B-cell lymphoma 2 is associated with advanced tumor grade and clinical stage, and reduced overall survival in young Chinese patients with colorectal carcinoma

JIASHENG WANG¹, GAN HE¹, QIANG YANG¹, LIAN BAI¹, BIN JIAN¹, QUGANG LI¹ and ZHONGFU LI²

¹Department of Gastrointestinal Surgery, Yongchuan Hospital, Chongqing Medical University, Chongqing 402160; ²Department of Gastrointestinal Surgery, Chongqing Emergency Medical Center, Center Hospital of Chongqing University, Chongqing 400014, P.R. China

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Abstract. The development of biomarkers that accurately and reliably detect colorectal cancer is a promising approach for colorectal cancer screening. Therefore, the objective of the present study was to evaluate the protein expression of α-methylacyl-CoA racemase (P504S/AMACR), tumor protein p53 (p53), B-cell lymphoma 2 (Bcl-2) and Ki-67/mindbomb E3 ubiquitin protein ligase 1 (MIB-1) in a population of Chinese patients with colorectal carcinoma. Colorectal tumors with matched normal tissue margins were collected from 148 surgical patients, and the demographic and clinical characteristics were collected. Immunohistochemical staining and western blot analysis of P504S/AMACR, p53, Bcl-2 and Ki-67/MIB-1 were conducted. Statistical analyses were used to compare protein expression in the colorectal tumors and matched normal tissue margins and to identify any associations between them and various clinicopathological parameters. Survival analyses were performed using the Kaplan-Meier method. In the present study, immunohistochemistry and western blot analysis revealed significantly higher expression of all four proteins in colorectal tumors compared with matched normal tissue margins (P<0.001). Spearman's rank correlation analysis revealed that Bcl-2 expression was negatively correlated with pathological grade and Tumor-Node-Metastasis (TNM) stage (-0.827 and -0.388, respectively; P<0.05). Bcl-2 expression was revealed to be a significant prognostic indicator of colorectal carcinoma [relative risk (95% CI), 0.703 (0.552-0.895); P<0.05]. The log-rank test revealed a significant association between low

Correspondence to: Dr Zhongfu Li, Department of Gastrointestinal Surgery, Chongqing Emergency Medical Center, Center Hospital of Chongqing University, 1 Jiankang Road, Yuzhong, Chongqing 400014, P.R. China E-mail: li_zhongfu@sina.com

Bcl-2 expression and reduced overall survival (P=0.039), as well as a significant association between older age (>55 years) and reduced overall survival (P<0.001) in Chinese patients with colorectal carcinoma. In conclusion, low expression of Bcl-2 is significantly correlated with advanced pathological grade and TNM stage and is a prognostic indicator of reduced overall survival in young Chinese patients with colorectal carcinoma.

Introduction

Colorectal cancer ranks as the third most common malignancy worldwide, with a global 5-year prevalence of ~3.2 million cases (1). Due to changes in diet and lifestyle, several increasingly affluent Asian countries (e.g., China, Japan, South Korea and Singapore) have displayed a 2- to 4-fold increase in the incidence of colorectal carcinoma over the past few decades (2). However, non-polypoidal lesions and *de novo* colorectal neoplasms (i.e., without preceding adenoma) are more common in Asian patient populations. The non-polypoidal lesions are less visible, which poses difficulties in disease screening by conventional imaging and endoscopy (2). Therefore, the development of alternative screening methods for colorectal carcinoma is particularly important.

Early detection of colorectal cancer in its localized or pre-invasive form is a more realistic approach to screening (3). Current screening strategies, including fecal occult blood testing and colonoscopy, have poor levels of patient acceptability (4). The development of peripheral blood protein biomarkers that are able to accurately and reliably detect colorectal cancer at its earliest stages appears to be promising alternative approach for colorectal cancer screening (5).

 α -methylacyl-CoA racemase (P504S/AMACR) is overexpressed in colorectal adenomas and cancer, and has been demonstrated to be associated with advanced distal colorectal adenoma (6) and tumor differentiation (7). Furthermore, an inverse correlation was demonstrated between p53 and B-cell lymphoma 2 (Bcl-2) expression in colorectal tumorigenesis, and abnormal activation of Bcl-2 inhibited apoptosis *in vivo* (8). The Ki-67 labeling index is positively correlated with poor survival in colorectal cancer patients (9). Therefore, the objective of the present study was to evaluate the protein

Key words: α-methylacyl-CoA racemase, tumor protein p53, B-cell lymphoma 2, Ki-67, mindbomb E3 ubiquitin protein ligase 1, colorectal carcinoma, biomarker

expression of four key genes, P504S/AMACR, p53, Bcl-2 and Ki-67/mindbomb E3 ubiquitin protein ligase 1 (MIB-1), in a population of Chinese patients with colorectal carcinoma. The present study aimed to shed light on the potential applicability of these four proteins as diagnostic biomarkers for colorectal carcinoma in Chinese patient populations.

Materials and methods

Patients and tumor samples. Frozen tumor tissues with matched normal tissue margins were collected from 148 Han Chinese patients (68 males and 80 females; with a mean age of 50 years old, and the ranging from 25-75 years old) that had undergone surgery at Yongchuan Hospital (Chongqing Medical University, Chongqing, China) from January 2015 to January 2016. For each patient, the following relevant demographic and clinical data were collected: i) Hospital/surgery location (city, country); ii) patient age (years); iii) patient sex; iv) risk factors [e.g., body mass index (BMI), smoking status and excessive alcohol consumption (i.e., a male consuming >4 drinks/day or >14 drinks/week; or a female consuming >3 drinks/day or >7 drinks/week)] (10); v) tumor characteristics, including pathological grade of tumor cells (G1-G4) (11), Clinical stage (T1-T4), samples were staged according to the American Joint Committee on Cancer TNM staging system (12); and vi) survival time post-surgery (months).

Immunohistochemical staining. Immunohistochemical staining was performed using an Ultrasensitive[™] SP kit (cat. no. KIT-0105M, Maixin Biotech Co., Ltd., Fuzhou, China) and diaminobenzidine (DAB; Fuzhou Maixin Biotech Co., Ltd., Fuzhou, China), according to the manufacturer's protocols. Briefly, 4-µm thick paraffin-embedded sections, which were fixed with 4% formaldehyde at 20°C for 24 h, were xylene-deparaffinized at 60°C for 1 hour, rehydrated in a descending alcohol series and rinsed in phosphate-buffered saline (PBS). Antigen retrieval was performed by placing the slides in boiling citric acid buffer (pH 6.0) for 5 min. The sections were subsequently incubated in 3% hydrogen peroxide in methanol at room temperature for 20 min to quench endogenous peroxidase activity. Non-immune serum albumin (5%; Beyotime Institute of Biotechnology, Haimen, China) was used to block non-specific binding at 20°C for 5 min. The tissue sections were then incubated at 37°C for 2 h with the following monoclonal primary antibodies: sheep anti-P504S/AMACR (cat. no. AF7508-SP; dilution, 1:500), rabbit anti-p53 (cat. no. AF 1355; dilution, 1:2,000), mouse anti-Bcl-2 (cat no. AF 810; dilution, 1:500) and sheep anti-Ki-67/MIB-1 (cat no. AF 7617; dilution, 1:800) (all from R&D Systems Inc., Minneapolis, MN, USA). Next was an incubation with biotinylated goat anti-rat IgG secondary antibodies (cat. no.; NL013, dilution, 1:1,000; R&D Systems) at 37°C for 30 min followed by a streptavidin horseradish peroxidase complex, according to the instructions of UltrasensitiveTM SP kit, with intermittent PBS rinses. Immunoreactivity was visualized with diluted DAB. Finally, sections were rinsed with distilled water, and counterstained at 37°C for 5 min with Mayer's hematoxylin and histomounted. As a negative control, the primary antibody was replaced with an equal amount of normal human/rabbit/rat/mouse IgG. The positive controls were cancer cell lines with known positive expression of P504S/AMACR, p53, Bcl-2 or Ki-67/MIB-1.

Evaluation of immunohistochemistry. Images of stained sections were captured in a series of 10 randomly selected high-power fields with a laser scanning confocal microscope (Olympus Corp., Tokyo, Japan) at magnification, x400. The sections were then scored by the proportion and staining intensity of positively stained tumor cells. The 'proportion score' of positively stained tumor cells was determined as follows: 0, no positive tumor cells; 1, <10% positive tumor cells; 2, 10-50% positive tumor cells; and 3, >50% positive tumor cells. The 'staining intensity' score was determined as follows: 0, no staining; 1, weak staining/light yellow; 2, moderate staining/yellow-brown; and 3, strong staining/brown. Immunohistochemical staining was scored independently by two investigators blinded to the clinicopathological findings. Cases with score discrepancies were re-reviewed simultaneously by the original two investigators and a senior pathologist until a consensus was reached.

The 'staining index' was then calculated as the staining intensity score multiplied by the proportion score. Using this method, a staining index was obtained for P504S/AMACR, p53, Bcl-2 and Ki-67/MIB-1 in colorectal specimens, with scores of 0, 1, 2, 3, 4, 6 or 9 for each protein biomarker. Subsequently, cut-off values (Table I) for immunoreactivity for each protein biomarker were based on measuring heterogeneity by the sequential association analysis for SPA progression. This cut-off point was used to distinguish between 'low' and 'high' expression of each protein biomarker.

Western blot analysis. Total protein was extracted from the colorectal cancer and matched normal tissue margin specimens with Cell Extraction Buffer (cat. no. FNN0011; Thermo Fisher Scientific, Inc., Waltham, MA, USA). Protein concentrations were determined using a bicinchoninic acid protein assay kit, and 30 μ g protein per sample was separated by 8% SDS-PAGE and transferred onto a methanol-activated nitrocellulose filter membrane (Bio-Rad Laboratories, Inc., Hercules, CA, USA). Prior to immunoblotting, membranes were blocked within 5% skimmed dry milk at 37°C for 2 h. Expression levels were normalized to β -actin (dilution, 1:1,000; cat. no. MAB8929; R&D Systems). The following primary monoclonal antibodies were diluted in buffer and incubated at 4°C overnight: Sheep anti-P504S/AMACR (cat. no. AF7508-SP; R&D Systems Inc.; dilution, 1:1,000), rabbit anti-p53 (cat. no. AF1355; R&D Systems Inc.; dilution, 1:2,000), mouse anti-Bcl-2 (cat no. AF810; R&D Systems Inc.; dilution, 1:1,000) and mouse anti-Ki-67/MIB-1 (cat no. AF7617; R&D Systems Inc.; dilution, 1:1,200). Following washing in TBST, membranes were incubated with a horseradish peroxidase-conjugated goat anti-rat secondary antibody (cat. no. ab7097; dilution, 1:1,000; Abcam, Cambridge, UK) for 1 h at room temperature. This experiment was repeated three times. Bands were detected using a SuperEnhanced chemiluminescence kit (cat. no. P1010; Applygen Technologies, Inc., Beijing, China).

Statistical analysis. All data were analyzed using SPSS 13.0 for Windows (SPSS, Inc., Chicago, IL, USA). χ^2 and Fisher's exact tests were used to compare P504S/AMACR,

Protein	Proportion of IHC staining	Area under ROC curve	Cut-off value, %	Sensitivity, %	Specificity, %
P504S/AMACR	6.53±4.51	0.62	8	60.2	84.2
p53	10.39±5.28	0.65	11	55.1	78.3
Bcl-2	18.30±13.20	0.72	21	66.7	80.3
Ki-67/MIB-1	10.20 ± 7.28	0.70	13	61.4	71.6

Table I. Results of ROC curve analysis of protein biomarkers and determination of the cut-off values.

ROC, receiver operating characteristic; IHC, immunohistochemistry; AMACR, α-methylacyl-CoA racemase; p53, tumor protein p53; Bcl-2, B-cell lymphoma 2.

Table II. Immunohistochemical analysis of four proteins in colorectal carcinoma tumors and matched normal tissue margins (control).

	Tissue	type, n		
Variable	Tumor	Control	χ^2 value	P-value
P504S/AMACR expression			88.80	<0.001ª
High	76	4		
Low	72	144		
Bcl-2 expression			97.93	<0.001 ^a
High	104	20		
Low	44	128		
p53 expression			76.85	<0.001ª
High	88	16		
Low	60	132		
Ki-67/MIB-1 expression			50.71	<0.001ª
High	72	16		
Low	76	132		

^aStatistically significant (P<0.05). AMACR, α-methylacyl-CoA racemase; Bcl-2, B-cell lymphoma 2; p53, tumor protein p53.

p53, Bcl-2 and Ki-67/MIB-1 protein expression (from the immunohistochemical 'staining index') in the colorectal tumors and matched normal tissue margins and various clinicopathological parameters (e.g., age, sex, BMI and smoking status). The Student's t-test was used to analyze the western blotting results. In order to confirm the correlations between P504S/AMACR, p53, Bcl-2 and Ki-67/MIB-1 protein expression and clinicopathological parameters, Spearman's rank correlation analysis was applied. To determine whether P504S/AMACR, p53, Bcl-2 and Ki-67/MIB-1 expression levels were independent prognostic factors in patients with colorectal carcinoma, prognostic relevance was evaluated using multivariate Cox regression analysis. P<0.05 was considered to indicate a statistically significant difference. Survival analysis was performed using the Kaplan-Meier method, followed by the log-rank test (13).

Results

P504S/AMACR, p53, Bcl-2 and Ki-67/MIB-1 protein levels are increased in colorectal tumor tissues compared with normal

tissue margins. Immunohistochemical analysis revealed that a significantly greater proportion of colorectal tumors exhibited a high expression of all four proteins (P504S/AMACR, p53, Bcl-2 and Ki-67/MIB-1), compared with matching normal tissue margins (Table II; Fig. 1). Additionally, western blot analysis revealed significantly higher expression of the same four proteins in colorectal tumors compared with matching normal tissue margins (all P<0.001; Fig. 2).

Higher expression of Bcl-2 was observed in patients >55 years of age, while a lower expression was observed in carcinomas at advanced pathological grades and TNM stages. Statistical analysis comparing P504S/AMACR, p53, Bcl-2 and Ki-67/MIB-1 protein expression in colorectal tumors with various clinicopathological parameters (e.g., age, sex, BMI and smoking) revealed that Bcl-2 expression was significantly associated with patient age (>55 vs. <45 years), pathological grade, and TNM stage (all P<0.05; Table III). Additionally, the Spearman's rank correlation analysis revealed that Bcl-2 expression, based on IHC score, was negatively correlated with pathological grade (r=-0.827; P<0.001) and TNM stage (r=-0.388; P=0.018).

Bcl-2 may serve as a prognostic indicator of colorectal carcinoma. To determine whether P504S/AMACR, p53, Bcl-2 and Ki-67/MIB-1 expression levels are independent prognostic indicators of colorectal carcinoma, The survival rate was considered to indicate prognosis, and was evaluated by multivariate Cox regression analysis. Only Bcl-2 expression displayed statistical significance as a prognostic indicator of colorectal carcinoma with a relative risk value of 0.703 (95% CI, 0.552-0.895; P<0.05; Table IV).

Older patients and those with a lower expression of Bcl-2 exhibit a reduced overall survival time. Based on the aforementioned results, survival analysis based on age and Bcl-2 expression was performed using the Kaplan-Meier method. The Kaplan-Meier curves are plotted in Fig. 3. Log-rank testing revealed that low expression of Bcl-2 in patients was associated with a reduced overall survival time when compared with patients with a higher expression of Bcl-2 (P=0.039), and that older age (>55 years) was associated with a reduced overall survival time (P<0.001).

Discussion

The aim of the present study was to evaluate the protein expression of four key genes, P504S/AMACR, p53, Bcl-2 and Ki-67/MIB-1, in order to assess their potential applicability as diagnostic biomarkers of colorectal carcinoma in a population of Chinese patients. Although a significantly higher expression of all four proteins was observed in colorectal tumors compared with the matching normal tissue margins (as determined by immunohistochemistry and western blot analysis), only Bcl-2 downregulation was significantly correlated with advanced pathological grade and TNM stage in these patients. Additionally, Bcl-2 downregulation was revealed to be a prognostic indicator of a reduced overall survival time.

Apoptosis is well-established as a highly conserved process for eliminating damaged cells in multicellular organisms (14,15). Therefore, defects in apoptotic pathways may result in the survival and proliferation of pre-malignant cells, which eventually leads to the development of human cancer (14,16). In particular, the Bcl-2 protein family has been identified as a key set of apoptotic regulators that conduct the following three activities: i) Promotion of cell survival (e.g., Bcl-2 and Bcl-xL); ii) initiation of cell death (e.g., Bcl-2-interacting mediator of cell death and Bcl-2-interacting domain); and iii) activation of apoptotic effector pathways (e.g., Bcl-2-associated X protein and Bcl-2 homologous antagonist killer) (13). With respect to Bcl-2 and tumorigenesis, elevated Bcl-2 gene expression has been positively correlated with a poorer patient prognosis in colorectal cancer, prostate cancer, bladder cancer, small cell lung cancer, breast cancer, melanoma and acute myeloid leukemia (17). Furthermore, a previous study demonstrated that higher Bcl-2 expression contributes to resistance to chemotherapy and radiation therapy (17).

In contrast with these conventional findings, the present study revealed that Bcl-2 expression was significantly correlated with advanced pathological grade and TNM stage in addition to reduced overall survival time in this Chinese colorectal carcinoma patient population. This phenomenon may be explained through colorectal tumor biology. In the colon, Bcl-2 has been demonstrated to be more highly



Figure 1. Immunohistochemistry of the four proteins in the colorectal tumors and matching normal tissue margins (control). Representative images of the expression of P504S, Bcl-2, p53 and Ki-67 in colorectal tumors and matching normal tissue margins are shown. P504S, α -methylacyl-CoA racemase; Bcl-2, B-cell lymphoma 2; p53, tumor protein p53.

expressed in adenomas than in carcinomas (18,19). As a result, colorectal tumors that retain their Bcl-2 expression tend to display adenoma-like characteristics and are often less aggressive, while tumors with reduced Bcl-2 expression tend to be more aggressive (20).

In terms of tumor characteristics and clinical outcomes, the results of the present study are in line with those of several previous studies in colorectal cancer patients. Biden et al (19) observed a significant negative association between Bcl-2 expression and microsatellite instability, as well as a significant positive association between Bcl-2 expression and patient survival. Ofner et al (21) associated Bcl-2 downregulation with increased tumor size, decreased lymphocytic infiltration and an increased likelihood of a poor clinical outcome. Baretton et al (22) demonstrated that Bcl-2-positive carcinomas are associated with a significantly longer disease-free survival time. Sinicrope et al (23) demonstrated that a high Bcl-2 expression is an independent predictor of improved relapse-free survival, but not overall survival, following adjustment for proliferative index, DNA ploidy and ethnicity. Bhatavdekar et al (24) observed a positive correlation between Bcl-2 expression and a poor survival outcome in patients with UICC/AJCC stage I and III

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	P5()4S/AMA	CR expre	ssion		Bcl-2 e	xpression			p53 ex	pression		K	i-67/MIB	-1 expres	sion
Characteristic	Low	High	χ^2	P-value	Low	High	χ^2	P-value	Low	High	χ^2	P-value	Low	High	χ^2	P-value
Age, years			0.10	0.873			19.75	0.026^{a}			3.02	0.385			0.13	0.858
<45	36	36			36	36			24	48			32	40		
>55	36	40			12	64			36	40			36	40		
Sex			0.06	0.9			0.22	0.815			0.43	0.742			0.08	0.886
Male	36	32			36	44			48	40			44	32		
Female	44	36			28	40			36	24			40	32		
$BMI, kg/m^2$			1.88	0.791			0.16	0.98			0.05	0.994			09.0	0.928
<18.5	28	20			24	24			24	28			20	28		
18.5-23.9	32	24			32	28			28	32			32	36		
≥24	20	24			20	20			16	20			16	16		
Smoking			0.01	0.97			1.08	0.603			0.14	0.85			0.06	6.0
Yes	24	40			28	44			20	32			36	32		
No	32	52			36	40			40	56			44	36		
Alcohol			0.01	0.957			0.35	0.769			0.75	0.666			0.43	0.742
Yes	28	36			48	44			52	44			48	40		
No	36	48			32	24			32	20			36	24		
Pathological grade			2.83	0.622			91.72	<0.001 ^a			3.59	0.313			3.35	0.262
G1	4	12			0	16			4	12			4	12		
G2	16	8			4	20			4	20			20	4		
G3	24	24			36	12			28	20			24	24		
G4	32	28			09	0			24	36			28	32		
TNM stage			1.82	0.813			36.47	0.005 ^a			2.54	0.572			1.35	0.885
T1	0	4			0	4			0	4			0	4		
T2	16	12			8	20			12	16			20	8		
T3	20	24			32	12			24	36			16	28		
Τ4	36	36			09	12			24	32			40	32		
^a Statistically significant Tumor-Node-Metastasis	(P<0.05).	Data are p	resented a:	the number	of patients	. AMACR,	α-methyla	cyl-CoA race	mase; Bcl-	.2, B-cell ly	mphoma	2; p53, tumoi	protein p.	53; BMI, bo	ody mass i	ndex; TNM,

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	Multivariate analysis	
Variable	Relative risk (95% confidence interval)	P-value
P504S/AMACR (low vs. high)	0.982 (0.843-1.114)	0.816
Bcl-2 (low vs. high)	0.703 (0.552-0.895)	0.004^{a}
p53 (low vs. high)	0.922 (0.731-1.162)	0.492
Ki-67/MIB-1 (low vs. high)	1.024 (0.763-1.375)	0.873

Table IV. Multivariate Cox regression analyses of four proteins as prognostic indicators of colorectal carcinoma.

^aStatistically significant (P<0.05). AMACR, α-methylacyl-CoA racemase; Bcl-2, B-cell lymphoma 2; p53, tumor protein p53.



Figure 2. Western blot analysis of the four proteins in the colorectal tumors and matching normal tissue margins (control). (A) Western blot analysis of the four proteins. (B) Expression levels of the four proteins normalized to β -actin. *P<0.001. P504S, α -methylacyl-CoA racemase; Bcl-2, B-cell lymphoma 2; p53, tumor protein p53.



Figure 3. Survival analysis using the Kaplan-Meier method. (A) The association between Bcl-2 expression and overall survival time in Chinese patients with colorectal carcinoma (log-rank P=0.039). (B) Subgroup analysis of overall survival time comparing younger (<45 years) and older (>55 years) Chinese patients with colorectal cancer (log-rank P<0.001). Bcl-2, B-cell lymphoma 2.

colorectal carcinoma. However, in contrast to these findings, other studies have revealed no significant association between Bcl-2 expression and prognosis in patients with colorectal cancer (25-28). Despite numerous clinical investigations, there remains no clear-cut association between Bcl-2 expression and clinical outcomes in patients with colorectal carcinoma.

As stated earlier, non-polypoidal lesions and *de novo* colorectal neoplasms (i.e., without preceding adenoma) occur more commonly in Asian patients (2). Due to the

lower incidence of adenoma-like colorectal tumors in Asian populations and the association between Bcl-2 and an adenomatous differentiated phenotype, it is likely that a lower Bcl-2 expression would be associated with less differentiated, more aggressive colorectal tumors and worse clinical outcomes in Asian patients, which is precisely what was observed in the Chinese patient population enrolled in the present study. Further studies in Chinese and other East Asian (e.g., Japanese and Korean) populations that correlate gross pathological examination (by surgery or endoscopy) with the histopathological and molecular observations are required to validate this hypothesis.

There are several limitations to the present study. To begin with, the sample size used was limited to only 47 patients; a larger sample size would have provided more reliable statistical results. Additionally, only three risk factors for colorectal carcinoma (BMI, smoking status and excessive alcohol consumption) were assessed in the present study. Other risk factors, including a current diagnosis of type-2 diabetes, a previous diagnosis of colorectal cancer, colorectal adenoma/polyps or inflammatory bowel disease (IBD), and a family history of colorectal cancer or colorectal adenoma/polyps, require further investigation in future studies. Furthermore, due to resource limitations, the expression levels of only four proteins, P504S/AMACR, p53, Bcl-2 and Ki-67/MIB-1, were analyzed in the present study. Other proteins that have been previously associated with IBD and colorectal malignancy, including adenomatous polyposis coli, retinoblastoma, deleted in colon cancer and v-Ki-ras2 Kirsten rat sarcoma viral oncogene homolog, also require further investigation in future studies (29). Finally, the present study did not analyze gross tumor morphology, epigenetic factors or the mRNA expression of the proteins examined, all of which may have provided additional insights.

In conclusion, significantly higher expression levels of P504S/AMACR, p53, Bcl-2 and Ki-67/MIB-1 were observed in colorectal tumors than in matched normal tissue margins. Bcl-2 downregulation was revealed to be significantly associated with advanced pathological grade and TNM stage in a Chinese patient population with colorectal carcinoma. Furthermore, it was demonstrated that Bcl-2 downregulation is a prognostic indicator of reduced overall survival time in these patients. Bcl-2 may be a novel target of colorectal carcinoma treatment in the Chinese population of patients.

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Availability of data and materials

All the data and materials are available upon reasonable request.

Authors' contributions

ZL contributed to the study design. GH, QY and LB contributed to data collection. JW contributed to the performance of experiments. BJ and QL contributed to data analysis.

Ethics approval and consent to participate

The present study was approved by the Ethics Committee (Institutional Review Board) of Yongchuan Hospital, Chongqing Medical University (Chongqing, China). All subjects recruited for the present study provided written informed consent prior to participation.

Consent for publication

All patients signed informed consent and agreed to the publication of the present research.

Competing interests

The authors declare that they have no competing interests. The funding body had no role in the study design, data collection and analysis, decision to publish or preparation of the manuscript.

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