

Upregulation of p38 pathway accelerates proliferation and migration of MDA-MB-231 breast cancer cells

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Abstract. Tumor cells capture the signaling pathways used by normal tissue to promote their own survival and dissemination and among them, the NF- κ B and MAPK pathways (ERK, JNK and p38). MAPK activation has ambiguous effects on tumor cell fate depending on cell type, cancer stage and the engaged MAPK isoforms. A synthetic peptide named LyeTx II, derived from the venom of the Brazilian spider *Lycosa erythrognatha*, was capable of increasing MDA-MB-231 aggressive breast cancer cell proliferation as indicated by MTT and BrdU (5-bromo-2'-deoxyuridine) incorporation assay and cell migration. A correlation has been established between the accelerated proliferation and migration observed in the presence of LyeTx II and the upregulation of p38 MAPK phosphorylation. The use of the selective inhibitor of p38 α / β (SB203580) abrogated the peptide effect in MDA-MB-231 cells. Besides, an augment of the canonical NF- κ B pathway activation considered as crucial in cancer progression was noted after cell incubation with LyeTx II. Importantly, activation of p38 and NF- κ B pathways was dependent on TAK1 activity. Together, these data suggest that TAK1-p38 pathway may represent an interesting target for treatment of aggressive breast cancers.

Introduction

Cancer is the second most common cause of death worldwide, preceded only by heart diseases (1). Breast cancer is one of the most common in women, with high morbidity. Generally,

treatment occurs via surgery, chemotherapy, radiotherapy and immunotherapy. However, despite the various therapeutic strategies that can be adopted, the mortality rate and the side effects of the treatment remain a challenge (2,3). Therefore, the search for new alternative treatments with less side effects is under investigation.

In such context, there is great interest in the pharmaceutical industry to target pathways that regulate cell proliferation and apoptosis in cancer. These pathways are initiated from various cell surface receptors, and may converge on the MAPK cascade, a module consisting of MAP kinase kinase (MEK) and MAPK (4-6). The MAPKs are serine/threonine protein kinases that participate in different intracellular processes such as proliferation, differentiation, cellular stress responses, and apoptosis (7,8). ERK1/2, JNK1/2/3 and p38 α / β / γ / δ constitute the main mammalian MAPKs studied in cancer area. ERK pathway is activated by mitogen factors, and this pathway is one of the most mutated in cancer often leading to an increase of cell proliferation and generally a decrease of apoptosis (9). On the other hand, p38 and JNK pathways are activated by stress factors, but their roles in cancer remain unclear, depending on cancer stage, cell type or MAPK isoforms (10-13).

Another pathway that called attention and has also been described as crucial in cancer progression is the NF- κ B pathway (14,15). The transcription factor NF- κ B controls many intracellular signals including cell cycle (e.g. cyclin D1), suppression of apoptosis (e.g. Bcl-2 and Bcl-xL) and inflammation (e.g. IL-6) (16). In response to a wide variety of stimuli, NF- κ B becomes active via canonical or non-canonical pathways in cancer. Generally, in the canonical pathway NF- κ B activation is preceded by the phosphorylation of IKK- α / β and I κ B α (17,18).

Nowadays, the major interest is to identify relevant molecular targets that may offer a specific therapeutic alternative treatment to cancer. In this context, MAPKs and NF- κ B pathways represent an interesting signaling network for investigation. In the present study, we used a peptide (named LyeTx II) derived from the venom of the spider *Lycosa erythrognatha*, that induced an exacerbated proliferation of MDA-MB-231 breast cancer cells, to evaluate the importance of the MAPKs and NF- κ B pathways in proliferation and migration of this breast cancer cell line considered as aggressive. According to our data, the proliferative and migratory

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Abbreviations: MAPK, mitogen-activated protein kinase

Key words: breast cancer, MDA-MB-231, MAPK, p38, proliferation, migration

effect induced by the peptide occurred mainly through upregulation of p38 pathway activation, without involvement of ERKs and JNKs. This study provides new insights into the role of p38 in aggressive breast tumor indicating that p38 may be a suitable target in this cancer type.

Materials and methods

Reagents. DMEM and fetal bovine serum (FBS) (Gibco) were from Thermo Fisher Scientific (Waltham, MA, USA). Thiazolyl Blue Tetrazolium Blue (MTT) and RIPA buffer were purchased from Sigma-Aldrich (St. Louis, MO, USA). p38 inhibitor SB203580 were obtained from InvivoGen (San Diego, CA, USA). Protease cocktail inhibitor and phosphatase cocktail inhibitor were acquired from Roche (Mannheim, Germany). Monoclonal antibodies against phospho-p44/42 (ERK1/2), phospho-p38 MAPK, phospho-JNK MAPK, β -actin and peroxidase-conjugated goat anti-mouse IgG and goat anti-rabbit IgG were obtained from Cell Signaling Technology (Danvers, MA, USA). Apoptosis, DNA Damage and Cell Proliferation Kit were obtained from BD Biosciences (Franklin Lakes, NJ, USA). Immobilon-P membranes and Luminat were acquired through Merck Millipore (Darmstadt, Germany).

Peptide information, synthesis and purification. LyeTx II is a 19 amino acid peptide, which includes four lysine residues conferring positive charge to it. LyeTx II synthesis and purification was performed at Professor J.M. Resende's Laboratory, from the Chemistry Department of UFMG, Brazil, according to methodology described by Santos *et al.* (19). The purity of LyeTx II was checked by mass spectrometry.

Cell culture conditions. The breast tumor cell lines MDA-MB-231 and MCF-7 were gifts from the Laboratory of Prof. A.M. Goes. MACL-1 and MGSO-3 cell lines were derived from primary tumor samples in the same laboratory (20). Cells were grown at 37°C in a humidified atmosphere of 5% CO₂ in DMEM supplemented with 10% heat inactivated FBS. For starvation conditions, cells were incubated with serum-free medium for 2 h before the western blot assay.

MTT assay. Breast cancer cells (10⁴/well) were plated in 24-well plates in DMEM with 10% serum. Cells were incubated for a further 96 h in the following conditions: medium alone, LyeTx II (from 0.1 to 100 nM), SB203580 (10 μ M). After 96 h, the medium was removed and cells were incubated with 100 μ l/well of MTT at 5 mg/ml for 2 h, at 37°C. After removing the medium, 200 μ l of DMSO was added to dissolve the formazan crystals and the absorbance was measured in a microplate reader at a wavelength of 595 nm. A value of 100% was assigned to untreated control cultures. Results were derived from at least three independent sets of triplicate experiments.

Western blotting. Cells seeded in 6-well plates (10⁶ cells/well) were serum-starved for 2 h. Then, cells were treated or not with LyeTx II (100 nM) in DMEM containing 10% FBS for the indicated time-points. 5z-7-oxozeaenol was added 30 min before the addition of LyeTx II. Cells were harvested and lysed in RIPA buffer supplemented with phosphatase inhibitor and

protease inhibitor cocktail according to the manufacturer's instructions. Protein lysates were separated by polyacrylamide gel electrophoresis on 12% gels, and electrotransferred to Immobilon-P membranes. Membranes were incubated with primary antibodies. After incubation with peroxidase-conjugated secondary antibody, protein expression was detected using Luminat reagent.

Measurement of cell proliferation by BrdU incorporation. This assay was realized using the Apoptosis, DNA Damage, and Cell Proliferation kit from BD Biosciences in accord with the manufacturer's instructions. Briefly, cells were treated or not with LyeTx II (100 nM) in DMEM containing 10% FBS and incubated with BrdU (10 μ M) for 8 h. When used, SB203580 was added 30 min before the addition of LyeTx II. After labeling, cells were fixed and permeabilized. After several washing, cells were treated with DNase (30 μ g/10⁶ cell) for 1 h. Following this treatment, cells were simultaneously stained with PerCP-CyTM5.5 anti-BrdU and Alexa-647 anti- γ H2AX for 20 min at room temperature. After washing, cells were suspended in staining buffer and analyzed by flow cytometry. The data were collected by the cell analyzing LSRFortessa (BD Biosciences - Immunocytometry Systems) using BD FACSDivaTM software (BD Biosciences) and analyzed with FlowJo (Tree Star) software.

Cell counting. Cells were seeded in a density of 1x10⁴ cells/well in 24-well plates. Cells were treated with LyeTx II (100 nM) in DMEM containing 10% FBS and incubated for 24, 48, 72 and 96 h. In each time, cells were harvested, stained with Tripa Blue (Gibco) and then counted in Neubauer plate. Number of cells = media of three counting x10⁴/ml.

Transwell assay. The 24-well Boyden chamber with 8 μ m pore size polycarbonate membrane Millipore (Millicell Hanging Cell Culture Inserts, Millipore, EUA) was used to analyze cell motility. For invasion assay, the membrane was pre-coated for 1 h with 300 μ l of free serum DMEM. Cells (10⁶) were seeded on the upper chamber with serum-free medium with or without LyeTx II (100 nM). When used, SB203580 (10 μ M) was added 30 min before LyeTx II addition also in the upper chamber. Medium 700 μ l (or until the volume reaches the upper chamber membrane) with 10% serum was added to the lower chamber as a chemoattractant. After 24 h of incubation, the non-motile cells at the top of the membranes were removed with cotton swabs, then the membranes were fixed and stained with 0.5% crystal violet (Sigma). Five visual fields of x200 magnification of each membrane were randomly selected and counted.

Results

Peptide LyeTx II enhances MDA-MB-231 cell population growth. Without initial information about LyeTx II activity, we predicted that this peptide, derived from the venom of the Brazilian spider *Lycosa erythrognatha*, could interact with the anionic membrane of the cancer cells due to its amino acid composition (four lysine residues conferring positive charge) and its secondary structure (α -helix). Therefore, we have evaluated the effect of the peptide on four different breast cancer

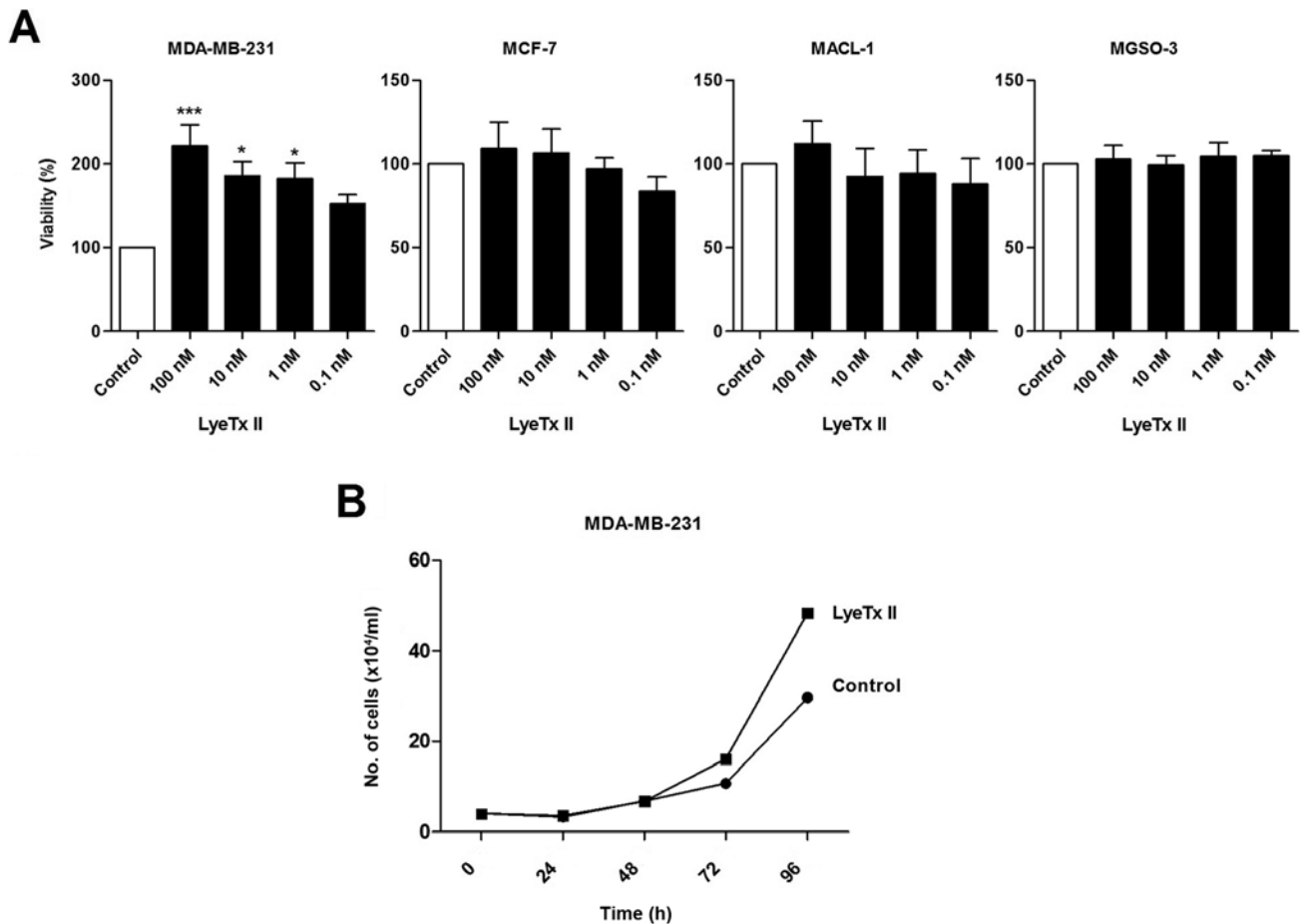


Figure 1. The peptide LyeTx II increases the proliferation of MDA-MB-231 cells. (A) Cell viability of breast cancer cell lines (MDA-MB-231, MCF-7, MACL-1, MGSO-3) was measured by MTT assay. * $p < 0.05$; *** $p < 0.01$. MTT assay was performed after 96 h of incubation in DMEM supplemented with 10% of FBS in the presence or not of LyeTx II at different concentrations. (B) Growth curve of MDA-MB-231 cells was realized in the presence of 100 nM of LyeTx II during 96 h and cell counting was performed at each 24 h. Statistical analyses were performed using one-way ANOVA followed by Bonferroni post test.

cell lines (MDA-MB-231, MCF-7, MACL-1 and MGSO-3). Cells were incubated with different peptide concentrations varying from 0.1 to 100 nM and an MTT assay was performed to evaluate cell viability (Fig. 1A). Of note, LyeTx II induced an increase of MDA-MB-231 cell viability after 4 days and this effect was more marked at 100 nM. No effect induced by the peptide was observed on the other breast cancer cell lines indicating that only the most aggressive cell line was responsive to LyeTx II. As MTT assay measures viable cell metabolism and not specifically cell proliferation, we confirmed our data by cell counting. According to Fig. 1B, an increased cell number was noted after 72 and 96 h of treatment with LyeTx II when compared to the control group.

Peptide LyeTx II enhances MDA-MB-231 cell proliferation. We sought to define whether the increased cell number observed in the presence of LyeTx II after 96 h was due to a direct activation of the proliferative machinery or liberation of secondary compounds. For this purpose, a BrdU assay was performed, that allowed us to correlate the proliferative effect induced by the peptide and the incorporation of BrdU that is a synthetic nucleoside analog of thymidine. Firstly, we pre-gated cells according to size (SSC) and granulometry (FSC). To select the population in division, H2AX in its phosphorylated

form (γ H2AX) has been used as a marker of the DNA doubled stranded breaks. The detection of the incorporation of BrdU and the presence of γ H2AX has been achieved by using antibody stained with PerCP-Cy5.5 and Alexa-647, respectively. As shown in the Fig. 2A, by double immunodetection, after 8 h of incubation with LyeTx II and BrdU, augmentation of the BrdU⁺/ γ H2AX⁺ population was noted when compared with the control cell population. The BrdU⁺/ γ H2AX⁺ population increased from 6.9 to 11.0% (Fig. 2B). Taken together, these data suggest that LyeTx II is able to interfere directly on the proliferative machinery of MDA-MB-231 cells. These findings led us to investigate the intracellular signaling pathways involved in the increased proliferation induced by LyeTx II.

p38 and NF- κ B pathway activation are upregulated in the presence of LyeTx II. Considering that MAPKs are frequently involved in the balance between cell proliferation and cell death, we sought to evaluate the phosphorylation state of the three main MAPKs (ERKs, JNKs and p38s) in the presence of LyeTx II (Fig. 3A). LPS (1 μ g/ml) and TNF- α (20 pg/ml) were used as positive control for the MAPK phosphorylation. At first, we observed that MDA-MB-231 cells did not use the JNK pathway, not in the presence of serum or in the presence of the peptide. However, we verified, that MDA-MB-231 cells

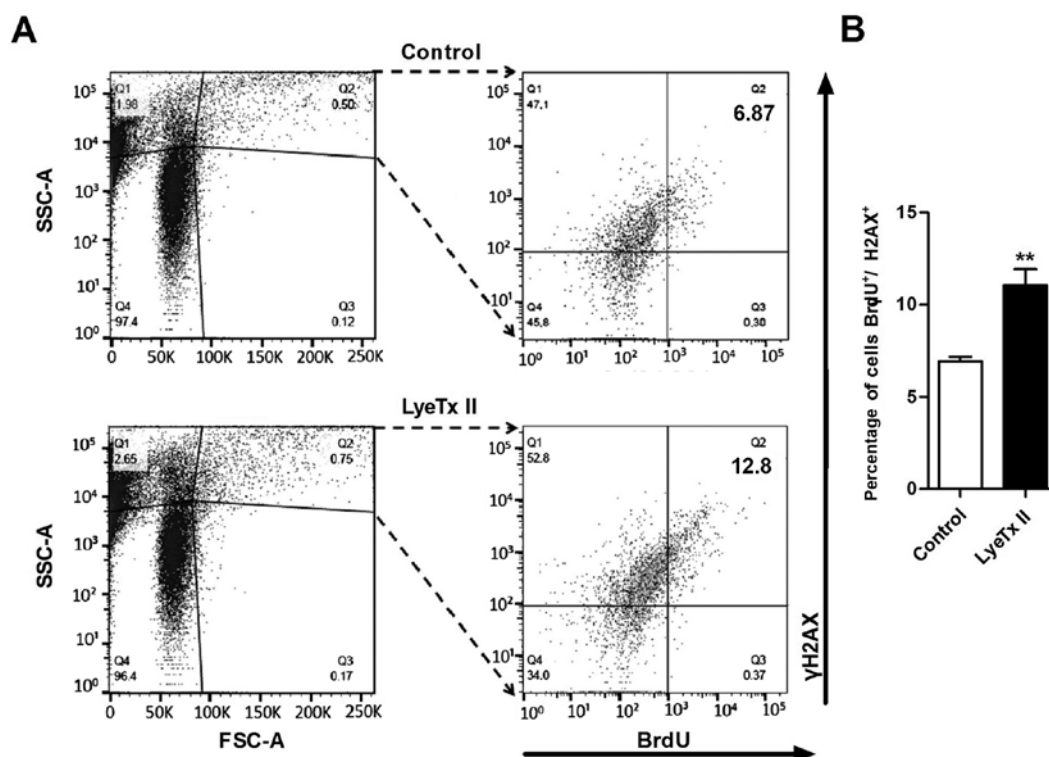


Figure 2. The peptide LyeTx II augments BrdU incorporation in MDA-MB-231 cells. MDA-MB-231 cells were incubated in DMEM with 10% of FBS and treated with LyeTx II (100 nM) and BrdU (10 μ M) for 8 h. Then, cells were fixed, permeabilized and stained with anti-BrdU and anti- γ H2AX and analyzed by FACS. Data are presented as (A) the representative dot plots of the frequency of individual data values and (B) the frequency of BrdU⁺/γH2AX⁺ cells (average \pm SD). Data are representative of two independent experiments in triplicate. Statistical analyses were performed using t-test. **p<0.01.

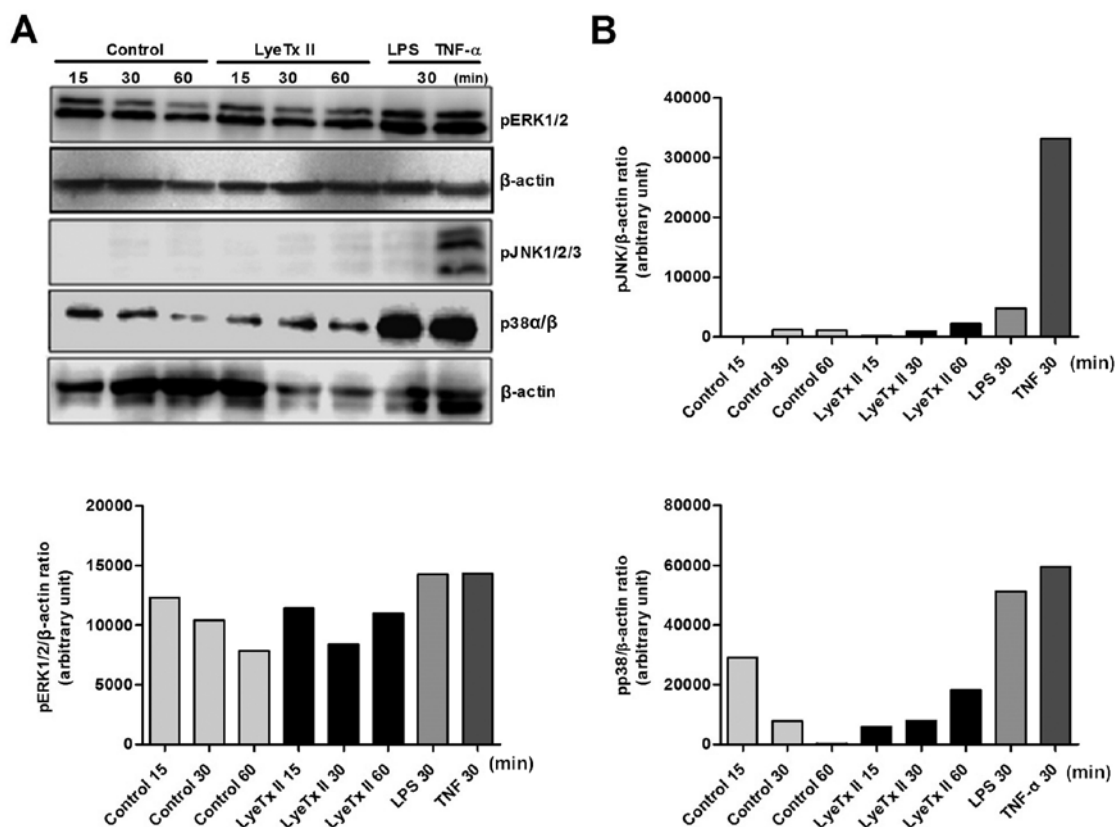


Figure 3. The peptide LyeTx II modulates p38 pathway activation in MDA-MB-231 cells. (A) MDA-MB-231 cells were incubated for 15, 30 and 60 min with LyeTx II at 100 nM in DMEM supplemented with 10% FBS (Control = medium 10% FBS). Then, cells were harvested and cell lysates were immunoblotted with the indicated antibodies. LPS (1 μ g/ml) and TNF- α (20 pg/ml) were used as positive controls. (B) The graphics represent the level of proteins quantified and normalized to the level of β -actin.

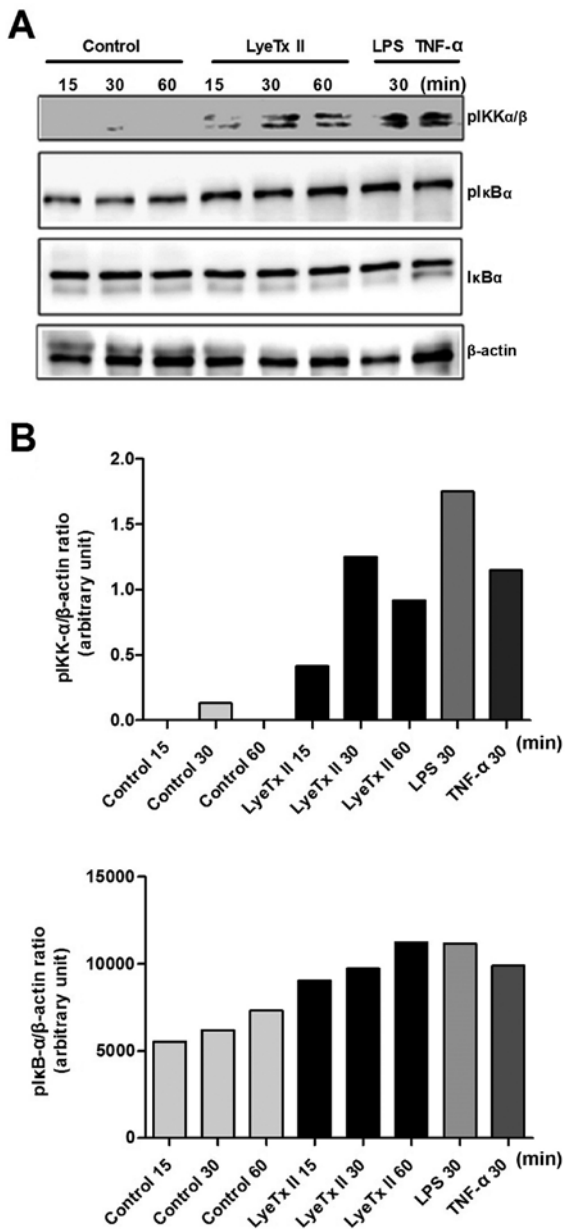


Figure 4. The peptide LyeTx II increases NF- κ B pathway activation in MDA-MB-231 cells. (A) MDA-MB-231 cells were incubated for 15, 30 and 60 min with LyeTx II at 100 nM in DMEM supplemented with 10% FBS (Control = medium 10% FBS). Then, cells were harvested and cell lysates were immunoblotted with the indicated antibodies. LPS (1 μ g/ml) and TNF- α (20 pg/ml) were used as positive control. (B) The graphics represent the level of proteins quantified and normalized to the level of β -actin.

were able to phosphorylate JNK1/2/3 under stress stimulation when TNF- α induced activation of the three isoforms of JNK. This constituted a direct evidence that LyeTx II did not require JNK activation to induce its proliferative effect. Concerning ERK1/2 pathway, no significant change in the kinetic phosphorylation of this MAPK was detected in the presence of LyeTx II. The most significant impact induced by the peptide was the modulation of p38 phosphorylation. We observed that the phosphorylation of p38 was upregulated during stimulation with LyeTx II, suggesting that the peptide effect might be related to p38 pathway activation. As shown in the Fig. 3A, phosphorylation of p38 remained elevated when compared to control group, even 60 min after peptide stimulation.

In parallel, we explored the NF- κ B pathway, another pathway well described as having an important role in cancer development. As shown in Fig. 4A, the peptide was able to enhance I κ B α phosphorylation as well as IKK- $\alpha\beta$, two of the main kinases involved in the canonical NF- κ B pathway activation. These data suggest a possible involvement of NF- κ B pathway in MDA-MB-231 cell proliferation under stimulation with LyeTx II.

Upregulation of p38 pathway is associated with the accelerated proliferation of MDA-MB-231 in the presence of LyeTx II. To evaluate whether the peptide effect was dependent on I κ B α and p38 activation, we used two selective inhibitors: BAY 11-7082 for NF- κ B pathway (21) and SB203580 for p38 pathway (22). It was not possible to define the involvement of NF- κ B pathway in the peptide activity, since I κ B α inhibitor induced MDA-MB-231 cell death in the concentration range used (2.5–20 μ M), that represent the concentrations able to inhibit I κ B α activity as previously reported (21). To correlate the peptide effect with the p38 pathway activation, we performed an MTT assay using SB203580 at concentration that did not affect the basal cell proliferation in our system. As shown in Fig. 5C, SB203580 was able to abrogate the proliferative effect induced by the peptide. To confirm the relationship between p38 activation and the proliferative effect induced by the peptide, we evaluated the BrdU incorporation in the presence of LyeTx II, SB203580 and SB203580 plus LyeTx II (Fig. 5A and B). Corroborating with the MTT assay, SB203580 was able to abolish the peptide effect maintaining the proliferative rate at a basal level. More precisely, in the presence of SB203580, the frequency of BrdU⁺ cells was significantly reduced in LyeTx II group comparing with LyeTx II group pretreated with SB203580. Collectively, the data confirmed that p38 plays a role in the increase of MDA-MB-231 cell proliferation when stimulated with LyeTx II.

p38 pathway upregulation is associated with the increased migration of MDA-MB-231 cells in the presence of LyeTx II. We hypothesized that accelerated proliferation and migration events might share the same signaling pathways in MDA-MB-231 cells. In such context, we evaluated whether the peptide could increase MDA-MB-231 cell migration by using the transwell assay. By placing the cells on one side of the membrane and using FBS as a chemoattractant on the other side, migration was determined by counting those cells that traversed the cell-permeable membrane. As shown in Fig. 6A and B, LyeTx II at 100 nM was also capable of enhancing cell migration in a significant way when compared to the control group. It is important to note that MDA-MB-231 cell line is considered as invasive, that explains the high number of cells that can pass through the membrane pores observed in serum conditions. The involvement of p38 pathway in MDA-MB-231 cell migration was confirmed when it was shown that SB203580 reduced the peptide effect on cell migration.

p38 and I κ B α phosphorylation induced by LyeTx II is dependent on TAK1. According to our data, LyeTx II was able to upregulate p38 and NF- κ B pathway activation, so we sought to determine possible crosstalk between these pathways. TAK1 has been previously described as a common upstream kinase

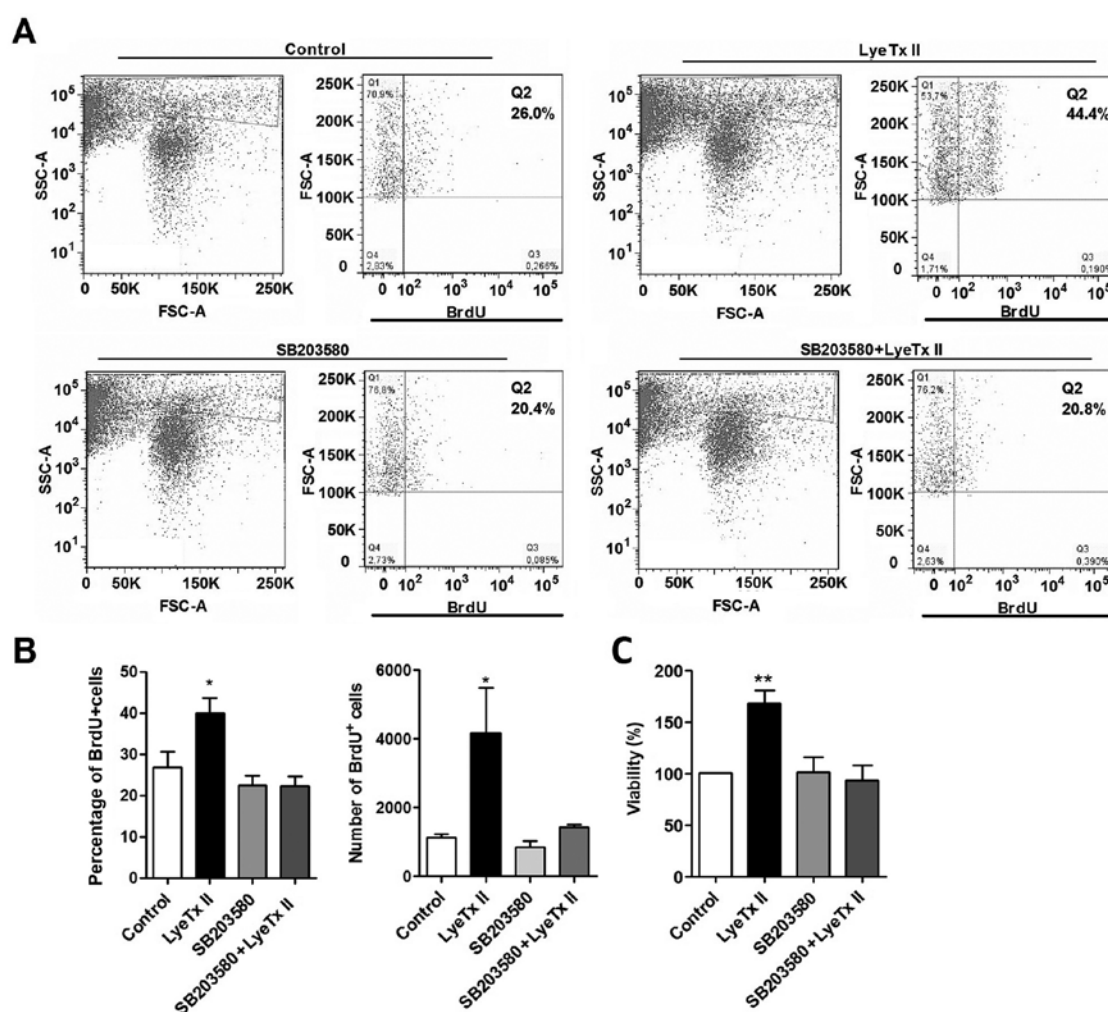


Figure 5. p38 inhibitor abrogates LyeTx II effect on MDA-MB-231 cell proliferation. (A) Cells incubated in DMEM supplemented with 10% of FBS were treated with LyeTx II (100 nM) and BrdU (10 μ M) for 8 h, pretreated or not with p38 inhibitor SB203580 (10 μ M) for 30 min. Then, cells were stained with fluorescent antibodies anti-BrdU and analyzed by FACS. Data are presented as (A) the representative dot plots of the frequency of individual data values and (B) the frequency and the number of BrdU⁺ cells (average \pm SD). (C) Cell viability was measured by MTT assay after incubation for 96 h with LyeTx II pretreated or not with SB203580 (10 μ M) for 30 min. Control group (NT) represents cells incubated with medium supplemented with 10% FBS. MTT data are representative of three experiments. Statistical analyses were performed using one-way ANOVA followed by Bonferroni post test. * $p < 0.05$; ** $p < 0.01$.

for both pathways (23). Further, other groups have previously described the involvement of TAK1 in the MDA-MB-231 cell survival (24,25). Thus, we explored the impact of TAK1 inhibition on the peptide effect. As shown in Fig. 7, when cells were stimulated with the peptide, they were still able to phosphorylate p38 and I κ B α in the presence of TAK1 at 300 nM. However, at 500 nM, TAK1 inhibitor reduced significantly p38 and I κ B α phosphorylation in LyeTx II-treated cells. Taken together, these data indicate that LyeTx II upregulates p38 and NF- κ B pathways in a TAK1-dependent manner.

Discussion

A peptide derived from the venom of the Brazilian spider *Lycosa erythrognatha* was used to correlate activated signaling pathways and aggressive breast cancer cell proliferation. There are several studies describing p38 as a proliferative agent (26,27); on the other hand, some studies have defined p38 as an apoptosis inducer (28-30). Herein, we clearly defined that p38 was involved in the peptide induced-MDA-MB-231 cell gain functions such as proliferation and migration.

Further, these data show a direct role for p38 in the cell cycle since the use of p38 inhibitor SB203580 abrogates the peptide effect on the BrdU incorporation in the first hours. In fact, it has been previously reported that p38 may regulate both the G2/M as well as G1/S cell cycle checkpoints (31-33). Different hypothesis may be advanced like the upregulation of cyclin D1 expression in a p38-dependent manner (34) or an additional mechanism that may contribute to augmentation of proliferation in the presence of the peptide. As p38 has also a well-known role in inflammation (35), we investigated a possible role of inflammatory mediators in the peptide effect. Since no augmentation of cytokine secretion was observed in the presence of LyeTx II, we were able to discard the correlation between the pro-inflammatory property of p38 and the peptide activity (data not shown). In addition, the use of rhTNF- α at a very low concentration (20 pg/ml) enhanced MDA-MB-231 proliferation (data not shown) and MAPK phosphorylation (including ERK and JNK) in a more pronounced way than the peptide. Taken together, these data support the idea that the proliferation effect induced by LyeTx II is independent on inflammatory processes.

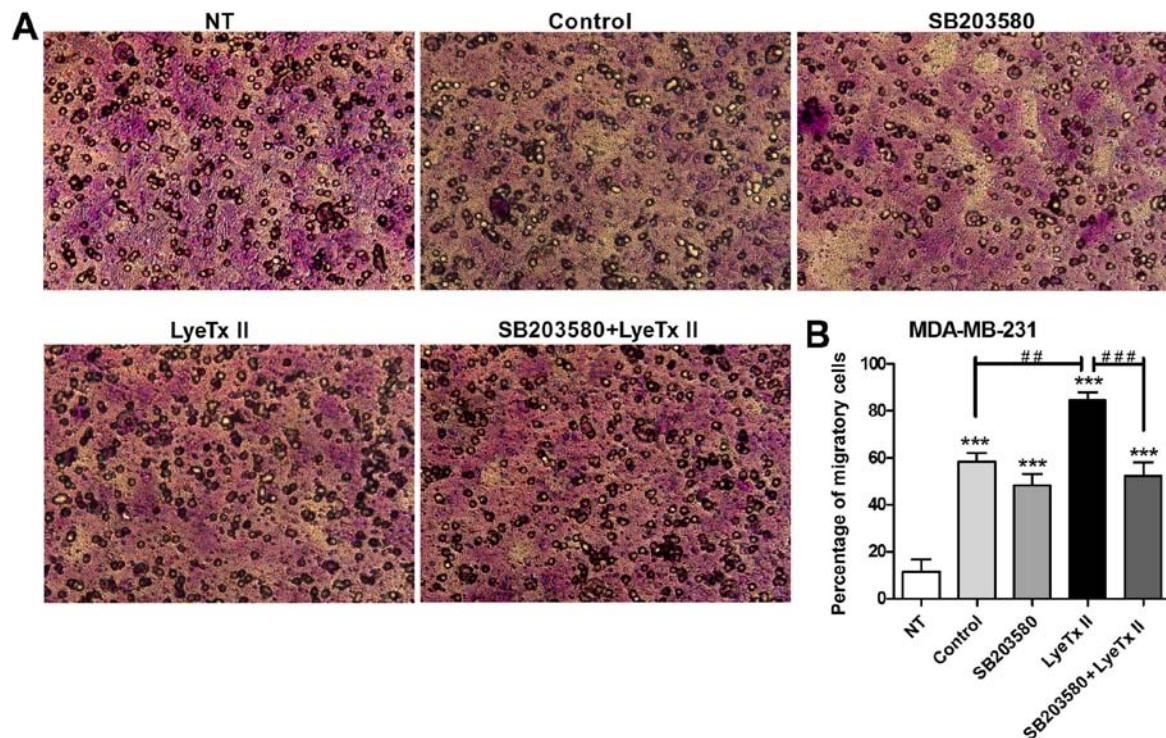


Figure 6. p38 inhibitor abrogates LyeTx II effect on MDA-MB-231 cell migration. (A) Cells were seeded on the upper chamber with serum-free medium in the presence or not of LyeTx II (100 nM). SB203580 (10 μ M) was added in the upper chamber 30 min before LyeTx II treatment. DMEM with 10% FBS was added to the lower chamber as a chemoattractant. After 24 h of incubation, cells that passed through the membrane were counted in 5 visual fields of x200 magnification of each membrane. (B) Graphic represents the media \pm SD of the percentage of cells that passed through the membrane after different treatments. Statistical analyses were performed using one-way ANOVA followed by Bonferroni post test. ***p<0.001 compared with non-treated (NT) group. **p<0.01; ###p<0.001.

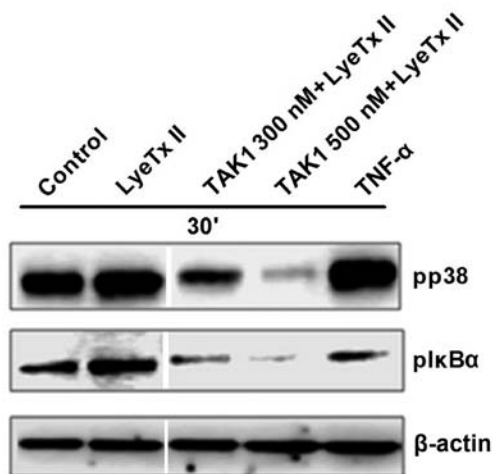


Figure 7. Impact of TAK1 inhibitor on p38 and IκBα phosphorylation induced by LyeTx II peptide. MDA-MB-231 cells were pretreated or not with TAK1 inhibitor [(5Z)-7-oxozeaenol] and incubated for 30 min with LyeTx II at 100 nM in DMEM supplemented with 10% FBS (Control = medium 10% FBS).

It has also been reported that p38 may play an important role in many steps of metastasis, such as invasion/migration, in pancreatic, hepatocellular and head and neck squamous carcinoma cell lines on the basis of the use of SB203580 and dominant-negative mutants (36). Herein we demonstrated that LyeTx II, through upregulation of p38 phosphorylation, enhanced MDA-MB-231 cell migration, reinforcing the

involvement of p38 in the augmentation of the aggressive character of the tumor cells.

We sought to define whether the increased proliferation was due to a combination of events involving proliferation and/or anti-apoptotic pathways. Given the role of the transcription factor NF-κB in prosurvival pathways (37), we analyzed the capacity of the peptide to upregulate the NF-κB pathway. At first, it was of interest to note that while MDA-MB-231 cells were capable of phosphorylating IKK-αβ and IκBα in the presence of serum, MCF-7 did not (data not shown). This is in accord with a model proposed by Lee *et al* (38) where ER⁺ breast cancer cells, such as MDA-MB-231, use constitutively NF-κB pathway and ER⁺ breast cancer cells, such as MCF-7, do not. The peptide activity fits this model by enhancing IKK-αβ and IκBα phosphorylation only in MDA-MB-231 cells. This suggests that LyeTx II, besides its proliferative activity, might play a pro-survival role through upregulation of NF-κB pathway. This might explain the significant impact of this peptide on MDA-MB-231 cell proliferation and migration at low concentration (approximately 100 nM). Peptides used in cancer studies act generally at concentration superior to 1 μ M (39,40).

Considering that LyeTx II modulated p38 and NF-κB pathways activation, we hypothesized that a common molecule could regulate both pathways in response to the peptide. TAK1 has been reported as an upstream kinase of p38 and NF-κB (41-43), so it appeared relevant in our model to correlate p38 and IκBα activation with TAK1 activity. Our data suggest that the phosphorylation of p38 and IκBα is dependent

on TAK1 activation. Noteworthy, it has been reported that p38 may exert a positive feedback on TAK1 phosphorylation (44), which may occur in our model since a sustained phosphorylation of p38 has been observed in presence of LyeTx II. Such feedback mechanism may contribute to the peptide activity.

An interesting finding in this study was the divergence of cellular response between the non-aggressive MCF-7 and aggressive MDA-MB-231 tumor cells to LyeTx II, probably due to the differential use of signaling pathways by cell lines. Comparative phosphoproteome analysis reveals differences in levels of various phosphoproteins in these two cell lines (45). This indicates that the sole presence or lack of receptors (ER, PR, HER2) may not be sufficient to classify tumor cells and to predict treatment.

In conclusion, we were able to associate a dysregulation of MAPK signaling pathways with a specific breast tumor cell function. We identified, by using the synthetic peptide LyeTx II, that upregulation of the p38 pathway in a TAK1-dependent manner led to an accelerated proliferation and migration rate in MDA-MB-231 cells. This study highlights the importance of using compounds derived from animal venoms that may contribute to identify new targets for breast cancer treatment. Furthermore, these data open new perspectives for the use of p38 and TAK1 inhibitors as a targeted-treatment in cancer.

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