### LABORATORY INVESTIGATION



# Cerebrospinal fluid VEGF levels and angiogenic capacity as potential prognostic markers in patients with gliomas: a pilot study

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Received: 29 July 2019 / Accepted: 3 October 2019 / Published online: 17 October 2019 © Springer Science+Business Media, LLC, part of Springer Nature 2019

## Abstract

**Introduction** Gliomas are tumors of the central nervous system. Despite new classifications, they are still divided in low and high-grade gliomas, being the latter of greater malignancy. The degree of malignancy is directly related with the angiogenic activity in tumoral tissues. We measured VEGF concentrations and angiogenic capacity in cerebrospinal fluid (CSF) from patients with high and low-grade gliomas. The purpose of this study was to find a biomarker that contributes in the differential diagnosis and prognosis of gliomas.

**Methods** CSF was obtained from 19 individuals: 8 with low-grade gliomas, 6 with high-grade gliomas and 5 controls. VEGF concentration in CSF was measured by ELISA and the angiogenic capacity was measured by chick chorioallantoic membrane (CAM) test.

**Results** The VEGF concentration was higher in patients with high-grade gliomas, compared to patients with low-grade gliomas and controls (2860 pg/mL±975 vs. 182.6±37.1 and 47.4±0.4, respectively). On the other hand, CSF from patients with high-grade gliomas generated a higher microvascular density (MVD) than patients with low-grade gliomas and controls (13.23±0.6 vessels/9000 $\mu$ m<sup>2</sup> vs. 9.3±0.3 and 7.92±0.2, respectively). Interestingly, there was not statistical differences in both VEGF levels and angiogenic capacity in patients with low-grade gliomas and controls.

Conclusion Together VEGF levels and angiogenic capacity in CSF can be used as a biological marker of gliomas malignancy.

Keywords Angiogenesis · Glioma · VEGF · Biomarker · CSF

## Introduction

The most common primary brain tumors in adults are gliomas. They can appear anywhere in the CNS but they primarily develop from glial stem cells in the brain [1, 2]. This type of tumors has a growing importance given that each year about 300,000 new cases are diagnosed in the world, and they represent the second cause of cancer mortality in patients under 35 years in the United States [3, 4]. Moreover,

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high-grade gliomas represent 80.7% of CNS malignant tumors [5].

Gliomas are tumors that present the same histological features as normal glial cells. They comprehend a heterogeneous group of diseases, including astrocytic, oligodendrocytic, ependymomas and mixed gliomas [6]. They range from lesions with a circumscribed growth pattern (WHO grade I pilocytic astrocytoma, pleomorphic xanthoastrocytoma)-showing a frequent mutation of B-Raf (BRAF) -, to diffuse gliomas (WHO grade II, III, and IV). Traditionally, diffuse gliomas were divided into low-grade lesions (WHO Grade II Astrocytoma, Oligodendroglioma, or Oligoastrocytoma) and high-grade lesions (WHO grade III Anaplastic Astrocytoma, Anaplastic Oligodendroglioma, Anaplastic Oligoastrocytoma, and WHO Grade IV Glioblastoma), based on anaplasia, mitotic activity, microvascular proliferation and necrosis [7]. The most aggressive variant is Glioblastoma (GBM) and has a median survival of 16 months in patients with the currently available treatments [3].

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The development of integrated genomic analysis and the discovery of the1p/19q codeletion, and the IDH 1 and 2, and H3K27M mutations, revolutionized the understanding of the disease, creating new molecular subtypes in the integrated histological-molecular 2016 WHO classification of gliomas [7]. The IDH1 and IDH2 mutations-both present in the majority of WHO grade II and III gliomas-defined a new subtype associated with a favorable prognosis. When these mutations were combined with the 1p/19g codeletion, as occurred in Oligodendrogliomas, they showed a better response to radiochemotherapy and were associated with a longer survival than diffuse gliomas, where these alterations have not been observed [8-11]. Therefore, the lower-grade glioma concept arises, including WHO grade II and grade III diffuse gliomas, and stratifies them according to their genetic profile [10]. Nevertheless, this new classification has not been massified in clinic, and classic WHO classification continues being the standard diagnosis [5].

In patients with low-grade gliomas, long-term following up is performed by serial MRI, as well as complementary radiotherapy in selected cases. In the case of GBM, the combination of radio and chemotherapy has shown significantly better results [12]. However, gliomas usually tend to increase their degree of malignancy over the time, and low-grade gliomas can become high-grade gliomas [13]. Despite aggressive surgery, radiotherapy and chemotherapy, high-grade gliomas remain being lethal. For progressing, these tumors stimulate the formation of new blood vessels through the process known as angiogenesis, driven mainly by vascular endothelial growth factor (VEGF) [14]. Therefore, glioma malignancy is directly related to its vascularization level. Angiogenic level of a tumor is a predictor of progression, recurrence and metastatic dissemination, which has been demonstrated in several studies of different types of cancer, where VEGF levels were significantly high [15-23].

The monitoring of the angiogenic potential of brain tumors is not simple and requires invasive procedures such as brain biopsies [24]. Moreover, pathological diagnoses based on biopsies may not be accurate due to tumor heterogeneity, vascularization, cellularity and necrosis, and serial biopsies are frequently required [25]. The reduction, recurrence or progression of a glioma responding to therapy is evaluated by serial MRI, but this technique is not capable to distinguish these tumors from the so-called pseudoprogression (MRI pattern that resembles tumor progression without clinical deterioration) [25, 26]. As a result of this, several genomic and epigenomic features have emerged as diagnostic, prognostic and predictor biomarkers of tumor classification and risk stratification, such as the isocytrate dehydrogenase (IDH) mutation, the 1p/19q codeletion, and the O6-methylguanine-DNA-methyltransferase (MGMT) methylation [27]. Nevertheless, more available and cost-effective biomarkers are needed. In this context, numerous potentially

diagnostic and prognostic biomarkers have been proposed, including VEGF, its soluble receptor sVEGFR1, basic fibroblast growth factor (bFGF), platelet derived growth factor (PDGF) and angiopoietin 2 (Ang2), all of which can be measured in blood or CSF. These biomarkers would allow developing a better tumor classification, to inform probable prognoses, and to facilitate patient management for elaborating a personalized treatment strategy [13, 28].

The aim of this pilot study is to contribute to the knowledge of tumor angiogenesis in low-grade and high-grade gliomas, by determining VEGF levels in CSF and angiogenic capacity of CSF from patients with high and low-grade gliomas.

## **Materials and methods**

## Subjects and collection of samples

The study was approved by the Ethic Committee of Hospital Clínico Universidad de Chile. An informed consent was obtained from participants before the enrollment. CSF samples were obtained from a total of 19 individuals, all without previous antiangiogenic treatment: 14 patients with gliomas, from which 8 has a low-grade gliomas (3 oligodendrogliomas II, 2 pilocytic astrocytoma, 1 subependymoma, 1 gemistocytic astrocytoma, 1 ependymoma) and 6 has a highgrade gliomas (5 glioblastoma, 1 anaplastic astrocytoma), and 5 control adults. Control CSF samples were obtained from joint injury hospitalized patients before spinal anesthesia. Approximately 2 mL of CSF were extracted from each individual by non-traumatic lumbar puncture. Samples were aliquoted and stored at -80° C until analysis.

## **Determination of VEGF concentration in CSF**

VEGF in CSF samples was quantified by ELISA, using the commercial kit Quantikine® Human VEGF Immunoassay (R & D systems, Minn, USA). Briefly, a microtiter plate pre-coated with a monoclonal antibody specific for human VEGF was incubated with 100  $\mu$ L of CSF sample. Bound VEGF was detected using a second polyclonal antibody for human VEGF conjugated with peroxidase and then, the correspondent substrate. Absorbance was measured at 450 nm. This ELISA kit detects the human isoforms VEGF<sub>165</sub> and VEGF<sub>121</sub>. The lowest detection limit was 31 pg/mL. All samples were tested by triplicated and the intra-assay variation coefficient was 5.5%.

#### Determination of the angiogenic effect of CSF

The procedure was performed in White Leghorn breed embryonated chicken as described by Sinning et al. [24]. 70 eggs of embryonated chicken were incubated in a chamber at 37°C. 48 h after, an air chamber was drilled to extract approximately 2 ml of albumin egg with a Pasteur pipette. Eggs were disinfected with an iodized alcohol solution and holes were covered with adhesive tape. Maintaining the same incubation position, a window of 2.5 cm wide and 2 cm high was performed at the dorsal level under sterile conditions, to expose the chicken chorioallantoic membrane (CAM).

After 8 days, a methylcellulose filter impregnated with 10 µL CSF sample was placed on each egg CAM to measure the angiogenic effect of CSF components. Each control or glioma patient sample was tested in quintuplicate. 48 h post incubation, CAMs were removed by surgical procedure, and fixed in a buffered 10% formalin solution. All 5 CAMs treated with CSF from the same patient were processed in the same histological block under conventional histological procedures, and serial sections of 5µm thick were stained with Alcian blue and Hematoxylin-eosin for histological study. Using an optical microscope with ×100 magnification, blood vessels (capillaries) were counted in five fields per membrane (each field corresponds to 9000  $\mu$ m<sup>2</sup>), which gives 25 fields for each control or glioma patient treated with CSF. For each CSF sample, mean and standard error of the blood vessel count were calculated. Angiogenic effect of the CSF sample was expressed as microvascular density (number of vessels present in 9000  $\mu$ m<sup>2</sup>) of CAM and the procedure was performed by blind counting.

## **Statistical analysis**

The VEGF concentration and the number of vessels per area were expressed as mean  $\pm$  SEM. Normal, high-grade and low-grade groups were analyzed by one-way ANOVA with Bonferroni Multiple Comparison test, using the Graph Pad Prism 5 software. Difference was considered significant when p < 0.05.

## Results

#### VEGF concentration in CSF

The VEGF concentration in CSF obtained from high-grade glioma patients was higher than that measured in low-grade glioma patients or control individuals (p=0.003). Additionally, the VEGF concentration in CSF obtained from low-grade glioma patients presented a great variation between samples, being 3 of them similar to controls and 5 of them about 5 times higher than controls. VEGF levels of control individuals were all similar (Fig. 1, Table 1).



**Fig. 1** Distribution of VEGF concentration in CSF from different study groups. HG patients with high-grade gliomas; LG patients with low-grade gliomas; Control healthy individuals; ns not significant; \*\* p = 0.003. VEGF levels were quantified by ELISA. See material and methods for details

#### Angiogenic capacity of CSF measured by CAM assay

Angiogenic capacity, expressed as microvascular density (MVD) of the CSF of high-grade glioma patients, was higher than that observed in low-grade glioma patients or control individuals (p < 0.0001 in both comparisons). There was no difference between low-grade glioma patients and control individuals (Fig. 2, Table 1).

Individual values of VEGF concentration and microvascular density after CSF treatment are summarized in Table 1.

## Discussion

In the present study, we found that VEGF concentration in CSF from patients with high-grade glioma is higher than that measured in low-grade glioma patients or control individuals. Accordingly, the angiogenic capacity of the CSF measured in a chick CAM model was also higher in patients with high-grade gliomas, compared to patients with low-grade glioma or control individuals. Our data provide evidence that supports the use of these parameters as biological markers of glioblastomas. In addition, CSF samples are relatively easier to obtain compared to biopsies and can be obtained serially to following-up each patient case.

In our study, VEGF levels in CSF were increased almost 50-times compared to control values in 4 of the 6 patients with high-grade glioma. These results are in agreement with the metanalysis reported by Chen et al. [29] that observed a correlation in the overexpression of VEGF with the pathological grade of gliomas, and with the observed by Sampath et al. [30], that reported higher VEGF concentrations in CSF

Patient	Group	Diagnosis	VEGF concentration (pg/mL)	MVD (vessels/9000µm <sup>2</sup> )
1	HG	Glioblastoma	277	$14.77 \pm 2.14$
2	HG	Glioblastoma	60	$12.77 \pm 2.16$
3	HG	Glioblastoma	2699	$12.13 \pm 1.78$
4	HG	Glioblastoma	3132	$14.87 \pm 3.54$
5	HG	Anaplastic astrocytoma	5628	$11.13 \pm 2.57$
6	HG	Glioblastoma	5260	$13.72 \pm 2.10$
HG group mean (mean $\pm$ S.E.)			$2860 \pm 975$	$13.23 \pm 0.60$
7	LG	Oligodendroglioma II	276	$7.90 \pm 2.07$
8	LG	Oligodendroglioma II	67	$8.63 \pm 2.46$
9	LG	Ependymoma	56	$9.37 \pm 2.24$
10	LG	Oligodendroglioma II	51	$8.30 \pm 1.86$
11	LG	Pilocytic astrocytoma	230	$10.31 \pm 1.26$
12	LG	Gemistocytic astrocytoma	248	$9.64 \pm 2.40$
13	LG	Subependymoma	237	$9.80 \pm 1.97$
14	LG	Pilocytic astrocytoma	294	$10.1 \pm 0.82$
LG group mean (mean $\pm$ SE)			$182.6 \pm 37.1$	$9.30 \pm 0.31$
15	С		46	$7.70 \pm 1.76$
16	С		48	$8.40 \pm 1.73$
17	С		46	$8.25 \pm 2.28$
18	С		47	$7.97 \pm 1.56$
19	С		48	$7.29 \pm 1.14$
Control group mean (mean $\pm$ SE)			$47.4 \pm 0.4$	$7.92 \pm 0.20$

Table 1 Data of individual VEGF levels and Microvascular Density (MVD) in CSF obtained from the different study groups

*HG* patients with high-grade gliomas, *LG* patients with low-grade gliomas, *Control* healthy individuals, *MVD* mean±standard error of 25 fields of CAM counted per patient treated with CSF, *SE* Standard Error



**Fig. 2** *Microvascular density (MVD), amount of blood vessels in*  $9000\mu m^2$  *in CAM treated with CSF from different study groups.* HG: group instilled with CSF obtained from patients with high-grade gliomas; LG: group instilled with CSF obtained from patients with low-grade gliomas; Control: group instilled with CSF obtained from healthy individuals. \*\*\* p < 0.0001. *ns* not significant. See material and methods for details

from patients with high-grade astrocytomas compared to patients with non-astrocytic tumors.

We observed that VEGF concentration in CSF from some low-grade patients was higher than controls, although not significant. Sampath et al. [30] and Ribom et al. [31] did not detect VEGF in CSF from low-grade glioma patients or control individuals. We observed low but detectable levels of VEGF in CSF from control individuals, using the same commercial kit that both authors used, although Sampath reported their results in a different unit of measurement than ours, so results are not comparable.

It is difficult to propose a biological marker for the detection of low and high-grade gliomas that would be easy to obtain and that deliver reliable results. Tumor growth is associated with local vascularization and for this reason VEGF levels have being studied in fluids such as blood and CSF. Nowacka et al. [32] reported that serum VEGFA values in patients with CNS tumors, are higher than in individuals without tumors. However, the concentration of VEGF in blood can be influenced by different factors so their levels do not reflect exclusively tissue production; Handzhiev et al. [33] analyzed concentration of VEGF and observed that lowgrade gliomas have higher VEGF levels than high-grade gliomas, both in pre- and postoperative collection from plasma. In the CNS, VEGF is released directly to the CSF so its concentration in this fluid is a reliable indicator.

Peles et al. [34] analyzed VEGF in both CSF and serum from patients with high and low-grade gliomas. They found differences in the VEGF concentration obtained from CSF but not in that obtained from serum, both in the two types of gliomas. Ribom studied VEGF levels in both fluids from patients with low-grade gliomas and the median of VEGF in serum from low-grade patients was 166.5 pg/mL, while in CSF they did not find detectable values. Serum VEGF was tested as a high grade glioma marker but without positive results when compared with control individuals [35]. Given the enormous variability of VEGF concentration, both in blood and CSF, obtained by different authors, it is necessary to consider another quantification tool of angiogenesis to discriminate between high and low-grade gliomas. It could be possible that, in some tumors, another factor besides VEGF may mediate tissular angiogenesis. To explore this idea, we combined the study of the quantity of angiogenic molecules with the determination of the biological activity in CSF.

By measuring the biological activity of CSF in a CAM bioassay, we observed a higher angiogenic capacity in eggs treated with CSF from high-grade glioma patients, compared to low- grade glioma patients and control individuals. Very few studies have evaluated the angiogenic capacity of CSF from patients with tumors. We previously reported that patients with high-grade gliomas have greater angiogenic capacity than individuals without tumors [24]. Now, using the same assay, we extended the study including low-grade glioma patients and we conclude that this increased capacity is seen only in high-grade gliomas. Given that in our results there were two discordant cases, future studies should consider a correlation between VEGF levels, tumor malignancy and the biological activity of each evaluated patient.

It is important to consider the potential relationship that exists between VEGF levels and angiogenic capacity in CSF. Although we did not study the vascularity of tumors from the patients, we did determine the angiogenic capacity of CSF from these patients using a CAM bioassay. This bioassay allowed checking the presence of some factors that could regulate the formation of blood vessels, as reported by Dai et al. [36] and Sinning et al. [24]. Our purpose was to investigate if some factors in CSF could also act on tumor tissue. The findings of Peles et al. [34] support our idea because they observed a correlation between VEGF or FGF-B levels in CSF and tumor vascularity in patients with astrocytic tumors.

An accurate diagnosis of cerebral tumors led to the determination of an appropriated clinical treatment. However, in some cases, biopsies are too difficult or risky procedures due to the location of the tumor. Efforts have been done to find biological markers for cerebral tumors, which can be obtained and measured in an easy way for guiding the diagnosis, prognosis and serve as a predictor of the efficiency of treatment. Tumoral development is associated with tissue vascularization, and then, it could be expected that angiogenic stimulators could be enhanced in CSF. Therefore, considering both factors, high levels of VEGF and angiogenic capacity in CSF could be together markers of high-grade gliomas, supporting the diagnosis with a procedure less aggressive than a biopsy, and considering that VEGF levels may vary depending on tumor progression or response to treatment; however limited by the fact that some treatments could be interfere with the results.

## Conclusion

VEGF levels in CSF and angiogenic capacity could together serve as a progression marker and glioma vascularization, being useful in the diagnosis and clinical evaluation of tumor progression. Future studies should focus on the measurement of these markers in other neoplastic and non-neoplastic diseases involving the CNS. In addition, more studies that correlate the levels of these markers with the outcome of patients and the response to treatment are needed. Thus, the measurement of the pro-angiogenic potential of CSF in gliomas could have important implications in clinical practice, because it would allow to determine the malignancy of tumors and to establish when a low-grade glioma becomes a high-grade glioma. Additionally, it would allow to perform postoperative following-up of patients, evaluate response to therapies, including antiangiogenic complementary therapies, through less expensive and invasive methods that could be implemented routinely.

**Funding** This study was funded by"Iniciación" I08/08–2 of the "Vicerrectoría de Investigación y Desarrollo" of the Universidad de Chile and the grant "Temas Libres" 324/08 of the "Oficina de apoyo a la investigación" of the Hospital Clínico de la Universidad de Chile.

### **Compliance with ethical standards**

**Conflict of interest** All authors declare that they have no conflict of interest.

Ethical approval All applicable international, national and/or institutional guidelines for the care and use of animals were followed. All procedures performed in the studies involving human participants were in accordance with the ethical standards of the institutional and/ or national research committee and with the 1964 Helsinki declaration and its later amendments or comparable ethical standards. Informed consent was obtained from all individual participants included in the study.

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