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Anterior Gradient 3 Promotes Breast Cancer Development and Chemotherapy Response

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Purpose

Anterior gradient 3 (AGR3) belongs to human anterior gradient (AGR) family. The function of AGR3 on cancer remains unknown. This research aimed to investigate if AGR3 had prognostic values in invasive ductal carcinoma (IDC) of breast cancer and could promote tumor progression.

Materials and Methods

AGR3 expression was detected in breast benign lesions, ductal carcinoma *in situ* and IDC by immunohistochemistry analysis. AGR3's correlations with clinicopathological features and prognosis of IDC patients were analyzed. By cell function experiments, collagen gel droplet-embedded culture drug sensitivity test and cytotoxic analysis, AGR3's impacts on proliferation, invasion ability, and chemotherapeutic drug sensitivity of breast cancer cells were also detected.

Results

AGR3 was up-regulated in luminal subtype of histological grade I-II of IDC patients and positively correlated with high risks of recurrence and distant metastasis. AGR3 high expression could lead to bone or liver metastasis and predict poor prognosis of luminal B. In cell lines, AGR3 could promote proliferation and invasion ability of breast cancer cells which were consistent with clinical analysis. Besides, AGR3 could indicate poor prognosis of breast cancer patients treated with taxane but a favorable prognosis with 5-fluoropyrimidines. And breast cancer cells with AGR3 high expression were resistant to taxane but sensitive to 5-fluoropyrimidines.

Conclusion

AGR3 might be a potential prognostic indicator in luminal B subtype of IDC patients of histological grade I-II. And patients with AGR3 high expression should be treated with chemotherapy regimens consisting of 5-fluoropyrimidines but no taxane.

> Key words Anterior gradient 3, Cancer, Prognosis

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Introduction

Breast cancer is one of the most common cancers in women and has been a major cause for cancer death of women [1]. In clinical work, breast cancer is classified into four molecule subtypes mainly based on status of estrogen receptor (ER), progesterone receptor (PR), Ki67, and human epidermal growth factor receptor 2 (HER2), consisting of luminal A, luminal B, HER2-overexpressing and triple-negative breast cancer (TNBC) [2]. Luminal A and B are ER and PR positive whereas the other two are negative. Among of them, luminal subtype accounts for approximately 70% of all breast cancer patients, TNBC accounts for about 19% and HER2-overexpressing accounts for the rest [2-4]. The presence of ER means that tumor may respond to endocrine therapy taking ER signaling pathways as targets. So endocrine therapy has been a major method to treat luminal patients, luminal A is much more sensitive to this method than luminal B [3] and patients with luminal A need no other treatment such as extra chemotherapy [4]. In luminal B, additional chemotherapy combined with endocrine therapy is very important. But the resistance of chemotherapy in this subtype is common, so recurrence rate is much higher than luminal A and new molecular targets should be developed to improve outcome of patients [5].

The proteins of human anterior gradient (AGR) family have three members, respectively named AGR1, AGR2, and AGR3. In higher vertebrates, AGR1 is lost due to decreased regeneration ability, but AGR2 and AGR3 still express [6]. However, physiological functions of AGR2 and AGR3 in vertebrates are poorly understood. In tumors such as breast [7-10], ovarian [11], prostate [12], esophageal [13], lung [14], and pancreas cancers [15], AGR2 has been acknowledged as an oncogene and a biomarker indicating poor prognosis of patients. Comparing to definite role of AGR2, AGR3's impacts on cancers are still unclear. Reports about AGR3 in cancers are few and contradictory.

In our clinicopathologic study, we performed immunohistochemistry (IHC) analysis to detect AGR3 expression in 51 cases of breast benign lesions, 62 cases of ductal carcinoma *in situ* (DCIS) and 336 cases of invasive ductal carcinoma (IDC). Our samples were randomly selected and covered all molecular subtypes and all histological grades. We confirmed that AGR3 mainly expressed in the IDC patients of luminal subtype of histological grade I-II and could predict poor prognosis of luminal B. AGR3 highly expressed IDC patients tend to occur bone metastasis and liver metastasis. In breast cancer cell lines, we found that AGR3 could promote proliferation and invasion ability of cancer cells which proved that AGR3 could accelerate progression of breast cancer from the perspective of cell functions. Besides, AGR3 high expression indicated a poor prognosis of breast cancer patients treated with taxane but a favorable prognosis treated with 5-fluoropyrimidines. Performing collagen gel dropletembedded culture drug sensitivity test (CD-DST) and cytotoxic analysis to achieve more results, we found that breast cancer cells with AGR3 high expression were resistant to taxane but sensitive to 5-fluoropyrimidines. All of these results above had never been reported before.

Materials and Methods

1. Patient selection and clinical information

Paraffin-embedded specimens were randomly selected from 51 cases of breast benign lesions, 62 cases of DCIS and 336 cases of IDC diagnosed between 2004 and 2009. All specimens came from the Department of Breast Cancer Pathology and Research Laboratory, Tianjin Medical University Cancer Institute and Hospital (Tianjin, China). All diagnosis of histopathology was confirmed according to World Health Organization (WHO) criteria. The median age of IDC patients was 52 years old which ranged from 28 to 89. None of them had received any oncotherapy before operation. Three hundred thirty-six cases of IDC patients included 240 cases of luminal subtype (71.4%), 28 cases of HER2-overexpressing subtype (8.3%), 62 cases of triple-negative subtype (18.5%), and six cases without information about molecular subtype (1.8%).

After operation, the chemotherapy regimens that IDC patients received included 118 cases of TE/TA (taxane+epirubicin/taxane+doxorubicin) regimen, 111 cases of CEF (cyclophosphamide, epirubicin, and 5-fluoropyrimidines)/ CAF (cyclophosphamide, doxorubicin, and 5-fluoropyrimidines) regimen, 78 cases of CMF (cyclophosphamide, methotrexate, and 5-fluoropyrimidines) regimen, nine cases of platinum-based regimen and seven cases just treated with taxane.

2. Patients' prognostic information

All IDC patients were followed up with 5-146 months. In this period, 14 cases (4.2%) developed tumor recurrence, 58 cases (17.3%) developed distant metastasis, and 25 cases (7.4%) died for cancer. And 56 cases developed cancer progression within 5 years.

In addition, 58 cases of patients with distant metastasis included 43 cases of bone metastasis, 17 cases of lung metastasis, 13 cases of liver metastasis, eight cases of brain metastasis, and 19 cases with multiple organic metastases.

3. Immunohistochemical staining

AGR3 expression was detected by IHC. All procedures were performed in Benchmark XT (Roche, Basel, Switzerland). Antigen retrieval was performed in citrate buffer at 121°C for 2 minutes 15 seconds. After retrieval, sections were steeped in 3% H_2O_2 buffer for 25 minutes and in 10% goat serum buffer for 25 minutes. Then, the sections were steeped in primary antibody against AGR3 (1:100, HPA053942, Sigma, St. Louis, MO) at 4°C for one night. Next, sections were steeped in anti-rabbit secondary antibody for 20 minutes and in horseradish peroxidase–conjugated streptavidin also for 20 minutes. Finally, sections were incubated with 3,3'-diaminobenzidinetera hydrochloride which was peroxidase substrate.

4. Evaluation of staining

All stained sections were reviewed independently by two pathologists under light microscopy. Evaluation of AGR3 depended on staining intensity and area. Intensity was graded like this: 0 (–) no staining, 1 (+) weak staining, 2 (++) moderate staining, and 3 (+++) strong staining. Then patients were categorized into two groups: group 1 (the grade of intensity was 2-3 and the staining area of tumor cells was 75%-100%) and group 2 (the grade of intensity was 1 or the staining area of tumor cells was 0%-74%). Additionally, group1 was defined as AGR3 high expression and group 2 was defined as AGR3 low expression.

Evaluation of ER and PR was based on the 2010 American Society of Clinical Oncology/College of American Pathologists (ASCO/CAP) guideline. Sections were regarded as positive cases if the staining area of tumor cells with nuclear staining was more than 1%.

5. Cell culture and reagents

MDA-MB-231 and T47D cell lines were bought from American Type Culture Collection (Manassas, VA) which had been checked by DNA profiling. MDA-MB-231 were cultured with Dulbecco's modified Eagle's medium and T47D were cultured with 1640 medium which were mixed with 10% fetal bovine serum (FBS). Both two cell lines were incubated in a 5% CO₂ incubator at 37°C.

6. Construction of lentiviral vector expressed with full length of AGR3

Full length of human AGR3 gene (GenBank accession No. NM_176813) was amplified in T47D cell line by polymerase chain reaction (PCR). The PCR primers were as follows: for-ward, 5'-GAGAGCTAGCGCCACCATGATGCTACACTC-

AGCT-3' and reverse, 5'-TCTCGGATCCTTAAGCGTAAT-CTGGAACATCGTATGGGTACATTAGCTCTGACT-GAATAAGTCTTAATGCTTT-3'. The production of PCR contained full length of AGR3 gene, $3 \times$ Flag label and HA label. Then the production was cloned into pCDH-CMV-MCS-EF1-Puro lentiviral vector. The sequences of the inserts were checked by gene sequencing technique. Lentiviral plasmid, packing plasmids Δ R and pVSVg were transfected into HEK-293T cells to make lentiviruses.

Then, MDA-MB-231 cells were infected by lentiviruses and screened by puromycin. After verification by western blot, stable clones were named as 3×Flag-AGR3-HA/MDA-MB-231.

7. Construction of lentiviral vector knocking down AGR3

Two different AGR3 specific RNA interference sequences were applied (siAGR3#1-#2). The sequences were as follows: siAGR3#1, 5'-gccttcacttcaaagaagtca-3'; siAGR3#2, 5'-gatga-catcacttgggtacaa-3'. Lentiviral plasmid, packing plasmids Δ R and pVSVg were transfected into HEK-293T cells to make lentiviruses. T47D cells were infected by lentiviruses and verified by western blot. Cells knocking down AGR3 successfully were named as siAGR3/T47D.

8. Western blot

Western blot was performed to detect AGR3 expression of breast cancer tissues and cells. The protein was lysed by lysis buffer (sodium dodecyl sulfate) on ice. After separating with sodium dodecyl sulfate polyacrylamide gel electrophoresis, protein was electro-transferred onto nitrocellulose membranes. The nitrocellulose membranes were incubated by primary antibodies. The primary antibodies contained AGR3 (ab201464, Abcam, Cambridge, UK), Flag (AF519–1, Beyotime, Shanghai, China), and β -actin (SC-47778, Santa Cruz Biotechnology, Santa Cruz, CA). Then the nitrocellulose membranes were treated with secondary antibodies.

9. Cell ATP/viability assay and sulforhodamine B assay

In cell ATP/viability assay, the cell lines were incubated in 24-well plates. Every type of cell line had six replicates. After 5 days, ATP level was measured by Cell Titer-Glo luminescent cell viability assay kit (catalog number G7571, Promega, Madison, WI). In sulforhodamine B (SRB) assay, cells were also incubated in 24-well plates for 5 days. After that, cells were fixed with 10% trichloroacetic acid and then stained with SRB (0.4%) buffer in 1% acetic acid. Lastly, added Tris-base (10 mM) into the plate to dissolve SRB and measured absorbance at 564 nm.

10. Migration assay

Migration assays were performed on cell lines using 24well transwell chambers (Corning, New York, NY). Cells with 200 μ L serum-free medium were added into the upper chambers and 600 μ L medium containing 5% FBS were added into the lower chambers. The cells were incubated at 37°C for 16 hours. Then, scraped the cells at upper layer of polyethylene membrane and stained the cells at lower layer with Giemsa solution. Lastly, taken photograph under light microscopy and quantified the cell number in randomly selected fields. All experiments were performed in triplicates.

11. Invasion assay

Invasion assays were also performed on cell lines using 24well transwell chambers (Corning). But the upper chambers were coated with prediluted extracellular matrix. Then cells with medium containing 0.5% FBS were added into the upper chambers and 600 μ L medium containing 10% FBS were added into the lower chambers. After incubating at 37°C for 24 hours, the invading cells were stained, counted and photographed under light microscopy as the procedure of migration assay. All experiments were performed in triplicates.

12. Cytotoxic analysis

To measure the cytotoxic effects of 5-fluoropyrimidine (5-FU) and taxane, 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide (MTT) assays were performed on cell lines. Briefly, cells were seeded at 24-well plates. After incubation to allow cells adhesion, cells were exposed to various concentrations of taxane or 5-FU (MDA-MB-231, taxane: 15 to 5×10⁴ ng/mL, 5-FU: 1 to 1.5×10⁴ µg/mL; T47D, taxane: 2.5 to 2.5×10^4 ng/mL, 5-FU: 0.1 to 5×10^3 µg/mL) with 500 µL/ well medium for 48 hours. Besides, drug-free medium was added to the control and blank wells. After exposure was finished, 500 μ L MTT (5 mg/mL) were added to each well. After 4-hour incubation, MTT solution was removed and 500 μ L dimethyl sulfoxide (Sigma) were added into the wells to dissolve the insoluble formazan crystals. After dissolution, 150 µL supernatant of every well was removed into the 96-well plates and the absorbance of the dye was measured at a wave length of 570 nm. All experiments were performed in triplicates and calculated the IC₅₀ value.

13. Wound healing assay

To measure the migration ability of the cell lines, wound healing assay was performed. The cells were seeded at 6-well plates. Then, a scratch was performed with a 10 μ L pipette

tip. After removing cell debris by phosphate-buffered saline, cells were incubated in a medium without FBS at 37°C. The migration distances were measured at 0, 6, 12, 18, and 24 hours. The cell-free area of the wounds was estimate under a microscope. All assays were performed in triplicates.

14. Statistical analyses

Using SPSS ver. 17.0 software package (SPSS Inc., Chicago, IL) to perform statistical analysis. Group comparisons were calculated by Mann-Whitney U test, Kruskal-Wallis test, ANOVA test, and chi-square test. Two variables' correlation was calculated by Spearman's rank correlation test. Using Kaplan-Meier method to calculated patients' overall survival (OS) survival and progression-free survival (PFS) time. p-values were computed by log-rank test. The relevant factors for patients' prognosis were identified by Cox proportional hazards regression model. *In vitro* experiments, using two-tailed Student's t test to detect if the difference between two groups had statistical meaning or not. In all statistical analyses, two-sided p < 0.05 was regarded as statistical significance.

15. Ethical statement

All information about patients in this study were obtained with informed consent. And this study was approved by the Ethic Committee of Tianjin Medical University Cancer Institute and Hospital. The approval number of ethic committee was bc2017019.

Results

1. AGR3 expression increased with the development of breast tumor malignancy

We detected AGR3 expression at proteomic level by IHC in 51 cases of breast benign lesion tissues, 62 cases of DCIS tissues and 336 cases of IDC tissues. The immunohistochemical staining of AGR3 was assessed and representative images of staining were shown in Fig. 1A. We found that AGR3 mainly located in cytoplasm of epithelial cells of mammary gland ducts in breast tissues. From benign lesions to DCIS and to IDC, AGR3 expression gradually evaluated (Fig. 1B). 7.8% (4/51) of benign lesions, 27.4% (17/62) of DCIS and 38.4% (129/336) of IDC specimens showed high expression of AGR3 (r_s =0.195, p < 0.001) (S1 Table). This indicated that AGR3 expression increased with the development of breast tumor malignancy.



Fig. 1. Anterior gradient 3 (AGR3) expression increased with the development of breast tumor malignancy and was negatively correlated with histological grade but positively correlated with estrogen receptor (ER)/progesterone receptor (PR) status. (A) Representative immunohistochemistry (IHC) images of AGR3 low expression group (upper) and AGR3 high expression group (lower). (B) AGR3 expression increased with progression of malignancy degree of lesions. AGR3 expression of benign lesions was low, ductal carcinoma *in situ* (DCIS) was higher and IDC was the highest. (C) Representative IHC images of AGR3 expression in different histological grades of IDC (left panel). Western blot analysis of AGR3 expression in the frozen breast tumor specimens consisting of normal tissues, grade I, grade II, and grade III. Every type of tissues had 6 cases. β -actin was used as a loading control (right panel). (*Continued to the next page*)



Fig. 1. (*Continued from the previous page*) (D) The expression of AGR3, ER, and PR were detected by IHC in serial paraffin sections. The upper panel was representative IHC images of AGR3 and ER. The lower was AGR3 and PR. (E) Representative IHC images of AGR3 expression in disease-free group and recurrence or metastasis group.

2. AGR3 mainly expressed in luminal subtype of IDC patients of histological grade I-II

In IDC, we analyzed the relationship between AGR3 expression and clinicopathological features of patients. We found that high expression of AGR3 was associated with high ER positivity (r_s =0.336, p < 0.001), high PR positivity

(r_s =0.225, p < 0.001), high risk of recurrence or metastasis (r_s =0.179, p=0.001), and low histological grade (r_s =-0.160, p=0.004) (Table 1). The relationships between AGR3 expression and patients' age (r_s =-0.099, p=0.069), tumor size (r_s = 0.041, p=0.457), lymph node metastasis (r_s =-0.007, p=0.901), distant metastasis (r_s =0.077, p=0.161), HER2 status (r_s =-0.098, p=0.074), and Ki-67 status (r_s =-0.009, p=0.874) were not obvi-

Table 1. AGR3 expression and pathological features of IDC patients

		AGR3 expre	ession, n (%)		
Pathological feature	No.	Low	High	Ľs	p-value
Age (yr)					
< 50	156	88 (56.4)	68 (43.6)	-0.099	0.069
≥ 50	180	119 (66.1)	61 (33.9)		
Histological grade ^{a)}					
Ι	14	7 (50.0)	7 (50.0)	-0.160	0.004**
II	252	147 (58.3)	105 (41.7)		
III	59	46 (78.0)	13 (22.0)		
Histological grade ^{a)}					
I-II	266	154 (57.9)	112 (42.1)	-0.159	0.004**
III	59	46 (78.0)	13 (22.0)		
Tumor size (cm) ^{a)}					
< 2	79	51 (64.6)	28 (35.4)	0.041	0.457
2-5	231	141 (61.0)	90 (39.0)		
> 5	23	13 (56.5)	10 (43.5)		
Lymph node metastases					
0	122	75 (61.5)	47 (38.5)	-0.007	0.901
1-3	100	62 (62.0)	38 (38.0)		
4-9	52	30 (57.7)	22 (42.3)		
>9	62	40 (64.5)	22 (35.5)		
Distant metastasis					
No	278	176 (63.3)	102 (36.7)	0.077	0.161
Yes	58	31 (53.4)	27 (46.6)		
ER status ^{a)}					
Negative	123	102 (82.9)	21 (17.1)	0.336	< 0.001
Positive	212	104 (49.1)	108 (50.9)		
PR status ^{a)}					
Negative	121	92 (76.0)	29 (24.0)	0.225	< 0.001
Positive	214	114 (53.3)	100 (46.7)		
HER2 status ^{a)}					
- to +	265	157 (59.2)	108 (40.8)	-0.098	0.074
++ to +++	69	49 (71.0)	20 (29.0)		
Ki-67 status ^{a)}					
Negative	96	59 (61.5)	37 (38.5)	-0.009	0.874
Positive	234	146 (62.4)	88 (37.6)		
Recurrence or distant metastasis ^{a)}					
No	261	173 (66.3)	88 (33.7)	0.179	0.001
Yes	69	31 (44.9)	38 (55.1)		

p-value was calculated by Spearman's rank correlation test. IDC, invasive ductal carcinoma; AGR3, anterior gradient 3; ER, estrogen receptor; PR, progesterone receptor; HER2, human epidermal growth factor receptor 2. **p < 0.01. ^aSome missing data.

ous (Table 1).

Fifty percent (7/14) of grade I, 41.7% (105/252) of grade II, and 22.0% (13/59) of grade III showed high expression of AGR3 (r_s =-0.160, p=0.004) (Table 1). Taking grade I and II as a group of low grade and grade III as a group of high grade, we found that 42.1% (112/266) of grade I-II showed high

expression of AGR3 and 22.0% (13/59) of grade III showed high expression of AGR3. The difference also had statistical meaning (r_s =-0.159, p=0.004) (Table 1). In addition, western blot analysis were performed in fresh tissues of IDC of different grades (6 cases of normal breast tissues, 6 cases of grade I, 6 cases of grade II, and 6 cases of grade III). Results

Mologular subtrac	No	AGR3 expres	sion, n (%)		n valuo
Molecular subtype	190.	Low	High	Is	p-value
Luminal A	68	37 (54.4)	31 (45.6)	-0.249	0.000***
Luminal B	172	90 (52.3)	82 (47.7)		
HER2-overexpressing	28	25 (89.3)	3 (10.7)		
TNBC	62	53 (85.5)	9 (14.5)		

Table 2. Relationship between AGR3 expression and molecular subtypes in IDC

p-value (luminal A vs. luminal B)=0.770, p-value (HER2-overexpressing vs. TNBC)=0.876, p-value (luminal A and luminal B vs. HER2-overexpressing and TNBC)=0.000; p-value was calculated by Spearman's rank correlation test. AGR3, anterior gradient 3; IDC, invasive ductal carcinoma; HER2, human epidermal growth factor receptor 2; TNBC, triple-negative breast cancer. ***p < 0.001.

also showed that AGR3 expression in grade I and II was significantly higher than grade III (Fig. 1C).

All results above indicated that AGR3 mainly expressed in IDC patients of histological grade I-II. In grade I-II, the relationship between AGR3 expression and clinicopathological features were also analyzed. High expression of AGR3 was also associated with high ER positivity (r_s =0.333, p < 0.001), high PR positivity (r_s =0.224, p < 0.001) and high rate of recurrence or metastasis (r_s =0.220, p < 0.001) (S2 Table). The correlations between AGR3 expression and patients' age (r_s = -0.099, p=0.105), tumor size (r_s =0.035, p=0.647), lymph node metastasis (r_s =-0.029, p=0.756), distant metastasis (r_s =0.089, p=0.149), HER2 status (r_s =0.113, p=0.066), and Ki-67 status (r_s =0.014, p=0.823) were also not obvious (S2 Table). But in grade III, all clinicopathological features had no correlation with AGR3 expression (S3 Table).

Then, we detected AGR3, ER, and PR expression in serial paraffin sections by IHC. The results also showed that AGR3 expression was positively correlated with ER and PR status. Representative images of staining were shown in Fig. 1D. Based on the important roles of ER and PR in molecular subtypes [2], further analysis was performed to confirm the relationship between AGR3 expression and molecular subtypes. We found that AGR3 expression in luminal subtype was much higher than non-luminal subtype in IDC. Fortyfive point six percent (31/68) of luminal A, 47.7% (82/172) of luminal B, 10.7% (3/28) of HER2-overexpressing, and 14.5% (9/62) of TNBC showed high expression of AGR3 (r_s = -0.249, p < 0.001) (Table 2).

Moreover, in IDC patients of grade I-II, AGR3 expression in luminal subtype was also much higher than non-luminal subtype. Forty-seven point four percent (27/57) of luminal A, 50.3% (72/143) of luminal B, 5.3% (1/19) of HER2-overexpressing, and 19.5% (8/41) of TNBC showed high expression of AGR3 (r_s =-0.218, p < 0.001) (S4 Table). But grade III had no such relationship (r_s =-0.241, p=0.136) (S5 Table). All of these results above indicated that AGR3 mainly expressed in luminal subtype of IDC patients of histological grade I-II.

3. High expression of AGR3 was a promoting factor for breast cancer patients to develop recurrence or distant metastasis

Table 1 showed that AGR3 expression was positively correlated with high risk of recurrence or metastasis. The representative images of IHC were shown in Fig. 1E. In order to get more accurate results, similar analysis was completed in different molecular subtypes. Additionally, we also analyzed the relationship between AGR3 expression and risk of recurrence or metastasis within 5 years.

In IDC, AGR3 expression of patients occurring recurrence or metastasis was much higher than disease-free patients (p=0.001); AGR3 expression of patients occurring recurrence or metastasis within 5 years was also higher than patients disease-free over 5 years (p=0.031) (Fig. 2A). Then, in luminal and luminal B subtype of IDC, AGR3 expression of patients occurring recurrence or distant metastasis was higher than disease-free patients (luminal, p < 0.001; luminal B, p=0.005); AGR3 expression of patients occurring recurrence or distant metastasis within 5 years was also higher than patients disease-free over 5 years (luminal, p=0.004; luminal B, p=0.023) (Fig. 2B and D). But in luminal A and non-luminal subtype including HER2-overexpressing and TNBC, AGR3 expression had no relationship with recurrence or metastasis (Fig. 2C and E-G).

In IDC of grade I-II, we got similar results. AGR3 expression of patients occurring recurrence or distant metastasis was higher than disease-free patients (p < 0.001); AGR3 expression of patients occurring recurrence or distant metastasis within 5 years was also higher than patients disease-free over 5 years (p=0.014) (Fig. 2H). Besides, the relationship between AGR3 expression and recurrence or metastasis still existed in luminal and luminal B subtype (Fig. 2I and K), but didn't exist in luminal A, non-luminal and TNBC (Fig. 2J, L,



Fig. 2. High expression of anterior gradient 3 (AGR3) was associated with high risk of recurrence or metastasis in luminal B subtype of invasive ductal carcinoma (IDC) patients of grade I-II. Compared the proportion of AGR3 highly expressed patients in the group developing metastasis or recurrence (MR group) and the group disease-free (DF group). Besides, compared the proportion of AGR3 highly expressed patients in the group developing metastasis or recurrence within 5 years (MR 5 years group) and the group disease-free over 5 years (DF 5 years group). The proportion of every group was as follows. (A) MR, 55.1%; DF, 33.7%; MR 5 years, 50.0%; DF 5 years, 33.9%. (B) Luminal of MR, 70.8%; luminal of DF, 40.9%; luminal of MR 5 years, 69.4%; luminal of DF 5 years, 42.2%. (C) Luminal A of MR, 56.9%; luminal A of DF, 37.7%; luminal A of MR 5 years, 72.7%; luminal A of DF 5 years, 40.0%. (D) Luminal B of MR, 68.6%; luminal B of DF, 42.2%; luminal B of MR 5 years, 15.0%; non-luminal of MR, 19.0%; non-luminal of DF, 10.4%; non-luminal of MR, 13.0%; HER2-overexpressing of DF, 0%; HER2-overexpressing of MR, 13.0%; HER2-overexpressing of DF, 0%; HER2-overexpressing of DF 5 years, 7.7%. (*Continued to the next page*)



Fig. 2. (*Continued from the previous page*) (G) Triple-negative breast cancer (TNBC) of MR, 23.5%; TNBC of DF, 9.1%; TNBC of MR 5 years, 18.7%; TNBC of DF 5 years, 0%. (H) Grade I-II of MR, 64.0%; grade I-II of DF, 36.5%; grade I-II of MR 5 years, 58.5%; grade I-II of DF 5 years, 37.0%. (I) Luminal of grade I-II in MR, 75.68%; luminal of grade I-II in DF, 43.13%; luminal of grade I-II in MR 5 years, 72.41%; luminal of grade I-II in DF 5 years, 70.0%; luminal A of grade I-II in MR, 72.72%; luminal A of grade I-II in DF, 40.91%; luminal A of grade I-II in MR 5 years, 70.0%; luminal A of grade I-II in MR, 76.9%; luminal B of grade I-II in MR, 30.8%; non-luminal of grade I-II in DF, 8.9%; non-luminal of grade I-II in MR 5 years, 17.0%; non-luminal of grade I-II in DF 5 years, 0%. (*Continued to the next page*)



Fig. 2. (*Continued from the previous page*) (M) TNBC of grade I-II in MR, 36.4%; TNBC of grade I-II in DF, 10.3%; TNBC of grade I-II in MR 5 years, 30.0%; TNBC of grade I-II in DF 5 years, 0%. (N) Grade III of MR, 30.8%; grade III of DF, 20.0%; grade III of MR 5 years, 20.7%; grade III of DF 5 years, 22.2%.

and M). The number of HER2-overexpressing specimens was too few, so the investigation could not be performed. And in IDC of grade III, AGR3 expression also had no relationship with recurrence or metastasis (Fig. 2N). All results above reminded that high expression of AGR3 indicated high risk of recurrence or metastasis in luminal B subtype of IDC patients of grade I-II.

In addition, we investigated the relationship between AGR3 expression and metastatic organs such as bone, lung, liver, and brain which were common metastatic organs of breast cancer patients. We found that AGR3 expression was correlated with bone metastasis (total IDC patients: $r_s=0.119$, p=0.029; grade I-II patients: r_s =0.216, p=0.031) and liver metastasis (total IDC patients: r_s=0.127, p=0.020; grade I-II patients: r_s=0.135, p=0.027) in IDC of grade I and II (Table 3). But there was no correlation between AGR3 expression and lung metastasis (total IDC patients: r_s=-0.015, p=0.788; grade I-II patients: r_s=0.009, p=1.0) or brain metastasis (total IDC patients: r_s =-0.003, p=0.958; grade I-II patients: r_s =0.020, p=1.0) (Table 3). In IDC of grade III, relationships between AGR3 expression and these four metastatic organs were not obvious (S6 Table). These results indicated that high expression of AGR3 could lead to high risk of bone and liver metastasis.

4. AGR3 was an independent prognostic factor in luminal B patients of grade I-II

Considering relationship between AGR3 expression and high risk of recurrence or metastasis. Kaplan-Meier analysis was performed in 330 cases of IDC with complete follow-up data to explore the value of AGR3 in prognosis.

In IDC, both OS and PFS of AGR3 high expression group

were significantly shorter than AGR3 low expression group (OS, p=0.035; PFS, p=0.004) (Fig. 3A). With further analysis in different molecular subtypes, we found that OS and PFS of AGR3 high expression group shortened in luminal subtype (OS, p=0.004; PFS, p=0.001) (Fig. 3B). Separately, in luminal A, AGR3 high expression could lead to shorter PFS (p=0.014) but had no influence on OS (Fig. 3C); in luminal B, both OS and PFS significantly shorten if AGR3 highly expressed (OS, p=0.010; PFS, p=0.014) (Fig. 3D). Whereas, in nonluminal and TNBC subtype, AGR3 expression had no influence on either OS or PFS (Fig. 3E and F). Number of HER2-overexpressing patients was too few, so Kaplan-Meier analysis could not be performed.

In IDC of grade I-II, we got similar results. OS and PFS also significantly shortened when AGR3 highly expressed (OS, p=0.027; PFS, p=0.001) (Fig. 3G). And AGR3's impacts on OS and PFS still existed in luminal and luminal B subtype (Fig. 3H and J), but didn't exist in luminal A, non-luminal or TNBC (Fig. 3I, K and L). Kaplan-Meier analysis still could not be performed in HER2-overexpressing subtype of grade I-II due to the small number of specimens.

Results above reminded that high expression of AGR3 indicated poor prognosis in luminal B subtype of IDC patients of grade I-II. In addition, in multivariate Cox regression analysis, we found that AGR3 expression was an independent predictor for both OS and PFS (Table 4).

5. AGR3 affected proliferation and invasion ability of breast cancer cells

After confirming the promoting role of AGR3 in clinical specimens, we detected AGR3 expression in breast cancer cell lines by western blot. In western blot analysis, we took

Measatu organs (ID) Low High to High to High to High to Polute Bone metastasis 293 187 (63.8) 106 (36.2) 0.119 0.029* 234 154 (65.8) 80 (34.2) 0.216 0.031* No 293 187 (63.8) 106 (36.2) 0.119 0.029* 234 154 (65.8) 80 (34.2) 0.216 0.031* No 319 196 (61.4) 123 (38.6) -0.015 0.788 257 149 (58.0) 108 (42.0) 0.009 1.000 Yes 17 11 (64.7) 6 (35.3) -0.015 0.788 257 149 (58.0) 108 (42.0) 0.009 1.000 Yes 17 11 (64.7) 6 (35.3) -0.012 0.788 255.6 $4 (44.4)$ -0.002 $7 (77.8)$ 1000 1000 $7 (77.8)$ 1000 $7 (77.8)$ 1000 1000 1000 1000 1000 1000 10002 252.2 100		No.	AGR3 expr	ession of IDC		-	No.	AGR3 expressio	n of grade I-II		-
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	Yes	8	5 (62.5)	3 (37.5)			4	2 (50.0)	2 (50.0)		

fresh tissues with AGR3 high expression as positive control. We found that AGR3 highly expressed in luminal cell line T47D, but low expressed in TNBC cell line MDA-MB-231 (Fig. 4A). We overexpressed full length of AGR3 in MDA-MB-231 and knocked down AGR3 in T47D (named as 3× Flag-AGR3-HA/MDA-MB-231 and siAGR3/T47D, respectively) to make sure the role of AGR3 *in vitro*. The overexpression and interference effects were detected by western blot (Fig. 4A). The interference sequence of siAGR3#1/T47D was used *in vitro* studies.

In ATP and SRB assays, proliferation ability of 3×Flag-AGR3-HA/MDA-MB-231 significantly increased comparing to 3×Flag-vector/MDA-MB-231; proliferation ability of siAGR3/T47D significantly reduced comparing to scr/T47D (Fig. 4B). Besides, we performed proliferation assays by using Nikon ECLIPSE Ti microscope to count cell numbers. The results were the same as ATP and SRB assays (Fig. 4B).

In Matrigel Boyden chamber assays and wound healing assays, migration and invasion ability of 3×Flag-AGR3-HA/ MDA-MB-231 significantly increased comparing to 3×Flagvector/MDA-MB-231; migration and invasion ability of siAGR3/T47D significantly reduced comparing to scr/T47D (Fig. 4C and D). All results of cellular functional experiments were consistent with results we got from clinical samples and indicated that AGR3 might improve the development of breast cancer by promoting tumor cells' proliferation and invasion ability.

6. AGR3 high expression indicated a poor prognosis of breast cancer patients treated with taxane but a favorable prognosis treated with 5-fluoropyrimidines

From AGR3 highly expressed IDC specimens, we choose 19 cases only receiving TE/TA chemotherapy regimen, 29 cases only receiving CEF/CAF chemotherapy regimen and 20 cases only receiving CMF chemotherapy regimen. These regimens were commonly used in the treatment of breast cancer patients. By performing Kaplan-Meier analysis, we found that PFS of TE/TA group were significantly shorter than CEF/CAF and CMF group (TE/TA vs. CEF/CAF, p=0.009; TE/TA vs. CMF, p=0.035). But PFS had no significant difference between CEF/CAF and CMF group (Fig. 5A). In addition, from AGR3 low expressed IDC specimens, we choose 57 cases only receiving TE/TA chemotherapy regimen, 48 cases only receiving CEF/CAF chemotherapy regimen and 24 cases only receiving CMF chemotherapy regimen. Kaplan-Meier analysis showed that both OS and PFS had no correlation with chemotherapy regimen in AGR3 low expressed IDC (Fig. 5B).

In AGR3 highly expressed IDC patients, CEF/CAF and CMF chemotherapy regimens could lead better prognosis than TE/TA chemotherapy regimen. We speculated that











grade I-II were shown, respectively. (K) OS and PFS of 58 non-luminal patients of grade I-II were shown, respectively. (L) OS and PFS of 40 TNBC patients of grade I-II were shown, respectively.

	OS (univari	ate)	OS (multivari	ate)	PFS (univaria	(te)	PFS (multivar	iate)
Variable	HR (95% CI)	p-value	HR (95% CI)	p-value	HR (95% CI)	p-value	HR (95% CI)	p-value
Age	0.911 (0.385-2.156)	0.832	1.071 (0.430-2.664)		0.676 (0.386-1.187)	0.173	0.759 (0.422-1.367)	
Tumor size	1.624(0.740-3.565)	0.227	1.017 (0.456-2.270)	ı	1.852 (1.096-3.130)	0.021	1.139(0.656-1.976)	
Lymph node metastasis	1.680(1.149-2.455)	0.007	1.757 (1.173-2.630)	0.006	1.651 (1.294-2.106)	0.000	1.679(1.303-2.164)	< 0.001
ER status	0.766 (0.316-1.857)	0.556	0.436 (0.147-1.295)	ı	0.739 (0.417-1.309)	0.300	0.559(0.282-1.109)	ı
PR status	0.907 (0.366-2.251)	0.834	0.780 (0.277-2.197)	ı	0.645 (0.366-1.137)	0.129	0.498 (0.261-0.950)	ı
AGR3 expression	2.695 (1.082-6.713)	0.033	4.161 (1.406-12.312)	0.010	2.531 (1.421-4.511)	0.002	3.856 (1.953-7.613)	< 0.001

 Table 4.
 Univariate and multivariate analysis for OS and PFS in IDC patients of grade I-II

terone receptor; AGR3, anterior gradient 3.

Qiao Xu, AGR3 Promotes Breast Cancer Development

AGR3 might change the sensitivity of one drug in these regimens. We divided all specimens into ten groups based on whether chemotherapy regimen consisted of taxane, epirubicin/doxorubicin, cyclophosphamide, methotrexate or 5fluoropyrimidines. These groups were respectively named as T/non-T group, E/non-E group, C/non-C group, M/non-M group and F/non-F group. We wanted to know whether presence or absence of one particular drug would change the prognosis of AGR3 highly expressed IDC patients by comparing "X" group to "non-X" group.

We found that, in AGR3 highly expressed IDC patients, PFS of T group was shorter than non-T group (p=0.009) (Fig. 5C) and PFS of F group was longer than non-F group (p=0.007) (Fig. 5D). But when comparing other groups, epirubicin/ doxorubicin, cyclophosphamide, and methotrexate had no impact on both OS and PFS (S7A, S7B and S7C Fig.). In AGR3 low expressed IDC patients, OS and PFS also had no difference during any comparison.

Then, more analysis was performed in different molecular subtypes. In luminal subtype of AGR3 highly expressed IDC patients, PFS of T group was also shorter than non-T group (p=0.006) (Fig. 5E) and PFS of F group was also longer than non-F group (p=0.014) (Fig. 5F). But in non-luminal subtype, both OS and PFS had no correlation with taxane and 5-fluoropyrimidine (S8A and S8B Fig.). And in AGR3 low expressed IDC patients, no matter luminal subtype or non-luminal subtype had no such relationship (Fig. 5E, 5F, S8A and S8B Fig.).

Besides, in IDC patients of grade I-II, PFS of T group was also shorter than non-T group (p=0.035) (Fig. 5G) and PFS of F group was also longer than non-F group (p=0.007) (Fig. 5H) when AGR3 highly expressed. And the impact of taxane and 5-fluoropyrimidine on PFS also existed in luminal subtype (Fig. 5I and J) but not in non-luminal subtype (S8C and S8D Fig.). In addition, grade III patients had no such relationship (S8E and S8F Fig.). All of these results above indicated that AGR3 high expression indicated a poor prognosis of breast cancer patients treated with taxane but a favorable prognosis treated with 5-fluoropyrimidines.

7. Breast cancer cells with AGR3 high expression were resistant to taxane but sensitive to 5-fluoropyrimidines

CD-DST assay had been widely applied in clinical work to test sensitivities of chemotherapy drugs. Based on the results of CD-DST, we chose other 161 cases of IDC specimens consisting of 44 cases sensitive to taxane, 39 cases resistant to taxane, 42 cases sensitive to 5-fluoropyrimidines and 36 cases resistant to 5-fluoropyrimidines. Then, we performed IHC to detect AGR3 expression in these specimens. We found that AGR3 expression of patients sensitive to taxane was much lower than patients resistant to taxane (p < 0.001) (Fig. 6A)



Fig. 4. Anterior gradient 3 (AGR3) promoted proliferation and invasion ability of breast cancer cells. (A) Western blot analysis of AGR3 expression in MDA-MB-231 and T47D cells (patient tissues with AGR3 high expression as positive control) (left panel). AGR3 expression was detected by primary flag antibodies in $3\times$ Flag-vector/MDA-MB-231 and $3\times$ Flag-AGR3-HA/MDA-MB-231 cells (middle panel). AGR3 expression was detected by primary AGR3 antibodies in T47D, scr/T47D and siAGR3/T47D cells (right panel). (B) Proliferation ability was detected by ATP and sulforhodamine B (SRB) assays in $3\times$ Flag-vector/MDA-MB-231 and $3\times$ Flag-AGR3-HA/MDA-MB-231 cells, respectively (upper panel in left and middle). Proliferation assay was repeated by using Nikon ECLIPSE Ti microscope to count cell number in $3\times$ Flag-vector/MDA-MB-231 cells (upper panel in right). Proliferation ability was also detected by ATP and SRB assays in scr/T47D and siAGR3/T47D cells, respectively (lower panel in left and middle). Proliferation ability was also repeated by using Nikon ECLIPSE Ti microscope to count cell number in $3\times$ Flag-vector/MDA-MB-231 cells, respectively (lower panel in right). Proliferation ability was also detected by ATP and SRB assays in scr/T47D and siAGR3/T47D cells, respectively (lower panel in left and middle). Proliferation assay was also repeated by using Nikon ECLIPSE Ti microscope to count cell numbers in scr/T47D and siAGR3/T47D cells (lower panel in right). Bars are mean±SD. All experiments were performed 3 times independently (*p < 0.05, **p < 0.01). (*Continued to the next page*)

but AGR3 expression of patients sensitive to 5-fluoropyrimidines was much higher than patients resistant to 5-fluoropyrimidines (p=0.004) (Fig. 6A). In breast cancer cell lines, 3×Flag-AGR3-HA/MDA-MB-231 and siAGR3/T47D were performed MTT and ATP assay to measure cell viability with various concentrations of tax-

















page)

MB-231 cells in MTT and ATP assay were much lower than 3×Flag-vector/MDA-MB-231 cells (MTT, p=0.025; ATP, p=0.009, lower panel). (Continued to the next



Fig. 6. (*Continued from the previous page*) (C) With the treatment of taxane, cell viability of siAGR3/T47D cells was much lower than scr/T47D cells (*p < 0.05, ** p < 0.01, upper panel). Taxane IC₅₀ values of siAGR3/T47D cells in MTT and ATP assay were much lower than scr/T47D cells (MTT, p=0.033; ATP, p=0.006, upper panel). With the treatment of 5-FU, cell viability of siAGR3/T47D cells was much higher than scr/T47D cells (*p < 0.01, ***p < 0.001, lower panel). 5-FU IC₅₀ value of siAGR3/T47D cells in MTT and ATP assay were much higher than scr/T47D cells (MTT, p=0.017, lower panel).

ane and 5-fluoropyrimidines. Moreover, the IC₅₀ values of taxane and 5-fluoropyrimidines in different cells were also detected. $3 \times Flag$ -vector/MDA-MB-231 and $3 \times Flag$ -AGR3-HA/MDA-MB-231 cells were treated with 15 to 5×10^4 ng/mL of taxane and $1 \cdot 1.5 \times 10^4$ µg/mL of 5-fluoropyrimidines for 48 hours. Both MTT and ATP assay showed that $3 \times Flag$ -AGR3-HA/MDA-MB-231 cells exhibited more resistant to taxane but more sensitive to 5-fluoropyrimidines than $3 \times Flag$ -vector/MDA-MB-231 cells (Fig. 6B). Scr/T47D and siAGR3/T47D cells were treated with 2.5 to 2.5×10^4 ng/mL of taxane and $0.1 \cdot 5 \times 10^3$ µg/mL of 5-fluoropyrimidines for 48 hours. Both MTT and ATP assay showed that siAGR3/T47D cells exhibited more sensitive to taxane but more resistant to 5-fluoropyrimidines for 48 hours.

All of these results above indicated that breast cancer cells

with AGR3 high expression were resistant to taxane but sensitive to 5-fluoropyrimidines. These results explained why AGR3 high expression could indicate a poor prognosis of breast cancer patients treated with taxane but a favorable prognosis treated with 5-fluoropyrimidines.

Discussion

AGR3's impact on cancers is still unclear. Reports about AGR3 in cancers are few and contradictory. In tumors, AGR3 protein was firstly identified in the membrane of breast cancer cell lines by proteomic screening [16]. But later, Loader's group confirmed that AGR3 overexpressed in the cytoplasm of breast cancer cells [8]. And Garczyk et al. [17] found that AGR3's mRNA significantly increased in breast cancer tissues of histological grade I-II, but not in grade III and AGR3 played a role of indicating poor prognosis in grade I-II which were consistent with our results. But they didn't perform more study about the correlation between AGR3 and molecular subtypes. They also found that combining expression of AGR3 and AGR2 in serum sample of breast cancer patients could diagnosed early breast cancer [17]. Besides, it had been demonstrated that AGR3 also overexpressed in ovarian [18] and prostate [19] cancer. In ovarian cancer, Gray et al. [20] found that AGR3 was up-regulated by a hormone (ER)independent mechanism and could mediate resistance to cisplatin. In prostate cancer, Vaarala et al. [21] found that AGR3 highly elevated in prostate cancer tissues and Bu et al. [22] found that both AGR2 and AGR3 genes were regulated by androgen and estrogen. Moreover, androgen receptor-binding sites in the promoter region of AGR2 and AGR3 were identified in the report of Bu et al. [22]. In these studies, researchers considered that AGR3 could promote cancer development and regarded AGR3 as an indicator of poor prognosis of cancer patients. However, some studies took opposite opinions. In breast cancer, Obacz et al. [23] found that AGR3 was associated with differentiation level, weak proliferation and favorable prognosis of patients. In ovarian cancer, AGR3 expression in serous borderline ovarian tumors and low-grade serous ovarian carcinomas was much higher than high-grade serous ovarian carcinomas and AGR3 was associated with longer patient survival [24].

The role of AGR3 remained uncertain and researchers even got opposite results in the same kind of tumor tissues. We thought it was due to the limit of sample size. With larger scale of samples, our study confirmed that AGR3 expression significantly increased with the development of breast tumor malignancy and had close relationships with molecular subtypes and histological grades. AGR3 mainly expressed in luminal subtype of IDC patients of histological grade I-II. Most importantly, in luminal B subtype, high expression of AGR3 was correlated with high risk of recurrence and metastasis and could predict poor prognosis of patients. Therefore, AGR3 might be a potential prognostic indicator of luminal B subtype in which patients were more likely to occur therapeutic resistance and poor outcome than luminal A [5]. As for why AGR3 was also overexpressed in luminal A but couldn't indicate prognosis of this subtype, we considered that it was due to the excellent sensitivity to endocrine therapy in luminal A. And we considered that AGR3's overexpression in luminal subtype of IDC was due to the transcription activating function of $ER\alpha$. Welboren et al. [25] had verified several ER α -binding sites existing on the promoter regions of AGR2 and AGR3. But the accurate mechanism

needed more proofs. Besides, through further analysis, we found that the majority of luminal patients were grade I-II (83.33%) but only 16.67% of luminal patients were grade III. This result probably explained why AGR3 mainly expressed in grade I-II rather than grade III.

Besides, we firstly found that taxane lead to worse outcome but 5-fluoropyrimidine lead to favorable outcome in AGR3 highly expressed IDC. Then, by performing CD-DST and cytotoxic analysis, we demonstrated that breast cancer cells with AGR3 high expression were resistant to taxane but sensitive to 5-fluoropyrimidines. As we known, in clinic therapy, luminal B patients were routinely treated with additional chemotherapy on the basis of endocrine therapy [5]. These indicated that luminal B patients with AGR3 high expression should be treated with chemotherapy regimens consisting of 5-fluoropyrimidines but not taxane.

As we known, taxane could cure cancer patients depending on its antimitotic characteristic. But ER stress had been identified as another mechanism of taxane cytotoxic effects. The ER stress mediated by taxane could induce apoptosis of cancer cells [26]. In this process, unfolded protein response (UPR) signaling pathway could weaken ER stress to maintain ER homeostasis and help cells survive [27]. Researchers had demonstrated that AGR2 could bind GRP78 which was a central trigger of UPR. The interaction of AGR2 and GRP78 would activate UPR signaling pathway to attenuate cell death induced by ER stress [28]. As homologous protein, AGR2 and AGR3 gene shared 71% sequence identity and lay adjacent to one another at chromosomal position 7p21 [22-24]. Therefore, in tumor cells, we speculated that AGR3 might also interact with GRP78 to activate UPR signaling pathway. Then, the UPR would attenuates ER stress mediated by taxane so that tumor cells could survive from the taxane treatment. Besides, Kim et al. [29] found that ER stress induced 5-fluoropyrimidines resistance in human colon cancer cells and Yun et al. [30] found that down-regulation of GRP78 could lead to enhanced sensitivity of cancer cells to cytotoxic effect of 5-fluoropyrimidines [30]. Therefore, we considered that AGR3 could promote sensitivity of breast cancer cells to 5-fluoropyrimidines also due to the interaction with GRP78. All of these speculations need a large number of experiments to prove.

Migration and invasion ability of tumor cells played important roles in the development of cancer. We reported for the first time that AGR3 could promote the proliferation and invasion ability of tumor cells which might be the biological basis of AGR3's prompting role in cancer.

In conclusion, our research confirmed that, in luminal B subtype of IDC patients of histological grade I-II, AGR3 high expression indicated poor prognosis and patients should be treated with chemotherapy regimens consisting of 5-fluoropyrimidines but not taxane. AGR3 could promote tumor

progression by improving tumor cells' proliferation ability, invasion ability, and chemotherapeutic drug resistance. AGR3 might be a potential prognostic indicator for classifying prognosis and therapy regimens in this population. But all of these conclusions need more experiments and analysis to confirm.

Electronic Supplementary Material

Supplementary materials are available at Cancer Research and Treatment website (https://www.e-crt.org).

Conflicts of Interest

Conflict of interest relevant to this article was not reported.

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