Tumor Necrosis Factor α Stimulates Her-2 Cleavage by Activated Caspase-8

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Key Words
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Abstract
Background/Aim: Her-2 over-expression has been correlated with a poor prognosis in patients with breast cancer. Now, we explored the effect of TNF-α treatment and/or NFκB activation on Her-2 expression in MCF-7 breast adenocarcinoma cells. Methods: Stably transfected MCF-7 cell lines with pcDNA3.0, IκBα MT, c-FLIP/control shRNA were established by FuGENE with the supplementation of G418 (500μg/ml). Western blot and Real-time PCR were applied to assess the expression levels of protein and mRNA of target gene. In addition, caspase-8 activity was evaluated by the incubation with a caspase-8 fluorogenic substrate, Ac-IEPD-AMC using a spectrofluorometer. Results: It was uncovered that Her-2 was a new substrate for caspase-8 and that tumor necrosis factor α (TNF-α) stimulation resulted in a caspase-8-dependent Her-2 cleavage in MCF-7 breast adenocarcinoma cells defective for nuclear factor κB (NFκB) activation. We demonstrated that the antiapoptotic transcription factor NFκB counteracted this cleavage through the induction of caspase-8 inhibitor, c-FLIP. Conclusion: we propose a novel mechanism in which NFκB functions as a new antiapoptotic factor by counteracting TNF-α-triggered Her-2 cleavage.

Introduction

Tumor necrosis factor α (TNF-α) can regulate proapoptotic and antiapoptotic signaling pathways through interaction with two receptors, TNF receptor 1 (TNF-R1) and TNF receptor 2 (TNF-R2) [1]. Following TNF-α triggering, TNF receptor is trimerized, and subsequently,
various signaling proteins were recruited to the receptor cytoplasmic domains [1-3]. Activation of TNF-R2 yields an anti-apoptotic and proinflammatory cascade [4], while TNF-R1 trimerization results in apoptosis through the recruitment and activation of caspase-8 as blocked the activation of the anti-apoptotic and proinflammatory transcription factor, nuclear factor κB (NFκB) [4-6]. Furthermore, NFκB target genes, including caspase inhibitors such as c-IAPs, XIAP, and c-FLIP; Bcl-2 family members; and other proteins like A20, counteract TNF-α-induced apoptosis [7, 8].

Human cancers are defined as an upregulated proliferation and a downregulated apoptosis. Among the proteins implicated in this dysregulation, Her-2 functions as a pivotal role in breast cancer cells [9-11]. Her-2 oncogene induces the expression of a transmembrane receptor protein structurally similar to epidermal growth factor receptor (EGFR). It was reported that Her-2 was over-expressed in approximately one-third of primary breast cancer, which was correlated with a poor prognosis and decreased overall or disease-free survival [9-11]. Her-2 over-expression is potential to enhance proliferation, metastasis, and prosurvival signals in breast cancer cell lines and to induce resistance to hormonal therapy [12, 13]. Moreover, anti-Her-2 monoantibody, trastuzumab has clinical activity either alone or combined with chemotherapy in Her-2-positive breast cancers [12]. Because association between Her-2 over-expression and NFκB currently is elusive, we explored the effect of TNF-α treatment and/or NFκB activation on Her-2 expression in MCF-7 breast adenocarcinoma cells. Our outcomes showed that TNF-α treatment led to intracellular Her-2 cleavage via a caspase-8-dependent signaling pathway in MCF-7 cells defective for NFκB activation. Therefore, our results provided the evidence to refer Her-2 as a brand-new caspase-8 substrate and demonstrated that NFκB can remarkably inhibit Her-2 degradation, that could be a novel mechanism for NFκB antiapoptotic role.

Materials and Methods

Reagents
Human recombinant TNF-α and actinomycin D were purchased from Sigma (St. Louis, U.S.A). Recombinant caspase-8 and caspase-8 inhibitors were purchased from Calbiochem (La Jolla, CA).

Cell culture and transfection
MCF-7 breast adenocarcinoma cells (generously supplied by Dr. Chen Huang, Medical College of Xi’an Jiao Tong University, Xi’an, Shaanxi Province, P.R. China) were cultured and passaged in RPMI 1640 medium supplemented with 10% stripped fetal bovine serum, 1% L-glutamine (200 mM), penicillin (100 IU), and streptomycin (100µg/ml). For the stably transfected cell lines (pcDNA3.0 and IκBα MT), culture medium was supplemented with G418 (500µg /ml, Roche). For DNA transfection, cells were plated at a concentration of 7×10⁵ cells per 35-mm-diameter culture dish and transfected 24 h later with FuGENE (Sigma, U.S.A), according to the protocol provided by the manufacturer (Roche, Switzerland).

Protein extraction and Western blot analysis
Whole cell extracts were obtained by resuspending the PBS-washed cellular pellets in 1% SDS. Then, lysates were boiled for 10 min, and protein amounts were quantified with Micro BCA Protein Assay reagent (Pierce, Rockford, IL) using a BSA standard solution curve as the manufacturer’s protocol. Protein extracts were separated on SDS-PAGE gels and blotted onto a polyvinylidenedifluoride (PVDF) membrane (Millipore, Bedford, MA). The membranes were blocked in Tris-buffered saline/Tween 20% buffer plus 5% nonfat dry milk, incubated for 2 h with first antibody, washed with Tris-buffered saline/Tween 20%, and incubated for 1 h with second horseradish peroxidase-conjugated antibody (DAKO, Glostrup, Denmark). The reaction was revealed with the enhanced chemoluminescence detection method (ECL kit; Amersham Pharmacia Biotech, Piscataway, NJ).
The antibodies were used for Western blot analysis: rabbit polyclonal anti-Her-2 (ab-2428), mouse monoclonal anti-actin (ab-3280), rabbit polyclonal anti-XIAP (ab-21278), rabbit polyclonal anti-c-FLIP (ab-8421), rabbit polyclonal anti-c-IAP-1 (ab-2399), and anti-c-IAP-2 (ab-23423) were obtained from Abcam (U.S.A).

Real-time quantitative PCR
Total RNA was extracted using RNeasy columns from Qiagen (Valencia, Spain) according to the manufacturer’s recommendations. After DNase treatment, RNAs were eluted and quantified using a spectrophotometer. 1.0 µg of RNA was reverse transcribed using the first-strand cDNA synthesis kit for reverse transcription-PCR (Roche). The quantitative PCR reaction system involved 2.0 µl of 20× diluted cDNAs, 2.0 µl of 10× SYBR Green PCR mix buffer, 1.6 µl of 25 mM MgCl₂ and 7.0 µM of each primer. The number of cycles was selected to allow linear amplification of the cDNAs. For quantitative PCR, GAPDH was selected as control. Quantification was performed with the Light Cycler PCR (Eppendorf). The primer sequences were as follows: GAPDH, 5′-ATGGGGAAGGTGAAGGTGGTC-3′ and 5′-TGATGGCATGGACTGTGG-3′; and Her-2, 5′-AGACGAAGCATACGTGA-3′ and 5′-GTACGAGCCGCACATC-3′.

Caspase-8 activity
To evaluate caspase-8 activity, pcDNA3 MCF-7 cells and IκBα MT cells were stimulated with TNF-α. Cell lysates then were incubated for 3 h at 37°C with a caspase-8 fluorogenic substrate, Ac-IEPD-AMC (Sigma, U.S.A). Subsequently, the fluorescence of generated cleaved product was assessed using a spectrophotometer (380 nm and 460 nm). The experimental procedure followed the manufacturer’s recommendations.

FLIP antisense experiments
c-FLIP shRNA, 5′-ACTTGTCCCTGCTCCTTGAA-3′; control shRNA, 5′-ATCACGTATCGTCGCTTCTC-3′ bearing phosphorothioate linkages were delivered into cells by FuGENE (Roche, U.S.A) at a final concentration of 10 µM for 8 h before TNF-α treatment.

Results

TNF-α-induced Her-2 degradation in NfκB-defective cells
To investigate the effects of TNF-α and NfκB on Her-2 expression, we used MCF-7 breast adenocarcinoma cells stably transfected with either an empty vector pcDNA3.0 (control cells) or a vector encoding an NfκB repressor, mutated IκBα at the sites of serines 32 and 36 (IκBα MT or MT cells), that inhibited NfκB activity and in turn led to enhanced apoptotic response to TNF-α. Both cell lines were stimulated with TNF-α for 24 h. Then, Her-2 expression was investigated by Western blot. An obvious decrease in Her-2 expression was detectable in MCF-7 cells with the mutated form of IκBα (Fig. 1). To further explore the transcriptional influence on decreased Her-2 protein, MCF-7 cells with pcDNA3.0 or IκBα MT were stimulated with TNF-α for 24 h. Then, the transcriptional level of Her-2 was tested by quantitative real-time PCR using Her-2-specific primers and GAPDH gene as control. No significant differences
Fig. 2. Her-2 mRNA synthesis and stability. A, pcDNA3.0 or IκBα MT MCF-7 cells were stimulated by tumor necrosis factor α (TNF-α, 100 units/ml) as indicated. Quantitative real-time PCR was performed using Her-2-specific oligonucleotides and glyceraldehyde-3-phosphate dehydrogenase (GAPDH) as control. B, control or MT cells were treated with actinomycin D (5 µg/ml) as indicated. Quantitative real-time PCR was performed as in 2A.

Fig. 3. Her-2 protein stability. IκBα MT cells were incubated with or without Phenylmethylsulfonyl fluoride (PMSF, 10 µg/ml) for 1 h before tumor necrosis factor α (TNF-α, 100 units/ml) time course stimulation. 10 µg of total cellular extract was subjected to anti-Her-2 Western blot analysis.

in Her-2 mRNA level were observed after both cell lines were stimulated by TNF-α (Fig. 2A). Subsequently, control and IκBα MT MCF-7 cells were preincubated with actinomycin D for 1 h to block RNA neosynthesis before a TNF-α stimulation. The degradation of Her-2 mRNA was identical in both cell lines (Fig. 2B). These outcomes showed that TNF-α-mediated Her-2 downregulation was not the consequence of decreased RNA transcription.

Her-2 cleavage by TNF-α-activated caspase-8 in NFκB-defective cells

In the light of the outcomes mentioned above, we postulated that downregulated Her-2 in response to TNF-α treatment might be a consequence of protease activation. To test such a hypothesis, we preincubated cells with protease inhibitors and tested their ability to counteract TNF-α-mediated Her-2 decay. Phenylmethylsulfonyl fluoride (PMSF, a serine protease inhibitor) did not prevent Her-2 decay in response to TNF-α in the cells overexpressing the mutated IκBα (Fig. 3). Because it was reported that Her-2 may be a caspase substrate, it was of interests to investigate this signaling pathway in our study. It was observed that endogenous caspase-8 activation following TNF-α addition in the IκBα MT cells, which was perfectly paralleled with Her-2 decay (Figs. 4A and B). Moreover, a specific caspase-8 inhibitor, z-IETD-fmk, was used to block TNF-α-induced Her-2 degradation in MCF-7 cells with IκBα MT (Fig. 5). Her-2 cleavage presented and was inhibited by the specific caspase-8 inhibitor. Taken together, these data indicated that caspase-8 was responsible for the TNF-α-induced Her-2 degradation in the NFκB-deficient cells.
Our report showed that caspase-8-induced Her-2 decay only occurred in the MCF-7 cells with the inhibition of NFκB activation or synthesis. Furthermore, we identified NFκB-regulated genes in control and MT cells treated with TNF-α after total cellular lysates were analyzed by Western blot analysis revealed with the specific antibodies. Accordingly, c-IAP-1, c-IAP-2, and XIAP were identical in the TNF-α-treated control or IκBα MT MCF-7 cells. However, c-FLIP-L was increased remarkably in the TNF-α-stimulated control cells as compared with MT cells (Fig. 6). Because c-FLIP-L is a well-known caspase-8 inhibitor and an NFκB target gene [14, 15]. Therefore, we further investigated whether modulation of c-FLIP-L affected Her-2 cleavage. Specific shRNA targeting c-flip mRNA efficiently attenuated c-FLIP-L expression in MCF-7 control cells, whereas control shRNA did not (Fig. 7). The inhibition of c-FLIP-L expression in control MCF-7 cells was associated with TNF-α-induced Her-2 degradation (Fig. 7). In addition, the opposite experiment was performed by re-introducing c-FLIP-L exogenous transient expression in MT cells. Ectopic c-FLIP-L expression partially inhibited TNF-α-mediated Her-2 cleavage (Fig. 8).
Discussion

The apoptosis depends on the activity of caspases family, a growing family of aspartyl-specific cysteine proteases that are essential for the execution of apoptotic cascade by the cleavage of broad-spectrum substrates, including DNA repair molecule like PARP, focal...
adhesion kinase, oncoproteins such as Bcl-2, and signaling proteins [16-19]. Because caspases are responsible for the cell apoptotic cascade, identification of their cellular substrates is essential and provides new insights into downstream events involved in apoptosis signaling. Our outcomes clearly defined Her-2 as a caspase-8 substrate. It was of interest that Her-2 molecule cleavage by activated caspase-8 occurred after TNF-α treatment, implying that this cleavage might be involved in proapoptotic signaling progression.

It was previously demonstrated that TNF-α treatment led to a decrease in Her-2 mRNA synthesis and expression level of Her-2 protein [20, 21]. We did not observe any effects of TNF-α on Her-2 transcription level. It was likely that these discrepancies resulted from the differences in the cell lines. However, it was the first time that we reported a caspase-dependent cleavage of Her-2 following TNF-α treatment.

In breast cancer cells, it was proven that oncoprotein Her-2 was able to promote proliferative, metastatic, and prosurvival events and correlated with a poor prognosis [9, 22]. Furthermore, Her-2 expression induced a resistance to chemoradiotherapy, and its inhibition by a specific monoclonal antibody, trastuzumab, yields the predominant clinical activity either alone or in combination with chemotherapy in the Her-2-positive breast cancers [23-26]. Moreover, it has been uncovered that Her-2 over-expression could induce resistance to TNF-α stimulation and that trastuzumab can restore the cytotoxic response [20, 21]. Our results raised a hypothesis that caspase-8-mediated Her-2 cleavage following TNF-α treatment could take part in the cell apoptosis. Wild-type Her-2 over-expression by the transfection of the plasmid bearing wild-type Her-2 is potential to significantly inhibit TNF-α-induced cell death, and a survival disadvantage was not displayed following the over-expression of an uncleavable Her-2, demonstrating that Her-2 cleavage leads to apoptosis.

Her-2-related signalings counteract cell apoptosis by inducing antiapoptotic pathways, such as Bcl-2 and Bcl-XL upregulation or by activating the Akt/NFκB prosurvival cascade [27-29]. NFκB is an extensively defined antiapoptotic factor, whose nuclear DNA binding is potently and rapidly induced by TNF-α treatment in almost all of the cancer cell lines [30, 31]. Constitutive NFκB activation has been observed in a wide variety of cancers and associated with a resistance to apoptosis owing to its target genes encoding for antiapoptotic molecules [30, 31]. In our study, TNF-α stimulation led to NFκB activation, and subsequent expression of one of its target genes, c-FLIP. c-FLIP is extensively defined as a caspase-8 inhibitor and therefore counteracts Her-2 cleavage triggered by TNF-α stimulation. c-FLIP-mediated cleavage inhibition did not occur when NFκB activity was inhibited by IκBα MT expression, demonstrating that it required the integrity of the NFκB pathway. We demonstrated that Her-2 cleavage might participate in TNF-α-induced apoptosis, indicating that NFκB-dependent blockage of Her-2 cleavage was a novel mechanism for NFκB antiapoptotic role.

In conclusion, we provide evidence, for the first time, that caspase-8 cleaves Her-2 protein in response to TNF-α stimulation, and we propose an original model in which NFκB exerts a new antiapoptotic function through c-FLIP-induced expression and subsequent inhibition of TNF-α-triggered cleavage of Her-2. Therefore, this antiapoptotic mechanism underlines the opportunity to evaluate the NFκB activation status and the Her-2 expression level in breast cancer cells and opens new insights to develop combined anticancer therapies for Her-2-overexpressing breast cancers. Because NFκB inhibitors currently are being evaluated as anticancer therapeutic agents, our results indicate a putative novel activity for these agents. Antibodies against Her-2 and NFκB inhibitors may be combined with chemotherapy or biological modifiers for the management of Her-2-overexpressing cancers.

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