Development of anaplastic lymphoma kinase (ALK) inhibitors and molecular diagnosis in ALK rearrangement-positive lung cancer

Eiji Iwama1,2
Isamu Okamoto3
Taishi Harada3
Koichi Takayama2
Yoichi Nakanishi2,3

1Department of Comprehensive Clinical Oncology, Faculty of Medical Sciences, Kyushu University, 2Research Institute for Diseases of the Chest, Graduate School of Medical Sciences, Kyushu University, 3Center for Clinical and Translational Research, Kyushu University Hospital, Fukuoka, Japan

Correspondence: Isamu Okamoto
Center for Clinical and Translational Research, Kyushu University Hospital, 3-1-1 Maidashi, Higashi ku, Fukuoka 812-8582, Japan
Tel +81 92 642 5378
Fax +81 92 642 5389
Email okamotoi@kokyu.med.kyushu-u.ac.jp

Abstract: The fusion of echinoderm microtubule-associated protein-like 4 with anaplastic lymphoma kinase (ALK) was identified as a transforming gene for lung cancer in 2007. This genetic rearrangement accounts for 2%–5% of non-small-cell lung cancer (NSCLC) cases, occurring predominantly in younger individuals with adenocarcinoma who are never- or light smokers. A small-molecule tyrosine-kinase inhibitor of ALK, crizotinib, was rapidly approved by the US Food and Drug Administration on the basis of its pronounced clinical activity in patients with ALK rearrangement-positive NSCLC. Next-generation ALK inhibitors, such as alectinib, LDK378, and AP26113, are also being developed in ongoing clinical trials. In addition, the improvement and validation of methods for the detection of ALK rearrangement in NSCLC patients will be key to the optimal clinical use of ALK inhibitors. We here summarize recent progress in the development of new ALK inhibitors and in the molecular diagnosis of ALK rearrangement-positive NSCLC.

Keywords: ALK, rearrangement, NSCLC, ALK inhibitor, targeted therapy, diagnosis

Background
Lung cancer is the leading cause of cancer deaths worldwide. Non-small-cell lung cancer (NSCLC) accounts for 85% of lung cancer cases, and has usually achieved an advanced stage by the time of diagnosis.1 Cytotoxic chemotherapy has been the mainstay of treatment for metastatic NSCLC, but its efficacy has plateaued in recent years. Further improvement in the clinical outcome of individuals with NSCLC will thus depend on the development of new treatment strategies, such as molecularly targeted therapies. In 2004, the identification of activating mutations of the epidermal growth-factor receptor (EGFR) gene in a subset of NSCLC patients led to a change in treatment of the disease.2,3 Treatment of patients with NSCLC positive for EGFR mutations with such EGFR tyrosine-kinase inhibitors (TKIs) as gefitinib and erlotinib was found to have a high response rate and to result in both prolonged progression-free survival (PFS) and improved quality of life compared with cytotoxic chemotherapy.4,5 The discovery of EGFR mutations and the efficacy of EGFR TKIs in selected patients thus opened a new era of personalized treatment for NSCLC.

Anaplastic lymphoma kinase (ALK) is a receptor tyrosine kinase whose gene was initially identified in a subset of individuals with anaplastic large-cell lymphoma. A reciprocal translocation between chromosomes 2 and 5 apparent in such patients6 was found to result in the formation of a fusion gene comprising the 5′ portion of the nucleophosmin gene and the 3′ portion of ALK encoding the kinase domain.7 In 2007, a fusion gene formed by ALK and the echinoderm microtubule-associated
protein-like 4 (EML4) gene was identified in the tumor of a 62-year-old Japanese man with adenocarcinoma of the lung, and was shown to possess pronounced oncogenic activity. This genetic rearrangement has since been found to occur in 2%–5% of NSCLC patients, predominantly in those with adenocarcinoma who are of younger age and never- or light smokers.9,10

The EML4–ALK fusion oncogene arises from a small inversion within the short arm of chromosome 2 that joins the 5′ region of EML4 (encoding the NH2-terminal portion of EML4, including its coiled-coil domain) to the 3′ region of ALK (encoding the COOH-terminal portion of ALK, including the tyrosine-kinase domain). It exists in multiple variants that encode the same intracellular tyrosine-kinase domain of ALK but different truncations of EML4.11,12 The most common variants are variant 1 (detected in 33% of patients), in which exon 13 of EML4 is fused to exon 20 of ALK (E13;A20), and variant 3a/b (detected in 29% of patients), in which exon 6 of EML4 is fused to exon 20 of ALK (E6a/b;A20).12 Two other rare fusion partners of ALK (tyrosine-kinase receptor-fused gene and kinesin family member 5B) in addition to EML4 have also been identified in individuals with NSCLC.

All of these ALK fusion proteins undergo ligand-independent dimerization mediated by the coiled-coil domain of the fusion partner, resulting in constitutive activation of the ALK tyrosine kinase.13,14 Such phosphorylation-mediated activation of the ALK fusion proteins results in activation of downstream signaling pathways – including the JAK–STAT, MEK–ERK, and PI3K–AKT pathways – that contribute to oncogenicity.15–17 TKIs that target the kinase activity of ALK (ALK TKIs) have been found to have pronounced antiproliferative and proapoptotic effects in EML4–ALK-positive lung cancer cells.14,18

**Crizotinib**
The first clinically available ALK TKI

The structure of crizotinib is shown in Figure 1. Crizotinib is an oral and potent small-molecule ALK TKI that was initially designed as an inhibitor of the tyrosine kinase MET. Crizotinib competes with adenosine triphosphate for binding to the tyrosine kinase pocket of ALK and thereby inhibits its tyrosine-kinase activity, leading to inhibition of downstream signaling and to anticancer effects. Crizotinib exerts proapoptotic activity, with a median effective concentration

![Chemical structures and molecular weight (MW) of crizotinib, alectinib, LDK378, and AP26113.](https://www.dovepress.com/)

---

**Figure 1** Chemical structures and molecular weight (MW) of crizotinib, alectinib, LDK378, and AP26113.
Most adverse events of crizotinib treatment appear to be mild (grade 1 or 2), with those that occur most frequently being visual effects, nausea, diarrhea, constipation, vomiting, and peripheral edema. Three warning adverse events – interstitial lung disease (ILD), hepatotoxicity, and prolongation of the QT interval – have been identified. Life-threatening or fatal treatment-related ILD was found to occur in 1.6% of patients. It remains unclear whether the risk factors for EGFR TKI-associated ILD, such as male sex, a history of smoking, and coincidence of interstitial pneumonia, also apply to crizotinib-associated ILD. It is thus important that patients treated with crizotinib be monitored for pulmonary symptoms and radiographic findings indicative of ILD, and the drug should be discontinued immediately on such a diagnosis. Elevated serum aminotransferase levels of grade 3 or 4 were detected in ∼7% of crizotinib-treated patients, with such elevation usually being asymptomatic and reversible on discontinuation of crizotinib. Although crizotinib-induced hepatotoxicity with a fatal outcome has been reported in <1% of treated patients, routine evaluation of liver function, including measurement of aminotransferase and bilirubin levels, should be performed.

Mechanisms of crizotinib resistance

Although treatment with crizotinib has a pronounced clinical benefit for patients with ALK rearrangement-positive NSCLC, such individuals inevitably develop drug resistance. Several mechanisms of crizotinib resistance have been described to date, including secondary mutation or copy-number gain of ALK, inadequate drug delivery, and activation of alternative signaling pathways, such as those mediated by EGFR or KIT (Figure 3).

Two secondary mutations of ALK associated with crizotinib resistance – L1196M and C1156Y – were first detected in the same patient, who relapsed after achieving a partial response to the drug. The L1196M substitution occurs at the gatekeeper position of ALK (a position that controls the binding of nucleotides and TKIs), and corresponds to the T790M substitution in EGFR and the T315I substitution in the Bcr–Abl fusion protein, both of which confer resistance to corresponding TKIs. Multiple additional mutations in the ALK kinase domain have since been identified in patients who develop resistance to crizotinib. In contrast, T790M accounts for the vast majority of secondary mutations of EGFR that confer resistance to EGFR TKIs (Figure 3).
Although the relative contributions of the different mechanisms to crizotinib resistance remain unclear because of the small numbers of patients examined, biopsy of tumors performed after the onset of acquired resistance has suggested that secondary mutations in the ALK kinase domain account for only ∼30% of such cases of resistance.27,30 This situation also differs from that for EGFR mutation-positive NSCLC, for which the T790M substitution has been detected in up to 60% of tumors with acquired resistance to EGFR TKIs (Figure 3).36 Repeated biopsy and molecular analysis of relapsed tumors will be required in clinical trials of treatment strategies designed to overcome acquired resistance.

### Clinical development of other ALK TKIs

Several new ALK TKIs are currently under development (Table 1).

**Alectinib (CH5424802)**

Alectinib (Chugai Pharmaceutical, Tokyo, Japan) is a potent and selective ALK inhibitor with a median inhibitory concentration for ALK activity of 1.9 nM, and with little or no inhibitory activity for other protein kinases examined (Figure 1).37 The specific potency of alectinib for ALK inhibition appears to be related to its one-hinge hydrogen bond, whereas other ALK inhibitors, including crizotinib,

**Table 1** Selected clinical trials of anaplastic lymphoma kinase (ALK) inhibitors for the treatment of ALK rearrangement-positive non-small-cell lung cancer

<table>
<thead>
<tr>
<th>Drug</th>
<th>Phase</th>
<th>Compared drug</th>
<th>Treatment setting</th>
<th>Status</th>
<th>Clinical trial number</th>
</tr>
</thead>
<tbody>
<tr>
<td>Crizotinib</td>
<td>III</td>
<td>PEM or DOC</td>
<td>Second line</td>
<td>Published23</td>
<td>NCT00932893</td>
</tr>
<tr>
<td></td>
<td>III</td>
<td>Platinum + PEM</td>
<td>First line</td>
<td>Ongoing25</td>
<td>NCT01154140</td>
</tr>
<tr>
<td>Alectinib</td>
<td>i/ii</td>
<td>ALK inhibitor-naive</td>
<td>ALK inhibitor-naive</td>
<td>Published18</td>
<td>AF-001JP</td>
</tr>
<tr>
<td></td>
<td>III</td>
<td>Crizotinib</td>
<td>ALK inhibitor-naive</td>
<td>Ongoing26</td>
<td>JapiCTI-132316</td>
</tr>
<tr>
<td>LDK378</td>
<td>III</td>
<td>Platinum + PEM</td>
<td>First line</td>
<td>Ongoing27</td>
<td>NCT01828099</td>
</tr>
<tr>
<td></td>
<td>III</td>
<td>PEM or DOC</td>
<td>Both platinum and crizotinib failure, third line</td>
<td>Ongoing44</td>
<td>NCT01828112</td>
</tr>
<tr>
<td>AP26113</td>
<td>i/ii</td>
<td>ALK inhibitor-naive or failure</td>
<td>ALK inhibitor-naive or failure</td>
<td>Ongoing45</td>
<td>NCT01449461</td>
</tr>
<tr>
<td>ASP-3026</td>
<td>I</td>
<td>ALK inhibitor-naive or failure</td>
<td>ALK inhibitor-naive or failure</td>
<td>Ongoing46</td>
<td>NCT01401504</td>
</tr>
<tr>
<td>X-396</td>
<td>I</td>
<td>ALK inhibitor-naive or failure</td>
<td>ALK inhibitor-naive or failure</td>
<td>Ongoing47</td>
<td>NCT01625234</td>
</tr>
<tr>
<td>CEP-37440</td>
<td>I</td>
<td>ALK inhibitor-naive or failure</td>
<td>ALK inhibitor-naive or failure</td>
<td>Ongoing48</td>
<td>NCT01922752</td>
</tr>
</tbody>
</table>

**Note:** Patient accrual completed.

**Abbreviations:** PEM, pemetrexed; DOC, docetaxel.
form two- or three-hinge hydrogen bonds. In contrast to crizotinib, alectinib also shows substantial inhibitory activity against the L1196M mutant of ALK, apparently because it is able to maintain an efficient (CH/π) interaction with position 1196 even after the substitution of methionine for leucine.

A Phase I/II first-in-human study (AF-001JP) performed with previously treated and crizotinib-naive patients with ALK rearrangement-positive advanced NSCLC was performed in Japan. The participants were deemed to be ALK fusion gene-positive if a positive result was obtained either by reverse-transcription polymerase chain reaction (RT-PCR) analysis or by both immunohistochemistry (IHC) and fluorescence in situ hybridization (FISH). In the Phase I portion of the study, 24 patients received alectinib with a dose escalation from 20 to 300 mg bid, with the latter being determined as the highest planned dose on the basis of the available safety information for the additive formulation in Japan. Given that DLTs were not observed, the maximum tolerated dose (MTD) was not identified in this study. The highest planned dose (300 mg bid) was thus judged to be acceptable as the recommended dose for the 46 patients enrolled in the Phase II portion of the trial. Of these 46 patients, 43 individuals (93.5%) achieved an objective response, and 44 (95.7%) achieved disease control. The median PFS was not determined by the time of publication. This excellent clinical activity was associated with mostly mild adverse events, with those of grade 3 being detected in only 17 (37.0%) patients and those of grade 4 or death in none. The most frequently reported treatment-related adverse events were dysgeusia and liver dysfunction, both of which were of grade 1 or 2 in almost all cases. The characteristic adverse events of crizotinib treatment, including visual effects and gastrointestinal disorders (diarrhea, vomiting, and nausea), occurred at a low rate in this study of alectinib. Application for approval of alectinib in Japan was submitted on October 7, 2013.

A Phase III clinical trial (JapicCTI-132316) comparing alectinib with crizotinib in terms of PFS for the treatment of patients with ALK rearrangement-positive NSCLC is ongoing in Japan. Major eligibility criteria include advanced or metastatic ALK-rearranged NSCLC (identified either by RT-PCR or by both IHC and FISH), no prior treatment with an ALK inhibitor, an Eastern Cooperative Oncology Group performance status of 0–2, and either no previous treatment or one line of prior treatment with chemotherapy (Figure 4).

A dose-finding Phase I study (AF-002JG, NCT01588028) was also performed for alectinib in the US. Key eligibility criteria for this study included advanced NSCLC with ALK rearrangement confirmed by FISH, as well as failed crizotinib treatment. No treatment-related dose reductions were necessary up to a dose of 600 mg bid. Two of seven patients experienced DLTs (headache of grade 3, and neutropenia of grade 3 requiring dose-holding for 7 days) at the dose of 900 mg bid. On the basis of these results, 600 mg bid was determined as the recommended dose of alectinib for a Phase II study in the US. The ORR was 54.5% across all cohorts of the Phase I study, indicating that alectinib possesses significant clinical activity in ALK rearrangement-positive patients who are refractory to crizotinib. A global single-arm Phase II study of alectinib in patients with ALK-rearranged NSCLC resistant to crizotinib is ongoing (NCT01801111). The FDA granted breakthrough-therapy designation for alectinib on the basis of the NCT01588028 data, with early approval being expected.

LDK378

LDK378 (Novartis, Basel, Switzerland) is also a potent and selective small-molecule ALK inhibitor (Figure 1). In a Phase I study, 59 patients received LDK378 with dose escalation from 50 to 750 mg once daily (qd). DLTs were observed in two of the 14 patients who received the drug at 400 mg qd, in two of the nine patients at 600 mg qd, and in one of the nine patients at 750 mg qd. DLTs included diarrhea, vomiting, nausea, dehydration, and elevated serum aminotransferase levels. The MTD was thus defined as a dose of 750 mg qd. Among 88 evaluable ALK rearrangement-positive NSCLC patients who received LDK378 at 400–750 mg qd, the ORR was 70%. In the subset of 64 patients who had experienced crizotinib failure, the ORR was 73%. These results thus suggest that LDK378 may be effective for the treatment of patients with ALK-rearranged NSCLC who have developed acquired resistance to crizotinib.

A Phase III clinical trial (NCT01828112) comparing LDK378 with chemotherapy (pemetrexed at 500 mg/m² or
Crizotinib failure, LDK378 versus chemotherapy (NCT01828112)\textsuperscript{44}

- *ALK*-rearranged (FISH)
- Platinum and crizotinib failure, third line

Randomize

LDK378 750 mg qd po

Primary end point: PFS

Pemetrexed or docetaxel

First line, LDK378 versus chemotherapy (NCT01828099)\textsuperscript{45}

- *ALK*-positive (IHC)
- First line

Randomize

LDK378 750 mg qd po

Primary end point: PFS

Platinum + pemetrexed

---

**Figure 5** Ongoing Phase III studies of LDK378 for the treatment of *ALK* rearrangement-positive non-small-cell lung cancer.

**Abbreviations:** ALK, anaplastic lymphoma kinase; FISH, fluorescence in situ hybridization; IHC, immunohistochemistry; qd, once daily; po, oral administration; PFS, progression-free survival.

Docetaxel at 75 mg/m\textsuperscript{2} for the treatment of *ALK*-rearranged NSCLC patients who have progressed after prior treatment with both crizotinib- and platinum-based chemotherapy is ongoing (Figure 5).\textsuperscript{44} In addition, a Phase III clinical trial (NCT01828099) comparing LDK378 with standard first-line chemotherapy (pemetrexed plus either cisplatin or carboplatin) in previously untreated *ALK* positive NSCLC patients assessed by IHC is also ongoing (Figure 5).\textsuperscript{45}

**AP26113**

AP26113 (Ariad Pharmaceuticals, Inc., Cambridge, MA, USA) is another highly selective small-molecule ALK inhibitor that shows activity against the L1196M mutant (Figure 1).\textsuperscript{46} A Phase I/II study of AP26113 (NCT01449461) is ongoing.\textsuperscript{47} In the Phase I portion of the study, 44 patients received AP26113 with dose escalation from 30 to 300 mg qd. The most common adverse events were fatigue, nausea, and diarrhea, most of which were of grade 1 or 2. One DLT (increased serum alanine aminotransferase level of grade 3) was observed in one of nine patients treated at a dose of 240 mg, and one DLT (dyspnea of grade 4) was observed in one of two patients at a dose of 300 mg. Although the MTD has not been defined, a recommended Phase II dose was identified as 180 mg qd on the basis of safety, efficacy, and pharmacokinetic data. In the Phase I portion of the study, 24 patients with *ALK*-rearranged NSCLC were evaluable for response. Fifteen of these patients achieved an ORR of 63%, including 12 of the 16 individuals who had progressed after previous crizotinib therapy (ORR 75%). In addition, four of five patients showed objective responses for metastases in the central nervous system.

**Other new ALK TKIs**

Other new ALK TKIs, such as ASP-3026 (Astellas Pharma, Tokyo, Japan), NMS-E628 (Nerviano Medical Sciences, Milan, Italy), X-396 (Xcovery, West Palm Beach, FL, USA), CEP-37440 (Teva Pharmaceutical Industries Ltd, Petah Tikva, Israel), TSR-011 (Tesaro, Inc., Waltham, MA, USA), and PF-06463922 (Pfizer, New York, NY, USA), are currently introduced into clinical trials.

**Molecular diagnosis of ALK rearrangement-positive NSCLC**

**FISH**

Break-apart FISH analysis was applied for detection of *ALK* rearrangement in clinical trials with crizotinib. In this approach, the 5’ and 3’ portions of the *ALK* gene are separately...
labeled with red or green fluorescent probes (Figure 6). If the signals of the two probes overlap, resulting in yellow fluorescence, then there is no translocation. If a translocation is present, the two probes are spatially separated, and each is detected as an isolated signal (red or green). Tumors are deemed positive for ALK rearrangement if 15% or more of the tumor cells show isolated signals. Such analysis detects ALK rearrangement regardless of the ALK fusion partner or the specific EML4–ALK variant. The break-apart FISH assay is a unique diagnostic approach approved for screening for ALK rearrangement in NSCLC by the FDA. FISH has several disadvantages, however. First, it is an expensive and low-throughput method that requires technical expertise. Second, false-negative results sometimes occur because of difficulty in interpretation of separated signals. And third, the use of FISH alone (without IHC or RT-PCR) for screening may give rise to false-positive results. Indeed, in one study, the ORR for crizotinib was only 48% among patients screened with FISH alone, but increased up to 81% among those screened with FISH in combination with IHC or RT-PCR.48

Figure 6 Schematic illustration for break-apart fluorescence in situ hybridization for detecting ALK rearrangements. Abbreviation: ALK, anaplastic lymphoma kinase.

RT-PCR

RT-PCR is a highly sensitive and specific method for the identification of ALK rearrangement.60,61 In addition, unlike FISH or IHC, it can determine both the fusion partner of ALK (from among those previously identified) and the EML4–ALK variant.62 RT-PCR requires high-quality ribonucleic acid (RNA) extracted from nonfixed or freshly frozen specimens, however. It is generally difficult to extract suitable RNA from the paraffin-embedded specimens used in daily clinical practice.
Table 2 Summary of the sensitivity and specificity of improved immunohistochemistry (IHC) procedures for the detection of anaplastic lymphoma kinase (ALK) in non-small-cell lung cancer specimens

<table>
<thead>
<tr>
<th>Reference</th>
<th>FISH-positive cases</th>
<th>Antibody</th>
<th>Cutoff point of IHC score</th>
<th>Detection system</th>
</tr>
</thead>
<tbody>
<tr>
<td>51</td>
<td>43/132</td>
<td>D5F3</td>
<td>97.7</td>
<td>SignalStain® Boost IHC Detection</td>
</tr>
<tr>
<td>52</td>
<td>44/161</td>
<td>D5F3</td>
<td>NE</td>
<td>EnVision+</td>
</tr>
<tr>
<td>53</td>
<td>63/196</td>
<td>D5F3</td>
<td>NE</td>
<td>OptiView/OptiView Amplification</td>
</tr>
<tr>
<td>54</td>
<td>7/594</td>
<td>D5F3</td>
<td>92.8</td>
<td>EnVision FLEX+</td>
</tr>
<tr>
<td>55</td>
<td>15/352</td>
<td>5A4</td>
<td>NE</td>
<td>Polymer+ Detection Kit (Leica)</td>
</tr>
<tr>
<td>56</td>
<td>25/262</td>
<td>5A4</td>
<td>80.0</td>
<td>Polymer+ Detection Kit (Leica)</td>
</tr>
<tr>
<td>57</td>
<td>22/153</td>
<td>5A4</td>
<td>100</td>
<td>EnVision+</td>
</tr>
<tr>
<td>58</td>
<td>10/101</td>
<td>ALK1</td>
<td>90.0</td>
<td>ADVANCE (Dako)</td>
</tr>
<tr>
<td>59</td>
<td>20/351</td>
<td>D5F3</td>
<td>97.8</td>
<td>Polymer+ Detection Kit (Leica)</td>
</tr>
</tbody>
</table>


Abbreviations: FISH, fluorescence in situ hybridization; NE, not evaluated.

EGFR mutation-negative

↓

ALK test

IHC

RT-PCR

Negative

Positive

Negative

Positive

FISH

Chemotherapy

ALK inhibitor

Figure 7 Proposed algorithm for testing for ALK rearrangement in patients with non-small-cell lung cancer.

Abbreviations: ALK, anaplastic lymphoma kinase; IHC, immunohistochemistry; RT-PCR, reverse-transcription polymerase chain reaction; FISH, fluorescence in situ hybridization.
Several new RT-PCR-based methods have recently been developed. MassARRAY is a nucleic acid-analysis platform for the detection of EML4–ALK that involves PCR amplification, single-base primer extension, and analysis by MALDI-TOF (matrix-assisted laser desorption ionization–time of flight) mass spectrometry.63 The region of EML4–ALK complementary deoxyribonucleic acid containing the fusion point is amplified by PCR, but given that the amplicons are relatively small (70–130 bp), the quality of RNA extracted from paraffin-embedded specimens is sufficient for the analysis. RT-PCR-based assays may thus come to be more convenient and a major tool for detection of ALK rearrangement if the use of paraffin-embedded tissue is validated.

Future perspectives

The identification of the EML4–ALK fusion gene has accelerated translational research and changed clinical practice for NSCLC, with crizotinib now being in clinical use as an ALK inhibitor for the treatment of patients with ALK rearrangement-positive NSCLC. Although crizotinib has an excellent initial therapeutic effect, all treated patients eventually develop resistance to this drug. The development of therapeutic strategies able to overcome crizotinib resistance, including those based on the administration of new ALK inhibitors, is thus warranted. In addition, given the existence of other drivers of NSCLC, such as EGFR mutations as well as reactive oxygen species 1 and RET fusion genes, it will be important to improve and validate methods for the detection of and screening for these various genetic changes, so that the appropriate drug can be prescribed.

Disclosure

The authors report no conflicts of interest in this work.

References


Treatment and diagnosis of ALK-rearranged lung cancer


