Clinicopathological characteristics of vascular endothelial growth factor expression in uveal melanoma: A meta-analysis

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Abstract. Angiogenesis is a potential prognostic factor that has been extensively investigated in patients with uveal melanoma (UM). Vascular endothelial growth factor (VEGF) expression is crucial in angiogenesis. However, there have been conflicting data regarding the clinicopathological data in UM. A meta-analysis was performed of all the germane literature to assess the clinicopathological characteristics of VEGF expression by combining separately estimated odds ratio (OR) values. Our combined results demonstrated that, according to the available studies, the expression of VEGF in UM was significantly higher compared to normal tissue [338 patients and 99 controls; OR=16.15, 95% confidence interval (CI): 8.65-30.12, P<0.00001]. When stratifying the studies by age (315 patients; OR=2.08, 95% CI: 1.19-3.62, P=0.01), cell type (423 patients; OR=0.54, 95% CI: 0.32-0.90, P=0.02), tumor size (222 patients; OR=0.30, 95% CI: 0.14-0.68, P=0.004) and scleral invasion (248 patients; OR=0.34, 95% CI: 0.15-0.78, P=0.01), significant clinicopathological information was provided. Our results indicated that VEGF expression in UM patients was significantly higher compared to that observed in controls. It was also significantly higher in patients who presented with scleral invasion and those who were aged <50 years. In addition, VEGF expression was higher in mixed-cell type and epithelioid-cell type UM and in patients with large-sized tumors.

Introduction

Uveal melanoma (UM) is a malignant tumor of the eye involving the iris, choroid or ciliary body (collectively referred

to as the uvea). It is the most common intraocular tumor, with an incidence of 5-6 cases per 1,000,000 individuals (1). There is a reported incidence of 4-6 cases per 1,000,000 individuals annually in the US and Europe (2). UM spreads through the blood and exhibits a high degree of malignancy.

Angiogenesis is a prognostic factor that has been extensively investigated in patients with UM (3). It involves the formation of new blood vessels from the endothelium of the existing vasculature. When a new tumor reaches a size of 1-2 mm, its further growth requires the formation of new blood vessels, which may lead to the development of metastases (4). As one of the most important biological markers, vascular endothelial growth factor (VEGF) is an endothelial cell mitogen and permeability factor that is potently angiogenic *in vivo* (5). VEGF induces the proliferation, differentiation and migration of vascular endothelial cells. VEGF is also required for haematopoiesis in malignant tumors, which favors primary tumor growth and metastasis (6).

Targeting constitutive VEGF and/or its receptors has been an attractive approach for cancer therapy. However, the most widely investigated clinicopathological characteristics associated with VEGF refer to variables including tumor size, largest tumor diameter (LTD), depth of invasion, lymph node status and vascular invasion. Although implicated in the pathogenesis of UM, the results on the correlation between VEGF and these factors have been conflicting and inconclusive (7-10).

It has not been elucidated whether the differences in these studies were due to the limited size of the samples or genuine heterogeneity. Therefore, in order to gain a full insight into the clinicopathological characteristics of VEGF expression in patients with UM, we enrolled data from studies of medical centers. The clinicopathological significance of our present analysis may enable a better understanding of the natural history of UM. In addition, the use of VEGF may be converted from candidate to the routine clinical setting as a predictor of the outcome of individual patients.

Materials and methods

Literature search. A search was conducted through PubMed, Embase, Cochrane Library, CNKI, VIP and Wanfang electronic databases, without language restrictions. The search was

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based on the terms 'melanoma', 'vascular endothelial growth factor' or 'VEGF', 'melanoma' and 'neovascularization'. The terms were also modified according to the different databases. The last query was updated on January 1, 2013. The references of the retrieved articles were cross-searched to identify any relevant studies that were overlooked during the electronic database search.

Inclusion criteria. The inclusion criteria for the primary studies were as follows: i) the articles included definitively diagnosed UM and normal eye tissue in humans; ii) all the eye samples with melanoma were obtained by surgery and immunohistochemical analysis was used to assess the expression of VEGF in the tumor samples; iii) the UM patients had not received immune therapy, radiotherapy or chemotherapy; iv) when multiple studies were published by the same authors or institutions, the most recent or informative was selected; and v) studies lacking clinicopathological data for meta-analysis, review articles without original data and single-case reports were excluded.

Methodological assessment. Our initial selection for all candidate studies was based on the careful screening of their abstracts by two independent reviewers (Meng Yang and Xiaocong Kuang), using a standardised data collection form, including the following items: name of first author, year of publication, ethnicity, patient gender, mean or median age, cell type, LTD, tumor size, scleral invasion, VEGF assessment method, cut-off value of VEGF positivity (%), number of readers, blinded reading (the investigator assessing the slides was blinded to the clinical information) and number of events in each category of VEGF.

We also screened the references from the relevant literature, including all the identified studies, without including additional reviews and editorials. The reference lists of the retrieved articles were also manually searched. Disagreements were resolved by consensus between the two readers. In the instance of a persistent disagreement, the final decision was made by a third expert investigator (Jianmin Li).

We did not set a predefined minimum number of patients or a minimum duration of median follow-up for a study to be included in our meta-analysis. We did not weigh each study by a quality score, since no such score has received general approval for use in a meta-analysis, particularly of observational studies, making the evaluation of its usefulness difficult. Our readers were not blinded to the studies; however, exclusions were always decided upon without knowledge of the global result of each study. When duplicate studies were retrieved, the study involving the highest number of patients from which data could be extracted (usually the latest) was included in our meta-analysis, in order to avoid overlapping between studies.

Statistical analysis. Three categories of stratified models were analyzed. The first stratified multivariate model was performed in order to confirm whether VEGF was highly expressed in UM patients compared with the controls. The second outcome of the meta-analysis was to assess the clinicopathological characteristics of VEGF expression, including patient gender, age, cell type, LTD, tumor size and scleral invasion.

According to the clinical characteristics, the following elements were combined: high and moderate VEGF expression; poor and no VEGF expression; mixed-cell and epithelioid-cell type tumors; tumors >15 mm in LTD; tumors <15 mm in LTD; small and medium tumor size; patients aged >50 years; and patients aged <50 years.

All statistical analyses were performed using RevMan statistical analysis software system, version 5.2. A two-tailed P<0.05 for the summary effect was considered to indicate a statistically significant difference. The heterogeneity of all the included studies was assessed by a statistical value I². When I^2 was <50%, the studies with an acceptable heterogeneity were considered and the fixed-effects model with the Mantel-Haenszel method was used; otherwise, a random-effects model with the DerSimonian and Laird method was adopted. The combined odds ratio (OR) was initially estimated using forest plots graphically. For each trial, the OR was estimated by a method depending on the data provided in the publication. The simplest method involved the direct retrieval of OR and its 95% confidence interval (CI) from the original article. If not available, we assessed the total numbers of events and the numbers of patients at risk in each group to estimate the OR.

The assessment of publication bias for each of the pooled study groups was performed mainly by the Egger's linear regression test. As a supplementary approach, the Begg's rank correlation was also applied to assess the potential publication bias. P<0.05 indicated that there was no publication bias in the study.

Results

Study selection. A total of 297 references were retrieved for initial reviewing using search strategies as previously described. A total of 195 citations were excluded from the analysis after the first screening based on the abstracts or titles. Following exclusion of the articles that were out of the scope of our meta-analysis, 39 potential studies were identified for detailed evaluation. Upon further review, a further 28 studies were eliminated due to the following reasons: 9 studies overlapped with others, 7 studies measured VEGF with methods other than immunohistochemistry and 12 studies lacked informative clinical data. Finally, 11 studies on the VEGF expression were included. The selection process for the studies included in this meta-analysis is summarized in Fig. 1. and the main characteristics of the eligible studies are summarized in Table I.

Correlation of VEGF expression between UM and controls. The combined results from all the studies demonstrated that the VEGF expression in patients with UM was significantly higher compared to that in controls in 7 studies (338 patients and 99 controls; OR=16.15, 95%CI: 8.65-30.12, P<0.00001). There was no statistical heterogeneity among the studies [I²=19%, degree of freedom (df)=6, P=0.28; Fig. 2A].

Correlation between VEGF expression and clinicopathological characteristics

Gender. The VEGF expression in UM according to patient gender was compared in 5 studies. VEGF expression was observed in 67 of the 102 male patients (66%) and in 62 of the 90 female patients (69%). There was no statistical

First author (publication year)	Language	Population	Study from PubMed	No. of patients (M/F)	Median age (years)	VEGF detection method	Cut-off for VEGF positivity (%)	Blinded reading ^b	No. of readers ^a	OR or HR estimate ^c	Results ^d	Refs.
Sheidow et al (2000)	English	Canada	Yes	25/22	62.1	Antibody	S	Yes	2	Data extrapolated	Positive	13
Clarijs et al (2001)	English	Netherlands	Yes	-/-	I	Antibody	I	I	I	Data extrapolated	Negative	20
Ugurel et al (2001)	English	Germany	Yes	63/62	I	Antibody	363.8 pg/ml	I	I	Reported in text	Positive	19
Boyd et al (2002)	English	Netherlands	Yes	14/16	63	Antibody	1	Yes	1	Data extrapolated	Negative	21
Wang et al (2003)	Chinese	China	No	33/32	46.1	Antibody	5	ı	7	Data extrapolated	Positive	22
Sahin <i>et al</i> (2007)	English	Turkey	Yes	10/8	39.5	Antibody	25	Yes	7	Data extrapolated	Positive	23
Zhao et al (2009)	Chinese	China	No	32/26	43	Antibody	S	Yes	7	Data extrapolated	Positive	24
Franco et al (2010)	English	Italy	Yes	23/30	61.8	Antibody	I	I	7	Data extrapolated	Positive	10
Xu et al (2011)	English	China	Yes	29/21	45.8	Antibody	S	Yes	7	Data extrapolated	Positive	25
Pang et al (2012)	Chinese	China	No	31/18	43	Antibody	I	I	7	Data extrapolated	Positive	26
Stitt et al (1998)	English	UK	Yes	15/10	59.7	Antibody	I	I	I	Data extrapolated	Positive	12
^a Readers of the histologic authors in the text. ^d Positi	al slides. ^b The r ve. inverse asso	readers of the slides sciation with VEGF	were unaware expression: ne	of the clinic: egative, no as	al patient out sociation. M	tcome. °Report I male: F fema	ed in the text, or prov le: VEGE vascular e	vided via mail	by the authors wth factor: O	s, or extrapolated from the R. odds ratio: HR. hazard	e data providec ratio: - missir	l by the

Table I. Main characteristics of the 11 studies included in the final meta-analysis



Flow chart of the literature search and inclusion strategy.

Figure 1. Selection process for the studies included in the meta-analysis.

heterogeneity among the studies ($I^2=0\%$, df=4, P=0.66). Thus, the fixed-effects model was used for statistical analysis. No association was observed between patient gender and VEGF expression (OR=0.69, 95% CI: 0.35-1.35, P=0.27).

Age. VEGF expression in UM was investigated according to patient age. The patients were divided into two groups aged >50 and <50 years. VEGF expression was observed in 116 of the 150 (77%) patients in the younger group and 100 of the 165 (61%) patients in the older group. No statistical heterogeneity was detected among the studies (I²=13%, df=6, P=0.33). Thus, the fixed-effects model was used. VEGF expression was significantly higher among patients aged <50 years (OR=2.08, 95% CI: 1.19-3.62, P=0.01).

Cell type. VEGF expression in UM according to cell type was compared in 7 studies. VEGF expression was observed in 99 of the 154 (64%) mixed- and epithelioid-cell type and in 164 of the 269 (61%) spindle-cell type tumors. The VEGF expression was lower in the spindle-cell type tumors compared to the other types (OR=0.54, 95% CI: 0.32-0.90, P=0.02). No statistical heterogeneity was detected among the studies (I²=0%, df =6, P=0.94; Fig. 2B).

LTD. The VEGF expression in UM according to LTD was compared in 5 studies. VEGF expression was observed in 122 of the 182 (67%) tumors with an LTD of >15 mm and in 69 of the 86 (80%) tumors with an LTD of <15 mm. There was statistical heterogeneity among the studies (I²=51%, df=4, P=0.09). Thus, the random-effects model was used for statistical analysis. No association was observed between LTD and VEGF expression (OR=0.43, 95% CI: 0.15-1.24, P=0.12).

Tumor size. VEGF expression in UM according to tumor size was compared in 4 studies. VEGF expression was observed in 47 of the 72 (65%) small- and medium-sized and in 121 of the 150 (81%) large-sized tumors. VEGF expression was significantly higher in large-sized tumors (OR=0.30,





0.40 [0.12, 1.32]

0.54 [0.32, 0.90]

Figure 2. Meta-analysis of (A) the association of vascular endothelial growth factor (VEGF) expression with uveal melanoma and comparison with controls and (B) the association between VEGF expression and tumor cell type.



Figure 3. Meta-analysis of (A) the association between vascular endothelial growth factor (VEGF) expression and tumor size and (B) the association between VEGF expression and scleral invasion.

95% CI: 0.14-0.68, P=0.004). There was no statistical heterogeneity among the studies (I²=26%, df=3, P=0.004; Fig. 3A).

Zhao XJ 2009

Total (95% CI)

Total events

16

99

25

154

27

164

33

269

20.8%

100.0%

Scleral invasion. VEGF expression in UM according to scleral invasion was compared in 5 studies. VEGF expression was observed in 119 of the 188 (63%) patients without and in 52 of the 60 (87%) patients with scleral invasion. VEGF expression was significantly higher in patients who exhibited scleral

invasion (OR=0.34, 95% CI: 0.15-0.78, P=0.01). There was no obvious statistical heterogeneity among the studies (I^2 =42%, df=4, P=0.14; Fig. 3B).

Sensitivity analysis and publication bias. The influence analysis revealed that no individual study significantly affected the pooled ORs and CIs. When each study was sequentially

removed and the meta-analysis was repeated with the remaining studies, the pooled OR remained essentially the same. However, the studies including analyses of VEGF expression according to patient gender, age, cell type, LTD, tumor size and scleral invasion, were <9. The Egger's linear regression and Begg's tests were not used to investigate publication bias.

Discussion

Since the VEGF was identified, there has been an increasing number of studies on the association between VEGF and cancer research. It was demonstrated that VEGF expression is strongly positive in a variety of human malignant tumors in the mRNA and/or the protein level (11). Pigment and blood vessels are abundant in uveal tissue. Tumor cells are mainly transferred through the blood; therefore, angiogenesis is crucial for the development of the tumor. However, currently available studies on VEGF expression in UM are sparse and the results obtained are inconsistent. Kvanta et al (5) reported that VEGF mRNA and protein expression were strongly positive in retinoblastoma, although there was no VEGF expression in UM. Stitt et al (12) investigated the expression of VEGF mRNA, VEGF protein and its receptor in retinoblastoma or UM and reported a strongly positive VEGF expression in the tumor; the VEGF expression rate was higher compared to that in the uninvolved normal retina, choroid and iris. Sheidow et al (13) reported that 37% (16/43) of the patients with choroidal melanoma developed distant metastases within the 10-year follow-up, but did not confirm the association of VEGF expression with distant metastasis, although the expression of VEGF in the choroidal melanoma cells was strongly positive (94% positivity rate).

To the best of our knowledge, this is the first comprehensive and detailed meta-analysis assessing the association between the expression of VEGF in UM and clinicopathological characteristics. The stratification of the baseline patient characteristics, including gender, age, cell type, LTD, tumor size and scleral invasion, our results demonstrated that VEGF expression is of clinicopathological value in UM, which may increase the predictive accuracy of prognosis in UM patients.

Our results demonstrated that high VEGF expression, as detected by immunohistochemistry, was confirmed in patients with UM according to evidence-based medicine. Notably, in the subgroup meta-analysis, we also observed that VEGF overexpression was correlated with a patient age of <50 years (P=0.008), which may explain its prognostic effect to some extent. Similar findings were also reported by other studies on age (14). However, further studies are required to assess the association of VEGF expression with age. We also observed that VEGF expression was significantly higher in mixed-cell and epithelioid-cell tumors and lower in spindle-cell tumors.

The association between tumor size and VEGF expression was not clearly determined. Our meta-analysis indicated that the VEGF expression was increased in large-sized tumors and decreased in small- and medium-sized tumors. We also demonstrated that VEGF expression was significantly higher in patients with scleral invasion. A collaborative ocular melanoma study in the USA investigated 1,091 enucleation specimens and reported a scleral invasion rate of 55% (15). The data mentioned previously questions the feasibility and safety of the implementation of local tumor resection and may explain the high recurrence rate following tumor resection in patients with UM (16). Those findings may also explain the increased formation of tumor microvasculature with the increasing degree and local infiltration of UM.

High VEGF expression induces the formation of a rich vascular network and nutritious environment, which is an active process that requires degradation of the extracellular matrix and increase in vascular permeability of blood and lymphatic vessels, favoring the progression of tumor cells into the blood and lymphatic vascular space (17). This may offer an explanation for the observed strong statistical association of VEGF overexpression with tumor invasion and metastasis (18).

The present study had several limitations. Firstly, although we did not detect significant publication bias among the studies, apart for erythrocyte sedimentation rate (ESR), it is uncertain whether the cases are comparably representative. Furthermore, the studies were observational and, therefore, more prone to biases compared with prospective randomized controlled studies. We detected publication bias for ESR (P=0.2) and there may be missing information which may reflect a negative or a more conservative association of ESR with DR4. More samples are required to validate the reliability of our conclusions.

In conclusion, our meta-analysis demonstrated that VEGF may be a marker that may enable earlier identification of high-risk patients and guide clinical decision-making regarding therapy and outcome. The evaluation of serum levels of VEGF expression may therefore play an important role in selecting melanoma patients for antiangiogenic therapy (19).

However, our conclusion should be interpreted with caution, since this analysis would be ideally performed on series of patient data. Further investigation into this subset of patients from other studies should assess the generalization of results prior to the implementation of VEGF in the routine clinical management of UM patients. Future prospective studies investigating the association of VEGF expression with survival or response to antiangiogenic therapy are required. The assessment of these angiogenic markers may be better standardised in future studies.

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