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### Prognostic Relevance of Carbonic Anhydrase-IX in High-Risk, Early-Stage Cervical Cancer: A Gynecologic Oncology Group Study

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#### Abstract

Objectives-To determine whether carbonic anhydrase-IX (CA-IX) was associated with progression-free survival (PFS) and overall survival (OS) in women with high-risk, early-stage cervical cancer treated with adjuvant pelvic radiotherapy with or without radiosensitizing chemotherapy.

Methods—CA-IX expression was detected using an immunohistochemistry assay and categorized as low when  $\leq$ 80% of tumor cells exhibited CA-IX staining and high when >80% tumor cells display CA-IX staining. Associations between CA-IX expression and clinical characteristics, angiogenesis marker expression and clinical outcome were evaluated.

Results—High CA-IX expression was observed in 35/166 (21.1%) of cases. CA-IX expression was not associated with age, race, stage, cell type, grade, positive margins, parametrial extensions, positive lymph nodes or lymphovascular space invasion but was associated with tumor size

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categorized as <2 cm, 2-2.9 cm, or  $\ge$ 3 cm (high expression: 4.7% vs. 23.2% vs. 32.5%, p=0.003) and cervical invasion confined to the inner two thirds compared with the outer third of the cervix (high expression: 6.1% vs. 23.7%, p=0.028). CA-IX expression was not associated with immunohistochemical expression of p53, CD31, CD105, thrombospondin-1 or vascular endothelial growth factor-A. Women with high versus low CA-IX expression had similar PFS (p=0.053) and significantly worse OS (p=0.044). After adjusting for prognostic clinical covariates, high CA-IX expression was an independent prognostic factor for PFS (hazard ratio [HR]=2.12; 95% confidence interval [CI]=1.13 -3.95; p=0.019) and OS (HR=2.41; 95% CI=1.24-4.68; p=0.009).

**Conclusions**—Tumor hypoxia measured by immunohistochemical expression of CA-IX is an independent prognostic factor for both PFS and OS in high-risk, early-stage cervical cancer.

#### Introduction

Cervical cancer is the second-most common cause of cancer-related deaths in women worldwide, causing an estimated 273,000 deaths annually worldwide [1] and 3,870 deaths annually in the United States [2]. Of the 19,339 cases registered with Surveillance Epidemiology and End Results (SEER) program between 1996 and 2003, 51% of cervical cancers were diagnosed as local disease, with a 5-year survival rate of 92% for these women [3]. Analysis of specimens obtained from women receiving surgical treatment for early-stage cervical cancer has identified several clinical and pathologic poor prognostic factors including increased age, African-American ethnicity, human papillomavirus (HPV) 18 infection, deep cervical stromal invasion, tumor size >2 cm, lymphovascular space invasion (LVSI), nodal metastases, microscopic tumor in uterine parametrial tissues, and positive surgical margins [4-7]. Though these prognostic factors have been well established, the biologic factors associated with recurrence and survival remain largely unknown.

In the 1990s, the antigen MN was identified [8]. MN is a 54 to 58 kDa transmembrane glycoprotein, and is a member of the carbonic anhydrase gene family, and is more specifically designated carbonic anhydrase IX (CA-IX) [9]. CA-IX is a biomarker of several types of human tumors, namely carcinomas of the cervix [10-16], kidney, esophagus and stomach, colon, head and neck, lung, and breast [17-22]. CA-IX expression in cancerous tissues and its absence in normal counterparts suggest a role in carcinogenesis [23]. Not only is CA-IX emerging as an important biomarker involved in carcinogenesis but its expression appears to be induced by hypoxia [11,23]. CA-IX expression is controlled by the transcription factor, hypoxia inducible factor-1 (HIF-1) and is up-regulated in hypoxic regions of tumor tissues. CA-IX expression was associated with microvessel density and hypoxia in head and neck squamous cell cancers [20] and with tumor stage and lymph node metastasis in cervical cancer [12]. The independent prognostic significance of CA-IX expression was recently reported in breast cancer patients [22]. The prognostic significance of CA-IX expression in carcinomas of the lung and cervix has also been examined [21,11]. In the case of carcinoma of the cervix, preliminary studies have shown that CA-IX expression is up-regulated in hypoxic regions of cervical tumors and is associated with a poor prognosis [11]. However, other reports have also indicated that there was no association of CA-IX expression and clinical outcome in cervical cancer transitional cell carcinoma of the bladder, or head and neck squamous cell carcinoma [24-26].

The purpose of this study was for the Gynecologic Oncology Group (GOG) to determine whether CA-IX was associated with progression-free survival (PFS) and overall survival (OS) in women with high-risk, early-stage cervical cancer treated with adjuvant pelvic radiotherapy with or without radiosensitizing chemotherapy on a multi-center randomized phase III trial [27]. The secondary objectives of this study were to examine the relationship between CA-IX and clinical characteristics as well as the expression of p53 and other biomarkers previously reported in this cohort, including CD31, CD105, thrombospondin-1 (TSP-1) and vascular endothelial growth factor-A (VEGF-A) [40].

#### **Materials and Methods**

#### **Eligibility Requirements**

Women who participated in a multi-center randomized phase III trial (Southwest Oncology Group 8797/GOG 109/Radiation Therapy Oncology Group 91-12) between 1991 and 1996, and provided a primary tumor specimen were eligible for this translational research study. Women were required to have undergone type III radical hysterectomy and pelvic lymphadenectomy for International Federation of Gynecology and Obstetrics (FIGO) stage IA2, IB, and IIA cervical cancer with pathologic findings of lymph node metastases, parametrial involvement, or positive surgical margins prior to treatment with adjuvant pelvic radiotherapy (RT) with or without radiosensitizing chemotherapy. Women with squamous carcinoma, adenosquamous carcinoma, and adenocarcinoma histologic subtypes were included. The results of this multi-center randomized phase III trial have been previously reported [27]. All women provided written informed consent and annual approval by the institutional review board at the GOG participating institutions in accordance with federal, state, and local requirements for the treatment protocol including primary tumor block collection for research, and approval for this translational research study was obtained from the GOG, the Cancer Therapy Evaluation Program at the National Cancer Institute and the University of California, Irvine Institutional Review Board.

#### **Treatment Regimens**

Eligible women were randomized within six weeks of surgery to receive 49.3 Gy  $\pm$  4.5 Gray delivered via a standard four-field box to the para-aortic nodes in women with positive high common iliac nodes with or without 70 mg/m<sup>2</sup> cisplatin given as a 2-hour intravenous infusion on day 1 and 1,000 mg/m<sup>2</sup> 5-FU per day for 4 consecutive days delivered as a 96-hour infusion every 21 days for a total of 4 cycles.

#### Follow Up and End Points

Follow-up consisted of physical examinations performed quarterly for 2 years, semiannually for 3 years, and annually thereafter. PFS was calculated as the time in months from date of enrollment to disease progression or death independent of cause, or date of last contact for those with no evidence of disease progression. OS was calculated as the time in months from date of enrollment to death independent of cause, or date of last contact for those who were still alive.

#### **Primary Tumor Specimens**

Formalin-fixed and paraffin-embedded (FFPE) pre-treatment radical hysterectomy specimens or large excisional biopsies were cut into 4 micron sections and fixed onto positively-charged glass slides. Histologic eligibility was verified by the GOG Pathology Committee, and the adequacy of FFPE primary tumor block for research was determined by the study pathologist (SYL) who examined hematoxylin and eosin-stained tissue specimens to confirm that at least 50% of each section consisted of malignant tissue.

#### Immunohistochemical Expression of CA-IX

Immunohistochemical staining of tissue sections with anti-CA-IX antibody was performed using a peroxidase technique with pressure cooking pretreatment [10]. Briefly, the specimens were de-paraffinized in xylene, rehydrated in graded alcohols, and rinsed in distilled, deionized water. Slides were placed in antigen retrieval citrate solution and pre-treated with pressure

cooking, and then incubated with a 3% (v/v) hydrogen peroxide solution for 10 minutes and 5% (v/v) normal horse serum in phosphate buffered saline (PBS) for 20 minutes. Specimens were incubated at ambient room temperature with an anti-CA-IX mouse monoclonal antibody (M75) with 1:10,000 dilution for an hour, a 1:200 dilution of biotinylated horse anti-mouse immunoglobulin G in PBS for 30 minutes, avidin-biotin peroxidase complex (ABC Elite, Vector Laboratories, Burlingame, CA) for 30 minutes, and 3',3'-diaminobenzidine chromogen solution for 10 minutes. Specimens were counter-stained with hematoxylin (Dako, Carpenteria, CA), rinsed in double distilled water, and mounted with Aquamount. Positive and negative controls were included in each run. CA-IX immunostaining was evaluated by one of authors (SYL) blinded to all clinical and outcome data. Specific immunohistochemical staining was defined by the presence of a brown reaction product on the plasma membrane under  $400 \times$ magnification (with 10× ocular lens). Faint staining of the cytoplasm was considered negative. The intensity of immunoreactivity was scored as strong (2+) and weak (1+). Strong positivity was defined as dark brown immunoreactivity that was easily identified at a low power magnification ( $40 \times$  or  $100 \times$ ). A reasonable degree of heterogeneity of CA-IX-positive staining was noted; therefore, based on the average percentage of strongly (2+) positive cells present in the entire tissue section, the CA-IX immunoreactivity was divided into four groups: (A) >80% tumor cells positive; (B) 40 to 80% of tumor cells positive; (C) 15-39% of tumor cells positive; (D) <15% tumor cells positive or no immunoreactivity was present. We further classified tumors that were uniformly hypoxic across the entire section (Group A) from those that exhibited limited hypoxia or were not hypoxic (Groups B, C and D) as illustrated in Figure 1.

#### Immunohistochemical Expression of VEGF-A, TSP-1, CD31, CD105 and p53

Semi-quantitative immunohistochemical (IHC) staining for vascular endothelial growth factor-A (VEGF-A, pro-angiogenesis factor), thrombospondin-1 (TSP-1, anti-angiogenesis factor), CD31 (pan endothelial marker), and CD105 (tumor-specific endothelial marker) was evaluated as previously described [28]. Briefly, p53 was detected using the N-terminal DO-1 anti-p53 clone that recognizes the major normal and mutant p53 isoforms but not the isoforms lacking the first 40 or 133 amino acids of full length p53 (Santa Cruz Biotechnology, Santa Cruz, CA). TSP-1 and p53 were categorized as negative or positive [28]. CD31 and CD105 microvessel density (MVD) "hotspots" were counted in three  $20 \times$  high-power fields and categorized as low (<110 and <28) or high ( $\geq$ 110 and  $\geq$ 28), respectively [28]. Tumoral histoscores (HS) were calculated for VEGF-A using the formula: [% cells positive × (intensity +1)] and categorized as low (<200) or high ( $\geq$ 200) [28].

#### **Statistical Analysis**

Biomarker and clinical data for this ancillary study were analyzed using SPSS version 14 (SPSS Inc., Chicago, IL) and SAS version 9.1 (SAS Institute, Inc., Cary, NC). CA-IX expression was evaluated and categorized as low when  $\leq$ 80% of tumor cells exhibited CA-IX staining and high when >80% tumor cells display CA-IX staining. Exact testing was used to test the hypothesis of independence between CA-IX expression and clinical characteristics or biomarkers [29, 30]. Estimates of the survival probabilities were calculated using the Kaplan-Meier product limit method [31], and the logrank test employed to test the null hypothesis of equality between strata [32,33]. Hazard ratios were estimated using Cox proportional hazard regression analysis [34], without or with adjustments for prognostic clinical covariates (age, race, depth of cervical invasion, parametrial extensions, positive lymph nodes and treatment) for PFS and OS. These covariates were selected for adjustment based on their documented prognostic value in cervical cancer [4-7].

#### Results

Of the 243 eligible and evaluable women who participated in the multi-center randomized phase III trial there were 180 women who were enrolled at GOG institutions and provided FFPE primary tumor tissue for translational research and 166 of these women were eligible for this translational research study. One hundred and fifty three of the specimens (92.2%) were from radical hysterectomies and 13 (7.8%) were from large excisional biopsies. Fourteen women were excluded for the following reasons: benign disease (n=1), low-risk, early stage disease (n=2), no FFPE tumor for testing (n=4), and inevaluable for CA-IX expression (n=7). The patient characteristics for the 166 women in this cohort are summarized in Table 1 and are representative of that observed in the entire cohort of women who participated in the phase III intergroup trial [27]. At the time of the final analyses, 110 women were alive with no evidence of disease, 5 were alive with disease progression, 43 died due to disease progression, 5 died due to a reason other than disease progression or treatment, and three died of unknown cause. Median follow-up for the 115 women who were still alive at the time of the final analysis was 105.9 (range: 2.7 to 184.8) months. The distribution of cases by treatment was as follows: 86 (51.8%) women were randomized to radiation and 80 (48.2%) women were randomized to chemoradiation.

Figure 1 displays representative immunostaining for CA-IX. High level expression was observed in 35/166 (21.1%) of cases when >80% tumor cells displayed strong CA-IX immunostaining (2+ intensity) that was distributed throughout the tumor nests in the entire tissue section. CA-IX expression was not associated with age, race, stage, cell type, grade, positive margins, parametrial extensions, positive lymph nodes or lymphovascular space invasion (Table 1) but was correlated with tumor size categorized as <2 cm, 2-2.9 cm, or  $\geq$ 3 cm (high expression: 5% vs. 23% vs. 33%; p=0.003). Tumors with invasion into the outer third of the cervix also had higher expression of CA-IX than those confined to the inner two thirds (high expression: 24% vs. 6%, p=0.028). CA-IX expression was not associated with immunohistochemical expression of p53, CD31, CD105, TSP-1 or VEGF-A (Table 2).

The Kaplan-Meier method was used to estimate PFS and OS for women with low compared with high CA-IX expression (Figure 2). Women with high versus low CA-IX expression had similar PFS (Figure 2A, p=0.053) and significantly worse OS (Figure 2B, p=0.044). The five-year OS was 78% vs. 61% for women with tumors with low or high CA-IX expression, respectively.

Unadjusted and adjusted Cox regression analyses were performed to examine the association between CA-IX expression and PFS or OS (Table 3). Women whose tumor exhibited high compared with low CA-IX expression had a similar risk of disease progression (hazard ratio [HR]=1.760; 95% confidence interval [CI]=0.985-3.144; p=0.056) and a significant increase in the risk of death (HR=1.840; 95% CI=1.007-3.362; p=0.047). After adjusting for prognostic clinical covariates, high CA-IX expression was an independent prognostic factor for PFS (HR=2.12; 95% CI=1.13-3.95; p=0.019) and OS (HR=2.41; 95% CI=1.24-4.68; p=0.009).

In follow up to these findings, an analysis was then performed to explore the relationship between CD31-MVD (detailed analysis previously reported in [28]) and CA-IX in this cohort to provide enhanced resolution regarding tumor vascularization and permeability within the tumor tissue from these patients, as CD31-MVD contributes a hot spot assessment of regions within the tumor that exhibit the highest degree of vascularization whereas CA-IX expression provides a score for whether or not the tumor is exhibiting uniform hypoxia throughout the entire tissue section. Categorized CD31-MVD was not correlated with categorized CA-IX expression (Table 2). Specifically, women who have tumors with high CD31-MVD (angiogenesis) and low CA-IX (hypoxia) had the best PFS (Figure 3A) and OS (Figure 3B).

Those with low CD31-MVD and high CA-IX had the worst PFS and OS (Figure 3). Those with either low CD31-MVD (angiogenesis) and low CA-IX (hypoxia) or high CD31-MVD and high CA-IX had intermediate PFS and OS (Figure 3). Inclusion of both biomarkers in an adjusted Cox regression model demonstrated that after adjusting for CD31-MVD, high CA-IX expression was associated with increased risk of death (Table 3). In addition, after adjusting for CA-IX expression, high CD31-MVD was associated with a reduced risk of progression and death (Table 3).

#### Discussion

Experimental and clinical studies have shown that hypoxic tumors are not only resistant to radiation and chemotherapy, but are also associated with genetic instability and increased risk of invasion, metastasis, and poor clinical outcome. The intrinsic biological aggressiveness of hypoxic tumors is explained, in part, by the up-regulation of a number of hypoxia-inducible genes mediated by the activation of the transcription factor, HIF-1. Investigators continue to search for reliable diagnostic, prognostic and predictive biomarkers that can assist in clinical treatment and management decisions and identify new therapeutic targets. CA-IX has recently emerged as one of the most promising endogenous markers of cellular hypoxia and has been validated directly or indirectly by various studies [11,35,36].

Although one cervical cancer study showed that CA-IX was not associated with clinical outcome [24], others demonstrate that high CA-IX expression was associated with worse survival in cervical carcinoma [11,14,15]. In this study, we confirm the observation that high CA-IX expression was associated with worse clinical outcome Our study defined high levels of CA-IX expression as more than 80% of tumor cells in entire tissue section exhibiting strong CA-IX immunoreactivity (2+ intensity) because this represents a uniform diffuse distribution of hypoxia throughout the carcinoma. Based on this definition, 21% (35/166) of the tumors exhibited high levels of CA-IX expression (uniform hypoxia). Women with high levels of CA-IX expression had an increased risk of disease progression and death. In addition, categorized CA-IX expression was an independent prognostic factor for PFS and OS after adjusting for clinical covariates with documented prognostic relevance in this cohort [6] and in other studies [4-5,7]. Differences in stage of disease, sample type and size as well as treatment regimens may influence, at least in part, the disparity in findings [24] compared with others [11,14,15] and that described herein.

CA-IX expression was not associated with age, race, stage, cell type, grade, positive margins, parametrial extensions, positive lymph nodes or lymphovascular space invasion (Table 1), consistent with prior publications[11,14]. However, CA-IX expression did correlate with categorized tumor size and the depth of cervical invasion. These findings are biologically plausible given that CA-IX expression will be induced via HIF-mediated transcription when the diffusional capacity of the tumor vasculature is exceeded as the tumor increases in size and invades deeper into the surrounding tissue [11,35-38].

It is well established that hypoxia results in the upregulation of genes that facilitate anaerobic metabolism and promote tumor vascularization (e.g. VEGF). Although the study did not show a correlation between CA-IX expression and several angiogenesis biomarkers including CD31-MVD, CD105-MVD, TSP-1 and VEGF-A or p53, an interesting association was observed between CD31-MVD, CA-IX expression and prognosis. Specifically, women who have tumors with high CD31-MVD (angiogenesis) and low CA-IX (hypoxia) had the best PFS and OS while those with low CD31-MVD (angiogenesis) and high CA-IX (hypoxia) had the worst PFS and OS (Figure 3). Those with either low CD31-MVD (angiogenesis) and low CA-IX (hypoxia) or high CD31-MVD and high CA-IX had intermediate PFS and OS (Figure 3). Adjusted Cox regression modeling demonstrated that CA-IX was a prognostic factor for OS

even after adjusting for CD31-MVD. Taken together, these findings extend our previous hypothesis [28] by clarifying that improved survival is observed in the well-vascularized and well-oxygenated cervical cancers with high CD31-MVD and low CA-IX expression.

In conclusion, a high level of CA-IX expression in tumors appears to be an independent prognostic indicator in women with high-risk, early stage cervical carcinoma and may provide a number of implications for cancer patient management such as serving as an indicator of the patient selection in combination with CD31-MVD and prognostic clinical covariates for possible hypoxia-modifying therapy, bio-reductive drugs administration or combination regimens that utilize immunomodulatory agents. Moreover, CA-IX is an emerging target for cancer therapy and functional imaging [39-41], further broadening the potential clinical value of the CA-IX biomarker in the detection, treatment and management of women with cervical cancer.

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#### Figure 1.

Representative examples of CA-IX expression in squamous cell carcinoma of the cervix: Low expression (A): Focal CA-IX expression (brown immunostain) limited to a few neoplastic cell clusters (long arrow), the majority of carcinoma nests are negative (short arrow). High expression (B): Diffuse, strong (2+) immunoreactivity is seen in >80% of the neoplastic cells (arrow) in a single section. The stroma ( $\blacklozenge$ ) and areas of necrotic tumor are negative. Original magnification ×100.

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#### Figure 2.

Kaplan-Meier plots for progression-free survival (A) and overall survival (B) for women categorized by low or high CA-IX expression.

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#### Figure 3.

Kaplan-Meier plots for progression-free survival (A) and overall survival (B) for women categorized by CA-IX and CD31-MVD expression.

		CA-IX E	xpression *	Exact test
Clinical Characteristics	Cases	Low Cases (%)	High Cases (%)	- p-value
Age **				0.121
<40	91	72 (79)	19 (21)	
40-49	37	32 (86)	5 (14)	
50-59	23	18 (78)	5 (22)	
60-69	12	6 (50)	6 (50)	
70-79	3	3 (100)	0 (0)	
Race and Ethnicity				0.498
Caucasian	104	79 (76)	25 (24)	
African American	29	24 (83)	5 (17)	
Hispanic	25	20 (80)	5 (20)	
Other $^{\dagger}$	8	8 (100)	0 (0)	
Stage				1.000
IB	158	124 (78)	34 (22)	
IIA	8	7 (88)	1 (12)	
Cell Type				0.810
Squamous Carcinoma	135	107 (79)	28 (21)	
Other Carcinoma $^{\dagger\dagger}$	31	24 (77)	7 (23)	
Grade				0.240
1	13	8 (62)	5 (38)	
2	80	63 (79)	17 (21)	
3	73	60 (82)	13 (18)	
Tumor Size				0.003
<2 cm	43	41 (95)	2 (5)	
2-2.9 cm	82	63 (77)	19 (23)	
≥3 cm	40	27 (68)	13 (32)	
Depth of Cervical Invasion				0.028
Inner two-thirds ‡	33	31 (94)	2 (6)	
Outer third	131	100 (76)	31 (24)	
Positive Margin				1.000
No	156	123 (79)	33 (21)	
Yes	10	8 (80)	2 (20)	
Parametrial Extensions				1.000
No	107	84 (79)	23 (21)	
Yes	59	47 (80)	12 (20)	
Positive Nodes				0.451
No	27	23 (85)	4 (15)	
Yes ‡‡	139	108 (78)	31 (22)	

## Table 1 Relationship between CA-IX expression and clinical characteristics

		CA-IX E	xpression *	Exact test
Clinical Characteristics	Cases	Low Cases (%)	High Cases (%)	p-value
Lymphovascular Space Invasion	1			0.378
No	41	30 (73)	11 (27)	
Yes	124	100 (81)	24 (19)	

Cases (row percentage  $\times$  100).

\* CA-IX expression was categorized as low when ≤80% of tumor cells exhibited CA-IX staining and high when >80% tumor cells display CA-IX staining.

\*\* Median age at enrollment for the entire cohort was 39.03 years.

 $^{\dagger}$ Other includes Asian/Pacific Islander (5), Filipino (2), and Native American (1).

 $^{\dagger\dagger}$  Other carcinoma includes 20 adenocarcinomas and 11 adenosquamous carcinomas

 ${}^{\ddagger}$  Five invaded the inner third and 28 invaded the middle third of the cervix.

 $\ddagger$  There were 68 women with one and 71 with two or more positive lymph nodes.

## Table 2Relationship between CA-IX and other biomarkers including p53, CD31, CD105, TSP-1,and VEGFA

	CA-IX E	kpression *	Exact test
	Low Cases (%)	High Cases (%)	p-value
p53			0.445
Negative	57 (75)	19 (25)	
Positive	68 (81)	16 (19)	
CD31-Microvessel Density			1.000
Low <110	81 (78)	23 (22)	
High $\geq 110$	44 (79)	12 (21)	
CD105-Microvessel Density			0.823
Low <28	36 (78)	10 (22)	
High $\geq 28$	75 (80)	19 (20)	
TSP-1			1.000
Negative	40 (80)	10 (20)	
Positive	87 (78)	25 (22)	
VEGF-A Histoscore			0.150
Low <200	39 (72)	15 (28)	
High $\geq 200$	86 (83)	18 (17)	

Cases (row percentage  $\times$  100).

\*CA-IX expression was categorized as low when ≤80% of tumor cells exhibited CA-IX staining and high when >80% tumor cells display CA-IX staining.

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# Table 3

Associations between CA-IX expression and progression-free survival or overall survival

Biomarker Expression $^{\dagger}$	Cn.	adjusted Cox M	odel	AG	ijusted Cox Mode	* [0	Uni	adjusted Cox M	odel	PA	justed Cox Mode	ž ľ
	HR	95% CI	d	H	95% CI	d	HR	95% CI	ď	HR	95% CI	d
CA-IX												
Low	1.00			1.00			1.00			1.00		
High	1.76	0.99-3.14	0.056	2.12	1.13-3.95	0.019	1.84	1.01-3.36	0.047	2.41	1.24-4.68	0.009
CA-IX*												
Low	1.00			1.00			1.00					
High	1.71	0.94-3.11	0.077	1.82	0.96-3.46	0.067	1.77	0.95-3.29	0.072	2.07	1.05-4.10	0.037
CD31-MVD*												
Low	1.00			1.00			1.00			1.00		
High	0.36	0.18-0.71	0.003	0.35	0.17-0.70	0.003	0.36	0.17-0.73	0.005	0.33	0.16 - 0.69	0.003

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microvessel density (MVD) <110 and high for tumors with a MVD ≥110.

 ${}^{\sharp}$ Adjusted for patient age, race, stage, depth of cervical invasion, parametrial extensions, positive lymph nodes and treatment.

\* CA-IX and CD31-MVD were both included in the Cox regression analyses for PFS and OS.