

Do EGFR tyrosine kinase inhibitors (TKIs) still have a role in EGFR wild-type pre-treated advanced non-small cell lung cancer (NSCLC)? – the shifting paradigm of therapeutics

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Provenance: This is an invited Editorial commissioned by Guest Section Editor Dr. Long Jiang, MD, PhD (Second Affiliated Hospital, Institute of Respiratory Diseases, Zhejiang University School of Medicine, Hangzhou, China).

Comment on: Tomasini P, Brosseau S, Mazières J, *et al.* EGFR tyrosine kinase inhibitors versus chemotherapy in EGFR wild-type pre-treated advanced nonsmall cell lung cancer in daily practice. *Eur Respir J* 2017;50:1700514.

Submitted Dec 22, 2017. Accepted for publication Jan 12, 2018.

doi: 10.21037/tlcr.2018.01.06

View this article at: <http://dx.doi.org/10.21037/tlcr.2018.01.06>

Introduction

EGFR mutations are well established as important oncogenic drivers that occur in 10–44% of primary lung adenocarcinomas occurring more frequently in women, Asians and non-smokers (1,2).

Gefitinib and erlotinib, tyrosine kinase inhibitors (TKIs), were developed and first applied clinically before the significance of *EGFR* mutations was established, in the era when lung cancer therapy was largely empirical. A link between the presence of activating *EGFR* mutations and sensitivity to gefitinib was established in 2004 in a phenotypically enriched population (1,2), however it was still several years before large randomized controlled trials (RCTs) confirmed the superiority of a “targeted” drug approach in patients with sensitizing *EGFR* mutations over standard chemotherapy. In this time, RCTs were also conducted evaluating the empirical use of erlotinib and gefitinib versus placebo as second or third line therapy in unselected patients with chemotherapy refractory non-small cell lung cancer (NSCLC) and in other clinical settings such as maintenance therapy. Despite modest benefits when used empirically in unselected patients, the dramatic effects of these agents in patients with sensitizing *EGFR* mutations redefined modern practice as it is today. Molecular selection

for *EGFR* mutations and anaplastic lymphoma kinase (*ALK*) gene rearrangements is now standard practice in advanced NSCLC while more generalised molecular characterisation is becoming increasingly popular. Patients lacking sensitizing *EGFR* mutations [*EGFR* wild-type (*EGFR*-wt) NSCLC] continue to be treated based on traditional empirical chemotherapy evidence.

In the era of immunotherapy and the advancing molecular landscape, there is a need to reassess the role of *EGFR* TKIs in *EGFR*-wt NSCLC.

Background for TKIs in *EGFR*-wt NSCLC

The empirical use of *EGFR* TKIs compared to placebo in treatment refractory second and third line unselected NSCLC patients was reported in the ISEL trial in 2005, with no improvement in survival observed with gefitinib (3), while statistically significantly improved survival was seen with erlotinib in the same year (4). However, these study populations were not thoroughly assessed for the presence of *EGFR* mutations. Thus, they can not be considered as *EGFR*-wt populations as the true rate of *EGFR* mutations, which may have biased the observed clinical benefits, was not known (Table 1).

In the large BR.21 trial, 731 patients were randomized to

Table 1 Timeline of clinical trials evaluating the efficacy of EGFR TKIs in an unselected or EGFR-wt NSCLC population

Trial title	Author(s)	Year	Trial phase	Treatment	Disease line and population	PFS (months)
ISEL	Thatcher <i>et al.</i>	2005	III	Gefitinib vs. placebo	Second; third-line chemotherapy refractory*; prior platinum mandatory; EGFR unselected	5.6 (g) vs. 5.1 (p); P=0.087
BR.21	Shepherd <i>et al.</i>	2005	III	Erlotinib vs. placebo	Second; third-line, prior combination chemotherapy and not eligible for further chemotherapy; EGFR unselected (EGFR expression testing optional)	2.2 (e) vs. 1.8 (p); P<0.001
ISTANA	Lee <i>et al.</i>	2008	III	Gefitinib vs. docetaxel	Second-line, platinum refractory or recurrence and candidate for further chemotherapy; EGFR unselected	3.4 (d) vs. 3.3 (g); P=0.0134
V-15-32	Maruyama <i>et al.</i>	2008	III	Gefitinib vs. docetaxel	Second; third-line, chemotherapy failure including platinum; EGFR unselected	2.0 (d) vs. 2.0 (g); P=0.335
INTEREST	Kim <i>et al.</i>	2008	III	Gefitinib vs. docetaxel	Second-line platinum refractory and non-refractory; EGFR unselected (EGFR copy number gain sub-group reported)	2.7 (d) vs. 2.2 (g); P=0.47
IPASS	Mok <i>et al.</i>	2008	III	Gefitinib vs. platinum-doublet chemotherapy	First line, EGFR-wt sub-group (EGFR mutated and unknown population also reported)	Not reported as absolute number, approx. 2 mo (g); significantly in favour of chemotherapy (HR =2.85; 95% CI, 2.05–3.98, P<0.001)
SATURN	Capuzzo <i>et al.</i>	2010	III	Erlotinib vs. placebo	Maintenance post first-line platinum chemotherapy; EGFR-unselected (EGFR testing mandatory with missing data)	2.9 (e) vs. 2.6 (p); P<0.0001
CTONG0806	Zhou <i>et al.</i>	2011	II	Gefitinib vs. pemetrexed	Second-line, platinum pre-treated; EGFR-wt	4.8 (p) vs. 1.6 (g); P<0.001
IFCT-GFPC	Pérol <i>et al.</i>	2012	III	Erlotinib or gemcitabine vs. observation	Maintenance post first-line platinum chemotherapy; EGFR-unselected (also included EGFR mutations)	3.8 (g) vs. 2.9 (e) vs. 1.9 (o); P=0.3867
TITAN	Ciuleanu <i>et al.</i>	2012	III	Erlotinib vs. docetaxel	Second-line; disease progression on platinum doublet chemotherapy* shared first line chemotherapy run-in phase with SATURN; EGFR-wt	2.0 (d) vs. 1.5 (e); P=0.089
TORCH	Gridelli <i>et al.</i>	2012	III	Erlotinib then second-line chemotherapy vs. chemotherapy then second-line erlotinib	First (and second-line), switch therapy; EGFR-wt and mutated	5.4 (c) vs. 2.2 (e) (P value not reported)
TAILOR	Garassino <i>et al.</i>	2013	III	Erlotinib vs. docetaxel	Second-line; recurrence or progression after failing platinum chemotherapy; EGFR-wt	2.9 (d) vs. 2.4 (e); P=0.02

Table 1 (continued)

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Trial title	Author(s)	Year	Trial phase	Treatment	Disease line and population	PFS (months)
DELTA	Kawaguchi <i>et al.</i>	2014	III	Erlotinib vs. docetaxel	Second; third-line; post chemotherapy (including platinum); EGFR-wt	3.2 (d) vs. 2.0 (e); P=0.09
HORG	Karampeazis <i>et al.</i>	2013	III	Erlotinib vs. pemetrexed	Second; third-line; disease progression after chemotherapy (platinum not mandatory in ≥ 65 years); EGFR unselected [subgroup (n=11/123) with EGFR mutation reported on]	3.6 (e) vs. 2.9 (p); P=0.136
LUX-Lung 8	Soria <i>et al.</i>	2015	III	Afatinib vs. erlotinib	Second-line; squamous histology, progression after prior platinum chemotherapy; EGFR unselected	2.4 (a) vs. 1.9 (e); P=0.043
IUNO	Cic�nas <i>et al.</i>	2016	III	Maintenance erlotinib vs. placebo and late erlotinib at disease progression in placebo arm	Second-line; prior platinum chemotherapy without progression; EGFR-wt	Maintenance-3 (e) vs. 2.9 (p); P=0.48; 'Early' vs. 'Late' erlotinib; P=0.4759

*, refractory defined as recurrent or progressive disease within 90 days of the last chemotherapy dose; PFS, progression free survival. g, gefitinib; p, placebo; e, erlotinib; d, docetaxel; o, observation; c, chemotherapy; a, afatinib.

erlotinib or placebo in the second and third line setting. All NSCLC histologies were included and *EGFR* testing was not required for enrolment. Overall survival (OS) favoured erlotinib [hazard ratio (HR) =0.73; 95% confidence interval (CI), 0.49–1.1]; however, ongoing treatment if any was not described (4). In further investigation, *EGFR* mutation testing was successful in 197 tumour specimens with a mutation identified in 40 patients (20.3%). Mutational status had no statistically significant association with objective response rate (ORR): 7% *EGFR*-wt patients achieved a response, compared with 16% of those with an *EGFR* mutation (P=0.37) (4,5). Furthermore, erlotinib resulted in an absolute improvement over placebo in progression free survival (PFS) of 0.4 months. Based on this trial, erlotinib was widely used in Europe, the USA and elsewhere as empirical therapy in refractory NSCLC patients.

In 2010 the phase III SATURN study evaluated erlotinib or placebo in 889 patients with NSCLC in the first-line switch maintenance setting. The study included patients with stable or responsive disease after standard first line chemotherapy. Although PFS was reported as 'significantly' longer with erlotinib than with placebo, the difference in median PFS was modest 12.3 weeks for patients on erlotinib versus 11.1 weeks placebo (HR =0.71; 95% CI, 0.62–0.82 weeks; P<0.0001).

Both the adenocarcinoma and squamous (SCC) sub-populations derived benefit. ORR was 11.9% with erlotinib versus 5.4% with placebo and OS reported significantly longer by 1 month (HR =0.77; 95% CI, 0.64–0.93 months, P=0.0063). Forty-four percent were *EGFR*-wt, 6% found to have an *EGFR* mutation, and the remainder having a missing or indeterminate result (6). The IFCT-GFPC phase III study in 2012 investigated whether continuation maintenance with gemcitabine or switch maintenance with erlotinib improved clinical outcomes compared with observation in patients with unselected NSCLC whose disease was controlled after cisplatin-gemcitabine induction chemotherapy. Both gemcitabine and erlotinib prolonged PFS versus observation with improvements in the median PFS by a modest 1.9 and 1.0 months respectively (7).

Moving back, between 2008 and 2010 three RCTs compared gefitinib with docetaxel in unselected patients with previously treated advanced NSCLC. In a Korean population, the ISTANA trial reported improved OS, and quality of life with gefitinib but inferior PFS. The prevalence of *EGFR* mutations has been reported as 20% in Korean patients, although it was not defined in this trial (8). In the Japanese V-15-32 study there was no difference between gefitinib and docetaxel with regard to OS or

PFS (9). The international INTEREST trial demonstrated non-inferiority with gefitinib versus docetaxel in a large study cohort, including patients with copy number gain in the *EGFR* gene (10).

The efficacy and safety of pemetrexed or gefitinib as second-line treatments for advanced *EGFR*-wt non-squamous NSCLC was investigated in the randomized phase II CTONG0806 trial in Chinese patients, reported in 2011. Pemetrexed showed significant improvement in PFS compared with gefitinib (4.8 vs. 1.6 months, $P < 0.001$) (11).

In the TAILOR and DELTA trials of 2013 and 2014 respectively, erlotinib failed to improve OS in comparison with docetaxel in *EGFR*-wt cohorts (12,13), whereas erlotinib and pemetrexed demonstrated similar OS ($P = 0.986$) in a third HORG phase III trial unselected for *EGFR* (14).

In the 2012 TITAN trial erlotinib was compared with chemotherapy in a population of *EGFR*-wt NSCLC patients with poor prognosis and progressive disease during or immediately after first-line chemotherapy. This trial showed no significant difference in OS between the two groups either (5.3 vs. 5.5 months, respectively; HR = 0.96; 95% CI, 0.78–1.19; log-rank $P = 0.73$) however the safety profile again favoured erlotinib (15).

Aforementioned, these trials largely tested *EGFR* TKIs in unselected patients, and rapid and durable responses were more often observed in patients phenotypically enriched for the presence of *EGFR* mutations; adenocarcinoma histology, Asian ethnicity, and a history of never or light smoking. In the landmark 2008 Iressa Pan-Asia Study (IPASS) however, among patients with *EGFR*-wt NSCLC, gefitinib compared to carboplatin and paclitaxel in the first line setting resulted in a significantly inferior ORR (1.1% vs. 23.5%, $P = 0.001$) and shorter PFS (HR = 2.85; 95% CI, 2.05–3.98; median PFS 1.5 and 5.5 months, respectively) (16).

In 2012 subsequent to IPASS, again in the first line setting, the phase III TORCH trial investigated erlotinib followed by cisplatin and gemcitabine at the time of disease progression compared to the standard sequence of cisplatin and gemcitabine followed by erlotinib at the time of disease progression in unselected patients. In the subgroup of patients with *EGFR*-wt NSCLC ($n = 236$), erlotinib followed by cisplatin and gemcitabine compared to the inverse sequence was associated with inferior OS (HR = 1.29; 95% CI, 1.58 to 2.71; median, 6.5 and 9.8 months, respectively) and inferior PFS (17).

In 2015 the LUX-Lung 8 study, which compared the second generation *EGFR* TKI afatinib versus erlotinib in

pre-treated patients with SCC, a small PFS benefit was observed favouring afatinib (median PFS 2.4 vs. 1.9 months respectively, HR = 0.82; $P = 0.043$) (18). There was no chemotherapy or observation control arm in this trial.

Finally, in 2016 the phase III IUNO trial re-assessed the benefit of maintenance erlotinib versus erlotinib at progression in advanced/metastatic NSCLC that had not progressed following four cycles of platinum-based chemotherapy. Median OS was 9.7 and 9.5 months with 'early erlotinib' and 'late erlotinib', respectively ($P = 0.82$). OS with maintenance erlotinib was not superior to second-line treatment in *EGFR*-wt patients. Maintenance treatment with erlotinib in patients with advanced/metastatic NSCLC without *EGFR*-activating mutations was concluded to be considered unfavourable (19) (Table 1).

Article summary

The Intergroupe Francophone de Cancérologie Thoracique (IFCT) Biomarkers France study undertook nationwide genetic tumour profiling in patients with NSCLC, demonstrating the utility of this approach in directing the most suitable therapeutic sequence. This large database was utilised to enable the trial under-review in this editorial (20).

In 2017, Tomasini *et al.* reported on this retrospective real-world study that recruited patients with advanced, largely non-squamous NSCLC who completed molecular testing by the 28 French National Cancer Institute accredited centres between April 2012 and April 2013. Patients were eligible if they did not harbour an *EGFR* mutation or *ALK* rearrangement. The molecular panel funded for routine testing included *EGFR*, *KRAS*, *BRAF*, *PI3KCA* mutations and *ALK* rearrangement. All patients who had received prior first line chemotherapy and second line chemotherapy at the time of progression were eligible and clinical outcome data were provided by the prescribing physician (21).

The primary outcome of interest from this dataset was second-line PFS and OS.

Data were cut June 1, 2015 and 17,640 NSCLC patients for whom molecular testing had been undertaken were potentially eligible with 1,278 patients eventually included after meeting inclusion criteria. Altogether, 410 patients were treated with a second-line *EGFR* TKI and 868 second-line chemotherapy. The median follow-up time was 11.4 (range, 10.3–12.4) months. There were more male patients included (68.8%) than female (32.1%).

Potentially confounding baseline characteristics include more non-smokers in the *EGFR* TKI group than in

the chemotherapy group (16.7% *vs.* 8.8%, respectively; $P < 0.001$) and fewer patients with *KRAS*-mutated tumours (24.9% *vs.* 33.8%; $P = 0.001$). There were more patients with ECOG performance status ≥ 2 and more elderly patients (≥ 65 years) in the EGFR TKI group than in the chemotherapy group (27.1% *vs.* 18.2%; $P = 0.001$ and 46.8% *vs.* 32.7%; $P < 0.001$, respectively).

After adjusting for differences in observed characteristics between treatment groups, median OS and PFS in patients treated with chemotherapy were longer than those with EGFR TKI: OS 8.4 *vs.* 5.0 months, respectively (HR =0.70; 95% CI, 0.6–0.8 months; $P < 0.0001$) and PFS 4.3 (3.88–4.83) *vs.* 2.8 (2.6–3.1) months, respectively (HR =0.66; 95% CI, 0.57–0.77 months; $P < 0.0001$).

In multivariate analyses response to first-line chemotherapy ($P < 0.001$) and smoking status ($P < 0.001$) were prognostic factors predicting a longer OS with EGFR TKI.

PFS and OS of *EGFR*-wt patients treated with second-line EGFR TKI were inferior to those encountered in patients receiving second-line chemotherapy, even when corrected for potential confounding characteristics.

Noticeably, in both groups of patients, those who derived a substantial benefit from first-line therapy also benefited from second-line therapy independent of what the second-line regimen was.

In trials comparing second-line EGFR TKI with either placebo or chemotherapy aforementioned, in patients with *EGFR*-wt tumours, *KRAS* status was often not routinely assessed therefore any potential imbalance between therapeutic arms with respect to *KRAS*-activating mutations unknown. In BR.21, INTEREST and TAILOR, *KRAS* mutated patients did not benefit from erlotinib (5,10,14). Non-smoking patients, *KRAS*-wt treated with EGFR TKIs derived a meaningful OS improvement in this study (HR =0.43, 95% CI, 0.28–0.66; $P < 0.001$); however, numbers may have been small and were not reported (21).

Current *EGFR*-wt TKI guidelines

In 2017, the National Comprehensive Cancer Network (NCCN) panel deleted the recommendation for erlotinib as switch maintenance therapy and as subsequent therapy in non-squamous wild-type NSCLC (22). This was based on preliminary results from the randomized IUNO trial and revised indication by the U.S. Food and Drug Administration (FDA) (19,23). The data showed that survival was not improved with erlotinib versus placebo.

On October 18, 2016, the U.S. Food and Drug

Administration modified the indication for erlotinib for treatment of NSCLC to limit use to patients whose tumours have specific *EGFR* mutations (23).

The 2016 European Society of Medical Oncology (ESMO) also updated their guidelines to remove erlotinib as switch maintenance therapy in NSCLC, however recommend erlotinib still represents a potential second/third-line treatment option in pre-treated patients with unknown or *EGFR*-wt status based on limited efficacy compared with chemotherapy [II, C] (24).

Immunotherapy

This trial recruited prior to immunotherapy being readily available second-line outside of a clinical trial. The first phase III trial demonstrating the superior efficacy of program death receptor-1 (PD-1) inhibitor nivolumab over docetaxel chemotherapy was reported in July 2015 (25); Checkmate-017 demonstrated a 3.2 month OS benefit in squamous NSCLC, further supported by Checkmate-057 with a 2.8 month survival benefit (26). Ongoing comparable data to support the efficacy of PD-1 inhibitor pembrolizumab and program death ligand-1 (PD-L1) inhibitor atezolizumab have followed and changed our practice and the landscape of treating advanced NSCLC as further compelling data emerge in the first line setting and in combination therapy (27,28).

Molecular therapy in oncogene addicted NSCLC

In the last 10 years, in part enabled by the availability of advancing diagnostics with next generation sequencing, personalized therapy has evolved exponentially as a number of oncogenes in NSCLC have been described, with differing characteristics, therapeutics and prognostic implications.

A platform for personalized medicine has developed, with many clinical trials investigating and reporting on the efficacy of treatment not only for *EGFR* mutant NSCLC, but *ALK*, *ROS1*, *MET* rearranged, *MET* exon 14 skipping mutation positive, *NTRK* fusion positive, *BRAF* and *HER2* mutant, *RET* amplified and *KRAS* mutant NSCLC. Most of these gene signatures were not known or reported in the *EGFR* TKI literature reviewed here.

The NCI-Match Trial (Molecular Analysis for Therapy Choice) is an example of a precision medicine treatment clinical trial in which patients are assigned to receive treatment based on the genetic changes found in their tumours through genomic sequencing (29).

Conclusions

Given the current landscape of treatment for metastatic NSCLC, and advances into the era of empirical immunotherapy and molecular therapeutics, there is little enthusiasm nor rationale for the further investigation of EGFR TKI therapy alone or in combination in *EGFR*-wt NSCLC.

Furthermore, the definition of ‘*EGFR*-wt’ has certainly evolved in recent years, so the past literature must be interpreted cautiously, understanding that study data have come from heterogeneous patient populations largely predating reflex modern molecular testing for the presence of an *EGFR* mutation or more complex molecular panels used widely today.

The several large scale clinical trials investigating the efficacy of EGFR TKIs used empirically in mostly unselected patients have resulted in some statistically significant results to argue in the past for their use. But with the improved negative patient selection in more recent trials (by using wider molecular profiling to exclude patients with oncogene drivers), and the safety and substantial clinical efficacy observed with immunotherapy in these better defined populations for empirical therapy; the clinical significance of the modest benefits seen with EGFR TKIs in this setting must be questioned. Further investigation in to this issue is not of clinical interest nor relevant in the current lung cancer treatment paradigm.

Acknowledgements

None.

Footnote

Conflicts of Interest: The authors have no conflicts of interest to declare.

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Cite this article as: Itchins M, Clarke S, Pavlakis N. Do EGFR tyrosine kinase inhibitors (TKIs) still have a role in EGFR wild-type pre-treated advanced non-small cell lung cancer (NSCLC)?—the shifting paradigm of therapeutics. *Transl Lung Cancer Res* 2018;7(Suppl 1):S39-S45. doi: 10.21037/tlcr.2018.01.06