Activity of Crizotinib in Patients with ALK-Aberrent Relapsed/Refractory Neuroblastoma: A Children’s Oncology Group Study (ADVL0912)

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ABSTRACT

Purpose: Anaplastic lymphoma kinase (ALK) aberrations are a promising target for patients with neuroblastoma. We assessed the activity of first-generation ALK inhibitor crizotinib in patients with no known curative treatments and whose tumors harbored an activating ALK alteration.

Patients and Methods: Twenty patients with relapsed/refractory ALK-positive neuroblastoma received crizotinib at the recommended phase II dose of 280 mg/m²/dose. A Simon two-stage design was used to evaluate the antitumor activity of crizotinib monotherapy. Response evaluation occurred after cycles 1, 3, 5, 7, and then every 3 cycles. Correlation of ALK status and response was a secondary aim of the study.

Results: The objective response rate for patients with neuroblastoma was 15% [95% confidence interval (CI): 3.3%–34.3%]: two with partial responses and 1 with a complete response. All three patients had a somatic ALK Arg1275Gln mutation, the most common ALK hotspot mutation observed in neuroblastoma and the only mutation predicted to be sensitive to ALK inhibition with crizotinib. Two patients had prolonged stable disease (10 and 13 cycles, respectively); both harbored an ALK Arg1275Gln mutation. Three patients with ALK Phe1174Leu mutations progressed during cycle 1 of therapy, and one patient with an ALK Phe1174Val received three cycles before disease progression. The two patients with ALK amplification had no response. The most common adverse event was a decrease in neutrophil count.

Conclusions: Despite limited activity seen in this trial, we conclude that this is more likely due to an inability to reach the higher concentrations of crizotinib needed to overcome the competing ATP affinity.

Introduction

Anaplastic lymphoma kinase (ALK) is a receptor tyrosine kinase (RTK) that is known to be oncogenically activated by point mutations within the tyrosine kinase domain, copy-number amplification, or chromosomal translocations (1–4). ALK mutations were first identified in both the germline of patients with familial neuroblastoma (4) and somatically in the sporadic form of the disease (1, 2, 4, 5). In addition, recurrent neuroblastoma demonstrates an increased frequency of ALK mutations (3, 6, 7). These mutations, as well as ALK amplifications (8), are a promising therapeutic target.

The discovery of ALK rearrangements in 3% to 5% of patients with non–small cell lung cancer (NSCLC; ref. 9) drove early-phase clinical studies of crizotinib, a first-in-class dual ALK/Met/ROS1 small-molecule tyrosine kinase inhibitor (10, 11). The dramatic response rates in NSCLC validated ALK as a therapeutic target and led to the expedited FDA approval of crizotinib in August 2011 for use in patients with ALK-rearranged lung cancer (12, 13). A pediatric phase I/II trial of crizotinib (ADVL0912, NCT00939770) established the recommended phase II dose (RP2D) as 280 mg/m²/dose twice daily (14). Pediatric patients with ALK fusions, most notably those with anaplastic large cell non-Hodgkin lymphoma (ALCL) and inflammatory myofibroblastic tumors (IMT), showed robust and durable complete antitumor responses in the majority of cases with proven ALK rearrangements (15). These data supported the portfolio of data required to obtain FDA breakthrough designation in May of 2018 for crizotinib in the context of relapsed ALCL (14, 15). This report describes the outcomes in children with relapsed or refractory ALK+ (activating mutation or amplification) neuroblastoma treated with crizotinib at the RP2D.

Patients and Methods

Patients

Patients >12 months and ≤ 21 years old who had relapsed or refractory ALK+ neuroblastoma were treated at the RP2D of 280 mg/m²/dose twice a day in cycles of 28-day duration. ALK status was reported by each enrolling institution using a Clinical Laboratory Improvement Amendments–certified FISH or DNA-sequencing...
Translational Relevance

Gain-of-function mutations in the ALK oncogene occur in 15% or more of newly diagnosed patients with high-risk neuroblastoma. This discovery positioned ALK as the first tractable molecular target for patients with this disease. However, crizotinib showed limited antitumor activity in this phase II trial for patients with relapsed ALK+ neuroblastoma. The preclinical mechanism underlying this observation revealed that two of the three hotspot mutations in ALK confer intrinsic resistance to crizotinib due to preferential affinity for ATP binding that could potentially be overcome by higher drug exposures. The observed responses occurred in patients with the most common germline and somatic hotspot mutations at residue R1275. Despite limited activity and lack of objective responses in patients harboring de novo resistant ALK mutations, we conclude that while this was possibly a limitation of the number of patients enrolled, this is more likely due to an inability to reach the higher concentrations of crizotinib needed to overcome the competing ATP affinity. Emerging data with the third-generation ALK inhibitor lorlatinib shows promise for patients with ALK-driven neuroblastoma.

Results

Patients

Twenty patients with relapsed/refractory ALK+ neuroblastoma, treated at the RP2D, were enrolled between March 2012 and July 2015; all were eligible and evaluable for response. Of the 20 patients, 6 patients were treated at the RP2D in phase I component of the study (14), and 14 patients were enrolled in the phase II expansion cohort. Patient characteristics are summarized in Table 1. All ALK mutations were somatic; no patients with germline mutations were enrolled on the expansion cohort of the phase I trial, cohort A2, and not previously reported. The borrowing of these patients with known ALK+ mutation status and treated at the RP2D was part of the study design, a strategy warranted when studying a rare subset of patients. The predefined criteria of at least one response among the stage I patients was met. The trial proceeded to stage II during which an additional 10 patients were entered. This 10+10 Simon two-stage design provides 88% power to reject the one-sided null hypothesis that the proportion of patients with a response is <0.05 in favor of the alternative hypothesis that the proportion is at least 0.25 with a type I error rate of 0.07. Correlation of the impact of ALK mutation or amplification status and response to crizotinib in patients with relapsed neuroblastoma known to harbor an ALK alteration was a secondary aim of the study.

Safety evaluation

The Common Terminology Criteria for Adverse Events version 4.0 was used to grade adverse events (AE). All patients who received at least one dose of crizotinib were evaluated. The relative frequency of each AE considered possibly, probably, or likely related to crizotinib was estimated as the proportion of all toxicity-evaluable cycles in which such toxicity was observed.

Pharmacokinetic studies

Plasma samples for pharmacokinetics were obtained at steady state and analyzed as described previously (16).

Statistical analyses

Participant demographics and clinical characteristics were summarized by ALK status. The percent of responders (CR or PR) with 95% confidence interval was estimated using methods described by Koyama and Chen (18).

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Toxicity

The most common grade 3 or 4 drug-related adverse event were hematologic and predominantly manifested as a decrease in neutrophil count, with one dose-limiting toxicity among all 20 patients (Supple-

Table 1. Characteristics of the patients at baseline.

<table>
<thead>
<tr>
<th>Characteristics</th>
<th>Value</th>
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<tbody>
<tr>
<td>Age at enrollment (years)</td>
<td>Median (min, max)</td>
</tr>
<tr>
<td></td>
<td>Female</td>
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<tr>
<td></td>
<td>Male</td>
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<tr>
<td>Race</td>
<td>White</td>
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<td>Black</td>
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<td>Asian</td>
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<td>Unknown</td>
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<tr>
<td>No. of prior therapies</td>
<td>1–2</td>
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<td></td>
<td>3–4</td>
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<td></td>
<td>5–6</td>
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<td>7+</td>
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<td>No. of patients with at least one prior therapy</td>
<td>Chemotherapy</td>
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<td></td>
<td>Radiotherapy</td>
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<td>Transplant</td>
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<td>Immunotherapy</td>
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<td></td>
<td>Prior therapy NOS</td>
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<tr>
<td>Disease status</td>
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<td>Measurable</td>
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<tr>
<td>Cohort</td>
<td>Phase I</td>
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<td>Phase II</td>
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<tr>
<td>ALK Aberration</td>
<td>Amplification</td>
</tr>
<tr>
<td></td>
<td>Arg1275Gln</td>
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<tr>
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<td>Arg1275Gln;Phel174Leu</td>
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<td></td>
<td>Phel174Leu</td>
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<tr>
<td></td>
<td>Phel174Val</td>
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<tr>
<td></td>
<td>Tyr1278Ser</td>
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</tbody>
</table>

Pharmacokinetics

Pharmacokinetic analysis was available for 13 of 20 patients, with a median age of 6 years (range, 2–14 years). The median steady-state concentration of crizotinib was 665 ng/mL (range, 298–1,040 ng/mL). Of the three patients with a response (1 CR, 2 PR), one patient with a PR had steady-state concentration data available (923 ng/mL). These values are similar to those previously reported from the phase I component of the clinical trial (16).

Discussion

Survival rates for the childhood cancer neuroblastoma have improved incrementally from 25% to 50% over three decades with dramatic escalation in therapy intensity. We and others discovered in 2008 that activating mutations in the tyrosine kinase domain of ALK are the cause of hereditary neuroblastoma, and that these mutations can also be somatically acquired (1, 2, 4, 5). This work drove the early-phase clinical trial of crizotinib in patients with relapsed/
refractory pediatric tumors, with emphasis on enrollment of ALK-driven cancers. ADVL0912 (NCT00939770) opened to accrual within 18 months of the initial discovery of recurrent ALK mutations in a neuroblastoma, thus rapidly translating these seminal preclinical molecular findings into the clinic. Notably, this is the first clinical demonstration that ALK is a tractable molecular target in neuroblastoma, as well as demonstration that the preclinical cellular, biochemical, and in vivo data accurately predicted the differential sensitivity of the ALK R1275 versus F1174 mutations, albeit the small number of patients treated.

Early preclinical studies showed that crizotinib had activity against neuroblastoma cell lines and xenografts harboring activating ALK mutations, with mutations at two of the three hotspots demonstrating intrinsic resistance to ALK kinase inhibition with crizotinib (19, 20). The relative resistance in F1174- and F1245-ALK neuroblastoma results from an increase in ATP-binding affinity (20) and less binding of crizotinib to the target. Preclinical experiments demonstrate that neuroblastomas harboring an F1174-ALK mutation may benefit from ATP-competitive ALK inhibitors such as crizotinib if an increase in dosage to overcome the relative affinity for ATP compared with R1275-mutated ALK was possible (19, 20). In phase I, we maximized the dose of crizotinib in pediatric patients to achieve higher plasma concentrations to overcome the increased ATP-binding affinity of de novo resistant ALK mutations. We ultimately defined a crizotinib RP2D in children that is nearly 200% of the adult MTD (14).

In the phase I portion, we previously reported 5 patients with neuroblastoma and known ALK mutations treated at dose levels other than the RP2D (100–365 mg/m²/dose twice daily), and 6 patients treated at the RP2D who are included in the phase II trial, 3 previously reported without response and 3 not previously reported upon. Of these patients, one had a CR and had a germline Arg1275Gln mutation (treated above the RP2D at 365 mg/m²/dose); two had prolonged stable disease at dose levels below the RP2D, one of whom had a germline Arg1275Gln mutation and resolution of MIBG-avid disease (14). The other three borrowed patients treated in the phase I trial at the RP2D have not been previously reported; they were enrolled on the A2 expanded cohort for patients with confirmed ALK alterations that could enroll at any time. In the phase II portion of the trial, where crizotinib was administered orally twice daily as monotherapy at the RP2D of 280 mg/m²/dose twice a day, 3 of 12 patients with an activating Arg1275Gln ALK mutation had objective responses (1 CR, 2 PR), 2 had prolonged stable disease, and 6 of the remaining 7 patients had early disease progression (Fig. 1; Table 2). No responses were seen in patients harboring ALK F1174 mutations or ALK amplifications.

At the 280 mg/m² dose level in the phase I portion of ADVL0912 (NCT00939770), the steady-state average plasma concentration (Cave) was 580 ± 173 ng/mL or 2,920 ± 380 nmol/L (mean ± SD; ref. 16). The protein binding of crizotinib is 90%. Therefore, the Cave steady-state free-drug concentration is estimated to be 129 nmol/L, which is equivalent to the IC50 for F1174 L ALK mutation (130 ± 10 nmol/L) and only marginally higher than the IC50 for R1275Q ALK mutation (85 ± 8 nmol/L). This suggests that 50% inhibition of ALK mutations may be insufficient to produce a therapeutic effect. (19, 20) In the phase II component of the study, the mean steady-state Cave was variable, ranging from 298 ng/mL (660 nmol/L) to 1,040 ng/mL (2,310 nmol/L); one patient with a PR had a steady-state concentration of 923 ng/mL (2,050 nmol/L). Pharmacokinetic data were not available for the other 2 patients who had a response. Thus, given the protein binding of crizotinib (90%), it appears likely that the free drug concentrations of crizotinib achieved in plasma even at the 280 mg/m² dose level, may be insufficient to inhibit ALK kinase activity. Early on in the era of precision medicine, blockade of BRAF-mutant melanoma, using tyrosine kinase inhibitors demonstrated that complete pathway inhibition is necessary for objective antitumor activity (21). Our preclinical data suggest the same is true for ALK-aberrant neuroblastomas and that the reduced sensitivity to crizotinib is caused by a heightened ATP-binding affinity in tumors with an F1174 of F1245 mutation. In the context of neuroblastoma-specific ALK mutations, crizotinib is likely just good enough to have some effect for only the R1275 mutations because ATP binding is weaker in this context (20), and
without complete abrogation of ALK signaling, the remaining kinase activity is likely sufficient to maintain tumor growth, even in the setting of an R1275 mutation.

In summary, crizotinib is active against a subset of Arg1275Gln ALK–mutated neuroblastomas, with possible prolonged utility in the setting of a germline mutation, a context where ALK is likely the initiating truncal mutation in the disease; however, crizotinib has no objective single-agent activity in neuroblastomas harboring other hotspot ALK mutations or amplification. High-risk neuroblastomas are characterized by extensive intratumoral and stroma-derived heterogeneity and harbor both preexisting and acquired subclonal populations, as well as spatial intratumoral heterogeneity and temporal clonal evolution, that confer therapy resistance, rendering early-phase studies challenging, even in the context of exploiting an oncogenic driver. We have shown that crizotinib in combination with chemotherapy can overcome the intrinsic resistance of ALK hotspot mutations by a synergistic interaction through the intact p53 pathway (22). These data provided the preclinical rationale for the ongoing COG phase III trial ANBL1531 (NCT03126916).

As the field has moved forward, lorlatinib has been identified as a highly specific and potent third-generation ALK inhibitor that overcomes de novo resistance and exerts unprecedented activity in neuroblastoma patient-derived xenografts harboring all hotspot mutations (23). Lorlatinib is now completing phase I (NCT03107988) evaluation in children and adults with relapsed/refractory ALK-driven neuroblastoma, and has confirmed antineuroblastoma activity with minimal observed toxicity (24). Lorlatinib will be studied in the upfront setting for patients with ALK-aberrant neuroblastoma, replacing crizotinib in the current phase III trial (NCT03126916) for patients with high-risk neuroblastoma. Thus, ALK inhibition is an important strategy for improving the outcomes of children with neuroblastoma.

As we learn more about how cancers evolve under the selective pressure of antitumor therapy, we are recognizing that this does not occur through a linear stepwise selection of genetic changes. Genomic alterations occurring during neuroblastoma progression and the resulting genomic heterogeneity and mutational diversity between intrapatient tissues remain poorly understood, but it is likely that ALK driver mutations can occur as a branched or truncal mutation, as seen by the increased frequency of ALK mutations at relapse (3, 6, 7), whereas MYCN amplification is always a truncal event and completely shared between lesions. The overall effect of these factors on the

**Figure 2.** Representative responses to crizotinib monotherapy in patients with activating Arg1275Gln ALK mutation. **A**, Patient 114. Diagnostic CT (top row) and [131I]-MIBG SPECT/CT images (bottom row) show response of an enlarged right aortocaval lymph node (arrows) to crizotinib therapy. Images obtained at baseline and after 5 and 13 cycles of therapy, respectively, show a partial response at RP5, and a complete response at RP13, with no residual lymphadenopathy or MIBG uptake at RP13. A similar pattern of response was also shown for a separate site of retroperitoneal disease (not shown). **B**, Patient 121. Whole-body MIBG planar images (outer panels), axial (inner left panels), and sagittal (inner right panels) fat-suppressed T2-weighted MRI images show partial response of measurable disease by MRI and MIBG-avid evaluable disease to crizotinib therapy. The presacral mass present at baseline (BL, arrow) has decreased significantly in size, with only a small focus of measurable disease (circle) remaining by RP10. Multiple sites of MIBG avid disease, including the presacral soft tissue mass, left parietal skull, entire vertebral column, proximal humeri, bony pelvis, proximal femurs, distal left femur, and proximal left tibia, have either resolved (left parietal skull) or are significantly decreased in signal intensity, with only faint uptake remaining at sites of baseline MIBG-avid disease.
efficacy of treatment is daunting; however, multiregional sequencing, single-cell sequencing, and serial profiling of plasma circulating tumor DNA are all emerging technologies that will allow us to more precisely dissect the clonal complexity and spatiotemporal genomic heterogeneity to develop more effective personalized therapies.

**Authors' Disclosures**

K. Wilner reports other from Pfizer Inc outside the submitted work. S.L. Berg reports grants from Children’s Oncology Group during the conduct of the study. P.C. Adamson reports grants from NCi and other from Pfizer during the conduct of the study. Y.P. Moss reports grants from Pfizer, UMI CA097452, UMI CA228823, Cookies for Kids’ Cancer Foundation, and Children’s Oncology Group Foundation during the conduct of the study, as well as personal fees from Pfizer outside the submitted work; in addition, Y.P. Moss has a patent for 14/607,573 licensed to The Children’s Hospital of Philadelphia. No disclosures were reported by the other authors.

**Authors’ Contributions**

J.H. Foster: Data curation, formal analysis, writing–original draft, writing–review and editing. S.D. Voss: Data curation, software, formal analysis, methodology, writing–original draft, writing–review and editing. D.C. Hall: Data curation, formal analysis, visualization, writing–original draft, writing–review and editing. C.G. Minard: Data curation, software, formal analysis. F.M. Balis: Data curation, methodology, writing–original draft, writing–review and editing. K. Wilner: Conceptualization, resources, writing–review and editing. S.L. Berg: Data curation, investigation, writing–review and editing. E. Fox: Data curation, formal analysis, funding acquisition, writing–review and editing. P.C. Adamson: Conceptualization, writing–original draft, writing–review and editing. S.M. Blaney: Conceptualization, data curation, methodology, writing–review and editing. B.J. Weigelt: Conceptualization, data curation, investigation, methodology, writing–original draft, writing–review and editing. Y.P. Moss: Conceptualization, resources, data curation, formal analysis, supervision, funding acquisition, validation, investigation, visualization, methodology, writing–original draft, project administration, writing–review and editing.

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