



Galectin-9 as a prognostic biomarker in small cell lung cancer

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Contributions: (I) Conception and design: Y Xu, X Zhang; (II) Administrative support: X Zhang; (III) Provision of study materials or patients: X Zhang; (IV) Collection and assembly of data: C Guan; (V) Data analysis and interpretation: Y Xu; (VI) Manuscript writing: All authors; (VII) Final approval of manuscript: All authors.

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Background: Galectin-9 has been reported to play an important role in the immunoregulation, survival, and growth of different cancers. Our study aims to uncover the clinical significance of galectin-9 in small cell lung cancer (SCLC) postoperatively.

Methods: We studied 48 patients with SCLC who underwent surgery from 2008 to 2014. Galectin-9 expression was assessed by immunohistochemistry on tissue and plot survival curves by the Kaplan-Meier curve. Prognostic analysis was conducted via integrating all of the independent indicators into a multivariate Cox analysis for overall survival (OS). Furthermore, we evaluate whether SCLC patients with high or low galectin-9 expression might benefit from chemotherapy or radiotherapy.

Results: High galectin-9 expression was significantly correlated with high levels of neuron specific enolase (NSE) expression ($P=0.04$). We found that galectin-9 ($P<0.001$), NSE ($P=0.006$) and postoperative therapeutic methods ($P=0.006$) were associated with patient survival time by Kaplan-Meier survival calculations and log-rank tests. SCLC patients with high galectin-9 expression had a relatively shorter OS than those with low galectin-9 expression. In addition, a multivariate analysis revealed high galectin-9 expression as a potential independent prognostic factor for OS [hazard ratio (HR) =6.21, 95% confidence interval (CI), 2.98–12.95, $P<0.01$], disregarding elevated NSE and therapeutic methods. Moreover, the benefit to patients undergoing chemotherapy or radiotherapy was superior in galectin-9 high expressed SCLC patients.

Conclusions: High expression of galectin-9 is an independent prognostic factor for OS in patients with SCLC. Evaluation of galectin-9 expression may predict the benefits from chemotherapy or radiotherapy.

Keywords: Small cell lung cancer (SCLC); galectin-9, the immune checkpoint; prognosis analysis

Submitted Feb 04, 2018. Accepted for publication May 04, 2018.

doi: 10.21037/tcr.2018.05.18

View this article at: <http://dx.doi.org/10.21037/tcr.2018.05.18>

Introduction

Small cell lung cancer (SCLC) is a deadly tumor that is a highly aggressive and widely metastatic cancer, with a 7% 5-year survival rate (1). SCLC is one pathology type of lung cancer which also includes lung adenocarcinoma, lung squamous carcinoma and large cell lung cancer. Although SCLC accounts for approximately 15% of lung cancer patients, it has the poorest prognosis of all lung cancers. Although there has been much effort in conducting comprehensive molecular research (2), there

have been no important therapeutic clinical advances for 30 years, leading SCLC to be designated as an intractable cancer (1). Furthermore, SCLC is a neuroendocrine tumor which has a relatively complex tumor microenvironment. Recently, immune checkpoint inhibitors play an important role in anti-tumor therapies for melanoma and renal cell carcinoma, bladder cancer, non-small cell lung cancer and others (3,4). However, there have been few studies to analyze the functions of immune checkpoints in the progression of SCLC.

The Galectin family has many biological functions, such

as immunological modification, tumor cell metastasis, the induction and maintenance of tumor angiogenesis, and so on (5). Galectin-9 is one member of the galectin family and is mainly expressed on macrophagocytes, antigen-presented cells and tumor cells, and it can ligand with T-cell immunoglobulins and mucin-domain containing-3 (TIM-3) which is expressed on lymphocytes so as to inhibit its function (6,7). Blocking the ligation between galectin-9 and TIM-3 may partially recover the function of the exhausted lymphocytes (8,9). However, there are many different, even contradictory results about the survival analysis of galectin-9 in different cancers.

In this study, we evaluated the expression of galectin-9 in clinical specimens of 48 SCLC patients by immunohistochemical analysis (IHC). The association between galectin-9 expression and clinicopathological characteristics, and overall survival (OS) by the Kaplan-Meier curve and cox survival analysis were also elucidated. Moreover, we evaluated the predictive potential of galectin-9 expression in SCLC patients with chemotherapy or radiotherapy.

Methods

Patients and specimens

This study was approved by the Shengjing Hospital of Chinese Medical University ethics committee (No. 2016PS256K), and we acquired exemptions via informed consent from patients for the use of abandoned paraffinic specimens. We collected pathological specimens from 48 patients with SCLC who underwent surgery at the China Medical University affiliated Shengjing Hospital, China, from 2008 to 2014, though three patients failed to follow-up. Most of the SCLC patients that received surgery by mistake did not carry out an effective pathological examination or puncture before surgery. And the nodules of minor established SCLC patients could have been resected, such patients were in the 1A stage (T1N0M0 in TNM stage) and not infiltrated in visceral pleura, with main bronchus, surrounding lymph nodes or distant organs and tumor sizes of less than 3 centimeters. Patients were diagnosed in stages by serum tumor markers: CEA, CY21-1, NSE, SCC; and imaginology examinations, positron emission tomography/computed tomography (PET/CT) or chest CT scans, bone ECT, and brain contrast-enhanced magnetic resonance imaging were conducted before surgery. The mean age of these patients was 56.6 years, with a range of 33 to 82 years. Most of the included SCLC patients had

received etoposide and cis-platinum (EP) chemotherapy, and some received etoposide and lobaplatin, epirubicin and cyclophosphamide (EC), cyclophosphamide, adriamycin and vincristine (CAV) and irinotecan and paraplatin (IP). Patients who had received neoadjuvant therapy or immune system related diseases were excluded. Histologic diagnoses were based on hematoxylin and eosin staining according to the World Health Organization 2004 criteria. We used neoplastic and paraneoplastic specimens (more than 5 centimeters apart from margin of the tumor) to carry out the following research.

Neuron specific enolase (NSE) is one of the most important markers used to evaluate the progress of SCLC patients (10). SCLC are patients always accompanied with hyponatremia, which is caused by SIADH (inappropriate secretion of antidiuretic hormone) or paraneoplastic syndrome and so on, and hyponatremia always predicts a poor outcome. Meanwhile, some SCLC patients with a state of hypercoagulability (high level D-dimer) also had worse prognoses. Among the above-mentioned data, NSE, Na⁺, and D-dimer were acquired in the serum of patients the day after their admission to the hospital, always about one week before surgery. Clinicopathological data collected for analysis included sex, tumor location, age of diagnosis, tumor size, node involvement (N), NSE expression, Na⁺ (sodium ion) level and D-dimer expression. Disease recurrence and survival were observed during the follow-up clinic or obtained through telephone correspondence. Follow-up was until death or December 2015.

Immunohistochemistry

IHC was performed on the resected SCLC tumor tissues using the primary antibodies, anti-galectin-9 antibody (1:800, 54330, CST) and CD8 (1:100, 17335-1-AP, Proteintech Group, China), according to a previously described protocol (11). All results were recorded by NIS-Elements F 3.0 (Japan) and computerized image systems (Nikon E800 microscope, Japan). All IHC results were analyzed by two pathologists independently in a blind manner. The intensity for staining was defined as follows: no staining was considered a negative result ("0"). Other positively stained sections were analyzed by color: weak staining ("1"), moderate staining ("2"), strong staining ("3"). This was followed by a calculation of the histoscore (H-score) according to the following formula: $H\text{-score} = 1 * (\% \text{ cell "1"}) + 2 * (\% \text{ cell "2"}) + 3 * (\% \text{ cell "3"})$. A mean H-score was obtained by averaging these five

representative fields (400× objective).

Statistical analysis

We carried out correlations between IHC expression and the clinic-pathological features by the Pearson's chi-squared test, and calculated survival situations of different groups by the Kaplan-Meier survival and log-rank tests. And Cox regression model was used to carry out multivariate analysis. The statistical analyses were performed using the SPSS software program, version 20.0 (SPSS, Chicago, IL, USA), which was considered statistically significant when P values are less than 0.05.

Results

The expression level and correlation of galectin-9 and clinic-pathological features

We evaluated galectin-9 expression in 48 human SCLC tissues by IHC. First, galectin-9 was abundant and expressed mainly in the cytoplasm and membrane of cancer cells (*Figure 1A*). There was no obvious difference between the galectin-9 expression of tumor tissue and para-tumor tissues, including bronchial alveolar cells and lymphoid tissues (*Figure 1B*). There were 24 SCLC patients over cut-off values, forming the galectin-9-high expression group, and 24 SCLC patients below cut-off values, forming the galectin-9-low expression group. Detailed clinic-pathological characteristics are shown in *Table 1*. We found that over-expressed galectin-9 on tumor specimens was associated with high levels of NSE expression ($P=0.04$). In this way, we thought NSE could be an advantageous factor in the selection of SCLC patients who may benefit more from anti-galectin-9 targeting therapy. However, there was no significant association between over-expressed galectin-9 and tumor size, N, Na+ expression, D-dimer expression or number of CD8⁺T lymphocytes present in SCLC tissue.

Prognostic values of galectin-9 in human SCLC

To determine the prognostic value of the expression levels of galectin-9, we used Kaplan-Meier survival calculations and log-rank tests for sex, tumor location, age of diagnosis, tumor size, node status (N), NSE, Na+, D-dimer, postoperative therapeutic method and galectin-9. We found that galectin-9 ($P<0.001$), NSE ($P=0.006$) and postoperative therapeutic methods ($P=0.006$) were associated with patient survival time. As shown in *Figure 1C*, patients with higher galectin-9

expression (galectin-9-high group) tended to have a shorter overall survival (OS) (15.64 ± 2.32 months) compared to that of the galectin-9-low group (28.13 ± 5.43 months) ($P<0.001$). Furthermore, we used a multivariate Cox regression model to analyze galectin-9, NSE and postoperative therapeutic methods to determine their prognostic values. In addition to elevated NSE [hazard ratio (HR) =2.27, 95% confidence interval (CI), 1.09–4.70, $P=0.03$], high galectin-9 expression (HR =6.21, 95% CI, 2.98–12.95, $P<0.01$) and different therapeutic methods (HR =0.57, 95% CI, 0.33–0.97, $P=0.04$) were independent predictors of poor overall survival in SCLC (*Table 2*).

In the study, 39 of 48 (81.25%) patients received chemotherapy or radiotherapy. To evaluate whether patients with high or low expressed galectin-9 would benefit from chemotherapy or radiotherapy, we studied the relationship between galectin-9 expression and the OS of SCLC patients with different therapeutic methods. We divided the 48 SCLC patients into two groups by galectin-9 cut-off expression into a galectin-9 high group and a galectin-9 low group. In the galectin-9 high expression group, the survival rates from surgery and additional chemotherapy/radiotherapy were a mere 1/6 and 1/17, respectively. In the galectin-9 low expression group, the survival rates from surgery and additional chemotherapy/radiotherapy were 0/3 and 11/19, respectively. There were 3/48 lost during follow-up. We then analyzed the survival of SCLC patients by therapeutic methods (surgery versus a combination of surgery and chemotherapy or radiotherapy). We found patients with chemotherapy or radiotherapy had a longer OS than patients with merely surgery ($P=0.006$, *Figure 1D*). There appeared to be an association between the OS of SCLC patients and different therapeutic methods in the galectin-9 high group rather than the galectin-9 low group (*Figure 1D*). A study about the correlation between galectin-9 expression, therapeutic methods and OS revealed that in the galectin-9 high group, the benefit observed in patients with chemotherapy or radiotherapy was superior to that of those patients without chemotherapy or radiotherapy ($P=0.042$, *Figure 1D*). Furthermore, there was no obvious relationship between survival and therapeutic methods in the galectin-9 low group. Therefore, we believe that patients with high galectin-9 expression might benefit more from chemotherapy or radiotherapy after surgery.

Discussion

Lung cancer is one of the deadliest cancers worldwide. In

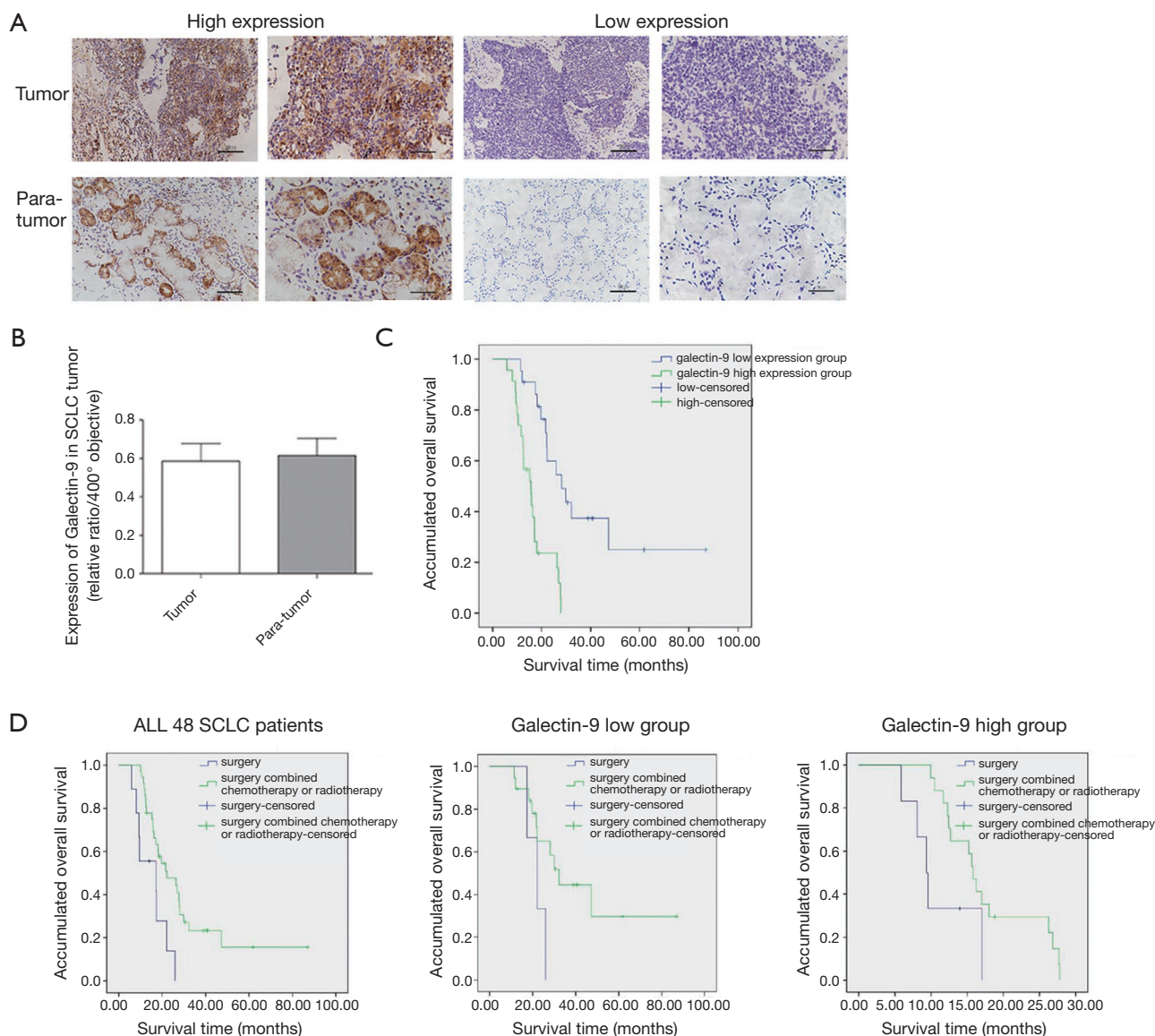


Figure 1 Galectin-9 expression and survival analysis in SCLC. (A) Representative immunohistochemistry results of galectin-9 of SCLC patients with high or low expression (staining methods: protein expression was detected by IHC with anti-galectin-9 antibody 200/400 \times object); (B) comparison between the galectin-9 expression of tumor tissue and para-tumor tissues including bronchial alveolar cells and lymphoid tissues. There were no obvious differences between tumor and para-tumor tissue; (C) survival analysis of galectin-9 high group and galectin-9 low group by Kaplan-Meier curve. patients with higher galectin-9 expression (galectin-9-high group) tended to have shorter overall survival (OS) (15.64 ± 2.32 months) compared to galectin-9-low group (28.13 ± 5.43 months) ($P < 0.001$); (D) survival analysis of different therapeutic methods by Kaplan-Meier curve in all SCLC patients, galectin-9 low group and galectin-9 high group. The benefit observed in patients with chemotherapy or radiotherapy was superior to the patients without chemotherapy or radiotherapy in galectin-9 high group ($P = 0.042$).

particular, SCLC is the poorest type of lung cancer with an unknown genetic mechanism and high invasion or metastasis. In our study, we revealed the prognostic value

of galectin-9 in SCLC. Pathological findings showed that patients with high galectin-9 levels were found to have a higher prevalence of NSE, which is a classic SCLC tumor

Table 1 Correlation between galectin-9 expression and patient characteristics

Characteristics	Low galectin-9	High galectin-9	P
Sex (male/female)	19/5	16/8	0.33
Location of tumor (left/right)	12/12	11/13	0.77
Age at diagnosis (year)	53.0±9.2	60.2±9.9	0.13
Tumor size (cm)	3.2±1.9	3.6±2.0	0.55
N status (N0/N1/N2)	6/7/11	8/5/11	0.73
NSE before operation (normal/elevated)	16/8	9/15	0.04
Na ⁺ before operation (normal/ elevated)	24/0	22/2	0.15
D-dimer before operation (normal/elevated)	20/4	22/2	0.38
Infiltrated CD8 ⁺ T lymphocytes in SCLC tissue (low/high)	9/15	15/9	0.08

The results of continuous variables are presented as mean ± SD (standard deviation).

Table 2 Multivariate Cox regression analysis of overall survival in 48 SCLC patients

Variables	Hazard ratio (95% CI)	P
Galectin-9 expression (high vs. low)	6.21 (2.98–12.95)	<0.01
CD8 T cells (more vs. less)	1.98 (0.57–6.84)	0.28
NSE (elevated vs. normal)	2.27 (1.09–4.70)	0.03
Therapeutic methods (surgery and chemotherapy or radiotherapy vs. surgery)	0.57 (0.33–0.97)	0.04

SCLC, small cell lung cancer.

biomarker. High galectin-9 expression was an independent prognostic factor of the OS of SCLC patients, disregarding NSE and different therapeutic methods. Furthermore, the benefit to patients undergoing chemotherapy or radiotherapy was superior in galectin-9 high expressed SCLC patients.

High-expression of galectin-9 in SCLC patients was found to be related to high NSE levels. On the one hand, NSE could be an advantageous factor in the selection of SCLC patients who may benefit more from anti-galectin-9 targeting therapy. On the other hand, mean NSE values were significantly higher in patients with extensive disease compared to those with limited disease. In this way, we speculate on whether galectin-9 could be related to tumor load, which may be one of the pieces of evidence suggesting that galectin-9 is a bad prognostic biomarker in SCLC.

The results of galectin-9 in survival analysis with various studies were different. Some researchers thought it was a protective factor in breast cancer (12), bladder urothelial carcinoma (13) and melanoma (14) by means of inducing cell aggregation and apoptosis, or decreasing

invasion and metastasis. However, others thought it was a disadvantageous risk factor in hepatocellular carcinoma (15), gastric cancer (16), pancreatic carcinoma (17) and lung adenocarcinoma (18) by means of up-regulating the interaction with TIM-3 to decrease the cytotoxic or proliferation of exhausted lymphocytes and so on. As for our results, an elevated galectin-9 expression was an independent disadvantage of a decreased OS in SCLC patients, and we thought this result may partially be related to the complex tumor microenvironment of SCLC. Furthermore, the other members of the galectin family, galectin-1 and galectin-3 probably play a role in the invasion, metastasis, anti-apoptosis and angiogenesis of cancer cells (19). Both galectin-1 and galectin-3 were associated with poor disease outcomes in lung cancer patients as previously reported (5,20).

Recently, there were some studies that found that galectin-9/TIM-3 functions as an immune checkpoint to inhibit the anti-tumor immune functions of lymphocytes, NK cells and so on (13). Zhou *et al.* found that antibodies against TIM3 restore responses of HCC-derived T cells

to tumor antigens (21). Kikushige *et al.* thought that TIM-3+ leukemic stem cells (LSCs) could also cause autocrine galectin-9 to react with TIM-3 to sustain self-renewal and promote Treg cell differentiation and maintenance (22). Furthermore, some studies have found that the survival time of animals with lung cancer, melanoma and glioma could be prolonged via blocking the reaction of galectin-9/TIM-3 (9,23,24). Aside from this, there have been many studies involved in the study of the immunoregulatory function of galectin-9. Wu *et al.* found that galectin-9 could react with CD44 to increase iTreg cell stability and function by the transcription factor Smad3, which participated in forming a feed-forward loop (25). In addition, galectin-9 can react with Dectin-1 which is expressed on macrophages and myeloid-monocytic cells to suppress their functions (17). However, Dectin-1 can protect against chronic hepatitis from hepatocarcinoma to induce hepatic fibrosis (26). Galectin-9 can also ligand with CD137 expressed on lymphocytes to activate their functions (27). So, the immuno-regulatory function of galectin-9 is complex and controversial.

In our study, we found that the survival time of patients with chemotherapy or radiotherapy was longer in galectin-9 high SCLC patients, though the survival rate was not obviously increased. Moreover, Liu and colleges also found the benefit associated with adjuvant chemotherapy was superior among galectin-9 low patients than among galectin-9 high patients in bladder urothelial carcinoma (13). We believe this result should be further proved in a larger group. Furthermore, many combination therapies with immune checkpoint inhibitors have been studied recently with great success. Kim *et al.* found that a combination of anti-TIM-3 on the basis of anti-PD-1 and stereotactic radiosurgery (SRS) could inhibit glioma growth and prolong the survival of mice (23). In this way, whether or not galectin-9 high expressed SCLC patients benefit more from a combination therapy of anti-galectin-9 immune therapy and chemotherapy or radiotherapy should feature in our subsequent studies. We think new immune checkpoint inhibitors with galectin-9 need to be further studied for the prolonging of the survival of SCLC patients.

Conclusions

Our results indicate that elevated galectin-9 expression is an independent indicator of a decreased OS in patients with SCLC. Evaluation of galectin-9 expression may predict the benefits from chemotherapy or radiotherapy in SCLC patients.

Acknowledgments

We are grateful to the pathology department of Shengjing Hospital affiliated to China Medical University in the acquisition of human tumor specimens. We thank members of the Medical Research Center of Shengjing Hospital, China Medical University.

Funding: This work was supported by grants from department of science and technology of Liaoning Province, Liaoning Province Finance Department (2014021032) and Chinese National Natural Science Foundation (61672146).

Footnote

Conflicts of Interest: All authors have completed the ICMJE uniform disclosure form (available at <http://dx.doi.org/10.21037/tcr.2018.05.18>). The authors have no conflicts of interest to declare.

Ethical Statement: The authors are accountable for all aspects of the work in ensuring that questions related to the accuracy or integrity of any part of the work are appropriately investigated and resolved. The study was conducted in accordance with the Declaration of Helsinki (as revised in 2013). This study was approved by the Shengjing Hospital of Chinese Medical University ethics committee (No. 2016PS256K), and we acquired exemptions via informed consent from patients for the use of abandoned paraffinic specimens.

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Cite this article as: Xu Y, Guan C, Zhang X. Galectin-9 as a prognostic biomarker in small cell lung cancer. *Transl Cancer Res* 2018;7(3):571-577. doi: 10.21037/tcr.2018.05.18