	4.4 \pm 2.1	3.2 \pm 0.9
WM 793 (melanoma)	5	3.6 \pm 4.2	4.8 \pm 4.2	45.6 \pm 14.6	43.7 \pm 4.5	48.6 \pm 6.8
Normal lung fibroblasts	2	4.1 \pm 0.1	4.1 \pm 2.3	3.2 \pm 1.3	0.9 \pm 1.2	1.0 \pm 1.0
Normal foreskin fibroblasts	2	3.9 \pm 1.2	3.2 \pm 2.0	2.9 \pm 1.0	2.6 \pm 1.4	2.1 \pm 1.1

*Mean percent specific lysis (+SD) at 50:1 M ϕ to target cell ratio.

‡ Mean percent specific lysis (+SD) with 1 μ g/ml LZ-TRAIL.

Means were calculated from experiments performed with M ϕ from the indicated number of donors.

I). Two normal human fibroblast cell types were also tested for sensitivity to the cytokine-stimulated M ϕ and were found to be resistant in all conditions from multiple donors.

Although FasL was not detected on the surface of the activated M ϕ (Fig. 1 A), activated M ϕ have been shown to release soluble FasL from intracellular stores (25). Therefore, to confirm that the observed tumoricidal activity was specific to TRAIL and not FasL, IFN- γ -stimulated M ϕ were pretreated with either TRAIL-R2:Fc (35) or Fas:Fc before adding the tumor cell targets. The TRAIL-R2:Fc reduced target cell death to control (unstimulated M ϕ effector) levels, whereas Fas:Fc did not alter the ability of the IFN- γ -treated M ϕ to mediate tumor lysis (Fig. 2 D). Finally, to determine whether M ϕ NO production contributed to the measured cytotoxic activity, M ϕ were stimulated as above but in the absence or presence of the NO synthase inhibitor, L-NMMA (44). The cytotoxic activity of the IFN- γ - and IFN- α -stimulated M ϕ was not decreased in the presence of L-NMMA compared with M ϕ stimulated in the absence of the inhibitor (Fig. 2, E and F). Similar results were observed with other tumor cell targets (data not shown). Furthermore, analysis of NO production by the M ϕ after 12 h stimulation, as measured by the accumulation of nitrite, revealed no increase in nitrite levels in the culture supernatants with any of the different stimuli compared with unstimulated M ϕ (data not shown). Collectively, these results confirm that the TRAIL expressed on M ϕ mediates the killing of tumor cells, demonstrating a novel mechanism of M ϕ -mediated apoptosis.

TRAIL-expressing M ϕ Induce Apoptotic Cell Death of Sensitive Tumor Cells. Although the release of 51 Cr from the tumor cell targets as measured in Fig. 2 indicates the amount of cell death, it does not discriminate between apoptotic and necrotic cell death. Previous reports have demonstrated

that TRAIL-induced cell death occurs through an apoptotic mechanism (30, 34, 38, 41). To confirm that the tumor cell death induced by the IFN-stimulated M ϕ was mediated through an apoptotic mechanism, the binding of FITC-conjugated annexin V to the tumor cells was analyzed. Annexin V preferentially binds to phosphatidylserine, a phospholipid component of the inner leaflet of the plasma membrane that is rapidly externalized during apoptosis (45, 46). Upon staining the OVCAR3 tumor cells after 6 h incubation with unstimulated or cytokine-stimulated M ϕ (E/T ratio 2:1) or soluble LZ-TRAIL, only those tumor cells incubated with IFN-stimulated M ϕ or LZ-TRAIL were positive for FITC-annexin V binding (Fig. 3), indicating that these cells were dying from the induction of apoptosis. Morphological changes (membrane blebbing and release of apoptotic bodies) were also observed using light microscopy (data not shown).

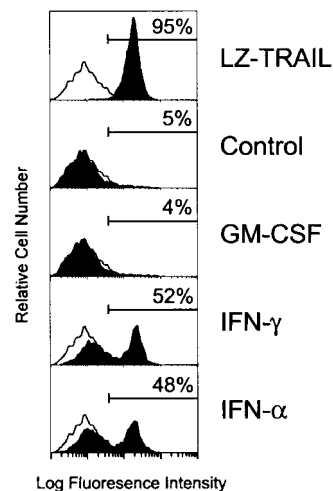


Figure 3. Phosphatidylserine externalization on OVCAR3 tumor cells during apoptosis induced by IFN-stimulated, TRAIL-expressing M ϕ . OVCAR3 tumor cells were cultured for 6 h in medium alone or in the presence of LZ-TRAIL (1 μ g/ml), unstimulated, or cytokine (GM-CSF, IFN- γ , IFN- α [100 ng/ml for 12 h])–stimulated M ϕ (E/T ratio 2:1). Cells were then stained with FITC-annexin V and analyzed by flow cytometry. The percent of FITC-annexin V positive tumor cells is indicated for each condition. Histograms represent 10^4 gated tumor cells. Similar results were seen with M ϕ from three other donors.

The Principal Mechanism of M ϕ Tumoricidal Activity Differs after IFN and LPS Stimulation. Having demonstrated that IFN-stimulated M ϕ express functional TRAIL, the differences between TRAIL and TNF-mediated tumoricidal activity by M ϕ were examined. The results presented in Fig. 1 show that TNF was only expressed on M ϕ after incubation with LPS. Thus, to demonstrate the biologic activity of the cell surface TNF detected after LPS stimulation, 2- and 12-h LPS-stimulated M ϕ were evaluated for the ability to kill the TNF-sensitive target cell, L929 (17). In direct correlation with the flow cytometric results, the 2-h LPS-stimulated M ϕ , but not the 12-h LPS-stimulated M ϕ , killed L929 target cells to levels comparable to soluble TNF (Fig. 4 A). When tested for sensitivity to LZ-TRAIL, the L929 cells were found to be resistant (data not shown). This killing was TNF-specific as demonstrated by the significant inhibition of the lysis of L929 cells upon the addition of TNFR:Fc, but not TRAIL-R2:Fc (Fig. 4 B). To further demonstrate that the cytotoxic activity of IFN- γ -stimulated M ϕ was mediated by TRAIL and not TNF, tumor cell lysis was measured in the presence of TRAIL-R2:Fc or TNFR:Fc. Only TRAIL-R2:Fc, and not TNFR:Fc, inhibited the IFN- γ -stimulated M ϕ from killing the tumor cell targets (Fig. 4 C). Thus, human M ϕ have multiple mechanisms for killing a variety of target cells depending on the activation mechanism.

TRAIL Receptor Expression on Human M ϕ Is Altered by IFN. Because TRAIL can interact with two death-inducing and two non-death-inducing receptors, the distribution of the four known TRAIL receptors on the M ϕ surface using receptor-specific mAbs was investigated. Unstimulated M ϕ expressed both TRAIL-R2 and -R3, whereas the levels of TRAIL-R1 and -R4 were at or below detection (Fig. 5 A). However, mRNA for each of the four TRAIL receptors could be detected by reverse transcription (RT)-PCR analysis (Fig. 5 B). The kinetics of TRAIL, TRAIL-R2, and TRAIL-R3 expression after IFN- γ stimulation were then measured at both the protein level by flow cytometry and the mRNA level by RT-PCR. Increased TRAIL expression could be detected by 2 h on the cell surface after addition of IFN- γ (Fig. 5 C), whereas RT-PCR analysis demonstrated that TRAIL mRNA levels increased by 1 h (Fig. 5 D). No consistent change in TRAIL protein or mRNA levels was detected after GM-CSF treatment compared with untreated M ϕ (Fig. 5, C and D).

Examination of the surface levels of TRAIL-R2 and -R3 during this same 8-h period revealed that IFN- γ -stimulated M ϕ downregulated TRAIL-R2 expression, whereas TRAIL-R3 was only slightly downmodulated (Fig. 5 C). In contrast, incubation with GM-CSF for 8 h resulted in a slight increase in TRAIL-R2 expression. Analysis of mRNA from IFN- γ -stimulated M ϕ revealed that TRAIL-R2 mRNA levels remained relatively constant over the 8-h period, whereas the TRAIL-R2 mRNA levels increased in M ϕ stimulated with GM-CSF over time (Fig. 5 D). No changes in TRAIL-R3 mRNA were observed with IFN- γ or GM-CSF incubation during this period of time (Fig. 5 D), nor were any changes in TRAIL-R1 or -R4 mRNA or

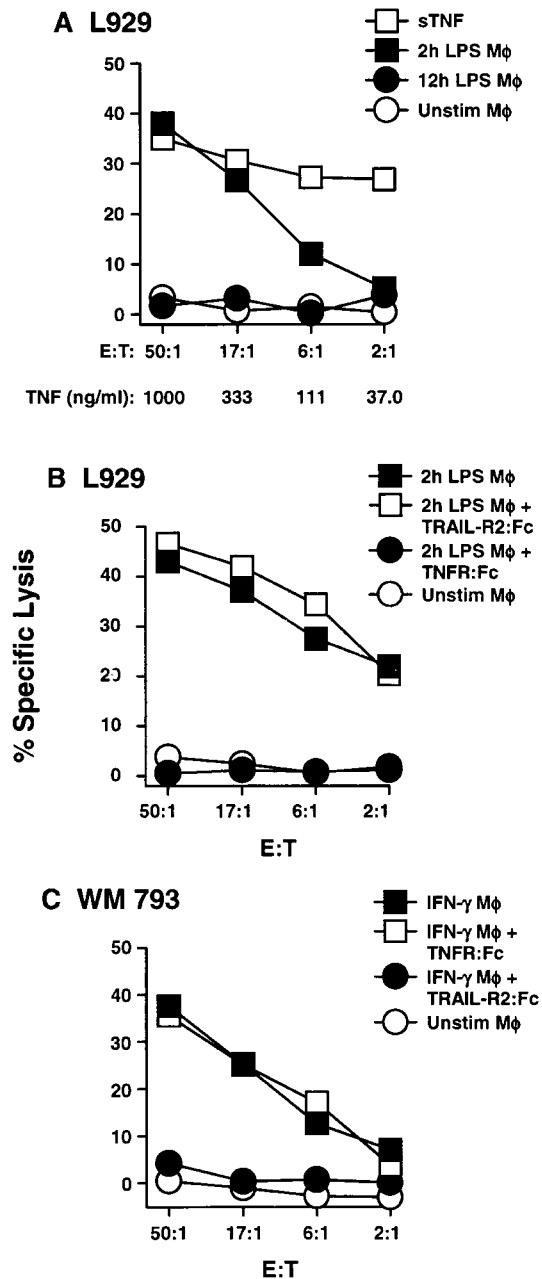


Figure 4. TNF-mediated apoptosis by human M ϕ occurs after stimulation with LPS but not IFN- γ . (A) M ϕ were incubated in the absence or presence of LPS for 2 or 12 h and then cultured for 8 h with ^{51}Cr -labeled L929 target cells at the indicated E/T ratios. As a positive control, soluble (s)TNF was added to target cells at the indicated concentrations. (B) Inclusion of TNFR:Fc (20 $\mu\text{g}/\text{ml}$) to 2-h LPS-stimulated M ϕ inhibited the killing of L929 target cells, whereas addition of TRAIL-R2:Fc (20 $\mu\text{g}/\text{ml}$) did not. (C) Killing of WM 793 tumor cells by M ϕ stimulated with IFN- γ for 12 h can be inhibited by TRAIL-R2:Fc (20 $\mu\text{g}/\text{ml}$), but not TNFR:Fc (20 $\mu\text{g}/\text{ml}$). Data points represent the mean of triplicate wells, and the experiments were repeated at least three times with similar results. For clarity, SD bars were omitted from the graphs, but were <10% of the value of all points.

protein detected (data not shown). Thus, TRAIL expression can be detected on M ϕ within 2 h after IFN- γ stimulation, paired with a concomitant loss in cell surface expression of the cognate death-inducing TRAIL-R2. Similar

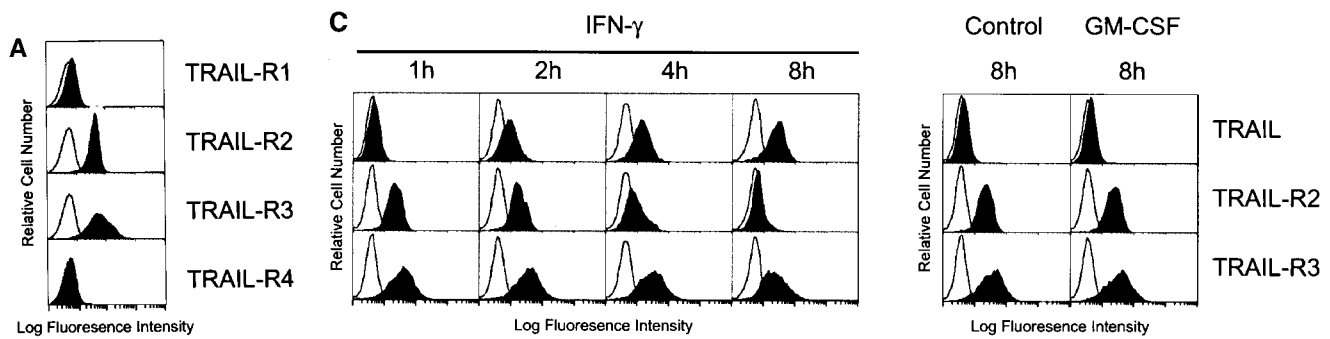


Figure 5. Surface analysis of IFN- γ -stimulated M ϕ reveals simultaneous increase in TRAIL expression with downregulation of TRAIL-R2 expression. (A) Flow cytometric analysis of TRAIL-R1, -R2, -R3, and -R4 expression on unstimulated M ϕ . Filled histograms represent staining by M271 (anti-TRAIL-R1 mAb), M413 (anti-TRAIL-R2 mAb), M430 (anti-TRAIL-R3 mAb), or M444 (anti-TRAIL-R4 mAb), and open histograms represent staining with isotype control mAb. (B) RT-PCR analysis of TRAIL receptor mRNA expression in normal M ϕ . (C) TRAIL, TRAIL-R2, and TRAIL-R3 expression after various times of IFN- γ stimulation. Filled histograms represent staining by M413 (anti-TRAIL-R2 mAb), M430 (anti-TRAIL-R3 mAb), or M181 (anti-TRAIL mAb), and open histograms represent staining with isotype control mAb. For comparison, M ϕ cultured in the absence or presence of GM-CSF for 8 h were also stained for TRAIL, TRAIL-R2, and TRAIL-R3. (D) RT-PCR analysis of TRAIL, TRAIL-R2, and TRAIL-R3 mRNA levels after M ϕ culture in the absence or presence of IFN- γ and GM-CSF. β -actin was used as a control over the same time course. Similar results were observed with M ϕ from two other donors.

results examining TRAIL and TRAIL receptor expression on IFN- α -stimulated M ϕ were also detected (data not shown).

IFN-stimulated M ϕ Are Resistant to TRAIL-induced Death. The loss of TRAIL-R2 expression suggested that peripheral blood M ϕ stimulated with IFN- γ would be resistant to TRAIL-mediated death. Thus, M ϕ were cultured in the absence or presence of GM-CSF or IFN- γ and then examined for sensitivity to LZ-TRAIL. M ϕ treated with GM-CSF displayed increased sensitivity to TRAIL-induced death compared with untreated M ϕ ; in contrast, IFN- γ treatment significantly decreased TRAIL-induced M ϕ death (Fig. 6 A). Similar results were obtained with IFN- α -stimulated M ϕ (data not shown). No significant changes in M ϕ sensitivity to TRAIL were seen with the other cytokines tested (IL-1, IL-2, IL-3, IL-4, IL-6, IL-7, IL-10, IL-12, and IL-15; data not shown) compared with untreated M ϕ . These results suggest that M ϕ stimulated with IFN- γ minimize TRAIL-mediated suicide or fratricide with the downregulation of TRAIL-R2 surface levels. However, it remains possible that additional mechanisms may contribute to the protection of the M ϕ from TRAIL-induced death.

The rendering of M ϕ resistant to TRAIL-mediated apoptosis after stimulation with IFN- γ suggested that the tumor cell targets used in Fig. 2 could be affected in a similar fashion upon culture with IFN- γ . OVCAR3 tumor cells were incubated in the absence or presence of IFN- γ for 12 h, and then tested for sensitivity to LZ-TRAIL or IFN- γ -stimulated M ϕ . In contrast to the M ϕ , the sensitivity of OVCAR3 tumor cells was not altered after incubation with IFN- γ (Fig.

6, B and C). Similar results were seen with other tumor cells (WM 793 and PC-3; data not shown). These results imply that not all cell types respond to IFN- γ by gaining resistance to TRAIL-induced death as observed with the M ϕ .

Discussion

M ϕ not only influence the activities of other immune and nonimmune cells in the body, but also function as effector cells under a variety of conditions (1). Activated M ϕ display potent tumoricidal activity against several different tumor cell types (7, 47-49). The results presented here demonstrate that one of the mechanisms by which M ϕ kill tumor cells is through expression of TRAIL. M ϕ stimulation with either IFN- γ or IFN- α resulted in the rapid expression of TRAIL on the cell surface, but not FasL or TNF. Because TRAIL mediates apoptosis in a high percentage (approximately two thirds) of hematopoietic and nonhematopoietic cell types (30, 36, 38, 42), M ϕ have the potential to mediate apoptosis of a broad range of tumor cell types via TRAIL. Expression of TRAIL appeared to be specific to the IFNs, as M ϕ stimulation with either IL-1, IL-2, IL-3, IL-4, IL-6, IL-7, IL-10, IL-12, IL-15, or GM-CSF resulted in no detectable TRAIL expression (data not shown). Interestingly, a concomitant loss of TRAIL-R2 (and to a lesser extent TRAIL-R3) expression was detected upon IFN stimulation, rendering the M ϕ resistant to TRAIL-mediated death. To our knowledge, this is the first demonstration of inducible TRAIL expression on a partic-

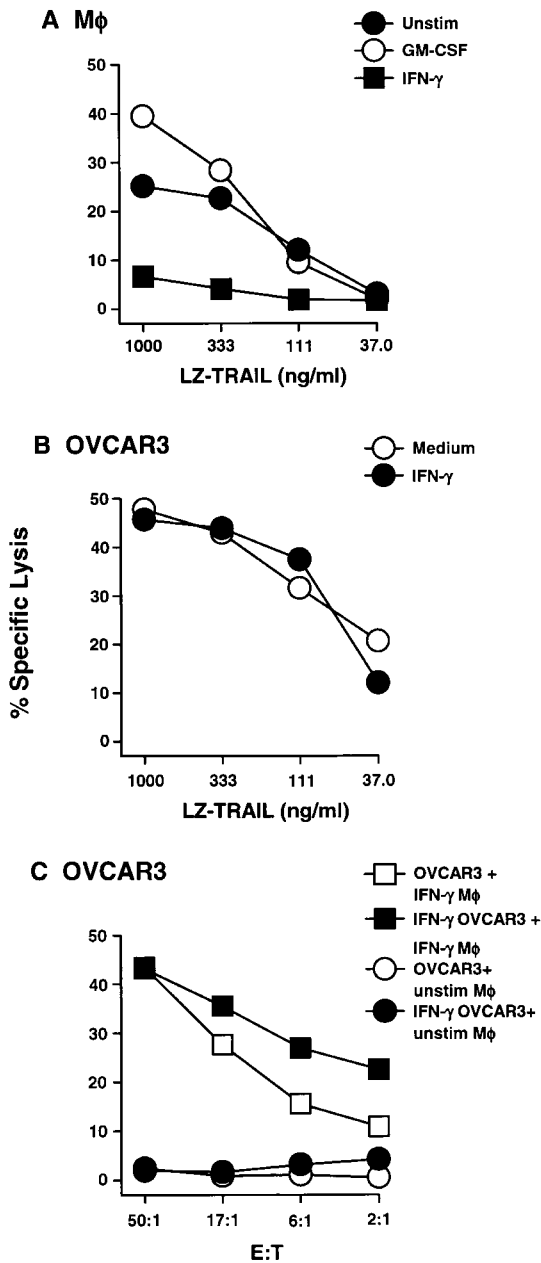


Figure 6. IFN- γ stimulation renders M ϕ resistant to TRAIL-induced death, but not tumor cell targets. (A) Peripheral blood M ϕ were incubated for 12 h in the absence or presence of GM-CSF and IFN- γ , then tested for sensitivity to LZ-TRAIL. (B) OVCAR3 tumor cells were incubated for 12 h in the absence or presence of IFN- γ , then tested for sensitivity to LZ-TRAIL. (C) OVCAR3 tumor cells were cultured for 12 h in the absence or presence of IFN- γ , then incubated with unstimulated or IFN- γ -stimulated M ϕ . Percent specific lysis was measured by ^{51}Cr release after 8 h, and each data point represents the mean of triplicate wells. For clarity, SD bars were omitted from the graphs, but were <10% of the value of all points. These experiments were repeated at least three times with similar results using M ϕ from at least three different donors.

ular human peripheral blood cell population, as well as modulation of the death-inducing TRAIL-R2 by a proinflammatory cytokine on the same cell population.

Although the expression of TRAIL on the IFN-stimu-

lated M ϕ is critical for the tumoricidal activity in our assay system, the sensitivity of the tumor cell to TRAIL-induced apoptosis is also an essential component of this phenomenon, as demonstrated by the fact that the tumor cell lines and normal cells that were resistant to TRAIL-mediated apoptosis were also resistant to TRAIL-expressing M ϕ . The identification of two TRAIL receptors with death-inducing ability and two without led to the initial hypothesis that the expression of TRAIL-R3 and/or -R4 conferred resistance to TRAIL-induced death (32, 33, 37). However, it is important to note that this hypothesis was formulated from reports examining the distribution of TRAIL receptor mRNA in several normal tissues and tumor cell lines and from experiments where TRAIL-R3 or -R4 was overexpressed in transfected cells. Most of the tumors used in this study express TRAIL-R3 and/or -R4 (38, 42). When the OVCAR3, WM793, and PC3 tumor cells were cultured with IFN- γ before incubation with LZ-TRAIL or IFN-stimulated M ϕ , the TRAIL receptor levels remained unchanged, and no significant change in the level of TRAIL sensitivity was observed (data not shown). Thus, the differences in sensitivity of the tumor cells to the TRAIL expressed on the M ϕ or the recombinant TRAIL added in solution are probably regulated by a variety of molecular mechanisms, both inside the cell and at the surface.

While our results focused on the tumoricidal activity of TRAIL-expressing M ϕ , previous reports have shown these cells can also produce cytotoxic inorganic oxidants, such as NO (10, 11). A role for NO in tumor cell killing has been documented for both human and mouse activated macrophages (4, 44, 50), where the toxicity of NO is mediated via mitochondrial damage, inhibition of DNA synthesis, and disruption of the tricarboxylic acid cycle, ultimately resulting in apoptosis (51, 52). Although murine macrophages release high levels of NO after either LPS or IFN- γ stimulation, studies with human peripheral blood M ϕ have reported contradictory findings (10). In some reports, M ϕ stimulated with either LPS or IFN- γ (or in combination) failed to release significant levels of NO (12–14, 44, 53), whereas others have reported that IFN- α stimulation results in a slight increase in NO production (54). In our studies, addition of the NO inhibitor L-NMMA to the cytotoxicity assays did not decrease the ability of the IFN- γ - and IFN- α -stimulated M ϕ to kill the tumor cell targets. Moreover, analysis of the culture supernatants for nitrites revealed no increase after 12 h stimulation with GM-CSF, IFN- γ , or IFN- α (data not shown). These observations, coupled with the fact that TRAIL-R2:Fc completely inhibited the tumoricidal activity of the M ϕ to background (unstimulated M ϕ) levels, imply that TRAIL is the primary mediator of the tumoricidal activity after IFN stimulation. Coexpression of TRAIL and TNF, and perhaps other unidentified death-inducing molecules, would theoretically increase both the cytolytic potential of the M ϕ and the range of different tumor targets susceptible to M ϕ -mediated death.

The importance of IFN in the management of spontaneously arising tumors was recently demonstrated in vivo us-

ing mice that lack sensitivity to IFN- γ (55). Compared with wild-type mice, the IFN- γ -insensitive mice develop tumors more rapidly and with greater frequency when challenged with a chemical carcinogen. Part of this "IFN effect" is via the interaction of the IFN with the tumor cells by enhancing the tumor cell tumorigenicity through heightened MHC class I expression. IFN- γ may also enhance an innate antitumor mechanism through the induction of TRAIL on cells of the M ϕ lineage (55). Although our data suggest that M ϕ would confer this antitumor activity, further studies are required to determine if other cell types, such as NK cells, neutrophils, and dendritic cells, are also able to express TRAIL after IFN stimulation (56).

Finally, in addition to a role in tumoricidal activity, these

data suggest that M ϕ TRAIL-expression may contribute to other physiologic and pathologic situations, such as the AICD of T cells during HIV infection. Recently, Katsikis et al. (57, 58) have demonstrated that activation-induced peripheral blood T cell apoptosis in HIV-infected individuals was Fas independent, and a potential role for TRAIL in this phenomenon was identified. It was observed that a blocking mAb to TRAIL could inhibit the AICD of T cells in a mixed population of HIV⁺ PBMCs (58). However, it was unclear which PBMC subset was expressing TRAIL and responsible for death of the T cells. Thus, the results presented here may provide an explanation for these experimental observations, as well as a basis for examining other activities mediated by activated M ϕ .

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Address correspondence to Neil A. Fanger, Department of Discovery Research, Immunex Corporation, 51 University St., Seattle, WA 98101. E-mail: nfanger@immunex.com

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