Novel Molecular Challenges in Targeting Anaplastic Lymphoma Kinase in ALK-Expressing Human Cancers

Abdulaheem Alshareef 1,2

1 Department of Clinical Laboratory Sciences, College of Applied Medical Sciences, Taibah University, Almedinah, Medina P.O. Box 41477, Saudi Arabia; amshareef@taibahu.edu.sa or al15@ualberta.ca
2 Department of Laboratory Medicine and Pathology, University of Alberta, Edmonton, AB T6G 2E1, Canada

Received: 26 August 2017; Accepted: 24 October 2017; Published: 28 October 2017

Abstract: Targeting anaplastic lymphoma kinase (ALK), a receptor tyrosine kinase receptor initially identified as a potent oncogenic driver in anaplastic large-cell lymphoma (ALCL) in the form of nucleophosmin (NPM)-ALK fusion protein, using tyrosine kinase inhibitors has shown to be a promising therapeutic approach for ALK-expressing tumors. However, clinical resistance to ALK inhibitors invariably occurs, and the molecular mechanisms are incompletely understood. Recent studies have clearly shown that clinical resistance to ALK inhibitors is a multifactorial and complex mechanism. While few of the mechanisms of clinical resistance to ALK inhibitors such as gene mutation are well known, there are others that are not well covered. In this review, the molecular mechanisms of cancer stem cells in mediating resistance to ALK inhibitors as well as the current understanding of the molecular challenges in targeting ALK in ALK-expressing human cancers will be discussed.

Keywords: anaplastic lymphoma kinase; ALK-expressing cancers; cancer stem cells; crizotinib; clinical resistance; tyrosine kinases; tyrosine kinase inhibitors

1. Introduction

Tyrosine kinases, such as ALK, are a very attractive therapeutic target for cancer treatment, especially on the basis of promising results from preclinical and early clinical studies [1]. Anaplastic lymphoma kinase (ALK) is a receptor tyrosine kinase that was initially discovered and characterized in a rare type of lymphoma called anaplastic large-cell lymphoma (ALCL) as an NPM-ALK fusion protein [2]. Specifically, the catalytic domain of the ALK protein was fused with the amino terminus of nucleophosmin (NPM), and it was found that the NPM-ALK fusion protein resulted in the constitutive activation of the ALK tyrosine kinase, thereby leading to deregulation of multiple cell signaling pathways and increased tumorigenicity (Figure 1) [3–5]. Subsequent studies of ALCL and other types of human cancer have revealed additional fusion partners of ALK and various types of ALK gene aberrations [6,7]. For example, the echinoderm microtubule-associated protein like 4 (EML4)-ALK fusion was identified in ~5% of non-small cell lung cancers (NSCLC) [8,9]. Amplified ALK or mutated ALK was identified in ~14% of neuroblastomas (NB), the most common and aggressive childhood malignancy [10–13]. To date, several ALK inhibitors are at various stages of clinical testing and the US Food and Drug Administration (FDA) [1]. Although most clinical results regarding ALK inhibitors are from patients with ALK-positive non–small-cell lung carcinoma (NSCLC), it is clear from preclinical studies that ALK inhibition is effective in all ALK-expressing cancers [14].
The ALK protein interacts and activates many essential adaptors involved in multiple signaling pathways, including PI3K, RAS/MEK/ERK, β-catenin, and JAK/STATs. Only four representative signaling pathways are shown here. EML4-ALK: echinoderm microtubule-associated protein like 4-anaplastic lymphoma kinase; NPM-ALK: Nucleophosmin-anaplastic lymphoma kinase; STAT: Signal transducer and activator of transcription; PI3K: phosphatidylinositol 3 kinase; ERK: extracellular signal-related kinase; JAK3: Janus kinase 3; Bcl2: B-cell lymphoma 2; Mcl1: Myeloid cell lymphoma 1; BAD: Bcl2-associated death promoter; mTOR: mammalian target of rapamycin; MEK: MAPK (Mitogen-activated protein kinase)/ERK (extracellular signal-regulated kinase); Sox2: (sex determining region Y)-box 2.

The data collected from clinical studies, especially for crizotinib (the first ALK inhibitor used in the clinic), were extremely promising [1]. In ALK+ NSCLC, for instance, comparing crizotinib with standard chemotherapy in the second-line setting resulted in an improved overall response rate (65% vs. 20%, respectively), a shorter response time (6.3 vs. 12.6 weeks), and an improved median progression-free survival (7.7 vs. 3.0 months) with crizotinib [15]. In ALK+ ALCL patients, crizotinib was administered to seven adults with resistant high-stage disease and resulted in a complete response (CR) in three patients and a partial response in one patient [16]. This later study was expanded and had a total of 11 patients (9 with ALCL) and a CR was observed in all 9 patients [17]. Furthermore, the Children’s Oncology Group-sponsored Phase 1 clinical trial (NCT00939770) with crizotinib in children with refractory ALK+ ALCL resulted in a CR in eight of the nine patients [18]. This Phase 1 clinical trial included 34 NB patients with recurrent or refractory cancer, and showed a wide range sensitivity to ALK kinase inhibition [18]. Specifically, only 2 out of 34 (6%) patients showed complete remission, 8 (23.5%) showed stable disease while 24 (71%) showed progressive disease [18].

Resistance to ALK inhibitors, including even second- or third-generation drugs used as a single therapy, is a ubiquitous problem in ALK-expressing cell lines as well as treated patients (Table 1 and Figure 2) [1]. Resistance to crizotinib, for instance, was initially reported in NSCLC [15,19] and inflammatory myofibroblastic tumors [20], followed by NB [18] and ALCL [17]. Previous reports have generally suggested two categories of mechanisms of resistance: (1) resistance mediated by mutations in the ALK kinase domain impairing binding of an inhibitor to an ALK protein, and/or (2) the activation of compensatory alternative oncogenic drivers such as MET, epidermal growth factor receptor (EGFR), KRAS, and c-KIT [1]. However, there is a lack of knowledge on the molecular basis of this resistance. In other words, almost all of the previous studies have focused on acquired resistance (which is caused by post-treatment changes such as alteration in drug targets and the activation of compensatory survival signaling pathways), while knowledge on intrinsic resistance (which includes the factors that exist
Resistance to targeted therapy has been reported to be mediated through multiple mechanisms [1]. Some of these mechanisms are known to exist among almost all tyrosine kinases, including ALK. For example, gene amplification, gene mutation, and upregulation of alternative signaling pathways; all of which have been shown to induce drug resistance [22,23,40]. Unlike ALK, there are several reported ways to overcome resistance to ALK inhibitors.

### Table 1. Summary of first and next generation ALK inhibitors.

| ALK Inhibitor | Other Names | FDA Approval (Month/Year) | Resistance Occurred [Reference] | Ways to Overcome Resistance [Reference] |
|---------------|-------------|---------------------------|---------------------------------|-----------------------------------------|
| Crizotinib    | PF-2341066 Xalkori® | Yes (08/2011) | Yes [15,23] | 1. Targeting PI3K/AKT/mTOR pathway [25,29] |
|               |             |                           |                                 | 2. Targeting Src [30]                     |
|               |             |                           |                                 | 3. Targeting HSP90 [31]                   |
|               |             |                           |                                 | 4. Targeting β-catenin [32]               |
| Ceritinib     | LDK-378 Zycadia® | Yes (04/2014) | Yes [33,34] | Alectinib [33,34] | Not performed |
| Alectinib     | RO5424802 Alectina® | Yes (12/2015) | Yes [35,36] | Ceritinib [35,37] | Not performed |
| Brigatinib    | AP26113 Alunbrig® | Yes (04/2017) | Yes [38] | - | Not performed |
| Lorlatinib    | PF-06463922 | No | Yes [39] | Crizotinib [39] | Not performed |

ALK: Anaplastic lymphoma kinase; FDA: Food and Drug Administration; PI3K: Phosphoinositide 3-kinase; mTOR: mammalian target of rapamycin; HSP90: heat shock protein 90.

**Figure 2.** The current active approaches to overcome resistance to ALK inhibitors. The most common approach mainly relies on second and third generation ALK inhibitors such as ceritinib, alectinib, and brigatinib. The less common approach relies on re-sensitizing resistant cells to ALK inhibitors by targeting other signaling pathways. X represents the inhibitory effect of the ALK inhibitor. Green triangle represents the addition of another ALK inhibitor. PI3K: Phosphoinositide 3-kinase; HSP90: heat shock protein 90.

### 2. Reported Mechanisms of Resistance

Resistance to targeted therapy has been reported to be mediated through multiple mechanisms [1]. Some of these mechanisms are known to exist among almost all tyrosine kinases, including ALK. For example, gene amplification, gene mutation, and upregulation of alternative signaling pathways; all of which have been shown to induce drug resistance [22,23,40]. Unlike ALK, there are several
mechanisms of resistance that have been reported in EGFR-expressing lung cancer cells and BCR-ABL (breakpoint cluster region-abelson)-expressing chronic myeloid leukemia (CML) cells. The existence of cancer stem cells (CSCs) is one of these mechanisms. For instance, stem cell population in lung cancer cells were shown to be relatively resistant to gefitinib, a tyrosine kinase inhibitor showing specificity to the epidermal growth factor receptor (EGFR) [41]. One study showed that acquired resistance to gefitinib was associated with a manifestation of stem cell–like properties in cancer cells [42]. Similarly, resistance to imatinib, a tyrosine kinase inhibitor showing specificity to BCR-ABL, has been well documented in leukemic stem cells.

Mechanistically, leukemic stem cells have been shown to be attributed to imatinib resistance through three independent mechanisms. First, leukemic stem cells were found to express a relatively high level of BCR-ABL [43,44]. Second, leukemic stem cells were found to be inefficient in maintaining the intracellular accumulation of imatinib, mainly due to the relatively low expression of the organic cation transporter-1 (OCT1), which is responsible for the cellular uptake of imatinib [45], as well as the relatively high expression of ABCB1 (ATP Binding Cassette Subfamily B Member 1) that mediates the efflux of imatinib [46]. Third, imatinib resistance was shown to be mediated by the activation of alternative signaling pathways such as Mitogen-activated protein kinase (MAPK), Notch, and hedgehog that maintain viability and growth despite continued suppression of BCR-ABL kinase activity [47–50].

The other mechanism of resistance that has been previously reported is histologic transformation which includes epithelial mesenchymal transition (EMT) and small cell transformation [40,51]. For instance, many studies have reported that histologic transformation, mainly EMT, could induce cancer cell resistance to EGFR-tyrosine kinase inhibitors (TKIs) [52–55]. Specifically, the transformation to SCLC as acquired resistance mechanism is observed in 14% of TKI-treated EGFR mutant pulmonary adenocarcinomas [56–58]. In ALK(F1174L)-driven NB cells, ALK inhibitor resistance was found to be associated with the induction of EMT [59]. Importantly, co-targeting EGFR (in lung cancer) or ALK (in NB) and the molecules that drive these pathways can reverse TKIs resistance [59–62]. A more recent study has suggested that EMT is only associated with, but does not drive resistance to, ALK inhibitors among EML4-ALK+ NSCLC [63].

Furthermore, it has been reported that due to a heterogeneous population of cancer cells, some of the population are not fully dependent on the activity of tyrosine kinases for their survival, which may be the reason behind the insensitivity of these cells to TKIs [64]. For instance, leukemia stem cells have been shown to utilize signaling pathways independent of BCR-ABL kinase activity for their maintenance and survival, and that these cells were insensitive to TKIs such as imatinib or dasatinib [65,66].

3. Role of Intra-Tumoral Heterogeneity in Dictating Resistance to ALK Inhibitors in ALK-Expressing Cancers

As mentioned previously, CSCs were shown to be a major contributing factor to disease relapses upon TKIs treatment. Oh et al., (2015) showed that targeting stemness with rapamycin, an mTOR inhibitor, synergized the crizotinib effect in EML4-ALK+ cells (lung cancer) in vitro and in vivo [67]. Additionally, they showed that the rapamycin treatment sensitized the crizotinib-resistant cell line to ALK inhibition [67]. A more recent study showed the synergistic effect of ALK and mTOR inhibitors in the treatment of NPM-ALK+ cells (ALCL) [68]. Using an mTOR inhibitor, namely Torin2, which is a selective mTORC1 inhibitor, was also shown to restore sensitivity to crizotinib in ALK-mutated (ALK(F1174L)) NB cells [28,69]. These studies have shed light into the importance of CSCs in ALK-expressing cancer cells; however, they did not mechanistically study CSCs as none of these studies purified and studied CSCs.

In our recent study, we found that CSCs derived from NB cells were significantly more resistant to crizotinib (under review). Of note, NB CSCs were purified based on their responsiveness to a Sox2 reporter, a strategy that has been used previously for several different cancer models [70–75]. Importantly, we concluded that the crizotinib resistant phenotype in CSCs can be attributed to their high β-catenin
expression since siRNA knockdown of β-catenin sensitizes CSCs to the crizotinib treatment. Our data suggested that combining β-catenin inhibitors and ALK inhibitors may be useful in treating NB patients. To the best of our knowledge, this latter study is the only study, to date, that has investigated the role of CSCs as a contributing factor to treatment failure and disease relapses to ALK inhibitors. Further studies need to be performed to study the CSCs and their role in mediating ALK TKIs resistance.

4. Role of ALK-Interacting Proteins in Mediating TKIs Resistance

Tyrosine kinases are known to bind to a large number of cellular proteins, thereby mediating their oncogenic effect [4]. To date, there is little known about whether some of these binding proteins might play a role in modulating resistance to TKIs. For instance, targeting β-catenin has been shown to cause an abrogation of tyrosine kinase resistance in the BCR-ABL CML model [76]. In a lung cancer model, inhibition of β-catenin was shown to enhance the anticancer effect of EGFR-TKI in EGFR-mutated cells [77]. Of note, β-catenin is the central mediator of Wnt/β-catenin signaling, and it can be localized either in the adherens junctions and is involved in cell–cell contacts, or in the nucleus where it is implicated in transcriptional regulation and chromatin modification [78]. The mechanism of how β-catenin mediates resistance to these TKIs.

Work from our laboratory using the NPM-ALK+ ALCL and ALK+ NB cells showed a physical interaction between ALK and β-catenin [32,79]. Computational analysis could provide a starting point for more in-depth mechanistic understanding and explanation for the reason behind ALK TKIs resistance and predicting the proteins that could mediate the blockage of TKI-ALK binding. To date, the fate of ALK—β-catenin interaction and other ALK-interacting proteins is still minimally understood, and requires more mechanistic studies.

5. CETSA as a Tool That Can Be Used to Predict the Resistance to ALK Inhibitors

The current read-out used to measure the effect of ALK inhibitors focuses on phenotypic assays where the response to an inhibitor is based on a functional readout such as changes in the phosphorylation status of downstream targets, or the impact on cellular viability [80]. While these functional readouts are very useful, they do not provide sufficient information regarding the resistance. We have recently shown that CETSA (Cellular Thermal Shift Assay), a recently described method that allows for the rapid and simple assessment of drug target engagement in a cellular context [81–83], is useful in predicting crizotinib sensitivity in ALK-carrying cancer cells [32]. Previous studies have shown that CETSA is as an excellent tool to evaluate the physical binding of an inhibitor to its target in intact cells [81–83]. A few studies have used the CETSA assay to evaluate crizotinib treatment [32,84,85]. The first report assessed the photosensitivity side effect of many kinase inhibitors, including crizotinib, on K562 (a BCR-ABL+ CML cell line) by combining the CETSA method with multiplexed quantitative mass spectrometry (MS) [84]. In the second report, the authors demonstrated a bond between the (S)-enantiomer crizotinib (which is not the clinically used ALK inhibitor; (R)-enantiomer) and MTH1 (MutT Homolog 1) and thereby worked as a suppressor of MTH1 activity [85]. In the third study, the authors identified a significant positive correlation between crizotinib-ALK binding and the observed IC₅₀, which provides a logical justification for the differential responsiveness [32]. Additionally, it proved that the CETSA assay was a very useful tool to predict crizotinib sensitivity in different ALK-carrying cancer types.

CETSA has increased in popularity as a tool to validate drug–target interaction [86,87]. For example, the proposed PARP-1 (Poly [ADP-ribose] polymerase-1) inhibitor iniparib reached Phase III clinical trials where it showed no efficacy, and was subsequently shown to lack activity against PARP-1 in living cells [86,87]. CETSA was used to compare the target engagement of PARP-1 for iniparib and olaparib [81], which is a well-established PARP-1 inhibitor in clinical development. Recently, the CETSA assay was used to assess the binding of these two PARP-1 inhibitors and showed that iniparib failed to induce a thermal shift, whereas olaparib binding induced a large thermal shift of PARP-1 [81]. Apparently, the mechanism of action of iniparib is not via physical binding to PARP-1;
instead, iniparib may kill cancer cells by unspecific modification of cysteine residues. All these reports highlight the significance of implementing CETSA as a tool that can prove the TKI binding to its target and thereby predict resistance in early stages of treatment.

6. Potential Significance of Precursor mRNA in Mediating Resistance to ALK TKIs

Cancer cells are generally known to develop multiple mechanisms to escape the signaling inhibition caused by TKI treatment. While the main mechanisms have been discussed earlier in this review, we recently postulated an additional mechanism where an ALK-intron retained transcript was detected in NB cell lines as well as patient samples (under review). This observation poses some intriguing questions with regard to the properties of ALK as an oncogene. In our hypothetical model, the ALK-intron 19 (ALK-I19) transcript was the final precursor to the fully spliced (FS)-ALK transcript, whose high expression is fundamental for sustaining NB growth and proliferation. Perhaps due to this inherent dependency on ALK, many NB cells may pre-synthesize the ALK-I19 transcript within the nucleus as a short-term storage system to bolster FS-ALK expression when it is critically required. This mechanism may be especially useful to the cancer cells to maintain homeostasis. In conditions of cellular stress, such as in normal tumor physiology like hypoxia and upon growth inhibiting drug treatments (i.e., chemotherapy), ALK-I19 may increase FS-ALK levels. A comparable stress-induced mechanism was identified for the ApoE gene in central nervous system neurons [88]. Upon injury of the neuronal cells, an intron retained transcript of ApoE pre-synthesized in the nucleus enhanced the cytoplasmic level of fully spliced ApoE [88]. This mechanism leads to the immediate production of proteins upon cellular stress. This phenomenon of a ‘buffer’ transcript could be relevant given the vital importance of ALK activity in many cancer types.

What molecular factors are responsible for this phenomenon? Answering this important question could aid the development of more effective therapeutics for ALK+ patients. For example, Intron 4-retaining CCND1 (cyclinD1) expressed in prostate and esophageal cancers was found to translate into a truncated cyclin D1 protein, which has oncogenic effects [89,90]. Indeed, RNA binding proteins, many of which participate in specific splicing complexes, are deregulated in cancers [91]. This disruption can lead to aberrant alternative splicing and boost tumorigenesis. Therefore, the examination of regulatory protein-networks involved in ALK protein synthesis and its role in mediating resistance to ALK TKIs is warranted.

7. Conclusions

This review presented data describing mechanisms of resistance to ALK TKI treatment and clearly showed that it is a multifactorial and complex mechanism (Table 1). There are at least four reasons supporting the notion that the mechanisms of resistance to ALK inhibitors are not exclusively attributed to one or two factors (e.g., specific ALK mutation). First, studies performed on crizotinib upon its discovery showed that a high concentration of crizotinib (defined as >300 nM) displayed off-target effects [92]; and the IC_{50} for U937, a histiocytic lymphoma cell line used as a negative control (as it expresses neither ALK nor c-Met) was 257 nM [93]. Second, NB cell lines carrying resistant ALK mutations such as ALK-F1174L, displayed drastically different IC_{50} to crizotinib (i.e., IC_{50}, 400 to 2000 nM)) [28]. Third, while some reports documented that crizotinib differential sensitivity in EML4-ALK-expressing cells was dependent on the EML4-ALK variant [94–96], other studies have shown no link between EML4-ALK fusion variants and crizotinib responses [97,98]. Fourth, crizotinib was shown to suppress the growth of ALK+ thyroid cancer cells; however, this potential therapeutic benefit was produced from non-Met/ALK-targeting effects [99]. All these facts highlight the importance of undertaking more research to understand the molecular basis of resistance to ALK TKIs.

The current active approach to overcome ALK-TKIs resistance mainly relies on second- and third-generation TKIs, with over 11 inhibitors being developed [100,101]. With increased experience in TKI resistance, the clinical response to the next-generation TKIs is commonly highly variable and unpredictable [101]. For instance, third-generation EGFR TKIs are being developed as part of a
strategy to overcome treatment resistance to first- and second-generation EGFR TKIs in lung cancer patients [102]; however, resistance to third-generation EGFR TKIs such as AZD9291 and HM61713 are also being arise [103]. A pervading theme regarding resistance to TKI therapy is its mediation by secondary mutations, which has not been resolved by introducing new generations of more TKIs. For example, despite the clinical efficacy of the first-, second-, and third-generation BCR-ABL inhibitors, resistance occurs invariably and more than 50 distinct point mutations encoding single amino-acid substitutions in the kinase domain of the BCR-ABL1 gene have been detected in patients with imatinib-resistant CML [64,104]. Adding more inhibitors does not work, as previously shown for other tyrosine kinase inhibitors. Therefore, it is crucial to perform further research to understand TKI resistance in ALK+ cancer patients.

Taken together, the evidence presented in this review depicts the importance of continuous investigation towards a deeper understanding of clinical resistance. Although there are copious and complex questions to be solved, the recent advances in both clinical and preclinical research, facilitated by the impressive developments in experimental methods and techniques, has generated much enthusiasm and hope for the future.

Conflicts of Interest: The authors declare no conflict of interest.

References
1. Vijayvergia, N.; Mehra, R. Clinical challenges in targeting anaplastic lymphoma kinase in advanced non-small cell lung cancer. Cancer Chemother. Pharmacol. 2014, 74, 437–446. [CrossRef] [PubMed]
2. Morris, S.W.; Kirstein, M.N.; Valentine, M.B.; Dittmer, K.G.; Shapiro, D.N.; Saltman, D.L.; Look, A.T. Fusion of a kinase gene, ALK, to a nucleolar protein gene, NPM, in non-Hodgkin’s lymphoma. Science 1994, 263, 1281–1284. [CrossRef] [PubMed]
3. Shiota, M.; Fujimoto, J.; Semb, T.; Satoh, H.; Yamamoto, T.; Mori, S. Hyperphosphorylation of a novel 80 kDa protein-tyrosine kinase similar to Ltk in a human Ki-1 lymphoma cell line, AMS3. Oncogene 1994, 9, 1567–1574. [PubMed]
4. Lai, R.; Ingham, R.J. The pathobiology of the oncogenic tyrosine kinase NPM-ALK: A brief update. Ther. Adv. Hematol. 2013, 4, 119–131. [CrossRef] [PubMed]
5. Amin, H.M.; Lai, R. Pathobiology of ALK+ anaplastic large-cell lymphoma. Blood 2007, 110, 2259–2267. [CrossRef] [PubMed]
6. Hallberg, B.; Palmer, R.H. Mechanistic insight into ALK receptor tyrosine kinase in human cancer biology. Nat. Rev. Cancer 2013, 13, 685–700. [CrossRef] [PubMed]
7. Grande, E.; Bolos, M.V.; Arriola, E. Targeting oncogenic ALK: A promising strategy for cancer treatment. Mol. Cancer Ther. 2011, 10, 569–579. [CrossRef] [PubMed]
8. Rikova, K.; Guo, A.; Zeng, Q.; Possemato, A.; Yu, J.; Haack, H.; Nardone, J.; Lee, K.; Reeves, C.; Li, Y.; et al. Global survey of phosphotyrosine signaling identifies oncogenic kinases in lung cancer. Cell 2007, 131, 1190–1203. [CrossRef] [PubMed]
9. Soda, M.; Choi, Y.L.; Enomoto, M.; Takada, S.; Yamashita, Y.; Ishikawa, S.; Fujiwara, S.; Watanabe, H.; Kurashina, K.; Hatanaka, H.; et al. Identification of the transforming EML4-ALK fusion gene in non-small-cell lung cancer. Nature 2007, 448, 561–566. [CrossRef] [PubMed]
10. Mosse, Y.P.; Laudenslager, M.; Longo, L.; Cole, K.A.; Wood, A.; Attiyeh, E.F.; Laquaglia, M.J.; Sennett, R.; Lynch, J.E.; Perri, P.; et al. Identification of ALK as a major familial neuroblastoma predisposition gene. Nature 2008, 453, 930–935. [CrossRef] [PubMed]
11. Janoueix-Lerosey, I.; Lequin, D.; Brugieres, L.; Ribeiro, A.; de Pontual, L.; Combaret, V.; Raynal, V.; Puisieux, A.; Schleiermacher, G.; Pierron, G.; et al. Somatic and germline activating mutations of the ALK kinase receptor in neuroblastoma. Nature 2008, 455, 967–970. [CrossRef] [PubMed]
12. George, R.E.; Sandra, T.; Hanna, M.; Frohling, S.; Luther, W.; II; Zhang, J.; Ahn, Y.; Zhou, W.; London, W.B.; McGrady, P.; et al. Activating mutations in ALK provide a therapeutic target in neuroblastoma. Nature 2008, 455, 975–978. [CrossRef] [PubMed]
13. Chen, Y.; Takita, J.; Choi, Y.L.; Kato, M.; Ohira, M.; Sanada, M.; Wang, L.; Soda, M.; Kikuchi, A.; Igarashi, T.; et al. Oncogenic mutations of ALK kinase in neuroblastoma. Nature 2008, 455, 971–974. [CrossRef] [PubMed]
14. Werner, M.T.; Zhao, C.; Zhang, Q.; Wasik, M.A. Nucleophosmin-anaplastic lymphoma kinase: The ultimate oncogene and therapeutic target. *Blood* 2017, 129, 823–831. [CrossRef] [PubMed]

15. Shaw, A.T.; Kim, D.W.; Nakagawa, K.; Seto, T.; Crino, L.; Ahn, M.J.; De Pas, T.; Besse, B.; Solomon, B.J.; Blackhall, F.; et al. Crizotinib versus chemotherapy in advanced ALK-positive lung cancer. *N. Engl. J. Med.* 2013, 368, 2385–2394. [CrossRef] [PubMed]

16. Gambacorti-Passerini, C.; Messa, C.; Pogliani, E.M. Crizotinib in anaplastic large-cell lymphoma. *N. Engl. J. Med.* 2011, 364, 775–776. [CrossRef] [PubMed]

17. Gambacorti Passerini, C.; Farina, F.; Stasia, A.; Redaelli, S.; Ceccon, M.; Mologni, L.; Messa, C.; Guerra, L.; Giudici, G.; Sala, E.; et al. Crizotinib in advanced, chemoresistant anaplastic lymphoma kinase-positive lymphoma patients. *J. Nat. Cancer Inst.* 2014, 106. [CrossRef] [PubMed]

18. Mosse, Y.P.; Lim, M.S.; Voss, S.D.; Wilner, K.; Ruffner, K.; Laliberte, J.; Rolland, D.; Balis, F.M.; Maris, J.M.; Weigel, B.J.; et al. Safety and activity of crizotinib for paediatric patients with refractory solid tumours or anaplastic large-cell lymphoma: A Children’s Oncology Group phase 1 consortium study. *Lancet Oncol.* 2013, 14, 472–480. [CrossRef]

19. Choi, Y.L.; Soda, M.; Yamashita, Y.; Ueno, T.; Takashima, J.; Nakajima, T.; Yatabe, Y.; Takeuchi, K.; Hamada, T.; Haruta, H.; et al. EML4-ALK mutations in lung cancer that confer resistance to ALK inhibitors. *N. Engl. J. Med.* 2010, 363, 1734–1739. [CrossRef] [PubMed]

20. Sasaki, T.; Okuda, K.; Zheng, W.; Butrynski, J.; Capelletti, M.; Wang, L.; Gray, N.S.; Wilner, K.; Christensen, J.G.; Demetri, G.; et al. The Neuroblastoma-Associated F1174L ALK Mutation Causes Resistance to an ALK Kinase Inhibitor in ALK-Translocated Cancers. *Cancer Res.* 2010, 70, 10038–10043. [CrossRef] [PubMed]

21. Roskoski, R., Jr. Anaplastic lymphoma kinase (ALK): Structure, oncogenic activation, and pharmacological inhibition. *Pharmacol. Res.* 2013, 68, 68–94. [CrossRef] [PubMed]

22. Azarova, A.M.; Gautam, G.; George, R.E. Emerging importance of ALK in neuroblastoma. *Semin. Cancer Biol.* 2011, 21, 267–275. [CrossRef] [PubMed]

23. Facchinetti, F.; Tiseo, M.; Di Maio, M.; Graziano, P.; Bria, E.; Rossi, G.; Novello, S. Tackling ALK in non-small cell lung cancer: The role of novel inhibitors. *Transl. Lung Cancer Res.* 2016, 5, 301–321. [CrossRef] [PubMed]

24. Shaw, A.T.; Engelman, J.A. Ceritinib in ALK-rearranged non-small-cell lung cancer. *N. Engl. J. Med.* 2014, 370, 2537–2539. [CrossRef] [PubMed]

25. Friboulet, L.; Li, N.; Katayama, R.; Lee, C.C.; Gainor, J.F.; Crystal, A.S.; Michellys, P.Y.; Awad, M.M.; Yanagitani, N.; Kim, S.; et al. The ALK inhibitor ceritinib overcomes crizotinib resistance in non-small cell lung cancer. *Cancer Discov.* 2014, 4, 662–673. [CrossRef] [PubMed]

26. Ajimizu, H.; Kim, Y.H.; Mishima, M. Rapid response of brain metastases to alectinib in a patient with non-small-cell lung cancer resistant to crizotinib. *Med. Oncol.* 2015, 32, 477. [CrossRef] [PubMed]

27. Kim, D.W.; Tiseo, M.; Ahn, M.J.; Reckamp, K.L.; Hansen, K.H.; Kim, S.W.; Huber, R.; West, H.; Groen, H.; Hochmair, M.J.; et al. Brigatinib in Patients With Crizotinib-Refractory Anaplastic Lymphoma Kinase-Positive Non-Small-Cell Lung Cancer: A Randomized, Multicenter Phase II Trial. *J. Clin. Oncol. Off. J. Am. Soc. Clin. Oncol.* 2017, 35, 2490–2498. [CrossRef] [PubMed]

28. Moore, N.F.; Azarova, A.M.; Bhattacharay, N.; Ross, K.N.; Drake, L.E.; Frumm, S.; Liu, Q.S.; Christie, A.L.; Sanda, T.; Chesler, L.; et al. Molecular rationale for the use of PI3K/AKT/mTOR pathway inhibitors in combination with crizotinib in ALK-mutated neuroblastoma. *Oncotarget* 2014, 5, 8737–8749. [CrossRef] [PubMed]

29. Umapathy, G.; El Wakil, A.; Witek, B.; Chesler, L.; Danielson, L.; Deng, X.; Gray, N.; Johansson, M.; Kvambrink, S.; Ruuth, K.; et al. The kinase ALK stimulates the kinase ERK3 to promote the expression of the oncogene MYCN in neuroblastoma. *Sci. Signal.* 2014, 7, ra102. [CrossRef] [PubMed]

30. Crystal, A.S.; Shaw, A.T.; Sequist, L.V.; Friboulet, L.; Niederst, M.J.; Lockerman, E.L.; Frias, R.L.; Gainor, J.F.; Amzallag, A.; Greninger, P.; et al. Patient-derived models of acquired resistance can identify effective drug combinations for cancer. *Science* 2014, 346, 1480–1486. [CrossRef] [PubMed]
31. Katayama, R.; Khan, T.M.; Benes, C.; Lifshits, E.; Ebi, H.; Rivera, V.M.; Shakespeare, W.C.; Iafrate, A.J.; Engelman, J.A.; Shaw, A.T. Therapeutic strategies to overcome crizotinib resistance in non-small cell lung cancers harboring the fusion oncogene EML4-ALK. Proc. Natl. Acad. Sci. USA 2011, 108, 7535–7540. [CrossRef] [PubMed]

32. Alshareef, A.; Zhang, H.F.; Huang, Y.H.; Wu, C.; Zhang, J.D.; Wang, P.; El-Sehemy, A.; Fares, M.; Lai, R. The use of cellular thermal shift assay (CETSA) to study Crizotinib resistance in ALK-expressing human cancers. Sci. Rep. 2016, 6, 33710. [CrossRef] [PubMed]

33. Sakamoto, H.; Tsukaguchi, T.; Hiroshima, S.; Kodama, T.; Kobayashi, T.; Fukami, T.A.; Oikawa, N.; Tsukuda, T.; Ishii, N.; Aoki, Y. CH5424802, a selective ALK inhibitor capable of blocking the resistant gatekeeper mutant. Cancer Cell 2011, 19, 679–690. [CrossRef] [PubMed]

34. Kodama, T.; Tsukaguchi, T.; Hiroshima, S.; Kodama, T.; Kobayashi, T.; Fukami, T.A.; Oikawa, N.; Tsukuda, T.; Ishii, N.; Aoki, Y. CH5424802, a selective ALK inhibitor capable of blocking the resistant gatekeeper mutant. Cancer Cell 2011, 19, 679–690. [CrossRef] [PubMed]

35. Katayama, R.; Friboulet, L.; Koike, S.; Lockerman, E.L.; Khan, T.M.; Gainor, J.F.; Iafrate, A.J.; Takeuchi, K.; Taji, M.; Okuno, Y.; et al. Two novel ALK mutations mediate acquired resistance to the next-generation ALK inhibitor alectinib. Clin. Cancer Res. Off. J. Am. Assoc. Cancer Res. 2014, 20, 5686–5696. [CrossRef] [PubMed]

36. Ou, S.H.; Milliken, J.C.; Azada, M.C.; Miller, V.A.; Ali, S.M.; Klempner, S.J. ALK F1174V mutation confers sensitivity while ALK I1171 mutation confers resistance to alectinib. The importance of serial biopsy post progression. Lung Cancer 2016, 91, 70–72. [CrossRef] [PubMed]

37. Ou, S.H.; Greenbowe, J.; Khan, Z.U.; Azada, M.C.; Ross, J.S.; Stevens, P.J.; Ali, S.M.; Miller, V.A.; Gitlitz, B. I1171 missense mutation (particularly I1171N) is a common resistance mutation in ALK-positive NSCLC patients who have progressive disease while on alectinib and is sensitive to ceritinib. Lung Cancer 2015, 88, 231–234. [CrossRef] [PubMed]

38. Gainor, J.F.; Dardaei, L.; Yoda, S.; Friboulet, L.; Leshchiner, I.; Katayama, R.; Dagogo-Jack, I.; Gadgeel, S.; Schultz, K.; Singh, M.; et al. Molecular Mechanisms of Resistance to First- and Second-Generation ALK Inhibitors in ALK-Rearranged Lung Cancer. Cancer Discov. 2016, 6, 1118–1133. [CrossRef] [PubMed]

39. Shaw, A.T.; Friboulet, L.; Leshchiner, I.; Gainor, J.F.; Bergqvist, L.; Brooun, A.; Burke, B.J.; Deng, Y.-L.; Liu, W.; Dardaei, L.; et al. Resensitization to Crizotinib by the Lorlatinib ALK Resistance Mutation L1198F. N. Engl. J. Med. 2016, 374, 54–61. [CrossRef] [PubMed]

40. Huang, L.; Fu, L. Mechanisms of resistance to EGFR tyrosine kinase inhibitors. Acta Pharm. Sin. B 2015, 5, 390–401. [CrossRef] [PubMed]

41. Huang, C.-P.; Tsai, M.-F.; Chang, T.-H.; Tang, W.-C.; Chen, S.-Y.; Lai, H.-H.; Lin, T.-Y.; Yang, J.C.-H.; Yang, P.-C.; Shih, J.-Y.; et al. ALDH-positive lung cancer stem cells confer resistance to epidermal growth factor receptor tyrosine kinase inhibitors. Cancer Lett. 2013, 328, 144–151. [CrossRef] [PubMed]

42. Shien, K.; Toyooka, S.; Yamamoto, H.; Soh, J.; Jida, M.; Thu, K.L.; Hashida, S.; Maki, Y.; Ichihara, E.; Asano, H.; et al. Acquired Resistance to EGFR Inhibitors Is Associated with a Manifestation of Stem cell-like Properties in Cancer Cells. Cancer Res. 2013, 73, 3051–3061. [CrossRef] [PubMed]

43. Barnes, D.J.; Palaiologou, D.; Panousopoulou, E.; Schultzis, B.; Yong, A.S.M.; Pattacini, L.; Goldman, J.M.; Melo, J.V. Bcr-Abl Expression Levels Determine the Rate of Development of Resistance to Imatinib Mesylate in Chronic Myeloid Leukemia. Cancer Res. 2005, 65, 8912–8919. [CrossRef] [PubMed]

44. Al-Achkar, W.; Wafa, A.; Moassass, F.; Klein, E.; Liehr, T. Multiple copies of BCR-ABL fusion gene on two isodicentric Philadelphia chromosomes in an imatinib mesylate-resistant chronic myeloid leukemia patient. Oncol. Lett. 2013, 5, 1579–1582. [CrossRef] [PubMed]

45. White, D.L.; Saunders, V.A.; Dang, P.; Engler, J.; Zannettino, A.C.; Cambarelli, A.C.; Quinn, S.R.; Manley, P.W.; Hughes, T.P. OCT-1-mediated influx is a key determinant of the intracellular uptake of imatinib but not nilotinib (AMN107): Reduced OCT-1 activity is the cause of low in vitro sensitivity to imatinib. Blood 2006, 108, 697–704. [CrossRef] [PubMed]

46. Jiang, X.; Zhao, Y.; Smith, C.; Gasparetto, M.; Turhan, A.; Eaves, A.; Eaves, C. Chronic myeloid leukemia stem cells possess multiple unique features of resistance to BCR-ABL targeted therapies. Leukemia 2007, 21, 926–935. [CrossRef] [PubMed]

47. Traer, E.; MacKenzie, R.; Sneed, J.; Agarwal, A.; Eiring, A.M.; O’Hare, T.; Druker, B.J.; Deininger, M.W. Blockade of JAK2-mediated extrinsic survival signals restores sensitivity of CML cells to ABL inhibitors. Leukemia 2012, 26, 1140–1143. [CrossRef] [PubMed]
Weisberg, E.; Wright, R.D.; McMillin, D.W.; Mitsiades, C.; Ray, A.; Barrett, R.; Adamia, S.; Stone, R.; Galinsky, I.; Kung, A.L.; et al. Stromal-mediated protection of tyrosine kinase inhibitor-treated BCR-ABL-expressing leukemia cells. *Mol. Cancer Ther.* 2008, 7, 1121–1129. [CrossRef] [PubMed]

Chu, S.; Holtz, M.; Gupta, M.; Bhatia, R. BCR/ABL kinase inhibition by imatinib mesylate enhances MAP kinase activity in chronic myelogenous leukemia CD34+ cells. *Blood* 2004, 103, 3167–3174. [CrossRef] [PubMed]

Aceves-Luquero, C.I.; Agarwal, A.; Callejas-Valera, J.L.; Arias-Gonzalez, L.; Esparis-Ogando, A.; del Peso Ovalle, L.; Bellon-Echeverria, I.; de la Cruz-Morcillo, M.A.; Galan Moya, E.M.; Moreno Gimeno, I.; et al. ERK2, but not ERK1, mediates acquired and “de novo” resistance to imatinib mesylate: Implication for CML therapy. *PLoS ONE* 2009, 4, e6124. [CrossRef] [PubMed]

Du, B.; Shim, J.S. Targeting Epithelial-Mesenchymal Transition (EMT) to Overcome Drug Resistance in Cancer. *Molecules* 2016, 21, 965. [CrossRef] [PubMed]

Kim, H.R.; Kim, W.S.; Choi, Y.J.; Choi, C.M.; Rho, J.K.; Lee, J.C. Epithelial-mesenchymal transition leads to crizotinib resistance in H2228 lung cancer cells with EML4-ALK translocation. *Mol. Oncol.* 2013, 7, 1093–1102. [CrossRef] [PubMed]

Uramoto, H.; Iwata, T.; Onitsuka, T.; Shimokawa, H.; Hanagiri, T.; Oyama, T. Epithelial-mesenchymal transition in EGFR-TKI acquired resistant lung adenocarcinoma. *Anticancer Res.* 2010, 30, 2513–2517. [PubMed]

Yauch, R.L.; Januario, T.; Eberhard, D.A.; Cavet, G.; Zhu, W.; Fu, L.; Pham, T.Q.; Soriano, R.; Stinson, J.; Seshagiri, S.; et al. Epithelial versus mesenchymal phenotype determines in vitro sensitivity and predicts clinical activity of erlotinib in lung cancer patients. *Clin. Cancer Res. Off. J. Am. Assoc. Cancer Res.* 2005, 11, 8686–8698. [CrossRef] [PubMed]

Thomson, S.; Buck, E.; Petti, F.; Griffin, G.; Brown, E.; Rammarine, N.; Iwata, K.K.; Gibson, N.; Haley, J.D. Epithelial to mesenchymal transition is a determinant of sensitivity of non-small-cell lung carcinoma cell lines and xenografts to epidermal growth factor receptor inhibition. *Cancer Res.* 2005, 65, 9455–9462. [CrossRef] [PubMed]

Sequist, L.V.; Waltman, B.A.; Dias-Santagata, D.; Digumarthy, S.; Turke, A.B.; Fidias, P.; Bergethon, K.; Shaw, A.T.; Gettinger, S.; Cosper, A.K.; et al. Genotypic and Histological Evolution of Lung Cancers Acquiring Resistance to EGFR Inhibitors. *Sci. Transl. Med.* 2011, 3, 75ra26. [CrossRef] [PubMed]

Fujita, S.; Masago, K.; Katakami, N.; Yatabe, Y. Transformation to SCLC after Treatment with the ALK Inhibitor Alectinib. *J. Thorac. Oncol.* 2016, 11, e67–e72. [CrossRef] [PubMed]

Ou, S.I.; Lee, T.K.; Young, L.; Fernandez-Rocha, M.Y.; Pavlick, D.; Schrock, A.B.; Zhu, V.W.; Milliken, J.; Ali, S.M.; Gitlitz, B.J. Dual occurrence of ALK G1202R solvent front mutation and small cell lung cancer transformation as resistance mechanisms to second generation ALK inhibitors without prior exposure to crizotinib. Pitfall of solely relying on liquid re-biopsy? *Lung Cancer* 2017, 106, 110–114. [CrossRef] [PubMed]

Debruyne, D.N.; Bhatnagar, N.; Sharma, B.; Luther, W.; Moore, N.F.; Cheung, N.K.; Gray, N.S.; George, R.E. ALK inhibitor resistance in ALKF1174L-driven neuroblastoma is associated with AXL activation and induction of EMT. *Oncogene* 2016, 35, 3681–3691. [CrossRef] [PubMed]

Wilson, C.; Nichols, K.; Bustos, D.; Lin, E.; Song, Q.; Stephan, J.P.; Kirkpatrick, D.S.; Settleman, J. Overcoming EMT-associated resistance to anti-cancer drugs via Src/FAK pathway inhibition. *Oncotarget* 2014, 5, 7328–7341. [CrossRef] [PubMed]

Hu, Y.; Zang, J.; Qin, X.; Yan, D.; Cao, H.; Zhou, L.; Ni, J.; Yu, S.; Wu, J.; Feng, J.T. Epithelial-to-mesenchymal transition correlates with gefitinib resistance in NSCLC cells and the liver X receptor ligand GW3965 reverses gefitinib resistance through inhibition of vimentin. *OncoTargets Ther.* 2017, 10, 2341–2348. [CrossRef] [PubMed]

Soucheray, M.; Capelletti, M.; Pulido, I.; Kuang, Y.; Paweletz, C.P.; Becker, J.H.; Kikuchi, E.; Xu, C.; Patel, T.B.; Al-Shahrour, F.; et al. Intratumoral Heterogeneity in EGFR-Mutant NSCLC Results in Divergent Resistance Mechanisms in Response to EGFR Tyrosine Kinase Inhibition. *Cancer Res.* 2015, 75, 4372–4383. [CrossRef] [PubMed]

Gower, A.; Hsu, W.H.; Hsu, S.T.; Wang, Y.; Giaccone, G. EMT is associated with, but does not drive resistance to ALK inhibitors among EML4-ALK non-small cell lung cancer. *Mol. Oncol.* 2016, 10, 601–609. [CrossRef] [PubMed]
64. Yang, K.; Fu, L.-W. Mechanisms of resistance to BCR-ABL TKIs and the therapeutic strategies: A review. Crit. Rev. Oncol. Hematol. 2015, 93, 277–292. [CrossRef] [PubMed]

65. Hamilton, A.; Helgason, G.V.; Schemionek, M.; Zhang, B.; Myssina, S.; Allan, E.K.; Nicolini, F.E.; Muller-Tidow, C.; Bhatia, R.; Brunton, V.G.; et al. Chronic myeloid leukemia stem cells are not dependent on Bcr-Abl kinase activity for their survival. Blood 2012, 119, 1501–1510. [CrossRef] [PubMed]

66. Corbin, A.S.; Agarwal, A.; Loriaux, M.; Cortes, J.; Deininger, M.W.; Druker, B.J. Human chronic myeloid leukemia stem cells are insensitive to imatinib despite inhibition of BCR-ABL activity. J. Clin. Investig. 2011, 121, 396–409. [CrossRef] [PubMed]

67. Oh, S.J.; Noh, K.H.; Lee, Y.-H.; Hong, S.-O.; Song, K.-H.; Lee, H.-J.; Kim, S.; Kim, T.M.; Jeon, J.-H.; Seo, J.H.; et al. Targeting stemness is an effective strategy to control EML4-ALK(+) non-small cell lung cancer cells. Oncotarget 2015, 6, 40255–40267. [CrossRef] [PubMed]

68. Redaelli, S.; Cecconi, M.; Antolini, L.; Rigolio, R.; Pirola, A.; Peronaci, M.; Gambacorti-Passerini, C.; Mologni, L. Synergistic activity of ALK and mTOR inhibitors for the treatment of NPM-ALK positive lymphoma. Oncotarget 2016, 7, 72886–72897. [CrossRef] [PubMed]

69. Berry, T.; Luther, W.; Bhatnagar, N.; Jamin, Y.; Poon, E.; Sanda, T.; Pei, D.; Sharma, B.; Vetharoy, W.R.; Hallsworth, A.; et al. The ALK<sup>F1174L</sup> Mutation Potentiates the Oncogenic Activity of MYCN in Neuroblastoma. Cancer Cell 2012, 22, 117–130. [CrossRef] [PubMed]

70. Zhang, H.F.; Wu, C.; Alshareef, A.; Gupta, N.; Zhao, Q.; Xu, X.E.; Jiao, J.W.; Li, E.M.; Xu, L.Y.; Lai, R. The PI3K/AKT/e-MYC Axis Promotes the Acquisition of Cancer Stem-Like Features in Esophageal Squamous Cell Carcinoma. Stem Cells 2016, 34, 2040–2051. [CrossRef] [PubMed]

71. Jung, K.; Gupta, N.; Wang, P.; Lewis, J.T.; Gopal, K.; Wu, F.; Ye, X.; Alshareef, A.; Abdulkarim, B.S.; Douglas, D.N.; et al. Triple negative breast cancers comprise a highly tumorigenic cell subpopulation detectable by its high responsiveness to a Sox2 regulatory region 2 (SRR2) reporter. Oncotarget 2015, 6, 10366–10373. [CrossRef] [PubMed]

72. Wu, F.; Zhang, J.; Wang, P.; Ye, X.; Jung, K.; Bone, K.M.; Pearson, J.D.; Ingham, R.J.; McMullen, T.P.; Ma, Y.; et al. Identification of two novel phenotypically distinct breast cancer cell subsets based on Sox2 transcription activity. Cell. Signal. 2012, 24, 1989–1998. [CrossRef] [PubMed]

73. Gelebart, P.; Hegazy, S.A.; Wang, P.; Bone, K.M.; Anand, M.; Sharon, D.; Hitt, M.; Pearson, J.D.; Ingham, R.J.; Ma, Y.; et al. Aberrant expression and biological significance of Sox2, an embryonic stem cell transcriptional factor, in ALK-positive anaplastic large cell lymphoma. Blood Cancer J. 2012, 2, e82. [CrossRef] [PubMed]

74. Liang, S.; Furuhashi, M.; Nakane, R.; Nakazawa, S.; Goudarzi, H.; Hamada, J.-L.; Iizasa, H. Isolation and characterization of human breast cancer cells with SOX2 promoter activity. Biochem. Biophys. Res. Commun. 2013, 437, 205–211. [CrossRef] [PubMed]

75. Iglesias, J.M.; Leis, O.; Pérez Ruiz, E.; Gumuzio Barrie, J.; Garcia-Garcia, F.; Aduriz, A.; Beloqui, I.; Hernandez-Garcia, S.; Lopez-Mato, M.P.; Dopazo, J.; et al. The Activation of the Sox2 RR2 Potentiates Phosphorylation Transcriptional Reporter in Human Breast Cancer Cell Lines is Dynamic and Labels Cells with Higher Tumorigenic Potential. Front. Oncol. 2014, 4. [CrossRef] [PubMed]

76. Heidel, F.H.; Bullinger, L.; Feng, Z.; Wang, Z.; Neff, T.A.; Stein, L.; Kalaitzidis, D.; Lane, S.W.; Armstrong, S.A. Genetic and Pharmacologic Inhibition of β-Catenin Targets Imatinib-Resistant Leukemia Stem Cells in CML. Cell Stem Cell 2012, 10, 412–424. [CrossRef] [PubMed]

77. Togashi, Y.; Hayashi, H.; Terashima, M.; de Velasco, M.A.; Sakai, K.; Fujita, Y.; Tomida, S.; Nakagawa, K.; Nishio, K. Inhibition of β-Catenin enhances the anticancer effect of irreversible EGFR-TKI in EGFR-mutated non-small-cell lung cancer with a T790M mutation. J. Thorac. Oncol. Off. Publ. Int. Assoc. Study Lung Cancer 2015, 10, 93–101. [CrossRef] [PubMed]

78. Zhang, X.; Hao, J. Development of anticancer agents targeting the Wnt/β-catenin signaling. Am. J. Cancer Res. 2015, 5, 2344–2360. [PubMed]

79. Anand, M.; Lai, R.; Gelebart, P. β-catenin is constitutively active and increases STAT3 expression/activation in anaplastic lymphoma kinase-positive anaplastic large cell lymphoma. Haematologica 2011, 96, 253–261. [CrossRef] [PubMed]

80. Chiarle, R.; Voena, C.; Ambrogio, C.; Piva, R.; Inghirami, G. The anaplastic lymphoma kinase in the pathogenesis of cancer. Nat. Rev. Cancer 2008, 8, 11–23. [CrossRef] [PubMed]
81. Martinez Molina, D.; Jafari, R.; Ignatoushchenko, M.; Seki, T.; Larsson, E.A.; Dan, C.; Sreekumar, L.; Cao, Y.; Nordlund, P. Monitoring drug target engagement in cells and tissues using the cellular thermal shift assay. *Science 2013*, *341*, 84–87. [CrossRef] [PubMed]

82. Jafari, R.; Almqvist, H.; Axelsson, H.; Ignatoushchenko, M.; Lundback, T.; Nordlund, P.; Martinez Molina, D. The cellular thermal shift assay for evaluating drug target interactions in cells. *Nat. Protoc. 2014*, *9*, 2100–2122. [CrossRef] [PubMed]

83. Jensen, A.J.; Martinez Molina, D.; Lundback, T. CETSA: A target engagement assay with potential to transform drug discovery. *Future Med. Chem. 2015*, *7*, 975–978. [CrossRef] [PubMed]

84. Savitski, M.M.; Reinhard, F.B.M.; Franken, H.; Werner, T.; Savitski, M.F.; Eberhard, D.; Molina, D.M.; Jafari, R.; Dovega, R.B.; Klaeger, S.; et al. Tracking cancer drugs in living cells by thermal profiling of the proteome. *Science 2014*, *346*, 1255784. [CrossRef] [PubMed]

85. Huber, K.V.; Salah, E.; Radic, B.; Gridling, M.; Elkins, J.M.; Stukalov, A.; Jemth, A.S.; Gokturk, C.; Sanjiv, K.; Stromberg, K.; et al. Stereoscopic targeting of MTH1 by (S)-crizotinib as an anticancer strategy. *Nature 2014*, *508*, 222–227. [CrossRef] [PubMed]

86. Liu, X.; Shi, Y.; Maag, D.X.; Palma, J.P.; Patterson, M.J.; Ellis, P.A.; Surber, B.W.; Ready, D.B.; Soni, N.B.; Ladror, U.S.; et al. Iniparib Nonselectively Modifies Cysteine-Containing Proteins in Tumor Cells and Is Not a Bona Fide PARP Inhibitor. *Clin. Cancer Res. 2012*, *18*, 510–523. [CrossRef] [PubMed]

87. Guha, M. PARP inhibitors stumble in breast cancer. *Nat. Biotech. 2011*, *29*, 373–374. [CrossRef] [PubMed]

88. Xu, Q.; Walker, D.; Bernardo, A.; Brodbeck, J.; Balestra, M.E.; Huang, Y. Intron-3 retention/splicing controls neuronal expression of apolipoprotein E in the CNS. *J. Neurosci. Off. J. Soc. Neurosci. 2008*, *28*, 1452–1459. [CrossRef] [PubMed]

89. Lu, F.; Gladden, A.B.; Diehl, J.A. An alternatively spliced cyclin D1 isoform, cyclin D1b, is a nuclear oncogene. *Cancer Res. 2003*, *63*, 7056–7061. [PubMed]

90. Comstock, C.E.; Augello, M.A.; Benito, R.P.; Karch, J.; Tran, T.H.; Utama, F.E.; Tindall, E.A.; Wang, Y.; Burd, C.J.; Groh, E.M.; et al. Cyclin D1 splice variants: Polymorphism, risk, and isoform-specific regulation in prostate cancer. *Clin. Cancer Res. Off. J. Am. Assoc. Cancer Res. 2009*, *15*, 5338–5349. [CrossRef] [PubMed]

91. Wong, J.J.; Au, A.Y.; Ritchie, W.; Rasko, J.E. Intron retention in mRNA: No longer nonsense: Known and putative roles of intron retention in normal and disease biology. *BioEssays News Rev. Mol. Cell. Dev. Boil. 2016*, *38*, 41–49. [CrossRef] [PubMed]

92. Zou, H.Y.; Li, Q.; Lee, J.H.; Arango, M.E.; McDonnell, S.R.; Yamazaki, S.; Koudriakova, T.B.; Alton, G.; Cui, J.J.; Kung, P.P.; et al. An Orally Available Small-Molecule Inhibitor of c-Met, PF-2341066, Exhibits Cytoreductive Antitumor Efficacy through Antiproliferative and Antiangiogenic Mechanisms. *Cancer Res. 2007*, *67*, 4408–4417. [CrossRef] [PubMed]

93. Christensen, J.G.; Zou, H.Y.; Arango, M.E.; Li, Q.; Lee, J.H.; McDonnell, S.R.; Yamazaki, S.; Alton, G.R.; Mroczkowski, B.; Los, G. Cytoreductive antitumor activity of PF-2341066, a novel inhibitor of anaplastic lymphoma kinase and c-Met, in experimental models of anaplastic large-cell lymphoma. *Mol. Cancer Ther. 2007*, *6*, 3314–3322. [CrossRef] [PubMed]

94. Heuckmann, J.M.; Balke-Want, H.; Malchers, F.; Peifer, M.; Sos, M.L.; Koker, M.; Meder, L.; Lovly, C.M.; Heukamp, L.C.; Pao, W.; et al. Differential protein stability and ALK inhibitor sensitivity of EML4-ALK fusion variants. *Clin. Cancer Res. Off. J. Am. Assoc. Cancer Res. 2012*, *18*, 4682–4690. [CrossRef] [PubMed]

95. Yoshida, T.; Oya, Y.; Tanaka, K.; Shimizu, J.; Horio, Y.; Kuroda, H.; Sakao, Y.; Hida, T.; Yatabe, Y. Differential Crizotinib Response Duration Among ALK Fusion Variants in ALK-Positive Non-Small-Cell Lung Cancer. *J. Clin. Oncol. Off. J. Am. Soc. Clin. Oncol. 2016*, *34*, 3383–3389. [CrossRef] [PubMed]

96. Woo, C.G.; Seo, S.; Kim, S.W.; Jang, S.J.; Park, K.S.; Song, J.Y.; Lee, B.; Richards, M.W.; Bayliss, R.; Lee, D.H.; et al. Differential protein stability and clinical responses of EML4-ALK fusion variants to various ALK inhibitors in advanced ALK-rearranged non-small cell lung cancer. *Ann. Oncol. Off. J. Eur. Soc. Med. Oncol. 2017*, *28*, 791–797. [CrossRef] [PubMed]

97. Voena, C.; Chiarle, R. The battle against ALK resistance: Successes and setbacks. *Expert Opin. Invest. Drugs 2012*, *21*, 1751–1754. [CrossRef] [PubMed]

98. Lei, Y.-Y.; Yang, J.-J.; Zhong, W.-Z.; Chen, H.-J.; Yan, H.-H.; Han, J.-F.; Yang, L.-L.; Wu, Y.-L. Clinical efficacy of crizotinib in Chinese patients with ALK-positive non-small-cell lung cancer with brain metastases. *J. Thorac. Dis. 2015*, *7*, 1181–1188. [PubMed]
99. Zhou, Y.; Zhao, C.; Gery, S.; Braunstein, G.D.; Okamoto, R.; Alvarez, R.; Miles, S.A.; Doan, N.B.; Said, J.W.; Gu, J.; et al. Off-target effects of c-MET inhibitors on thyroid cancer cells. *Mol. Cancer Ther.* **2014**, *13*, 134–143. [CrossRef] [PubMed]

100. Awad, M.M.; Shaw, A.T. ALK inhibitors in non-small cell lung cancer: Crizotinib and beyond. *Clin. Adv. Hematol. Oncol.* **2014**, *12*, 429–439.

101. Maione, P.; Sacco, P.C.; Sgambato, A.; Casaluce, F.; Rossi, A.; Gridelli, C. Overcoming resistance to targeted therapies in NSCLC: Current approaches and clinical application. *Ther. Adv. Med. Oncol.* **2015**, *7*, 263–273. [CrossRef] [PubMed]

102. Miller, V.A.; Hirsh, V.; Cadranel, J.; Chen, Y.M.; Park, K.; Kim, S.W.; Zhou, C.; Su, W.C.; Wang, M.; Sun, Y.; et al. Afatinib versus placebo for patients with advanced, metastatic non-small-cell lung cancer after failure of erlotinib, gefitinib, or both, and one or two lines of chemotherapy (LU-X-Lung 1): A phase 2b/3 randomised trial. *Lancet. Oncol.* **2012**, *13*, 528–538. [CrossRef]

103. Wang, S.; Tsui, S.T.; Liu, C.; Song, Y.; Liu, D. EGFR C797S mutation mediates resistance to third-generation inhibitors in T790M-positive non-small cell lung cancer. *J. Hematol. Oncol.* **2016**, *9*, 59. [CrossRef] [PubMed]

104. Quintas-Cardama, A.; Kantarjian, H.; Cortes, J. Flying under the radar: The new wave of BCR-ABL inhibitors. *Nat. Rev. Drug Discov.* **2007**, *6*, 834–848. [CrossRef] [PubMed]