



Prognostic role of beclin-1 in locally advanced non-small cell lung cancer in patients receiving docetaxel-platinum induction chemotherapy

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Background/Aims: The outcome of local treatment for advanced non-small cell lung cancer (NSCLC) remains poor, with therapies such as induction chemotherapy (IC) yielding conflicting results. This study aimed to assess the clinicopathologic and prognostic significance of the excision repair cross-complementation group 1 (ERCC1), beclin-1, and glucose-regulated protein of molecular mass 78 (GRP78) in patients with locally advanced NSCLC receiving docetaxel-platinum IC, along with efficacy and safety.

Methods: This is a retrospective observational cohort study. We reviewed medical records of 31 NSCLC patients receiving docetaxel-platinum IC, and conducted immunohistochemical staining of ERCC1, beclin-1, and GRP78.

Results: Response rate was 67.8% with 10.7 months of median relapse-free survival (RFS) and 23.1 months of median overall survival (OS), and no treatment-related death was reported. High expression of ERCC1, beclin-1, and GRP78 was identified in 67.7%, 87.1%, and 67.7%, respectively. Expression of ERCC1 and GRP78 did not reveal statistical significance in survival, whereas high beclin-1 expression revealed longer OS (7.6 months vs. 23.2 months; log-rank $p = 0.024$). In multivariate analysis, histologic differentiation (hazard ratio [HR], 3.48; $p < 0.001$), stage (HR, 8.5; $p = 0.024$), and adjuvant treatment (HR, 16.1; $p = 0.001$) were related to RFS, and in OS, stage (HR, 5.4; $p = 0.037$), adjuvant treatment (HR, 8.6; $p = 0.004$), and beclin-1 expression (HR, 8.2; $p = 0.011$) were identified as significant prognostic factors.

Conclusions: Our findings suggest that high beclin-1 expression predicts longer survival in locally advanced NSCLC and docetaxel-platinum IC is a treatment option that deserves consideration.

Keywords: BECN1 protein, human; Carcinoma, non-small-cell lung; Induction chemotherapy

INTRODUCTION

Lung cancer is a leading cause of cancer deaths. Non-small cell lung cancer (NSCLC) accounts for many lung

cancers, with up to one-third of patients with stage III of the disease. Despite advances in treatment modality, locally advanced NSCLC presents a therapeutic challenge. In terms of extent and size of the primary tumor, and

nodal status, stage III NSCLC is a heterogeneous disease. The accepted treatment for locally advanced NSCLC is a multi-modal strategy including chemotherapy, radiation therapy, and surgery. However, with outcomes remaining poor, the most appropriate treatment remains controversial.

Induction chemotherapy (IC) in locally advanced NSCLC has revealed conflicting data [1,2]. IC may eradicate micrometastasis and assess tumor sensitivity to chemotherapy, and this information may be valuable for follow-up chemotherapy at the adjuvant and palliative stages. IC can diffuse more efficiently into the tumor, making it more advantageous than postradiation or surgery, and most importantly, lead to downstaging, thereby resulting in a higher percentage of curative resections. Conversely, there are drawbacks, including a delay in definitive local treatment and the potential for increased morbidity because of chemotherapy-induced toxicities. Therefore, many studies have focused on determining a more accurate prediction of tumor response, to select patients that would benefit most from IC.

The excision repair cross-complementation group 1 (ERCC1) enzyme plays a key role in the nucleotide excision repair system. It recognizes and removes platinum-induced DNA adducts; thus, its expression is associated with platinum resistance [3-5].

Hypoxia, hypoglycemia, and acidosis are main characteristics of tumor microenvironment, that lead to chronic endoplasmic reticulum (ER) stress. Glucose-regulated protein of molecular mass 78 (GRP78), a member of heat shock protein family, is a molecular chaperone in ER. Under ER stress, GRP78 activates a pro-survival pathway using an autophagy mechanism, and this leads to the protection and survival of cancer cells and chemoresistance [6,7]. Autophagy is a lysosomal pathway involved in cellular defense, with beclin-1 playing a key role in this mechanism [8]. Recently, several studies have been focusing on the role of autophagy in cancer. Autophagy has been suggested to be involved in taxane cytotoxicity, although this correlation is under investigation [9-13].

This purpose of this study was to assess the clinicopathologic and prognostic significance of ERCC1, beclin-1, and GRP78 in patients with locally advanced NSCLC receiving docetaxel-platinum IC, along with efficacy and safety of the treatment.

METHODS

Patients and treatment

We reviewed the medical records of NSCLC patients diagnosed and treated at Seoul St. Mary's Hospital January 2003 to June 2012. Patients that satisfied the following eligibility criteria were enrolled in this study: patients with histologically proven locally advanced NSCLC, an Eastern Cooperative Oncology Group (ECOG) performance status of ≤ 1 , available biopsied tissue acquired before IC, and those who received docetaxel-platinum IC for curative local treatment including surgery and concurrent chemoradiation treatment (CCRT). Cancer staging was evaluated according to the 7th edition of the American Joint Committee on Cancer (AJCC). This study was approved by the Institutional Review Board (IRB) of the Catholic Medical Center (XC13TIM10010K). IRB approved a waiver of informed consent. All patients were treated with 75 mg/m² of docetaxel and 75 mg/m² of cisplatin or carboplatin area under curve (AUC) 5 on day 1 every 3 weeks.

Evaluation of tumor toxicity and response

Toxicity was assessed according to the Common Terminology for Adverse Events version 3.0 (CTCAE v.3). Following completion of IC, tumor response was evaluated according to the Response Evaluation Criteria in Solid Tumors (RECIST 1.1).

Immunohistochemical staining and scoring

Immunohistochemical (IHC) staining was conducted using 4- μ m sections of paraffin-embedded tissue, biopsied for initial diagnosis. After deparaffinization in 100% xylene and rehydration over a series of decreasing concentrations of ethanol, using standard protocols, slides were subjected to antigen retrieval using the heat induced epitope retrieval method in 0.01 M citrate buffer (pH 6.0). Endogenous peroxidase activity was neutralized by immersion in 3% hydrogen peroxide in methanol for 10 minutes, followed by a 4°C overnight incubation with mouse monoclonal anti-ERCC1 (1:200, GeneTex, Irvine, CA, USA), rabbit monoclonal anti-beclin-1 (1:200, Epitomics, Burlingame, CA, USA), and rabbit polyclonal anti-GRP78 (1:1,000, Abcam, Cambridge, UK). After washing, the sections were incubated with polymer-conjugated horseradish peroxidase for

10 minutes at room temperature. After incubation, the sections were counterstained with hematoxylin. Two pathologists with no clinical information about the patients interpreted all immunostained slides. We scored expression levels by measuring the intensity of positive staining and graded as low and high.

Statistical analysis

Pearson chi-square test and Fisher exact test were used to assess the association between the expression of molecules (ERCC1, beclin-1, and GRP78) and clinicopathologic factors. Relapse-free survival (RFS) was defined as the time from the start of IC to relapse or death for patients dying without relapse. Overall survival (OS) was defined as the time between start of IC and death. We assessed the relationship between survival (RFS, OS) and the expression of molecules (ERCC1, beclin-1, and GRP78) using Kaplan-Meier methods. Univariate and multivariate Cox regression analyses were used to identify prognostic factors. We used the SPSS/PC+ version 18.0 (SPSS Inc., Chicago, IL, USA), and a *p* value of < 0.05 was considered statistically significant.

RESULTS

Patient characteristics

Thirty-one patients were enrolled in this study. Baseline characteristics are presented in Table 1. Mean age at diagnosis was 64 years (range, 43 to 75) and 27 patients (87.1%) were men. ECOG performance status was reported as grade 1 in 25 patients (80.6%), and 26 patients (83.9%) were former or current smokers. Histology revealed squamous cell carcinoma in 20 patients (64.5%), adenocarcinoma in 10 patients (32.3%), and large cell carcinoma in one patient. Twenty-eight patients (90.3%) had stage III disease, and three patients (9.7%) had stage IIB disease. Twenty-eight patients (90.3%) received docetaxel-cisplatin IC and three patients (9.7%) received docetaxel-carboplatin. The median number of IC cycles was three (range, 2 to 4).

Response, operability, and survival outcomes

After two cycles of IC, three patients (9.7%) demonstrated complete response, 18 patients (58.1%) partial response, nine patients (29%) stable disease, and one pa-

tients (3.2%) revealed progressive disease.

Following IC, 18 patients (58.1%) underwent surgery, eight patients (25.8%) received CCRT, two patients (6.5%) received radiotherapy alone, two patients (6.5%) were treated with second-line chemotherapy or best supportive care, and one patient was lost to follow-up. Of 18 patients undergoing surgical resection, 15 patients (83.3%) achieved Ro resection, and 12 patients (66.7%) revealed pathologic downstaging. Two patients died within 30 days postsurgery (massive hemoptysis and sepsis), and one patient died of radiation pneumonitis after defini-

Table 1. Patient characteristics (n = 31)

Characteristic	Value
Male sex	27 (87.1)
Age, yr	64.4 ± 7.1
Performance status	
0	6 (19.4)
1	25 (80.6)
Smoking status	
Never-smoker	5 (16.1)
Former-smoker	14 (45.2)
Current-smoker	12 (38.7)
Pathology	
Squamous cell	20 (64.5)
Adenocarcinoma	10 (32.3)
Large cell	1 (3.2)
Histologic differentiation	
Moderate	20 (64.5)
Poor	11 (35.5)
Stage	
IIB	3 (9.7)
IIIA	21 (67.7)
IIIB	7 (22.6)
Regimen	
Docetaxel-cisplatin	28 (90.3)
Docetaxel-carboplatin	3 (9.7)
No. of IC cycles	
2	15 (48.4)
3	15 (48.4)
4	1 (3.2)
RDI	89.4 ± 11.7

Values are presented as number (%) or mean ± SD.

IC, induction chemotherapy; RDI, relative dose intensity.

Table 2. Toxicities of docetaxel-platinum induction chemotherapy in locally advanced non-small cell lung cancer

	All grades	Grade 3-4
Hematologic toxicities		
Neutropenia	23 (74.2)	21 (67.7)
Anemia	11 (35.5)	1 (3.2)
Thrombocytopenia	5 (16.1)	1 (3.2)
Non-hematologic toxicities		
Asthenia	27 (87.1)	1 (3.2)
Myalgia	15 (48.2)	0
Anorexia	26 (83.9)	1 (3.2)
Nausea	18 (58.1)	2 (6.5)
Vomiting	10 (32.3)	2 (6.5)
Diarrhea	11 (35.5)	2 (6.5)
Constipation	4 (12.9)	0
Alopecia	22 (71)	0
Mucositis	12 (38.7)	2 (6.5)
Neuropathy	6 (19.4)	0
Rash	5 (16.1)	0

Values are presented as number (%).

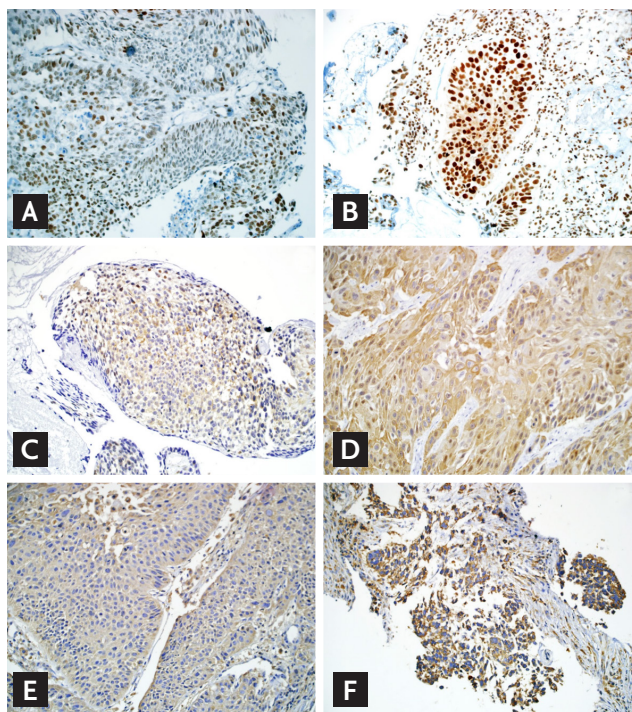


Figure 1. Expression by immunohistochemistry (×200). (A) Low excision repair cross-complementation group 1 (ERCC1). (B) High ERCC1. (C) Low beclin-1. (D) High beclin-1. (E) Low glucose-regulated protein of molecular mass 78 (GRP78). (F) High GRP78.

CCRT. Adjuvant CCRT was administered to six patients after surgical resection and adjuvant chemotherapy to nine patients.

Median follow-up duration was 18.8 months (range, 3 to 130), and median RFS and OS was 10.7 months (95% confidence interval [CI], 5.5 to 15.8) and 23.1 months (95% CI, 15.1 to 31), respectively.

Toxicities

Over the course of chemotherapy, asthenia, anorexia, neutropenia, alopecia, nausea, and myalgia were frequently observed (Table 2). However, all toxicities were transient and were managed with supportive care. Treatment-related death did not occur. The most common grade 3/4 toxicity was neutropenia (67.7%) and febrile neutropenia occurred in a patient. Grade 3/4 neutropenia was managed with granulocyte-colony stimulating factor (G-CSF), with or without antibiotics. Most of the patients recovered from neutropenia within 3 days and we did not observe delay of chemotherapy from prolonged toxicities of > 1 week. The mean duration from last chemotherapy treatment to surgery was 1.1 ± 0.3 months (range, 20 to 66 days).

Expression of ERCC1, Beclin-1, and GRP78

High expression of ERCC1, beclin-1, and GRP78 were identified in 21 patients (67.7%), 27 patients (87.1%), and 21 patients (67.7%), respectively (Fig. 1). High ERCC1 expression was found frequently in low T staging (T1, 2) compared to high T staging (T3, 4) with a relative risk of 2 (p = 0.021). Correlations between expression of the molecules and other clinicopathologic features or tumor response were not statistically significant (Table 3). In addition, relationships between expression of ERCC1, beclin-1, and GRP78 were not identified (Table 4).

Predictive and prognostic role of ERCC1, Beclin-1, and GRP78

In the Kaplan-Meier analysis of RFS, the expression of ERCC1, beclin-1, and GRP78 did not reveal statistical significance (median RFS in patients with low vs. high expression; ERCC1: 6.9 months vs. 10.7 months, log-rank p = 0.775; beclin-1: 6.3 months vs. 10.7 months, log-rank p = 0.185; GRP78: 6.9 months vs. 10.7 months, log-rank p = 0.314). In the analysis of OS, patients with high expression of beclin-1 showed longer OS (median

Table 3. Correlation between expression of molecules (ERCC1, beclin-1, and GRP78) and clinicopathologic characteristics

	ERCC1		Beclin-1		GRP78	
	High	p value	High	p value	High	p value
All cases	21		27		21	
Smoking						
Never-smoker:smoker	3:18	1	5:22	1	5:16	0.147
Pathology						
SqCC:non-SqCC	14:7	1	20:11	1	12:9	0.262
Histologic differentiation						
MD:PD	14:7	1	19:8	0.115	13:8	1
Stage						
T1, 2:T3, 4	16:5	0.021	19:12	0.619	14:7	0.447
No:N1, 2, 3	3:18	0.739	4:23	1	4:17	0.277
IIB, IIIA:IIIB	17:4	0.652	21:6	1	16:5	1
Tumor response						
CR + PR:SD + PD	15:6	0.685	18:9	1	14:7	1

Values are presented as number.

ERCC1, excision repair cross-complementation group 1; GRP78, glucose-regulated protein of molecular mass 78; SqCC, squamous cell carcinoma; MD, moderately differentiated; PD, poorly differentiated; CR, complete response; PR, partial response.

Table 4. Correlations between expression of ERCC1, beclin-1, and GRP78

	ERCC1			GRP78		
	High	Low	p value	High	Low	p value
Beclin-1						
High	18	9	1	18	9	1
Low	3	1		3	1	
GRP78						
High	16	5	0.222			
Low	5	5				

ERCC1, excision repair cross-complementation group 1; GRP78, glucose-regulated protein of molecular mass 78.

Table 5. Prognostic factors for survival by multivariate analysis

Factor	RFS		OS	
	HR	p value	HR	p value
Differentiation (PD vs. MD)	34.8	< 0.001		
Stage (IIIB vs. IIB, IIIA)	8.5	0.118	5.4	0.037
Nodal stage (N1, 2, 3 vs. No)				
Definitive treatment (CCRT vs. surgery)				
Adjuvant treatment (no vs. yes)	16.1	0.001	8.6	0.004
Beclin-1 (low vs. high)			8.2	0.011

RFS, relapse-free survival; OS, overall survival; HR, hazard ratio; PD, poorly differentiated; MD, moderately differentiated; CCRT, concurrent chemoradiation.

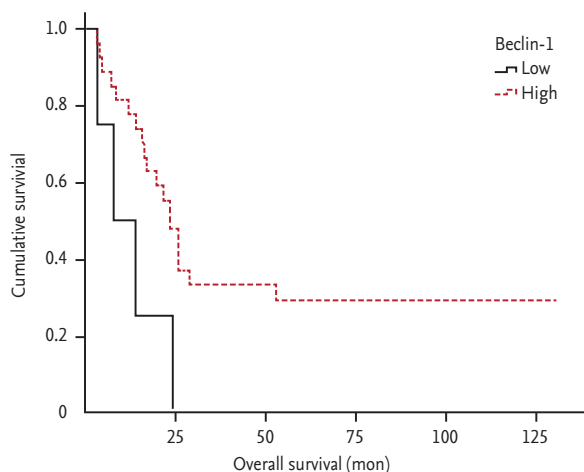


Figure 2. Overall survival of patients with non-small cell lung cancer according to beclin-1 expression.

OS in patients with low vs. high expression; ERCC1: 23.1 months vs. 19.5 months, log-rank $p = 0.647$; beclin-1: 7.6 months vs. 23.2 months, log-rank $p = 0.024$; GRP78: 19.5 months vs. 23.2 months, log-rank $p = 0.547$) (Fig. 2). Univariate Cox regression analysis revealed that histologic differentiation (moderate vs. poor; hazard ratio [HR], 17.8 and $p < 0.001$) was significantly related to RFS and beclin-1 expression (low vs. high; HR, 3.6 and $p = 0.012$) and adjuvant treatment after definitive treatment (no vs. yes; HR, 4.4 and $p = 0.035$) were related to OS. Multivariate analysis conducted with factors exhibiting $p < 0.2$ at univariate analysis (histologic differentiation, stage, nodal stage, type of definitive treatment, additional adjuvant treatment, and expression of beclin-1). Identified significant factors were histologic differentiation, stage, and adjuvant treatment for PFS, and stage, adjuvant treatment and expression of beclin-1 for OS (Table 5).

DISCUSSION

This study was conducted to find predictive and prognostic biomarkers in patients with locally advanced NSCLC receiving docetaxel-platinum IC, in company with efficacy and safety of IC. Our results suggested beclin-1 as a prognostic biomarker, and IC as a safe and effective treatment option.

In our study, patients with high expression of beclin-1 revealed longer OS, while significant predictive role of its expression in terms of IC such as RFS or tumor re-

sponse was not identified. However, high beclin-1 expression revealed a tendency towards longer RFS. Since only four patients (12.9%) revealed low expression of beclin-1, further study with an expanded cohort and standardizing method of beclin-1 assessment is needed. Autophagy maintains intracellular homeostasis which has conflicting roles in cancer cells, anti-tumors and pro-tumors. Beclin-1 is essential for autophagosome formation, and in human cancer, its downregulation and tumor suppressive role have been reported [12,14-16]. In cervical and duodenal cancer, low beclin-1 expression was associated with poor prognosis and in cervical cancer, the expression of beclin-1 in tumor tissue was lower compared to adjacent normal tissue [15,16]. In breast cancer, autophagy enhanced paclitaxel-induced cell death and autophagy-related markers are downregulated in docetaxel-resistant breast cancer cell lines [17]. Meanwhile, by contrast with aforementioned studies and our results, in endometrial and pancreatic cancer, high beclin-1 expression was associated with poor prognosis, suggesting a pro-tumor role of beclin-1 [18,19]. In NSCLC, autophagy and beclin-1 have been studied as well and conflicting results were reported. Wang et al. [12] and Kim et al. [20] reported consistent results with our study that low beclin-1 expression was associated with aggressive behavior and poor prognosis. Whereas several preclinical studies with NSCLC cell lines reported different results supporting pro-tumor role of autophagy, revealing contribution of autophagy to taxane resistance. In this context, autophagy was proposed as a therapeutic target to overcome chemoresistance [11,13]. In our study and the abovementioned studies, Wang et al. [12] and Kim et al. [20] reported a majority of patients had squamous cell histology (64.5%, 51.9%, and 51.9%, respectively), while the NSCLC cell lines in the preclinical studies revealing a pro-tumor role of autophagy were all adenocarcinoma cell lines [11-13]. Hence, in NSCLC, the role of autophagy may be different according to histology. The different roles of autophagy according to histologic subtype were not identified in our study, so further study is needed in an expanded cohort and preclinical setting with squamous histology cell lines. Beclin-1 expression did not have a significant predictive role in terms of IC such as RFS or tumor response; however, high beclin-1 expression revealed a tendency towards longer RFS.

In our study, high expression of GRP78 revealed longer RFS and OS compared to low expression group with no statistical significance. Emerging data indicates that ER stress and autophagy are involved in human cancer, specifically in carcinogenesis, disease aggressiveness, metastasis, and sensitivity/resistance to anti-tumor treatment [6-10]. Kim et al. [20] reported that expression of GRP78 and beclin-1 in NSCLC correlates with low tumor stage and longer survival, and that their expressions correlated with each other. Conversely, Sun et al. [21] reported that GRP78 mRNA expression was significantly higher (71.9%) in stage III/IV NSCLC, like our result of 71.7%. Additionally, in a small interfering RNA experiment, they have revealed that GRP78 promotes metastasis and anti-apoptosis.

Besides the undefined roles of autophagy and ER stress in NSCLC, several factors may contribute to results with lack of statistical significance [9,10,17]. First, we focused on locally advanced disease, mainly IIIB. We had limited sample numbers ($n = 31$), and there is concern in serving biopsied tissue as a representative sample for whole tumor.

Being not in accord with previous studies, neither predictive nor prognostic role of ERCC1 was identified in our study. A recent study suggested that IHC analysis with commercially available 8F1 antibodies did not specifically detect the unique functional ERCC1 isoform, thereby limiting its use in the clinic towards therapeutic decision making [22].

In the present study, the response rate of docetaxel-platinum IC was higher (67.8%) than those of previous studies employing paclitaxel-carboplatin IC (40% to 55%), and median OS (23.1 months) was similar to those of previous reports (OS, 16 to 29 months). Considering that about a quarter of patients in this study had stage IIIB disease, while previous reports focused on stage IIIA disease, the result of this study is notable [23-25]. In terms of toxicities, the present study revealed a higher incidence of grade 3/4 neutropenia (78.4%) than that with paclitaxel-carboplatin or gemcitabine-cisplatin IC [23,25]. Ethnic differences of pharmacogenomics in docetaxel metabolism may contribute to the serious hematologic toxicities. However, these were transient and successfully managed in all patients using G-CSF and isolation care. In addition, these toxicities did not lead to delay in the follow-up definitive local treatment, and we ob-

served no treatment-related death. The addition of adjuvant treatment after definitive treatment was identified as a viable prognostic factor in RFS and OS. Hence, IC followed by surgery and adjuvant treatment could be an optimal sequence. And the addition of immunotherapy and detection of cell free tumor DNA after definitive treatment might play roles in the treatment of locally advanced NSCLC [26].

Our findings suggested beclin-1 expression as a favorable prognostic biomarker in locally advanced NSCLC. Further study is required to validate biomarkers related to autophagy and ER stress in NSCLC.

KEY MESSAGE

1. Our study suggested the possibility of beclin-1 expression as a favorable prognostic marker in locally advanced non-small cell lung cancer (NSCLC) receiving docetaxel-platinum induction chemotherapy.
2. Further extended study is required to validate biomarkers related to autophagy and endoplasmic reticulum stress in NSCLC.

Conflict of interest

No potential conflict of interest relevant to this article was reported.

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