

# Autocrine regulation of *mda-7/IL-24* mediates cancer-specific apoptosis

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**A noteworthy aspect of melanoma differentiation-associated gene-7/interleukin-24 (*mda-7/IL-24*) as a cancer therapeutic is its ability to selectively kill cancer cells without harming normal cells. Intracellular MDA-7/IL-24 protein, generated from an adenovirus expressing *mda-7/IL-24* (*Ad.mda-7*), induces cancer-specific apoptosis by inducing an endoplasmic reticulum (ER) stress response. Secreted MDA-7/IL-24 protein, generated from cells infected with *Ad.mda-7*, induces growth inhibition and apoptosis in surrounding noninfected cancer cells but not in normal cells, thus exerting an anti-tumor “bystander” effect. The present studies reveal a provocative finding that recombinant MDA-7/IL-24 protein can robustly induce expression of endogenous *mda-7/IL-24*, which generates the signaling events necessary for bystander killing. To evaluate the mechanism underlying this positive autocrine feedback loop, we show that MDA-7/IL-24 protein induces stabilization of its own mRNA without activating its promoter. Furthermore, this posttranscriptional effect depends on *de novo* protein synthesis. As a consequence of this autocrine feedback loop MDA-7/IL-24 protein induces sustained ER stress as evidenced by expression of ER stress markers (BiP/GRP78, GRP94, GADD153, and phospho-eIF2 $\alpha$ ) and reactive oxygen species production, indicating that both intracellular and secreted proteins activate similar signaling pathways to induce apoptosis. Thus, our results clarify the molecular mechanism by which secreted MDA-7/IL-24 protein (generated from *Ad.mda-7*-infected cells) exerts cancer-specific killing.**

bystander antitumor activity | endoplasmic reticulum stress | reactive oxygen species | mRNA stabilization | cancer-specific killing

**M**elanoma differentiation-associated gene-7 (*mda-7*) was first identified as a gene associated with terminal differentiation of metastatic human melanoma cells (1, 2). Based on its structure, chromosomal location, and biochemical properties, *mda-7* has been classified as a member of the IL-10 family of cytokines that includes IL-10, IL-19, IL-20, IL-22, and IL-26 and has been redesignated IL-24 (3, 4). When expressed at supraphysiological levels, by means of an adenoviral (Ad) expression system (*Ad.mda-7*), MDA-7/IL-24 induces growth suppression and apoptosis in a broad spectrum of human cancer cells, including those from melanoma, malignant glioma, fibrosarcoma, and carcinomas of the breast, cervix, colon, rectum, liver, lung, ovary, and prostate, without exerting any deleterious effects on their normal counterparts (5–13). A phase I trial evaluating *Ad.mda-7* (INGN 241) activity by intratumoral injection in patients with advanced solid tumors was performed, indicating that *mda-7/IL-24* was safe and could induce as much as 70% apoptosis in tumors after a single injection of recombinant virus and that multiple injections elicited clinical responses (14–16). The successful transition of *Ad.mda-7* into the clinic in a phase I clinical trial reinforces the hypothesis that *mda-7/IL-24* is safe and affords remarkable potential as a cancer gene therapeutic. Moreover, secreted MDA-7/IL-24 protein, generated from *Ad.mda-7*-infected cells, promotes antiangiogenic,

immunostimulatory, radiosensitizing and “bystander” antitumor activities (6, 11, 17, 18).

*mda-7/IL-24* expression is detected in human tissues and cells associated with the immune system such as spleen, thymus, peripheral blood leukocytes, and normal melanocytes (19). Secreted MDA-7/IL-24 stimulates monocytes and specific populations of T lymphocytes and promotes proinflammatory cytokine production. When expressed at low, presumably physiological levels, MDA-7/IL-24 binds to currently recognized MDA-7/IL-24 receptor complexes consisting of two sets of heterodimeric chains, IL-20R1/IL-20R2 or IL-22R1/IL-20R2 (20–22). Most human tissues express the IL-20R1/IL-20R2 complex. IL-22R is found in a few tissues lacking IL-20R2, such as adult and fetal liver, colon, small intestine, and pancreas. A functional set of cell surface receptors can also be found in the majority of human tumor cells (23). Upon ligand binding, both receptors induce activation of STAT3 (20–22). However, our previous studies demonstrate that activation of the JAK/STAT pathway is dispensable for *Ad.mda-7*-induced apoptosis, and cell death triggered by intracellular MDA-7/IL-24 protein occurs through a receptor-independent mechanism (23). Intracellular MDA-7/IL-24 localizes to the endoplasmic reticulum (ER) and induces an ER stress response, also known as unfolded protein response (UPR), thus eliciting tumor-specific apoptosis (24).

A highly conserved UPR signal transduction pathway is activated by ER stress caused by misfolded protein accumulation (25). The UPR can be triggered by unfolding proteins in the lumen of the ER, resulting in *de novo* synthesis of ER proteins (such as the “glucose-regulated proteins” BiP/GRP78 and GRP94) that assist in protein folding. Cell death is an inevitable consequence of persistent ER stress, and ER stress can lead to apoptosis through both mitochondria-dependent and -independent pathways. Production and integration of apoptotic signals may occur in the ER, generating the death response. Mechanisms involved in this process include PERK/ $\alpha$ -subunit of eukaryotic translation initiation factor-2 (eIF2 $\alpha$ )-dependent induction of the proapoptotic transcription factor GADD153, BAK/BAX-regulated Ca<sup>2+</sup> release from the ER, and cleavage and activation of procaspase 12 (25).

*Ad.mda-7* infection of cancer cells induces growth arrest and DNA damage-inducible (GADD) gene family, classically associated with the stress response, including the ER stress pathways (9, 26). Induction of GADD genes and further upstream events such as activation of p38<sup>MAPK</sup> and PKR are promoted by *mda-7/IL-24* in a transformed cell-specific manner, and these events occur independently of the glycosylation of MDA-7/IL-24. *Ad.mda-7* also

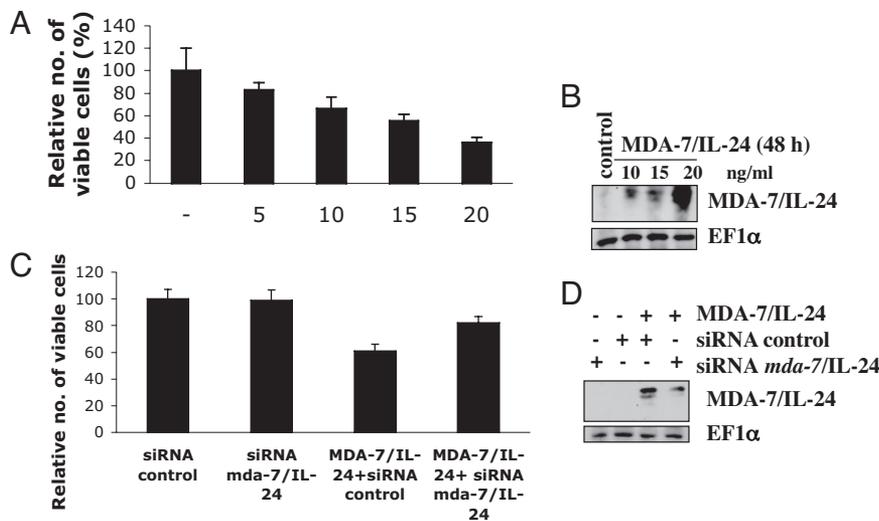
Author contributions: M.S. and P.B.F. designed research; M.S., Z.-z.S., and I.V.L. performed research; P.G. and P.D. contributed new reagents/analytic tools; M.S. and P.B.F. analyzed data; and M.S., D.S., and P.B.F. wrote the paper.

The authors declare no conflict of interest.

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**Fig. 2.** Induction of endogenous *mda-7/IL-24* is required for growth-inhibitory activity of recombinant MDA-7/IL-24. (A) DU-145 cells were treated with MDA-7/IL-24 (5, 10, 15, 20 ng/ml) for 72 h, and cell viability was evaluated by standard 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyl tetrazolium bromide (MTT) assay. (B) DU-145 cells were treated with MDA-7/IL-24 (5, 10, 15, 20 ng/ml) protein for 48 h, and expression of MDA-7/IL-24 and EF-1 $\alpha$  protein was analyzed by Western blotting. (C) HeLa cell viability assay showing a significant decrease in the number of dead cells in cultures treated with *mda-7/IL-24* siRNA plus MDA-7/IL-24 compared with MDA-7/IL-24 plus control siRNA. (D) HeLa cells were transfected with control or *mda-7/IL-24* siRNA (100 nmol/liter) and untreated or treated with recombinant MDA-7/IL-24 protein, and expression of MDA-7/IL-24 protein was analyzed by Western blotting 48 h later.

expression in both DU-145 and P69 cells without affecting the expression of housekeeping genes GAPDH and  $\beta$ -actin. These findings suggest that exogenous MDA-7/IL-24 protein induces endogenous *mda-7/IL-24* expression in normal and cancer cells.

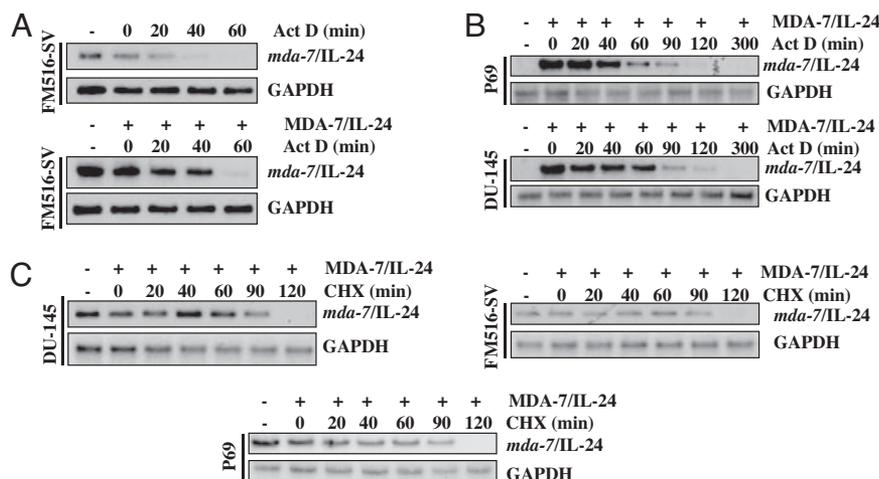
**Autocrine Feedback Loop Is Required for Growth-Inhibitory effect of MDA-7/IL-24.**

A direct correlation was evident with level of induction of endogenous MDA-7/IL-24 protein and reduction in tumor cell viability upon MDA-7/IL-24 treatment (Fig. 2 A and B). Exposing DU-145 cells to increasing concentrations of MDA-7/IL-24 resulted in a nonlinear dose-dependent induction of endogenous MDA-7/IL-24 protein. Similarly, MDA-7/IL-24 exposure also reduced viability of DU-145 cells in a concentration-dependent manner. To assess the importance of autoregulation of *mda-7/IL-24* in mediating growth-inhibitory activity, an siRNA approach was used to knock down expression of MDA-7/IL-24. HeLa cells were chosen for these studies because of high transient transfection efficiency. HeLa cells were transiently transfected with *mda-7/IL-24* siRNA or control siRNA and then challenged with recombinant MDA-7/IL-24 protein (10 ng/ml). Western blotting confirmed the ability of *mda-7/IL-24* siRNA to knock down protein expression (Fig. 2D). A significant reduction in the MDA-7/IL-24 protein level was observed with *mda-7/IL-24* siRNA but not with control siRNA. As a corollary, *mda-7/IL-24* siRNA, but not control siRNA, significantly protected cancer cells from

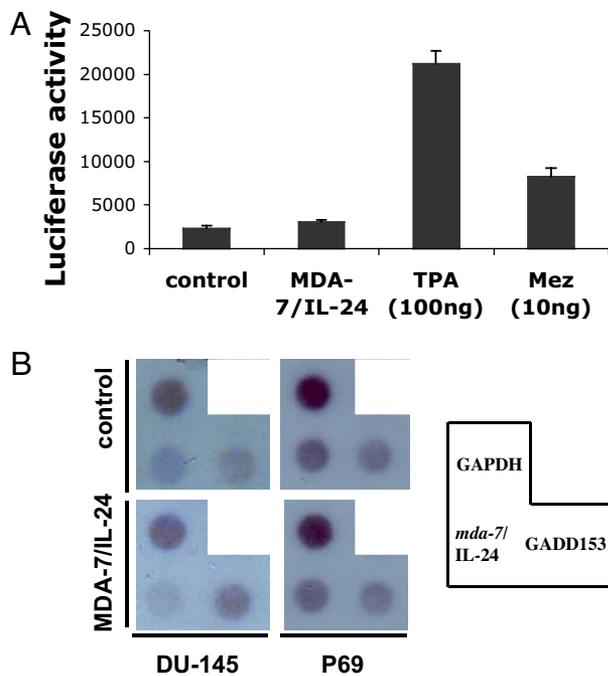
MDA-7/IL-24-induced reduction in viability, indicating a direct correlation between endogenous expression of MDA-7/IL-24 and recombinant MDA-7/IL-24-mediated killing (Fig. 2C).

**Recombinant MDA-7/IL-24 Increases *mda-7/IL-24* mRNA Stability.**

To explore the molecular mechanism by which MDA-7/IL-24 protein induces endogenous *mda-7/IL-24* expression, we analyzed *mda-7/IL-24* promoter function and *mda-7/IL-24* mRNA stability because other known inducers of *mda-7/IL-24* work predominantly by stabilizing *mda-7/IL-24* mRNA. Indeed, the 3'-UTR region of *mda-7/IL-24* mRNA contains several AU-rich elements (ARE), conferring rapid mRNA turnover, and the robust induction of *mda-7/IL-24* mRNA during terminal differentiation of melanoma cells involves stabilization of its mRNA rather than direct transcriptional control. FM516-SV cells, which endogenously express *mda-7/IL-24* mRNA, were used to determine whether treatment with MDA-7/IL-24 altered the half-life of this mRNA. FM516-SV cells were treated with vehicle or MDA-7/IL-24 protein (20 ng/ml) for 48 h followed by actinomycin D (Act D) (0.5  $\mu$ g/ml). Cells were then incubated further and harvested at different times points. Total RNA was isolated, and *mda-7/IL-24* mRNA expression was examined by Northern blotting to measure mRNA decay rates. Without any treatment, the half-life of *mda-7/IL-24* mRNA was <20 min (Fig. 3A Upper). MDA-7/IL-24 treatment extended the half-life of the mRNA to  $\approx$ 40 min



**Fig. 3.** Recombinant MDA-7/IL-24 protein increases stability of endogenous *mda-7/IL-24* mRNA that requires *de novo* protein synthesis. (A) FM516-SV cells were untreated or treated with MDA-7/IL-24 (20 ng/ml) for 48 h and then with Act D (0.5  $\mu$ g/ml). At the indicated times after treatment, total RNA was extracted and subjected to Northern blotting for the indicated mRNAs. (B) P69 and DU-145 cells were treated with MDA-7/IL-24 for 48 h and then with Act D (0.5  $\mu$ g/ml). At the indicated times after treatment, total RNA was extracted and subjected to Northern blot analysis for the indicated mRNAs. (C) P69, FM516-SV, and DU-145 cells were treated with or without MDA-7/IL-24 protein for 48 h and then with CHX (10  $\mu$ M). At the indicated times after treatment, total RNA was extracted and subjected to Northern blot analysis for the indicated mRNAs.

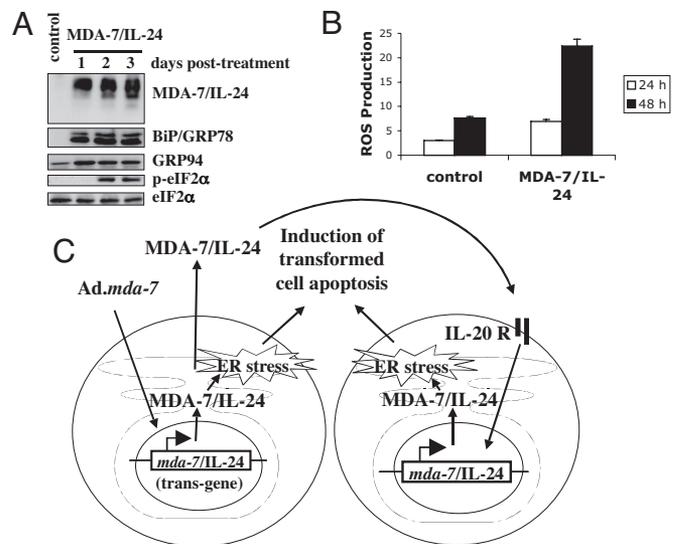


**Fig. 4.** Induction of *mda-7/IL-24* by MDA-7/IL-24 protein occurs posttranscriptionally. (A) HeLa cells were transfected with *mda-7*-Prom-luc and treated with MDA-7/IL-24 protein (20 ng/ml), TPA, or MEZ. Luciferase activity was measured after 48 h. (B) Nuclei were prepared from P69 and DU-145 cells. The isolated nuclei were used to label preinitiated RNA transcription with [ $\alpha$ - $^{32}$ P]UTP *in vitro*, and the purified RNA was hybridized to a dot blot carrying an equivalent amount of cDNA probes. The transcription rate of GAPDH served as control.

(Fig. 3A Lower). In DU-145 and P69 cells, no basal expression of *mda-7/IL-24* mRNA was detected. Combination of MDA-7/IL-24 and Act D treatment revealed that, similar to FM516-SV cells, the half-life of *mda-7/IL-24* mRNA was also  $\approx 40$  min in these cells (Fig. 3B).

We next analyzed the role of new protein synthesis in regulating MDA-7/IL-24-induced *mda-7/IL-24* mRNA expression. FM516-SV and DU-145 cells were treated with cycloheximide (CHX, 10  $\mu$ M) for different periods of time after treatment with MDA-7/IL-24 protein for 48 h. The cells were harvested, total RNA was extracted, and Northern blotting was performed by using a radiolabeled *mda-7/IL-24* cDNA probe (Fig. 3C). CHX treatment increased the *mda-7/IL-24* mRNA half-life to  $\approx 120$  min (Fig. 3C). These findings indicate that MDA-7/IL-24 protein increases the stability of *mda-7/IL-24* mRNA. Inhibition of protein synthesis further increased the mRNA half-life, indicating that new proteins binding to the mRNA and reducing stability may play a critical role in maintaining low (or undetectable) basal expression of *mda-7/IL-24* mRNA. Our data demonstrate that protein synthesis is necessary for promoting MDA-7/IL-24-mediated *mda-7/IL-24* message stability.

**Induction of *mda-7/IL-24* mRNA by MDA-7/IL-24 Protein Occurs Independently of *mda-7/IL-24* Promoter Activation.** To analyze potential transcriptional induction of *mda-7/IL-24* mRNA by MDA-7/IL-24 protein, we performed *mda-7/IL-24* promoter analysis by using a luciferase reporter-based approach. A plasmid was generated containing  $\approx 2.2$  kb of the *mda-7/IL-24* promoter sequence upstream of a luciferase gene (*mda-7*-Prom-luc). HeLa cells were transfected with *mda-7*-Prom-luc along with a  $\beta$ -galactosidase expression plasmid and treated with MDA-7/IL-24 protein (20 ng/ml) for 48 h. Luciferase and  $\beta$ -galac-



**Fig. 5.** Recombinant MDA-7/IL-24 protein triggers ER stress and ROS production. (A) Cells were treated with recombinant MDA-7/IL-24 protein (20 ng/ml), and changes in BiP/GRP78, GRP94, and phospho-eIF2 $\alpha$  proteins were evaluated by using Western blot analyses. (B) Cells were untreated or treated with MDA-7/IL-24 protein (20 ng/ml) for 24 and 48 h, and ROS production was determined as described. Results are average from at least three independent experiments. (C) Model illustrating the possible molecular mechanism of cancer cell-selective apoptosis induction by recombinant MDA-7/IL-24 protein.

tosidase activity was measured, and luciferase activity was normalized by  $\beta$ -galactosidase activity (Fig. 4A). Phorbol 12-tetradecanoate 13-acetate (TPA) and mezerein (MEZ), known inducers of *mda-7/IL-24* expression, significantly increased *mda-7/IL-24* promoter activity. MDA-7/IL-24 protein treatment did not augment promoter activity beyond the basal level. Nuclear run-on assays substantiated the results obtained by promoter analysis (Fig. 4B). MDA-7/IL-24 did not elevate the transcription of *mda-7/IL-24* mRNA in isolated nuclei from DU-145 or P69 cells, whereas the transcription of GADD153, a known downstream target of MDA-7/IL-24, was augmented only in DU-145. These results confirm that *mda-7/IL-24* is induced by MDA-7/IL-24 protein at a posttranscriptional level.

**MDA-7/IL-24 Protein Stimulates the UPR Pathway and Generates ROS.** We reported that Ad.*mda-7* induces an UPR pathway (17, 18, 24). The robust expression of MDA-7/IL-24 protein and the subsequent killing effect suggest that similar to Ad.*mda-7*, MDA-7/IL-24 may induce an UPR pathway. To determine whether chaperone proteins were activated after MDA-7/IL-24 treatment, we measured steady-state levels of specific proteins (BiP/GRP78, and GRP94) up-regulation of which frequently correlates with UPR. We also determined the phosphorylation of eIF2 $\alpha$ , a key downstream event of UPR that mediates inhibition of protein translation (Fig. 5A). In addition, we measured *de novo* synthesis of GADD153, a proapoptotic transcription factor (Fig. 4B). Enhanced BiP/GRP78 and GRP94 protein levels were apparent at 1 or 2 days after MDA-7/IL-24 treatment, indicating selective modulation of specific chaperone proteins (Fig. 5A). In addition, MDA-7/IL-24 treatment induced phosphorylation of eIF2 $\alpha$  (Fig. 5A).

We determined the time course of mitochondrial changes (ROS generation) after treatment of DU-145 cells with MDA-7/IL-24 protein. Cells were treated with MDA-7/IL-24 protein, collected at 24 and 48 h, and stained for ROS production with dichlorofluorescein diacetate (DCFH-DA). Fig. 5B shows that MDA-7/IL-24 protein increased ROS production similar to Ad.*mda-7* infection.



## Materials and Methods

**Cell lines, Viability and Apoptosis Assays, and ROS Production.** Cancer cells, DU-145, A549, PC-3, HO-1, and HeLa, and normal cells, FM516-SV immortalized normal human melanocytes, P69 immortalized normal human prostate epithelial cells, and PHFA, were cultured as described (11–13). MTT assays and apoptosis assays were performed by flow cytometry (11–13, 18). For analysis of ROS production, cells were stained with DCFH-DA and analyzed by flow cytometry (12).

**Protein Purification.** Ad.MDA-7/IL-24 with a His<sub>6</sub> tag on the C terminus (Ad.His-*mda-7*) was generated by standard protocols. Immortal primary human fetal astrocytes (IM-PHFA) (9) were infected with Ad.His-*mda-7* (100 pfu per cell) for 2 h, washed, and then cultured in complete growth medium. IM-PHFA were used because they are well infected and generate high levels of MDA-7/IL-24 protein that is secreted without being killed (9). After 96 h, medium was collected and centrifuged to remove any cell debris. Supernatant was mixed with a nickel-nitrilotriacetic (Ni-NTA) acid slurry and incubated overnight to allow binding of MDA-7/IL-24 to the Ni-NTA beads. After 24 h, the Ni-NTA slurry-containing medium was allowed to pass through a column to collect the beads. The beads were washed with sodium phosphate buffer (pH 6.0), and MDA-7/IL-24 protein was eluted in sodium phosphate buffer (pH 9.0) containing 250 mM imidazole. The protein was dialyzed against DMEM to remove imidazole. The size and purity of the recombinant MDA-7/IL-24 protein were determined by Coomassie blue staining gel and Western blot analysis (6).

**Northern Blot Analysis and RT-PCR.** Total RNA was isolated by using the RNeasy kit (Qiagen). Reverse transcription (RT) was performed on 5  $\mu$ g of total RNA with an oligo(dT) primer (18, 23). cDNA corresponding to 5 ng of total RNA was amplified for 35 cycles by PCR with specific primers for *mda-7/IL-24* (23). For Northern blotting, 15  $\mu$ g of total RNA was denatured, electrophoresed in a 1.2% agarose gel with 3% formaldehyde, and transferred onto a nylon membrane (11). The blots were probed with an [ $\alpha$ -<sup>32</sup>P]dCTP full-length human *mda-7/IL-24*

cDNA probe and then stripped and reprobed with an [ $\alpha$ -<sup>32</sup>P]dCTP human *gapdh* probe (11).

**Western Blot Analysis.** Protein extracts were prepared with RIPA buffer containing a mixture of protease inhibitors as described (6, 11). Fifty micrograms of protein was applied to a 12% SDS/PAGE and transferred to nitrocellulose membranes. The membranes were probed with polyclonal antibodies to MDA-7/IL-24, eIF2 $\alpha$ , BiP/GRP78, p-eIF2 $\alpha$ , and GRP94.

**siRNA Assay.** HeLa cells were transfected with 100 nM *mda-7/IL-24* siRNA (Dharmacon, Inc.) or control siRNA (38). After 24 h of siRNA transfection, cells were washed, and each plate was split in two and untreated or treated with MDA-7/IL-24 protein. Cells were harvested 48 h later, counted, and cell lysates were prepared and analyzed for MDA-7/IL-24 protein expression by Western blotting (6).

**Nuclear Run-On Assays.** Nuclear run-on assays for *mda-7/IL-24*, *gapdh*, and *gadd153* were performed as described in ref. 39.

**Promoter Analysis.** A plasmid containing  $\approx$ 2.2 kb of *mda-7/IL-24* promoter sequence upstream of luciferase gene (*mda-7*-Prom-luc) was generated (35). HeLa cells were transfected with *mda-7*-Prom-luc along with a  $\beta$ -galactosidase expression plasmid by using Lipofectamine 2000 (Invitrogen) according to the manufacturer's protocol. The cells were treated with MDA-7/IL-24 protein for 48 h, and luciferase and  $\beta$ -galactosidase activity was measured (35). Luciferase activity was normalized by  $\beta$ -galactosidase activity.

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