

Assessment of K-ras Gene Mutation after Neoadjuvant Chemotherapy in Patients with Stage III_A non-small Cell Lung Cancer

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Abstract Objective To assess the state of K-ras gene mutation after neoadjuvant chemotherapy in patients with stage III_A non-small cell lung cancer (NSCLC). **Methods** Twenty four patients with stage III_A NSCLC received treatment of operation after two cycles neoadjuvant chemotherapy. Histopathologic response in resection specimens was observed. Assessment of K-ras genes mutation in resection specimens was performed by polymerase chain reaction single strand conformation polymorphism (PCR-SSCP) analysis. **Results** Clinical effective rate of chemotherapy was 41.7% (10/24). Surgery resection rate was 95.8% (23/24). The rate of marked histopathologic response (tumor regression grade IV-III) for chemotherapy was 43.5%. Nine mutation (39.1%) of K-ras gene in resection specimens of 23 patients were found. The rate of K-ras mutation (20.0%) in patients who had tumor regression grade IV-III was significantly lower than those (53.8%) in patients with histologic response grade II-I for chemotherapy ($P<0.05$). The rate of K-ras mutation was 20.0% in patients who achieved complete resection of tumor, as compared with 75.0% in patients who had incomplete resection of tumor ($P<0.05$). **Conclusion** The lower rate of K-ras gene mutation in patients who had marked histologic response (grade IV-III) for chemotherapy may be a result of effective neoadjuvant chemotherapy, eliminating tumor compartments with mutation K-ras gene. The presence of K-ras gene mutation may not be of impact on effect of neoadjuvant chemotherapy and patients prognosis. Farther study of state of K-ras gene mutation after neoadjuvant chemotherapy in patients with NSCLC is needed.

Key Words Non-small cell lung cancer; Neoadjuvant chemotherapy; K-ras gene; Mutation

Approximately 25% to 30% of non-small cell lung cancer (NSCLC) patients present with stage III disease, which is defined as locally advanced tumor confined to the chest without distant metastasis. Stage III_A patients undergoing surgery had a 5-year survival of 15%~20%^[1]. Extrathoracic disease recurrence makes the possibility of surgical cure unlikely in this group. Combined-modality therapy employing systemic (chemotherapy) and local (surgery or radiotherapy) approaches has shown favorable results in patients with stage III disease. The results of numerous randomized trials demonstrate that neoadjuvant (induction) chemotherapy followed by surgery may improve outcome compared to that achieved with surgery alone^[2-5]. A randomized phase III trial published by Rosell et al shown that the low survival rate in the surgery-alone group may be associated with the higher incidence of K-ras gene point mutation in this group (42%) vs the neoadjuvant chemotherapy group (15%)^[2]. Hence, it is possible that K-ras gene mutation in the group of surgery alone is responsible for the observed outcome differences rather than a beneficial effect of the neoadjuvant chemotherapy itself. This study

was designed to treat the patients with stage III_A NSCLC by neoadjuvant chemotherapy followed surgery, then, to examined the mutational state of K-ras gene in resection specimens. The purpose of the study was to assess the possible impact of K-ras gene mutation on neoadjuvant chemotherapy in patients with stage III_A NSCLC.

MATERIAL AND METHODS

Patients

During the period from February 1999 to April 2002, 24 patients with stage III_A NSCLC were entered into our study. Among them, 16 were males, 8 females, mean age 58.4 ± 10.3 years old ranging from 48 to 69 years. All patients were diagnosed by histologic examination of biopsy material or cytologic examination of percutaneous needle aspiration specimen or bronchoscopic brushing or washings. Among them 13 were adenocarcinoma, 8 squamous cell carcinoma, 2 adenosquamous cell carcinoma, large cell carcinoma 1. Initial clinical evaluation included a medical history, physical examination, chest X-ray, CT scan of chest, brain and abdomen, pulmonary function, electrocardiogram, measurement of blood biochemical analysis. All patients were staged III_A NSCLC according to the International

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Staging System for Lung Cancer, 1997 revision^[6]. Eligibility criteria included a favorable medical condition (a Karnofsky performance score of 80 or higher); sufficient bone marrow reserve (leukocytes $>4000 \times 10^9$ and thrombocytes $>100000 \times 10^9$); normal renal and hepatic function, adequate pulmonary function (a predicted postoperative FEV1 greater than 1.0L).

Treatment, response assessment, and follow-up

All patients received neoadjuvant chemotherapy regimen consisting of mitomycin (8mg/m² day 1), vindesine (3mg/m² days 1, 8), and cisplatin (60 mg/m² day 1), or etoposide (80 mg/m² days 1-4) and cisplatin (60 mg/m² day 1) at three to four weeks intervals for two cycles. Intensive antiemetic therapy and hydration was administered as required.

Approximately 4 weeks after completion of neoadjuvant chemotherapy, clinical response was evaluated with a second CT scan of chest. Conventional response definitions were used for this CT scan-based clinical response assessment. A complete response (CR) was defined as the total disappearance of all radiographic evidence of tumor. A partial response (PR) was defined as any response less than complete, but with greater than 50% reduction in the sum of the products of the crossed diameters of all measurable lesions. Patients with less tumor shrinkage were considered stable disease (SD). Progressive disease (PD) was defined as greater than a 25% increase in the sum of the products of the crossed diameters of all measurable lesion. A thoracotomy was performed in four to five weeks after 2 cycles neoadjuvant chemotherapy. After postoperative recovery, patients were referred for adjuvant chemotherapy. A chemotherapy regimen identical to the pretreatment regimen was given for 4 to 5 cycles. Chemotherapy-related toxicity was observed and recorded.

Follow-up examination were taken every 3 months for the first 2 years and all 6 months thereafter.

Histopathologic assessment

The degree of histopathologic response was determined by assessing the extent of tumor necrosis, as described by Junker et al^[7], somewhat revised by us. Grade IV: complete tumor regression with no evidence of vital tumor tissue in the section of surgical specimen; Grade III: morphologic evidence of therapy-induced tumor regression with $<20\%$ residual tumor cell in the section of surgical specimen presenting focal disease; Grade II: tumor regression with at least 20% residual tumor cell in the section of specimen presenting more than focal microscopic disease; Grade I: no tumor regression or on-

ly spontaneous tumor regression in the section specimen.

Detection of K-ras gene mutation with PCR-SSCP

DNA preparation Resection tumor specimens were finely minced and suspended in 500 μ l DNA lysis solution. The mixtures were incubated at 55°C for 2h, heated at 95°C for 10min, and centrifuged briefly. After phenol/choroform extraction, the DNAs were precipitated with 50 μ l isopropanol. DNAs were washed once with 70% ethanol and resuspended in 100 μ l of filtered H₂O. Finally, DNA content were determined by fluorimetry.

PCR-SSCP and nucleotide sequencing PCR-SSCP sequencing analysis was performed according to the technique described by Cho et al^[8]. DNA was amplified by polymerase chain reaction (PCR). Reaction were generally performed in a volume of 50 μ l and were comprised of 0.2 μ g DNA sample, 200 μ M dNTPS, 5 μ l PCR reaction buffer, 1.5mM MgCl₂ and 2.5 μ Taq polymerase, two primers (K-ras / 12, 13; 5' - ATTATAAG-GCCTGCTG - 3', 5' - CCTGCACCAGTAATATGC - 3'), the samples were overlaid with paraffin oil and subjected to 35 cycles of amplification. The standard temperature profile of the cycles was as followed: denaturation at 94°C for 1min, annealing at 55°C for 1 min, and extension at 72°C for 30 seconds. The SSCP analysis was performed as followed: PCR products were diluted in loading buffer (95% formamide, 20mM EDTA, 10mM NaOH, 0.05% bromophenol blue, 0.05% xylene cyanol blue), boiled for 5 minutes, and rapidly cooled on ice. Samples were run at 150 Volt for 5h on 6% acrylamide gels with 10% glycerol at 4°C. Autoradiography against radiographic film with intensifying screens was performed at - 70°C.

Statistical analysis

Comparisons between patients with K-ras mutation and without K-ras mutation on the median survival were made on the student t test. Difference in the K-ras mutation and 3 - year survival rate were evaluated by the Chi square test. A *p* value of less than 0.05 was considered as statistically significant.

RESULTS

Clinical response and toxicity

All 24 patients received the planned 2 cycles of neoadjuvant chemotherapy. Among them, one patient (4.2%) had CR, 9 (37.5%) had PR, and 13 patients (54.2%) demonstrated SD, only 1 patient had evidence

of PD. Total response rate (CR+PR) was 41.7%. The main chemotherapy-related toxicity was gastrointestinal reactions (45.8%), such as nausea, vomiting and diarrhea, neutropenia (25.0%), thrombocytopenia (16.7%), and abnormal hepatic function tests (12.5%). It was entirely manageable to chemotherapy-related toxicity without treatment interruption.

Surgical result

All 24 patients underwent operation in 4 weeks after chemotherapy. Twenty three patients (95.8%) received surgical resection, 15 cases (62.5%) had complete resection, 8 (33.3%) had incomplete resection, in one case, the tumor was not resectable because of encasement of the great vessels. Operation included lobectomy (14 cases), lobectomy plus other segmental lobe resection (3 cases), lobectomy plus bronchoplastic sleeve resection (3 cases), and complete pneumonectomy (3 cases).

Histopathological assessment tumor specimens in 23 patients who received surgical resection shown that 1 (4.3%) was histologic response grade IV, 9 (39.1%) were grade III, 10 (43.5%) shown grade II. In 3 specimens (13.0%), no evidence of chemotherapy-induced tumor histological response (grade I) was found. The rate of marked histopathological response (grade IV-III) was 43.5% (10/23), among them, adenocarcinoma were 4, squamous cell carcinoma 5, adenosquamous cell carcinoma 1. All 10 patients with histopathologic response grade IV and III achieved complete resection. Only 5 of 13 patients with tumor regression grade II - I had complete resection.

Detection of K-ras gene mutation.

PCR-SSCP analysis is based on the fact that the electrophoretic mobility of single stranded nucleic acids depends not only on size but also on its sequence in non-denaturing polyacrylamide gels. In 23 resection tumor specimens, 9 K-ras mutation (39.1%) were confirmed by direct nucleotide sequencing. All 9 mutation were at codon 12 of the K-ras gene. Mutation were present in 6 of 12 adenocarcinomas (50.0%), 2 of 8 squamous cell carcinomas (25.0%), 1 of 2 adenosquamous cell carcinomas (50.0%). Rate of K-ras mutation was 20.0% (2/10) in tumor regression grade IV-III as compared to 53.8% (7/13) in histopathologic response grade II - I ($p < 0.05$) (table 1). Rate of K-ras mutation in patients who achieved complete resection was 20.0% (3/12), compared with 75.0% (6/8) in patients who had incomplete resection ($p < 0.05$) (table 2).

Table 1. Comparison of K-ras gene mutation between tumor regression grade IV-III and grade II - I.

K-ras mutation status	n	grade IV-III (%)	grade II - I (%)
K-ras mutation	9	2(20.0)	7(53.8)*
no K-ras mutation	14	8(80.0)	6(46.2)

$p < 0.05$, vs grade IV-III

Table 2 Comparison of K-ras mutation between patients with complete resection and incomplete resection.

K-ras mutation status	n	complete resection	incomplete resection
		(%)	(%)
K-ras mutation	9	3(20.0)	6(75.0)*
no K-ras mutation	14	12(80.0)	2(25.0)

$p < 0.05$, vs complete resection

Follow-up and survival

After neoadjuvant chemotherapy and surgery, all 23 patients were followed up. The median follow-up period was 40.5 months, the median survival duration were 23.5 months, 3-year survival rate was 30.4%. The median survival duration for patients with K-ras gene mutation were 15.6 months as compared to 35.8 months in without K-ras gene mutation ($p < 0.05$). The 3-year survival rate was 22.2% (2/9) for patients with K-ras gene mutation, 35.7% (5/14) for patients without K-ras gene mutation, the difference of 3-year survival rate between these patients were not statistically significant ($p > 0.05$).

DISCUSSION

The concept of neoadjuvant chemotherapy, surgical resection performed after chemotherapy, has gained widespread acceptance for patients with IIIA NSCLC because the results from numerous trials have demonstrate the feasibility of this approach and have suggest that survival rates are improved [1-5]. Neoadjuvant chemotherapy is considered as a cytoreductive therapy administered prior to surgical resection. The intent of cytoreductive therapy is to downstage primary tumor and thence increase the resectability rate [9]. Early administration of chemotherapy may eradicate micrometastasis, decrease relapse, improve overall survival, and allow accurate pathologic assessment of response to chemotherapy. Two randomized trials had indicated that the survival rate of patients with stage IIIA disease is significantly better in association with neoadjuvant chemotherapy and surgical resection than that with resection alone [2-5]. In a trial Rosell et al performed by mutation K-ras gene was found in 3 out of 20 patients (15%) in the chemotherapy-surgery group as opposed to 10 out of 24 (42%) in the surgery alone group [2]. We

want to know whether K-ras mutation status influences the effect of chemotherapy or chemotherapy could reverse the expression of K-ras gene, and prognostic impact of a K-ras gene mutation on patients with stage III_A NSCLC receiving chemotherapy followed by surgery.

The ras gene family comprises three genes: K-ras, H-ras and N-ras encoding membrane-bound proteins p21^{ras}. By a point mutation in codons 12 (80% of all K-ras mutations), 13 and 60, the oncogene acquires transforming potential. The amino acid change alters the resulting P21^{ras} protein configuration and stimulates cell growth and differentiation autonomously^[10]. Early studies suggested that K-ras gene mutations were an important negative prognostic factor, and were associated with tumor progression and shortened survival in patients with NSCLC. However, several more recent and larger studies cast doubt on this^[11]. Therefore, these analyses on the prognostic impact of K-ras gene mutation are still controversial.

Our study showed that K-ras gene mutation occurred more frequently in patients with a minor extent of histopathologic response (grade II - I) compared with patients with histopathologic response grade IV - III, and in patients with incomplete resection as compared to patients with complete resection. It seems to be difficult to confirm that an inhomogeneous distribution of mutated K-ras genes have multidrug resistance contributed to these different results, or the lower K-ras mutation rate in patients with marked histopathologic response (grade IV - III) was a result of effective neoadjuvant chemotherapy, eliminating tumor compartment with mutated K-ras gene. Experimental study found that several NSCLC cell lines were tested for resistance against different cytotoxic agents without any association between ras gene mutation and resistance^[12]. In our study, absence of tumor cell or the presence of significant areas of tumor necrosis were seen throughout resection specimens in patients with histopathologic response grade IV - III. Hence, lower rate of K-ras gene mutation in patient with tumor regression grade IV - III could be a result of effective chemotherapy eliminating tumor cells with mutated K-ras gene. Shorter median survival in patients with K-ras gene mutation in addition to could be impact of K-ras gene mutation itself, other important factor was the rate of surgical complete resection in patients without K-ras mutation were markedly higher than that in patients with K-ras mutation. The survival benefit in patients with complete resection after neoadjuvant chemotherapy had been confirmed^[9].

In conclusion, our study data and the results of the current analysis have indicated that the neoadjuvant

chemotherapy is acceptable to eligible patients with stage III_A NSCLC. The rate of K-ras gene mutation in surgical specimens with tumor regression grade IV - III was significantly lower than that with histopathologic response grade II - I, it may be a result of effective preoperative chemotherapy eliminating tumor cell with mutated K-ras gene. There are no sufficient data to conclude that K-ras gene mutation makes an impact on neoadjuvant chemotherapy in patients with NSCLC. The further evaluation of K-ras gene mutation in study on neoadjuvant chemotherapy is needed.

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