EDITORIAL

New targets bring hope in squamous cell lung cancer: neurotrophic tyrosine kinase gene fusions

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Neurotrophic tyrosine kinase genes encode for the Trk-family proteins TrkA, TrkB, and TrkC, which have an important role in the development of the nervous system; however, they have been identified as oncogenic fusions in solid tumors (NTK-1, NTRK-2, and NTRK-3) and are associated with poor survival in lung cancer. These three new fusions can be detected by fluorescent *in situ* hybridization or next-generation sequencing in less than 5% of the lung tumors. There are several ongoing clinical trials of NTRK oncogenes in lung cancer and other tumors. The agents entrectinib (RXDX-101), a multi-kinase small molecule inhibitor that selectively inhibits NTRK1, NTRK2, and NTRK3, ROS1 and ALK, and LOXO-101, an ATP-competitive pan-NTRK inhibitor, have shown responses in patients with lung cancer with an acceptable toxicity profile. Although these oncogenic fusions are not very prevalent, the high prevalence of lung cancer makes these findings very relevant and suggests the feasibility of these oncogenes as targets in lung cancer. New data from Ozono and collaborators presented in this issue suggest that BDNF/TrkB signal promotes proliferating migratory and invasive phenotypes and cellular plasticity in squamous cell carcinoma (SCC) of the lung but that it also represents a druggable target that may bring hope to squamous lung cancer patients. *Laboratory Investigation* (2017) **97**, 1268–1270; doi:10.1038/labinvest.2017.91

as epidermal growth factor receptor (EGFR) and anaplastic lymphoma kinase (ALK) rearrangements. New, unusual suspects, including BRAF, RET, ROS1, among others are bringing new hope to patients and caregivers. One of the new kids on the block is neurotrophic tyrosine kinase (NTRK). NTRK genes encode for the Trk-family proteins: TrkA, TrkB, and TrkC (encoded by NTRK1, NTTRK2, and NTRK3). The NTRK family plays a role in the development of the nervous system with the modulation of growth, proliferation, repair, maintenance and apoptosis.^{1,2} However, NTRK fusions are also present in solid tumors as oncogenic fusions responsible for growth of cancer cells, proliferation, and survival; the presence of these three oncogenes is associate with poor survival in lung cancers.^{3,4} NTRK1 oncogenic fusions have been detected by fluorescent in situ hybridization (FISH) or next generation sequencing in about 3% of NSCLC,

he therapeutic landscape of lung cancer

has been changing dramatically since the

introduction of therapuetic targets such

while NTRK2 and NTRK3 fusions in about 1% of NSCLC of all types.^{3,5} Several therapeutic agents are currently in development for these oncogenes; one of them is entrectinib (RXDX-101)^{6,7} which is a multi-kinase small molecule inhibitor that selectively inhibits NTRK1, NTRK2, and NTRK3, ROS1 and ALK, as shown in vitro and in vivo in several tumor models. In mice xenografts of colorectal cancer cells with the TrkA fusion TPM3-NTRK1, treatment with entrectinib caused tumor regression.8 Other TRKA inhibitors such as CEP-701 and ARRY-470 were also successful against lung cancer cells expressing NTRK1 fusions.³ Another important agent is LOXO-101, an ATP-competitive pan-NTRK inhibitor that has shown antitumor activity against lung adenocarcinomas.9 AZD7451 was tested on colorectal cancer cell lines harboring tropomyosin (TPM)/NTRK1 fusion (KM12 cell line) and the H460 (an LCNEC cell line exhibiting NTRK2 expression). Proliferation was inhibited in KM12 and H460. An analysis of the H460 cells demonstrated decreased levels of phosphorylation

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¹Phase I-Early Clinical

of NTRK2 after treatment with AZD7451.¹⁰ These and other preclinical studies in colorectal cell lines suggested a role for targeting NRTK-dependent cancer cell populations because these are oncogenic derivers of the cancer tumors.

There are several clinical trials ongoing for NTRK oncogenes, for example, the agent entrectinib first showed activity in colorectal cancer harboring a TPM3-NTRK1 fusions.8 The phase I studies for ALKA-372-001 and STARTRK-1 were dose-escalation trials of entrectinib for patients with advanced or metastatic solid tumors harboring TrkA, ROS1, or ALK molecular alterations. The results of both trials with 67 patients enrolled (38 were NSCLC), were presented at the 2015 ASCO.^{11,12} Among 11 patients received the recommended phase 2 dose of entrectinib 10 responses were reported including 3 of 3 patients harboring NTRK translocations obtained a partial response. In another publication, Farago et al¹³ screened 1378 cases of NSCLC using PCR and confirmed by FISH the presence of NTRK1 gene rearrangements. Two patients (0.1%) were found to have NTRK1 gene rearrangements: one was a TPM3-NTRK1 rearrangement, and in the second case, a fusion transcript containing sequence from SQSTM1 (sequestosome 1, which has previously been described as a fusion partner with ALK in NSCLC) and NTRK1 was observed. The patient presenting the SQSTM1/NTRK1 fusion participated in a phase 1 clinical trial with entrectinib and he achieved a partial response. There is an ongoing trial called STARTRK-2 (NCT02568267) which is a phase 2 basket study evaluating entrectinib for the treatment of solid tumors harboring NTRK1/2/ 3, ROS1, or ALK gene rearrangements currently recruiting patients in the US.

LOXO-101 is an ATP-competitive pan-Trk inhibitor, which has demonstrated encouraging activity on NTRK pre-clinical studies. A phase 1 trial (NCT02122913) is currently recruiting patients.¹⁴ The study has enrolled six patients with NTRK fusions, including one NSCLC (TPR-NTRK1) two mammary analog secretory carcinoma (MASC) (ETV6-NTRK3), one thyroid carcinoma (ETV6-NTRK3), one GIST (ETV6-NTRK3), and one soft tissue sarcoma (STS) (LMNA-NTRK1). All three patients evaluable for efficacy (STS, MASC, and GIST) reported partial responses to LOXO-101.15 Doebele et al published a case report reporting a PR and symptom improvement in a patient with a metastatic STS harboring the LMNA-NTRK1 gene fusion, participating in the LOXO-101 phase 1 trial.⁹ A

phase 2 basket study of LOXO-101 in patients NTRK fusion positive solid tumors, including CNS tumors is ongoing (NCT02576431). New drugs are currently under investigation to target tumors harboring activating NTRK molecular alterations in early phase I studies including patients with advanced solid tumors like the multi-target inhibitors XL-184, MGCD516, and DCC-2701 that are under evaluation in advanced cancers positive for different molecular alterations, including also NTRK (NCT02219711, NCT02228811). Among these, DCC-2701 (Altiratinib), a MET, TIE2, VEGFR2, and TRK kinase inhibitor currently in phase 1 clinical development has recently received orphan drug designation for glioblastoma by the US Food and Drug Administration (FDA). AZD7451 is a tyrosine kinase inhibitor that selectively targets NTRK1, NTRK2, and NTRK3 kinases, being considered then a pan NTRK kinase inhibitor which has shown promising pre-clinical activity in NTRK1 fusion positive lung cancer celllines.¹⁰ Finally, there is the TrkA/ALK inhibitor TSR-011 (NCT02048488), the pan-Trk/ROS1 inhibitor DS6051b (NCT02279433), and the pan-Trk inhibitor PLX-7486 (NCT01804530).

The study by Ozono *et al*¹⁶ in this issue of Laboratory Investigation hypothesizes that BDNF/ TrkB signal promotes proliferating migratory and invasive phenotypes and cellular plasticity in squamous cell carcinoma (SCC) of the lung. The investigators conclude that the BDNF/TrkB signaling pathway contributes to the aggressive behavior of lung SCC by promoting cell invasion and proliferation, and suggest that this pathway may be a new therapeutic target for lung SCC. We agreed with the authors' conclusions. This is the first study to demonstrate, using a Trk inhibitor, that the inhibition of TrkB suppresses the migration, invasion and proliferation activities of lung SCC specifically. There are previous studies of Trk in colon cancer and lung cancer, but this paper is devoted exclusively to SCC, a type of cancer that is in urgently in need of new targets because most of the common actionable genes for lung cancer (EGFR, ALK, ROS-1, RET) are generally found in adenocarcinomas of the lung. These results may pave the way for a target therapy for lung SCC.

We know that NTRK fusions are not very common (<5% of the lung cancers) and that might tamper some enthusiasm, however due to the fact that lung cancers are very prevalent and numerous, even this number of patients represent a very significant group of patients in need of therapy in the US and worldwide.¹⁷

DISCLOSURE/CONFLICT OF INTEREST

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