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Association of Beclin 1 expression with response to neoadjuvant chemoradiation therapy in patients with locally advanced rectal carcinoma

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Abstract

Beclin 1 is an essential regulator of autophagy that is induced in response to cellular stress and serves to maintain cell survival in established tumors. We recently demonstrated that *Beclin 1* suppression can sensitize colorectal cancer cells to radiation-induced DNA damage and apoptosis. Therefore, we hypothesized that the level of Beclin 1 expression may be associated with radiation sensitivity *in vivo*. We determined the association of Beclin 1 expression in pre-treatment rectal cancer tissues with response to neoadjuvant chemoradiation in surgical resection specimens. Consecutive stage II and III (n=96) rectal adenocarcinoma patients were treated with neoadjuvant chemoradiation followed by surgical resection with curative intent. Beclin 1 was analyzed by immunohistochemistry and the expression level was dichotomized at the median value with categorization into low and high groups. We identified 56 (58.3%) and 40 (41.7%) patients with high vs low level Beclin 1 expression, respectively. Patients with high vs low Beclin 1 expression were significantly less likely to be downstaged after chemoradiation treatment [45% (25/55) vs 58% (22/38); $p=0.02$]. In a multivariable analysis adjusted for age, sex, histological grade and baseline TNM stage, the impact of Beclin 1 expression on tumor downstaging remained statistically significant ($p=0.03$). The association of the level of Beclin 1 expression with the rate of tumor downstaging after chemoradiation is consistent with *in vitro* data, and suggests that Beclin 1 may be a predictive biomarker for the efficacy of chemoradiation in rectal cancer patients.

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CONFLICT OF INTEREST

The authors declare no conflict of interest.

Keywords

Beclin 1; rectal cancer; chemoradiation; autophagy; biomarker

Beclin 1 is an essential regulator of autophagy where it is required for autophagosome formation and maturation.¹ During autophagy, cytosolic proteins and organelles are engulfed into autophagosomes that fuse with lysosomes where they are degraded and recycled.² Autophagy is upregulated in tumor cells in response to metabolic, hypoxic or cytotoxic stress to maintain cell survival.³ Allelic loss of *Beclin 1* and defective autophagy were shown to sensitize cells to metabolic stress and to activate the DNA damage response in association with aneuploidy in immortalized murine epithelial cells and in mammary tumors.^{4, 5} Beclin 1 interacts with multiple proteins including the product of *UV-irradiation-resistance-associated gene (UVRAG)* to form core complexes that regulate autophagy. In a recent study in colorectal cancer (CRC) cells, we found that Beclin 1 and UVRAG expression can protect against radiation-induced DNA double strand breaks with a resultant decrease in tumor cell apoptosis.⁶ These data suggest that Beclin 1 overexpression in CRCs can contribute to radiation resistance. Conversely, suppression of Beclin 1 or inhibition of autophagy has been shown to sensitize CRC cells to radiation⁷ or chemotherapy⁸ which establishes autophagy as a therapeutic target in this malignancy.³

Nearly one-third of new diagnoses of CRC are due to cancers of the rectum.⁹ Patients whose tumors show transmural invasion of the rectal wall (T_{3,4}) or have any depth of invasion with regional lymph node metastases (T_xN_{1,2}) routinely receive concurrent chemotherapy and radiation prior to surgical resection. Preoperative or neoadjuvant chemoradiation is preferred to postoperative treatment since it allows for tumor downstaging that can improve resectability and local control, as well as reducing toxicity.^{10, 11} Tumor downstaging is defined as a decrease in the pathologically-determined tumor (T) and lymph node (N) categories and is commonly utilized as a measurement of treatment response. Downstaging is determined by comparing preoperative staging data with endoscopic ultrasonography and/or imaging with the surgical resection specimen. In rectal cancer patients, tumor downstaging by chemoradiation has been shown to predict long-term outcomes.¹² In this study, we tested the hypothesis that the level of Beclin 1 expression in pretreatment rectal cancer tissues is associated with the response to neoadjuvant chemoradiation therapy. To date, a predictive biomarker for the efficacy of chemoradiation therapy in rectal cancer is lacking^{13–15}, and such a biomarker has the potential enable personalized therapy including novel therapeutic strategies.

MATERIALS AND METHODS

Patient cohort

Consecutive patients with locally advanced rectal adenocarcinoma (T₃₋₄, N₀ or T_x, N₁₋₂) whose pre-treatment tumor tissues were available from our biorepository in the Center for Cell Signaling in Gastroenterology were included. All patients were treated with neoadjuvant chemoradiation and underwent surgical resection with curative intent between 1997 and 2009 at the Mayo Clinic in Rochester, MN. Pre-treatment tumor staging was

performed in all patients using endoscopic ultrasonography, abdominal/pelvic computed tomography, and chest imaging. Neoadjuvant chemoradiation therapy consisting of infusional 5-fluorouracil (5-FU)[225 mg/m²] and external beam radiation therapy (50.4 Gy in 28 fractions). Four to six weeks after completion of chemoradiation, all patients underwent surgical resection with negative surgical margins (R0 resections) being achieved in all. The study was approved by the Mayo Clinic Institutional Review Board.

Assessment of Treatment Response and Tumor Downstaging

Pathological examination of surgical resection specimens was performed in accordance with the TNM classification where T stage, number of involved regional lymph nodes, and histologic grade were determined. We classified the pathological treatment response following chemoradiation in each patient into the following categories: pathological complete response (pCR), residual microscopic disease, or residual macroscopic tumor. Tumor downstaging was determined by a comparison between pretreatment TNM staging and re-staging by pathological examination of the surgical specimen.¹⁶

Beclin 1 expression and scoring in tumor specimens

Tissue sections (4–6 μm) from formalin-fixed, paraffin-embedded tumors were deparaffinized and antigen retrieval was performed in a preheated 0.1mM EDTA, pH 8.0 buffer for 30 min. After blocking endogenous peroxidase activity, slides were incubated (30 min) with a primary anti-Beclin 1 rabbit polyclonal antibody (ABCAM, Cambridge, MA; diluted 1: 250). After rinsing in TBST wash buffer (DAKO, Carpinteria, CA), a secondary incubation was performed (15 min) using a DUAL+/HRP labeled polymer (K4061, DAKO). Slides were subsequently placed in 3,3'-diaminobenzidine (5 min) and counterstained with a modified Schmit's hematoxylin. Negative controls omitted the primary antibody, but included all other procedural steps. Each slide contained a unique number that enabled blinding with respect to clinical data.

Beclin 1 protein expression was scored in tumor biopsy specimens obtained prior to chemoradiation therapy using criteria determined a priori. Tumors were considered positive if > 5% of tumor cells stained for Beclin 1 with a staining intensity 1+. Intensity was graded as follows: 0=not detectable, 1=weak; 2=moderate; 3=strong. Staining extent was defined as the percentage of positive tumor cells and scored: 0: <5%; 1: 5–24%; 2: 25–49%; 3: 50–74%; 4: 75%. Staining intensity and extent were multiplied to produce a weighted score¹⁷ for each tumor whose values were dichotomized at the median for subsequent analyses. All specimens were analyzed by a gastrointestinal pathologist (TTW) without knowledge of clinical information.

Statistical analysis

For Beclin 1 data, categorical variables were analyzed using the Chi-square test and continuous variables using the t-test. The paired differences for the staging variable were compared between Beclin 1 expression levels (low vs high) using a 2-sample t-test. A multivariable linear regression model was developed for predicting the staging paired differences between Beclin 1 expression levels after adjusting for age, gender, histologic grade, and baseline TNM stage. Descriptive statistics and graphical methods were used to

summarize the data. All statistical tests were two-sided with a significance level defined at $p < 0.05$. Data were analyzed using JMP 10.0 (SAS Institute).

RESULTS

Patient Characteristics

Among 96 patients with locally advanced (T_{3-4} , N_0 or T_{any} , N_{1-2}) rectal cancer, tumor location included the distal rectum (0 to 5 cm from anal verge) ($n=38$; 39.6%), midrectum (5–10 cm) ($n=32$; 33.3%) and proximal rectum (>10 cm) ($n=26$; 27.1%). Forty-seven patients (49.0%) underwent a low anterior resection and 49 (51.0%) underwent an abdominoperineal resection. Mean patient age was 58.7 years (range, 25–92 years) and 69.8% were men. Characteristics of the study population stratified by the dichotomized pre-treatment Beclin 1 expression level in tumor tissue are shown in Table 1.

Beclin 1 protein expression was analyzed in carcinoma tissues prior to chemoradiation. Cytoplasmic expression (Fig. 1A) was dichotomized at the median value of the weighted score and Beclin 1 scores at the median were included in the high expression group. High vs low level Beclin 1 expression was detected in 56 (58.3%) and 40 (41.7%) tumors, respectively (Table 1). Beclin 1 expression was not significantly associated with the clinicopathological variables such as age, gender, tumor grade, or pre-treatment TNM stage (all $p > 0.05$) (Table 1).

Beclin 1 Expression and Response to Chemoradiation

Patients with high vs low Beclin 1 expression were significantly less likely to achieve a pathologic response to neoadjuvant chemoradiation, defined as pCR or microscopic residual disease [8 (14.3%) vs 16 (66.7%); $p=0.02$]. An analysis of tumor downstaging based upon the TNM staging system revealed that patients whose tumors had high vs low Beclin 1 expression were significantly less likely to be downstaged after chemoradiation treatment [45% (25/55) vs 58% (22/38); mean (median) downstaging: -0.64 (0) vs -1.16 (-1); ($p=0.02$)] (Fig. 1B). In a multivariable analyses that adjusted for age, gender, histologic grade, and baseline TNM stage, the association of the Beclin 1 expression level with tumor downstaging was maintained ($p=0.03$) (Table 2). Specifically, tumor downstaging by chemoradiation was significantly predicted by both the Beclin 1 expression level and by the TNM stage at baseline in contrast to other clinicopathological variables (Table 2).

DISCUSSION

The treatment of choice in patients with locally advanced rectal carcinoma is preoperative chemoradiation therapy followed by surgical resection.^{10, 11} In patients receiving such therapy, studies have suggested that the degree of tumor downstaging is clinically important in that it can be used for treatment monitoring and as a prognostic parameter.¹⁸ In this regard, tumor downstaging has been proposed as a surrogate marker for favorable long-term outcome in rectal cancer patients.^{19, 20} A range of pathological response to chemoradiotherapy occurs that includes pCR (ypT0N0), with no viable tumor cells left (currently 10% to 25% of patients), to microscopic residual disease, to virtually no tumor regression or even tumor progression during therapy.²¹ Based upon preclinical data, we

evaluated the level of Beclin 1 expression as a potential predictive marker of pathological response and downstaging by neoadjuvant chemoradiation. Consistent with *in vitro* data⁶, we found that a high level of Beclin 1 expression in pre-treatment rectal carcinoma tissues was associated with a significantly reduced rate of tumor downstaging by chemoradiation therapy compared to tumors with a low level of Beclin 1 expression. This finding for Beclin 1 remained statistically significant in a multivariable analysis after adjustment for relevant covariates. We also found that the pathological tumor response to chemoradiation was significantly reduced in tumors with high vs low level Beclin 1 expression. Together, these data suggest that Beclin 1 overexpression can contribute to treatment failure and may, therefore, enable tumor progression. Further support for these data derives from a study whereby the expression of Beclin 1 was associated with poor survival after chemoradiation in patients with esophageal squamous cell carcinoma compared to tumors lacking Beclin 1.²² Since all patients received 5-FU and concurrent radiotherapy, we were unable to determine the effect of Beclin 1 expression on 5-FU alone. Preclinical studies have shown that siRNA knockdown of *Beclin 1* or *Atg5* can sensitize human colorectal cancer cells to 5-FU.²³ In patients with stages II and III colon carcinoma treated with 5-FU-based adjuvant chemotherapy, Beclin 1 overexpression was associated with reduced survival.²⁴ While conflicting data have been reported regarding the prognostic impact of Beclin 1 in CRC, a recent meta-analysis found that patients whose CRCs had high Beclin 1 expression had a poor prognosis.²⁵

Radiation has been shown to promote autophagy as a mechanism of stress tolerance in cultured tumor cell lines. However, the role of autophagy or its key regulator Beclin 1 in radiation resistance in human cancers remains unclear. Beclin 1-deficient cells have been reported to display increase susceptibility to radiation-induced cell death.²⁶ Furthermore, knockdown of *Beclin 1* and *autophagy related 5 (ATG5)* genes were shown to increase radiosensitization induced by a mTOR/PI3K inhibitor (NVP-BEZ235).⁷ In CRC cell lines, we recently reported that Beclin 1 expression can confer resistance to radiation-induced DNA double strand breaks that can be reversed by siRNA knockdown of *Beclin 1* to promote cell death.⁶ These observations are clinically relevant in that the autophagy inhibitor chloroquine has been shown to strongly promote radiation-induced cell death in highly radioresistant cancer stem cells, and is currently being studied in clinical trials in cancer patients.²⁷

Strengths of our study are uniform preoperative staging procedures that included endoscopic ultrasonography in all patients performed at a major tertiary referral center. Study limitations including the relatively small sample size and retrospective design. We recognize the inherent limitations related to comparison of pre-therapeutic imaging and surgical pathology findings post chemoradiation that include the potential for overstaging rectal cancers by endoscopic ultrasonography²⁸. Our series included a higher than expected number of rectal cancers with poor differentiation. However, Beclin 1 expression did not differ by histologic grade in this study nor in most prior reports in this malignancy.²⁵ Other limitations include interpretation of IHC by one expert pathologist, although scoring criteria were established a priori. It will be important to confirm the association of the dichotomization of Beclin 1 expression used in our study with clinical outcome. A formal method to assess neoadjuvant treatment response by grading histologic changes in the resected specimen has been

proposed²⁹, although correlation with disease prognosis awaits further study and external validation is needed.

In conclusion, we found that the level of Beclin 1 expression in rectal carcinomas was significantly associated with pathological response and rates of tumor downstaging following neoadjuvant chemoradiation therapy. If validated prospectively, our findings support a role for utilizing Beclin 1 as a predictive biomarker for the efficacy of neoadjuvant chemoradiation therapy in patients with rectal cancer.

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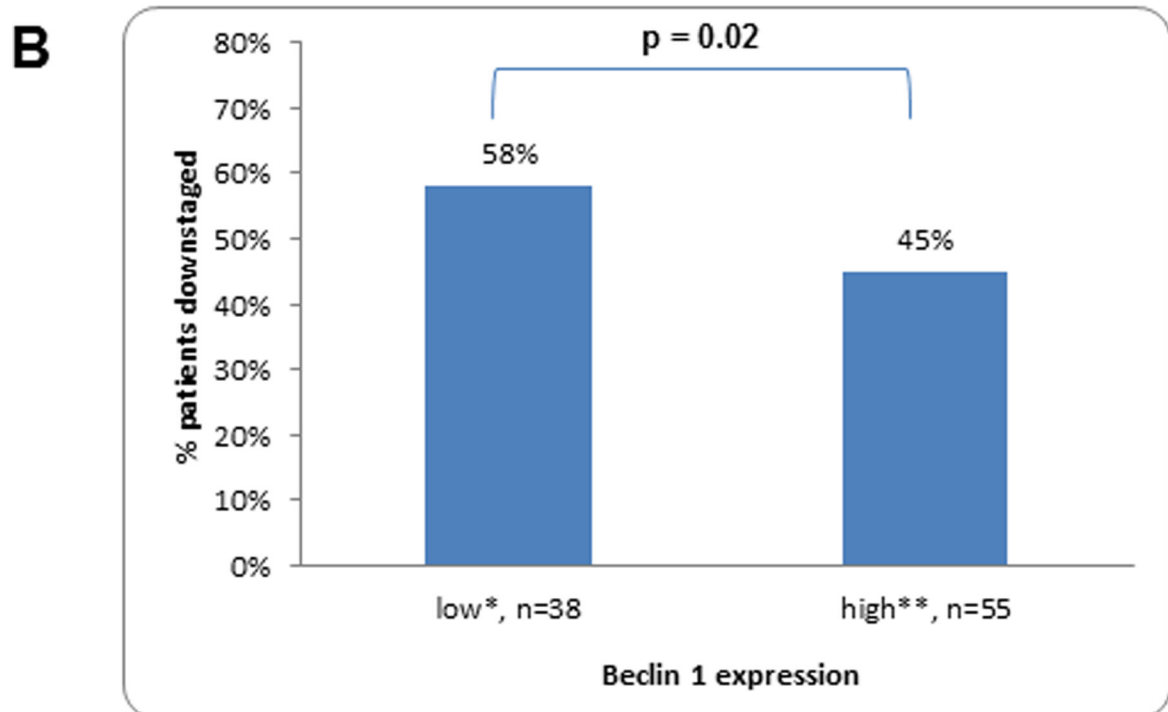
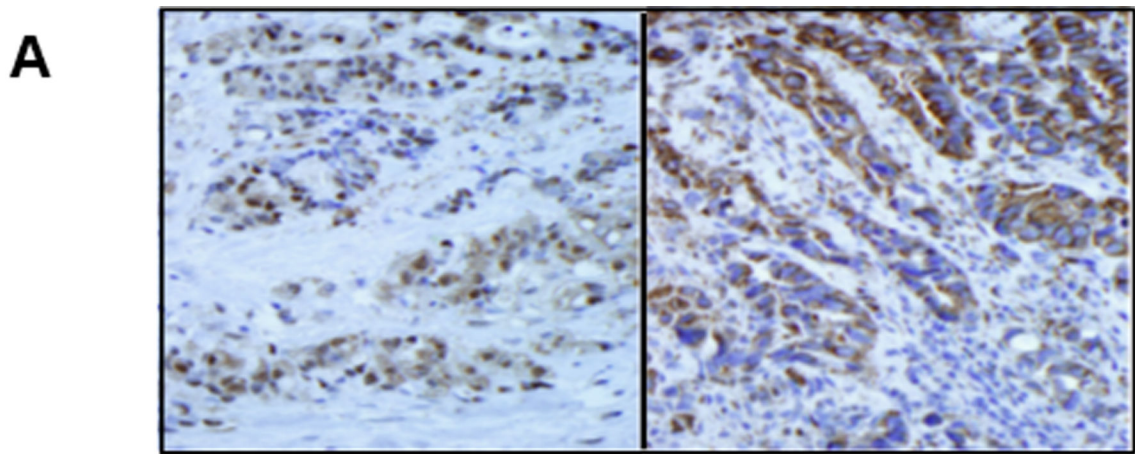
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Novelty and Impact of Work?

The essential autophagy protein Beclin 1 was recently shown to protect against radiation-induced DNA damage with a resultant decrease in colorectal cancer cell apoptosis. In locally advanced rectal cancer patients, we found that high vs low level Beclin 1 expression in pretreatment tumor tissues can predict the response to chemoradiation therapy with a significantly reduced rate of tumor downstaging. These data suggest the potential utility of Beclin 1 expression as a predictive biomarker.



* Missing data (n=2); ** missing data (n=1)

Figure 1.

A) Immunostaining for Beclin 1 protein expression was performed in rectal carcinomas. Representative tumors with low (*left*) vs high (*right*) level expression of Beclin 1 in the tumor cell cytoplasm (see Methods) are shown from pre-treatment biopsies ($\times 400$). **B)** Percentage of patients with rectal cancers who are downstaged after neoadjuvant chemoradiation therapy according to the level of Beclin 1 expression in pre-treatment tumor biopsies.

Table 1

Patient Characteristics

	All patients (%)	Low Beclin 1 expression	High Beclin 1 expression	P value*
No. of patients	96 (100)	40 (100)	56 (100)	
Mean age, range (years)	58.7 (25–92)	59.9 (38–92)	57.8 (25–88)	0.41**
Gender				
Male	67 (69.8)	25 (62.5)	42 (75.0)	0.19
Female	29 (30.2)	15 (37.5)	14 (25.0)	
Histologic grade				
Well/moderate	26 (27.1)	13 (32.5)	13 (23.2)	0.31
Poor	70 (72.9)	27 (67.5)	43 (76.8)	
Pre-treatment TNM staging				
T3-4, NO	19 (19.8)	7 (17.5)	12 (21.4)	0.69
T1-4, N1-2	74 (77.1)	31 (77.5)	43 (76.8)	
Unknown***	3 (3.1)	2 (5.0)	1 (1.8)	

* 2-sided Chi-square p value

** 2-sample t-test p value

*** Treated as missing for Chi-square test

Table 2

Multivariate Regression Model Predicting for Changes in TNM Stage From Baseline

Variable	Estimate* (standard error)	p value
Beclin 1 level expression	-0.49 (0.23)	0.03
Gender	-0.06 (0.25)	0.82
Histologic grade	-0.14 (0.25)	0.58
TNM stage (at baseline)	0.57 (0.28)	0.04
Age	0.005 (0.009)	0.60

* Estimate is the differences in the means between the groups after adjusting for the other variables. For example, for the Beclin 1 expression variable, the -0.49 estimate is the low Beclin 1 expression mean minus the high Beclin 1 expression mean after adjusting for the other variables (-1.01 - (-0.52)).

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