RESEARCH ARTICLE

Loss of Inhibitor of Growth (ING-4) is Implicated in the Pathogenesis and Progression of Human Astrocytomas

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Keywords

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Abstract

Inhibitor of growth 4 (ING-4) is a tumor suppressor gene that interacts with nuclear factorkappaB (NF-κB) and represses its transcriptional activity. Several lines of evidence suggest that the tumor suppressor gene ING-4, the transcription factor NF-KB and its target genes matrix metalloproteases MMP-2, MMP-9 and urokinase plasminogen activator (u-PA) are critically involved in tumor invasion. The aim of the present study was to investigate immunohistochemically the expression pattern of ING-4, NF- κ B and the NF- κ B downstream targets MMP-2, MMP-9 and u-PA in human astrocytomas from 101 patients. We found that ING-4 expression was significantly decreased in astrocytomas, and ING-4 loss was associated with tumor grade progression. Expression of p65, a NF- κ B subunit, was significantly higher in grade IV than in grade III and grade I/II tumors, and a statistical significant negative correlation between expression of ING-4 and expression of nuclear p65 was noticed. MMP-9, MMP-2 and u-PA were overexpressed in human astrocytomas. Of note, astrocytomas of advanced histologic grades (grade III, IV) displayed significantly higher expression levels of these proteins compared to tumors of lower grades (grade I, II). Collectively, our data suggest an essential role for ING-4 in human astrocytoma development and progression possibly through regulation of the NF-kB-dependent expression of genes involved in tumor invasion.

INTRODUCTION

Astrocytomas are the most common primary central nervous system tumors. The most prominent characteristic of these tumors is their ability to invade adjacent normal brain parenchyma by degrading components of extracellular matrix. The highly invasive nature of malignant astrocytomas often prevents curative surgical resections and complicates effective delivery of therapy, leading to serious clinical deficits and poor prognosis. Tumor invasion is a complex biologic process involving specific well-orchestrated events, namely adhesion, cell motility and degradation of extracellular matrix (25). Recent molecular studies suggest the implication of inhibitor of growth ING-4 (inhibitor of growth family, member 4) in astrocytomas invasion process (13). ING-4 is a member of the ING tumor suppressor gene family. The ING family involves five evolutionarily conserved proteins (ING-1-ING5) characterized by a highly conserved plant homeodomain finger motif in their carboxy-terminal end (7). Previous studies have shown that this motif is found in transcription factors (ie, AF10, AF17, Trx) that modulate chromatin structure (1) and that ING proteins interact

with multiprotein complexes containing histone acetyltransferase and deacetylase (12). These data indicate that ING proteins participate in transcriptional regulation through chromatin remodeling (12). ING-4 is located in chromosomal locus 12p13 and encodes for a 249-amino acid protein containing a C-terminal plant homeodomain finger motif and two nuclear localization signals. ING-4 has been shown to induce cell cycle arrest and p53-dependent apoptosis, to suppress cell migration and angiogenesis and to enhance chemosensitivity (16, 21). Additionally, in vitro studies in cell lines have suggested that ectopic expression of ING-4 decreases cell spreading and migration (42). In vitro data link the tumor suppressor function of ING-4 to nuclear factor-kappaB (NFκB) signaling pathway (13). NF-κB is a collective name for dimeric transcriptional factors comprising the Rel family of proteins that involve RelA (p65), c-Rel, RelB, NF-KB1 (p50) and NF-KB2 (p52). The most common heterodimer in activated cells is the NF-kB1-RelA that contains the p65 and p50 proteins (24). In quiescent cells, NF-kB remains in the cytoplasm bound to its inhibitory component IKB. Activation of NF-KB occurs after IKB phosphorylation, ubiquitination and subsequently degradation. This results in NF- κ B translocation to the nucleus, where it binds to specific binding sites of the promoters of target genes involved in a variety of processes including proliferation and survival (2, 20). In addition, previous studies have shown that ING-4 overexpression suppresses tumor invasion by decreasing the expression of both matrix metalloproteases (MMP)-2 and MMP-9 (26).

Many of the genes transcriptionally regulated by NF- κ B [ie, MMP-2, MMP-9 and urokinase-type plasminogen activators (u-PA)] are known to be involved in the process of invasion (5, 35). Accordingly, inhibition of NF- κ B activity suppresses invasion and metastasis through u-PA, MMP-2 and MMP-9 down-regulation (18, 32, 36). However, extensive *in vivo* evidence supporting a role of ING-4–NF- κ B–MMPs and u-PA signaling axis in human astrocytoma invasion process is lacking.

In the present study we examined the expression pattern of ING-4, p65, MMP-2, MMP-9 and u-PA in a well-characterized set of human astrocytomas of all grades and in normal brain. In addition, we investigated their implication in the pathogenesis and progression of astrocytic tumors and correlated their expression levels with clinicopathologic parameters such as age, sex and tumor grade. Finally, we assessed the potential of role of ING-4 as a biomarker for the prediction of the grade of astrocytic neoplasms.

MATERIALS AND METHODS

Tissue specimens

Formalin-fixed paraffin-embedded tissue samples from 101 patients with astrocytomas of all grades were obtained from the Department of Pathology of "Agios Andeas" and "Nikaia" General Hospitals, of Patras and Athens, Greece. The study was approved by the Committee on Research and Ethics and the Scientific Committee of the University Hospital of Patras, Greece. The mean age of the patients was 56 years ranging from 12 to 80 years, whereas the male-to-female ratio was 0.8. For statistical purposes we categorized patients' age into three groups: group A (10-39 years), group B (40-69 years) and group C (70-100 years). All cases were reviewed by two experienced pathologists (H.P. and J.V.), who assigned a histologic grade according to the latest World Health Organization Classification (28). Our cohort included four (3.9%) pilocytic (grade I), seven (6.9%) grade II, 18 (17.8%) grade III and 72 (71.3%) grade IV astrocytomas. Because of the relatively small number of grade I and grade II astrocytomas of our series, theses tumors were grouped together for the statistical analyses. We used normal brain cortex obtained from 15 autopsy cases as controls.

Immunohistochemical staining

Paraffin-embedded sections of tumor tissue fixed in 10% buffered formalin from 101 patients were used for immunohistochemistry. Four-micrometer serial sections were obtained from each specimen on SuperFrost[®] (Menzel-Gläser, Braunschweig, Germany) plus glass slides. In brief, after deparaffinization and rehydration in graded alcohols, immunohistochemistry was performed using the two-step peroxidase technique with a peroxidase-conjugated polymer (DAKO Envision kit, DAKO, Carpinteria, CA). The following primary antibodies were used: anti-p65 mouse monoclonal (Santa Cruz Biotechnology, Santa Cruz, CA), anti-MMP-2 (Chemicon, Hampshire, UK), anti-MMP-9 mouse monoclonal (Novocastra, Newcastle, UK), anti-u-PA mouse monoclonal (Calbiochem, Darmstadt, Germany), anti-ING-4 polyclonal (Proteintech, 10617-1-AP-Proteintech Group, Inc., Chicago, IL, USA).

Sections were counterstained with Harris' hematoxylin, dehydrated and mounted permanently. For each antiserum, all sections from different studied cases were immunostained concurrently. Negative controls were performed in all cases by omitting the primary antibodies. We have attained optimal reproducible results for each antibody, employing dilutions/incubation time and pretreatment conditions.

Immunohistologic evaluation

Each slide was independently evaluated by two experienced pathologists (E.P. and J.V.) and one investigator (G.K.). The intensity of the immunostaining for all the antibodies was scored as follows: 0, negative staining; 1, weak staining; 2, moderate staining and 3, strong staining. The distribution of immunostaining was scored according to the following assumption: 0, less than 10% of neoplastic cells display positive immunoreactivity; 1, 10-30% of neoplastic cells display positive immunoreactivity; 2, 30-80% of neoplastic cells display positive immunoreactivity; and 3, >80% of neoplastic cells display positive immunoreactivity for the examined protein. The total score combined the intensity and the distribution of immunostaining and was calculated according to the following assumption: 0: intensity and distribution of staining 0; 1: intensity 1+, distribution 2+/3+ or intensity 2+, distribution 1+; 2: intensity 2+, distribution 2+/3+ or intensity 3+, distribution 1+; and 3: intensity 3+, distribution 2+/3+. The immunohistochemical profile of all the tested antibodies was similar in all the areas of the examined tissue sections. No differences were observed between the invasive front and the central core of the examined tumors.

Statistical analysis

Statistical analysis was performed using SPSS for Windows, release 12.0 (SPSS Inc., Chicago, IL). To test the significance of differences among groups of clinicopathologic parameters, ordinal data were analyzed with non-parametric Kruskal–Wallis or Mann–Whitney tests, whereas correlations between expression of proteins (immunohistologic scores) were evaluated by Spearman rank order correlation coefficient. Tumor grade was modeled using binary logistic regression analysis. All ranking tests were performed with correction for ties. The significance level was defined as P < 0.05.

RESULTS

ING-4 protein expression

The results of immunohistochemical investigation are summarized in Table 1. Glial cells of normal brain tissue and tissue adjacent to neoplastic brain parenchyma showed strong nuclear expression of ING-4. In contrast, only 28 of 101 (27.7%) cases of astrocytomas cells showed nuclear ING-4 immunopositivity (Figure 1). In 21 of 101 (20.8%) of tumor cases, ING-4 displayed weak and in seven of 101 cases (6.9%) strong immunoreactivity. Loss of ING-4 expression correlated significantly with tumor grade. More specifically, the expression of ING-4 was significantly lower in grade IV than in grade III tumors, and in grade III than in grade I/II tumors

	Gliomas	Total	ING-4 expression	ession			Nuclear p6	Nuclear p65 expression			Cytoplasm	Cytoplasmic p65 expression	sion	
		101	0 n (%) 6 (8.7)	1 n (%) 19 (27.5)	2 n (%) 24 (34.8)	3 n (%) 20 (29)	0 n (%) 7 (6,9)	1 n (%) 38 (37.6)	2 n (%) 45 (44.5)	3 n (%) 11 (10.8)	0 n (%) 9 (8.9)	1 n (%) 25 (24.7)	2 n (%) 36 (35.6)	3 n (%) 31 (30.6)
Age	<20 20-50	5 32	4 (80) 22 (68.8)	1 (20) 8 (25)	0 (0) 2 (6.3)	(0) 0	0 (0) 2 (6.3)	3 (60) 13 (40.6)	2 (40) 15 (46.9)	0 (0) 2 (6.3)	0 (0) 2 (6.3)	2 (40) 7 (21.9)	2 (40) 14 (43.8)	1 (20) 9 (28.1)
Sex	~ >50	64 45	47 (73) 33 (73.3)	12 (18.8) 8 (17.8)	5 (7.8) 4 (8.9)	(0) 0	5 (7.8) 3 (6.7)	22 (34.4) 15 (33.3)	28 (43.8) 22 (48.9)	9 (14.1) 5 (11.i)	7 (6.9) 5 (11.1)	16 (25) 13 (28.9)	20 (31.3) 14 (31.1)	21 (32.8) 13 (28.9)
	:	56	40 (71.4)	13 (23.2)	3 (5.4)	(0) 0	4 (7.1)	23 (41.1)	23 (41.1)	6 (10.7)	4 (7.1)	12 (21.4)	22 (39.3)	18 (32.1)
Grade	Grade 1/II Grade III	11	2 (18.2) 13 (72.2)	7 (63.6) 2 (11.1)	2 (18.2) 3 (16.7)	(0) 0	1 (9.1) 4 (4.22)	8 (72.7) 8 (44.7)	2 (18.2) 5 (27.5)	0 (0) 1 (5.6)	1 (9.1) 4 (22.2)	7 (63.6) 6 (33.3)	3 (27.3) 6 (33.3)	0 (0) 2 (11.1)
	Grade IV	72	58 (80.6)	12 (16.7)	2 (2.8)	(0) 0	2 (2.8)	22 (30.6)	38 (52.8)	10 (13.9)	4 (5.6)	12 (16.7)	27 (37.5)	29 (40.3)

(P < 0.001, for both correlations). No significant difference was observed in ING-4 expression levels between grade I and grade II neoplasms. The expression levels of ING-4 did not reveal any significant difference between different age groups, as well as between male and female patients. By using binary logistic regression analysis we found that loss of ING-4 immunoreactivity could predict a high-grade astrocytoma of our series with an overall accuracy of 89%.

NF-kB (p65) protein expression

NF- κ B (p65) was detected in both the cytoplasm and the nucleus of 94 of 101 (93.1%) of the examined astrocytomas (Table 1). On the contrary, this protein was not detected in normal brain cells, under the conditions employed. Cytoplasmic p65 immunostaining was found in 92 of 101 (91.1%) tumors while nuclear p65 immunolocalization was observed in 94 of 101 (93.1%). Both nuclear and cytoplasmic immunoreactivity was detected in 92 of 101 (91.1%) astrocytomas. p65 showed both nuclear and cytoplasmic immunolocalization (Figure 1D-F). Importantly, its cellular levels (both nuclear and cytoplasmic) were significantly higher in grade IV compared to grade III and in grade III compared to grade I and II tumors (P < 0.001, for all correlations). We did not observe any significant difference among the p65 immunoexpression levels of grade I and grade II tumors. There was no significant correlation between p65 expression levels and patient gender and age.

MMP-9 protein expression

Cytoplasmic expression of MMP-9 was observed in 93 of 101 (92.1%) of the examined tumors (Table 2), while in normal brain tissue MMP-9 immunostaining was absent. MMP-9 expression levels correlated significantly with tumor grade. Indeed, grade IV astrocytomas demonstrated significantly higher expression of MMP-9 compared to grade III and grade III compared to grade I/II astrocytomas (P < 0.001) (Figure 2D–F). We did not detect significant differences in the MMP-9 expression levels between grade I and grade II neoplasms. No significant correlation was observed between expression levels of MMP-9 and patient age or gender.

MMP-2 protein expression

In normal brain tissue, immunostaining for MMP-2 was negative. With regard to astrocytomas 99 of 101 (98%) of the cases had cytoplasmic MMP-2 expression (Figure 2A-C). There was a statistical significant correlation between MMP-2 expression and tumor grade. MMP-2 expression was significantly elevated in grade IV compared to grade III and in grade III in comparison to grade I/II gliomas (P < 0.001) (Table 2). No significant difference was observed in MMP-2 expression levels among grade I and grade II astrocytomas. MMP-2 immunoexpression levels did not differ among different patient age groups or gender.

u-PA protein expression

The results of immunohistochemical evaluation are summarized in Table 2. Immunohistochemical analysis revealed cytoplasmic

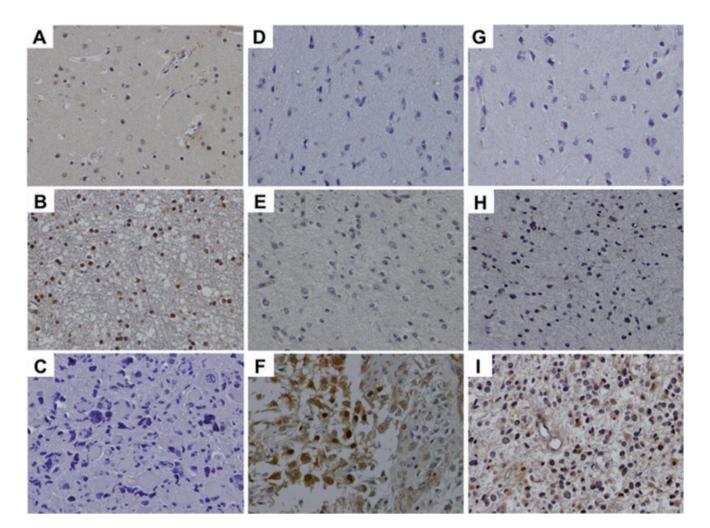


Figure 1. Loss of ING-4 expression in a glioblastoma (C). Note the increased nuclear expression of ING-4 in normal brain parenchyma (A) and low-grade astrocytoma (B). Elevated NF- κ B (p65) expression is evident in a high- and in a low-grade astrocytoma (F and E, respectively).

immunoreactivity for u-PA in 89 of 101 (88.1%) of gliomas, while normal brain did not show expression of u-PA, under the conditions employed. The expression levels of u-PA were significantly elevated in tumors of higher grades (grades III and IV) in comparison to lower grade astrocytomas (P < 0.001). In fact, low-grade astrocytomas display positive immunoreactivity in seven of 11 (63%) of the cases, whereas anaplastic astrocytomas and glioblastoma display positive cytoplasmic immunoreactivity in 16 of 18 (88%) and 66 of 72 (91%) of the examined tumors, respectively (Figure 1G–I). We did not observe any significant difference in u-PA expression levels between grade I and grade II brain tumors. u-PA immunostaining levels were not correlated with patient gender and age.

Correlations between the expression levels of the examined proteins

In our study there was a statistical significant negative correlation between expression of ING-4 and of p65 (r = -326, P = 0.001 for

Note that normal brain parenchyma is negative for p65 (D). A case of glioblastoma revealing augmented cytoplasmic u-PA expression (I). A low-grade astrocytoma (H) shows weak u-PA immunostaining, whereas normal brain does not show any immunoreactivity for this protein (F).

total p65, r = -290, P = 0.003 for nuclear p65 and r = -0.323, P = 0.001 for cytoplasmic p65). MMP-9 protein expression levels were strongly associated with p65 expression (r = 0.415, P < 0.001for total p65, r = 0.399, P < 0.001 for nuclear p65 and r = 0.393, P < 0.001 for cytoplasmic p65). Interestingly statistically significant negative correlation was found between MMP-9 and ING-4 expression levels in glioma cells (r = -0.264, P = 0.008). There was no significant correlation between MMP-2 and ING-4 expression; nevertheless, a trend toward a negative correlation was observed (r = -0.186, P = 0.062). However, the expression levels of MMP-2 were strongly associated with the cellular levels of p65 (r = 0.300, P = 0.002 for total p65, r = 0.270, P = 0.006 for nuclearp65 and r = 0.261, P = 0.008 for cytoplasmic p65). In addition, MMP-2 staining was robustly and positively correlated with MMP-9 (r = 0.393, <0.001). There was no significant correlation among u-PA and ING-4, although a trend toward a negative correlation was noted (r = -0.192, P = 0.054). The expression levels of u-PA were strongly correlated with the cellular levels of p65 (r = 0.277, P = 0.005 for total p65, r = 0.248, P = 0.12 for nuclear

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Gliomas		Total	MMP-9 expression	kpression			MMP-2 e	MMP-2 expression			u-PA expression	ssion		
		101	0 n (%) 8 (7.9)	1 n (%) 23 (22.7)	2 n (%) 32 (31.6)	3 n (%) 38 (37.6)	0 n (%) 2 (1.9)	1 n (%) 17 (16.8)	2 n (%) 46 (45.5)	3 n (%) 36 (35.6)	0 n (%) 12 (11.8)	1 n (%) 22 (21.7)	2 n (%) 55 (54.4)	3 n (%) 12 (11.8)
Age	<20 20–50	32 32	1 (20) 4 (12.5)	1 (20) 12 (37.5)	0 (0) 4 (12.5)	3 (7.9) 12 (37.5)	0 (0) 1 (3.1)	2 (40) 7 (21,9)	2 (40) 13 (40.6)	1 (20) 11 (34.4)	2 (40) 5 (15.6)	1 (20) 8 (25)	2 (40) 15 (46.9)	0 (0) 4 (12.5)
	>50	64	3 (4.7)	10 (15.6)	28 (43.8)	23 (35.9)	1 (1.6)	8 (12.5)	31 (48.4)	24 (37.5)	5 (7.8)	13 (20.3)	38 (59.4)	8 (12.5)
Sex	Σı	45	3 (6.7)	10 (22.2)	14 (31.1)	18 (40)	(0) 0	9 (20)	22 (48.9)	14 (31.1)	3 (6.7)	11 (24.4)	26 (57.8)	5 (11.1)
-	- - -	56	5 (8.9)	13 (23.2)	18 (32.1)	20 (35.7)	2 (3.6)	8 (14.3)	24 (42.9)	22 (39.3)	9 (16.1)	1 (19.6)	29 (51.8)	7 (12.5)
Grade	Grade 1/II Grade III	18	5 (45.5) 2 (11.1)	6 (33.3) 6 (33.3)	1 (9.1) 5 (27.8)	0 (0) 5 (27.8)	1 (9.1) 1 (5.6)	/ (63.6) 5 (27.8)	2 (18.2) 7 (38.9)	T (9.1) 5 (27.8)	4 (36.4) 2 (11.1)	6 (54.5) 4 (22.2)	1 (9.1) 8 (44.4)	0 (0) 4 (22.2)
	Grade IV	72	1 (1.4)	12 (16.7)	26 (36.1)	33 (45.8)	(0) 0	5 (6.9)	37 (51.4)	30 (41.7)	6 (8.3)	12 (16.7)	46 (63.9)	8 (11.1)

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p65 and r = 0.265, P = 0.007 for cytoplasmic p65), MMP-9 (r = 0.427, P < 0.001) and MMP-2 (r = 0.358, P < 0.001).

DISCUSSION

In the present study we demonstrated significantly reduced levels of ING-4, a candidate tumor suppressor gene, in human astrocytomas compared to normal brain tissue. This finding indicates that down-regulation of this protein might be involved in the pathogenesis of human astrocytic tumors, a notion that is in concert with recent molecular studies on human breast cancer cell lines and head and neck squamous cell carcinomas having shown loss of ING-4 heterozygosity (15). This hypothesis is further supported by previous reports suggesting that genomic alterations of the chromosomal locus 12p13, which involves ING-4 gene, occur frequently in human neoplasias (9, 15, 26).

Our results have demonstrated that decreased ING-4 expression in human astrocytomas correlated significantly with tumor grade progression, with lower expression levels of ING-4 observed in cases of high-grade neoplasms.

These in vivo finding parallels data from previous in vitro studies on glioblastoma and multiple myeloma cell lines showing a negative correlation between ING-4 expression and tumor grade (10, 13). Taken together, these results indicate that loss of ING-4 might be implicated not only in the development but also in the progression of astrocytic tumors to more aggressive phenotypes.

A remarkable volume of studies has shown that NF-kB regulates the expression of genes that govern major cancer-related processes, namely apoptosis, angiogenesis, metastasis, proliferation, tumor growth and survival (33).

In this study we have shown that NF- κ B is highly expressed in astrocytomas as compared with normal human brain tissue. These results provide further evidence that NF-kB is probably involved in the development of astrocytic tumors in humans.

In addition, we revealed a positive and significant correlation between NF-kB expression levels and tumor grade. Indeed, grade III and IV astrocytomas showed remarkably elevated NF-KB immunoreactivity compared to grade I and II neoplasms, suggesting that NF-kB might also contribute to histologic aggressiveness of astrocytomas. Similar immunohistochemical analyses have shown that NF-kB/p50 is also highly expressed in human astrocytic neoplasms and that its expression is robustly correlated with tumor grade (23). This points toward a critical role of the NF-κB family members in astrocytoma pathobiology.

ING-4 has been proposed to regulate tumorigenesis by modulating NF- κ B signaling (13). In the present study we demonstrated an inverse relationship between ING-4 and NF-KB protein expression in human astrocytomas, indicating that the ING-4-NF-kB axis might be involved the pathobiology of these neoplasms. In accordance with our data, it has been recently shown that one of the upstream regulators of the NF-kB is ING-4 in experiments performed on the glioblastoma cell line U87MG. That study showed that ING-4 expression was significantly suppressed at both the RNA and the protein levels and that ING-4 binding to RelA subunit of NF-kB acts as a negative regulator of NF-KB activity (13). In addition, functional biochemical studies have revealed that ING-4 suppresses brain tumor angiogenesis, and thus progression, through physical interaction and transcriptional repression of NF- κ B (13). We further assessed the cellular

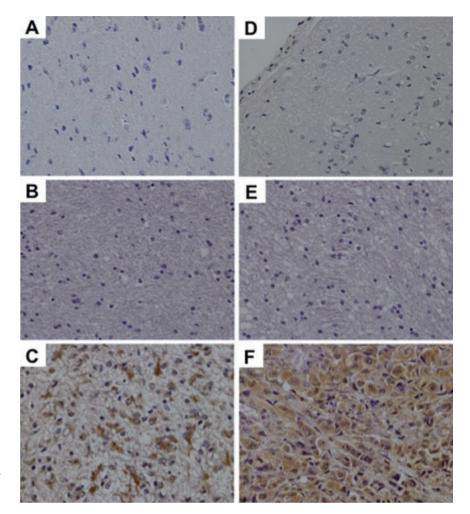


Figure 2. Normal brain tissue shows negative immunoreactivity for MMP-2 (A) and MMP-9 (D). Weak cytoplasmic expression of MMP-2 (B) and MMP-9 (E) in a low-grade astrocytoma. Strong cytoplasmic expression of MMP-2 (C) and MMP-9 (F) in a glioblastoma.

levels of ING-4 as biologic marker that could distinguish between high- and low-grade astrocytomas. By applying binary logistic regression analysis we revealed that loss of ING-4 expression could reliably predict a high-grade tumor of our series with an overall accuracy of 89%. This distinction might contribute to the individualization of prognosis and eventually treatment of astrocytoma patients.

Tumor invasion is a complex biologic process involving adhesion, motility and degradation of extracellular matrix (25). Many of the genes regulated by NF-KB, such as MMPs and u-PA, are known to be involved in invasion. MMP-2, -9 and u-PA have the ability to promote ECM remodeling, since they are key components of the tumor-associated proteolysis. Hence, many studies have proposed that these molecules could probably use therapeutic targets in oncology (14). We demonstrated increased expression levels of u-PA in human astrocytomas. More importantly, u-PA cellular levels exhibited significant correlation with progression of the astrocytomas to higher histologic grade. This is not a surprising finding, since there are reports indicating that u-PA activity is augmented in brain tumors in vivo (39). Additionally, studies on other human malignancies, such as lung, breast and colon carcinomas have clearly demonstrated a significant correlation between the production of u-PA and tumor invasiveness and metastatic propensity (19, 44). A considerable number of proteins, including NF- κ B, serve as essential regulators of u-PA (11, 40). In fact, molecular studies have shown that u-PA gene harbors NF- κ B binding site within its promoter region (37). Consistently with previous molecular studies that have linked NF- κ B with the regulation of u-PA (22, 27, 45), we observed that NF- κ B and u-PA immunoexpression levels were strongly and positively associated to each other, suggesting that in human astrocytomas u-PA may be influenced by NF- κ B.

In line with evidence suggesting that u-PA functions as an important physiologic activator of MMPs (29) we found a strong correlation between u-PA and MMP-2/MMP-9 expression in human gliomas, indicating that u-PA is probably involved in activation of MMPs in brain astrocytic tumors (38).

Similar to the plasminogen activator system, members of the MMP family of proteins are involved in extracellular matrix degradation and linked to various steps of tumor invasion and metastasis (3, 6, 30, 41). Among them MMP-2 and MMP-9 seem to have a central role in astrocytoma invasive potential (8, 31). In the present study we demonstrated increased expression of MMP-2 and MMP-9 in the majority of the astrocytomas of our sample, while this immunoreactivity was absent in normal brain tissue, suggesting a role of these enzymes in astrocytoma initiation. In addition, we observed augmented MMP-2 and MMP-9 expression levels in grade IV in comparison to grade III and grade I/II gliomas. Increased immunoexpression of MMP-2 and MMP-9 in human gliomas has also been reported by other study groups (8, 34), further strengthening the hypothesis that these proteins participate in the progression of astrocytic neoplasms. Since MMPs are activated by u-PA, a synergetic role of these enzymes in tumor progression and invasion has been considered (29).

NF- κ B also plays an important role in regulating tumor cell infiltration as it is essential for MMPs expression (4). In the current study we observed positive correlation between the expression levels of the evaluated MMPs and NF- κ B. This is consistent with earlier studies showing that the expression of several MMPs is regulated by a NF- κ B (17, 43).

Collectively, our findings provide evidence that the downregulation of ING-4 in human gliomas may induce MMP-2 and MMP-9 expression possibly via enhancement of NF- κ B. Clearly, more studies are required to clarify the role of these proteins in the pathobiology of human astrocytomas. If ING-4 proves to be essential in the pathogenesis of astrocytomas, drugs that could selectively modulate the activity of this protein might constitute a novel alternative therapeutic approach for the treatment of central nervous system tumors.

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