Efficacy of EGFR Tyrosine Kinase Inhibitors in Non-Small-Cell Lung Cancer Patients with/without EGFR-Mutation: Evidence Based on Recent Phase III Randomized Trials

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Background: EGFR mutation might be a predictive factor for applying EGFR-tyrosine kinase inhibitors (EGFR-TKIs, including gefitinib, erlotinib and afatinib) in non-small-cell lung cancer (NSCLS) patients. Thus, it is necessary to pool previous trials to compare the effect of EGFR-TKIs versus cytotoxic chemotherapy in EGFR mutation positive (mut+) and negative (mut–) patients.

Material/Methods: This study identified 8 first-line and 9 second-line phase III trials in databases. Hazard ratio (HR) was pooled to assess the risk of progression-free survival (PFS), and overall survival (OS), while odds ratio (OR) was pooled to assess objective response, disease control, and toxicity of EGFR-TKIs verses chemotherapy.

Results: In EGFR mut+ patients, EGFR-TKIs were associated with significantly lower risk of disease progression in the first-line setting, but this trend was only observed in the gefitinib group, not in the erlotinib group in the second-line setting. In EGFR mut– patients, gefitinib and erlotinib had significantly higher risk of disease progression in first-line and second-line setting, respectively. Compared with chemotherapy, the effects of EGFR-TKIs on OS in both first-line and second-line settings were not evident. Regarding toxicity, EGFR-TKIs had significantly higher risk of rash and lower hematological toxicity compared with chemotherapy.

Conclusions: All of the 3 EGFR-TKIs and gefitinib alone regimens had better effects in prolonging PFS in EGFR mut+ patients in first-line and second-line setting, respectively, but chemotherapy seemed more effective in EGFR mut- patients than EGFR-TKIs. Therefore, accurate identification of EGFR mutation status is useful to decide on an appropriate regimen for treatment of NSCLC patients.

MeSH Keywords: Carcinoma, Non-Small-Cell Lung • Genes, erbB-1 • Meta-Analysis

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Background

The EGFR signaling pathway plays a critical role in regulating the development and progression of non-small-cell lung cancer (NSCLC) [1,2]. Activation of this signal pathway might stimulate cancer cell proliferation and invasion and inhibit tumor cell apoptosis [2,3]. Gefitinib and erlotinib are the first generation of small and reversible molecular EGFR-tyrosine kinase inhibitors (TKIs) inhibiting the EGFR/ERBB1 receptor [4], while afatinib is the second generation of TKI inhibiting the EGFR/HER-1/ERBB1 and HER-2/ERBB2 receptors [5]. These three agents have been extensively used in patients with NSCLC [5].

Accumulating evidence shows that NSCLC patients with somatic mutations in certain genes have varied risks of lung cancer [6]. Somatic mutation in the region of EGFR genes encoding tyrosine kinase had higher response rate to EGFR-TKIs [7,8]. Therefore, EGFR-TKIs-based regimens, which were originally limited to use in patients who failed in previous platinum-based chemotherapy, have been gradually applied as first-line strategy for selective NSCLC patients who might have greater response to EGFR-TKIs. About 30–35% of NSCLC patients in Asia [9] and 10–15% of NSCLC patients in Europe [10] have EGFR activate mutation, in which the deletions in exon 19 and substitution of leucine-858 with arginine (L858R) in exon 21 of the EGFR kinase domain were the most common types [10,11].

Several phase III randomized controlled trials (RCTs) have been conducted to compare EGFR-TKIs (gefitinib, erlotinib, or afatinib) versus the standard doublet chemotherapy (platinum plus a new third-generation agent) as first-line treatment or EGFR-TKIs (gefitinib or erlotinib) versus a single new third-generation chemotherapy agent in second-line treatment for patients with advanced NSCLC. Most of the first-line treatments reported that EGFR-TKIs were associated with higher response rate and better progression-free survival (PFS) versus cytotoxic chemotherapy. In second-line treatment, the effect of EGFR-TKIs was not significant or was even associated with worse effect compared with cytotoxic chemotherapy. However, the small size of individual studies might not have sufficient statistical power to estimate the possible difference in experimental and control arms. Therefore, this study aimed to assess the efficacy of EGFR-TKIs versus chemotherapy in progression-free survival (PFS) and/or overall survival (OS) among patients with or without EGFR mutation.

Material and Methods

Search strategy

Trials were searched in MEDLINE, the Cochrane Central Register of Controlled Trials (CENTRAL), and EMBASE databases by using the following search terms and strategy: (“lung neoplasms” OR “non-small-cell lung cancer” OR “NSCLC”) AND (“gefitinib” OR “erlotinib” OR “afatinib” OR “tyrosine kinase inhibitor”) AND (“lung neoplasms” OR “non-small-cell lung cancer” OR “NSCLC”) AND (“randomized” OR “trial” OR “RCT”). The time range of the search was restricted to January 2000 to April 2014. No language restriction was applied for searching. Reference lists of included trials and other relevant reviews or meta-analyses were manually searched to avoid missing eligible trials.

Criteria for inclusion and exclusion

RCTs meeting the following criteria simultaneously were included for this meta-analysis: (1) phase III RCTs; (2) patients >18 years; and (3) studies compared EGFR-TKIs monotherapy vs. any chemotherapy in first-line or second-line trials for NSCLC patients. Trials were included regardless of publication status, date of publication, and language. Trials with a combination of chemotherapy and EGFR-TKIs in the experiment arm or merely with placebo in control arm were excluded.

Data extraction

Data extraction from original trials was independently performed by 2 authors (WQZ and TL). Disagreement was resolved by referring to original studies with a third author (HL) through group discussion. Data extracted include first author, year of publication, country/region in which the trials were conducted, regimen design in experiment and control group, and clinicopathological data including EGFR mutation, progression-free survival (PFS), overall response, disease control rate, and...
Table 1. Characteristics of first-line trials included.

| Study           | Country/region | Treatment               | Total no. pts | No. of EGFR+ pts (%) | No. of EGFR– pts (%) | No. of EGFR unknown pts (%) | Progression-free survival HR (95% CI) | Objective response OR (95% CI) | Disease control OR (95% CI) | Overall survival HR (95% CI) |
|-----------------|----------------|-------------------------|---------------|----------------------|----------------------|-------------------------------|--------------------------------------|-------------------------------|-----------------------------|-------------------------------|
| WJTOG3405 (2010) | Japan          | Gefitinib vs. Cis+D     | 172           | 172 (100)            | 0                    | 0                             | 0.49 (0.34–0.71)                  | 3.4 (1.6–7.4)                 | 3.8 (1.2–12.5)              | 1.64 (0.75–3.59)             |
| NEJ002 (2010)   | Japan          | Gefitinib vs. Car+Pa    | 228           | 228 (100)            | 0                    | 0                             | 0.32 (0.24–0.43)                  | 6.3 (3.6–11.2)                | 2.1 (1.0–4.6)               | 0.89 (0.63–1.26)             |
| IPASS (2009)    | East Asian     | Gefitinib vs. Car+Pa    | 1217          | 261 (21)             | 176 (15)             | 780 (64)                      | 0.74 (0.65–0.84)                  | 1.59 (1.25–2.01)              | 0.70 (0.54, 0.92)           | 0.90 (0.79–1.03)             |
| First-SIGNAL (2012) | Korean       | Gefitinib vs. Cis+G   | 309           | 43 (14)              | 54 (17)              | 212 (69)                      | 1.20 (0.94–1.53)                  | 1.46 (0.93–2.28)              | 0.56 (0.34, 0.94)           | 0.93 (0.72–1.20)             |
| Optimal (2011)  | China          | Erlotinib vs. Car+G     | 154           | 154 (100)            | 0                    | 0                             | 0.16 (0.10–0.26)                  | 8.6 (4.1–18.2)               | 5.8 (1.6–21.3)              | 1.07 (0.79–1.45)             |
| EURTAC (2012)   | Europe         | Erlotinib platinum+G/  | 173           | 173 (100)            | 0                    | 0                             | 0.37 (0.25–0.55)                  | 7.9 (3.8–16.4)                | 2.0 (1.0–3.9)               | 1.04 (0.65–1.66)             |
| LUX-Lung 3 (2012) | Global       | Aftatinib vs. Cis+Pe    | 354           | 354 (100)            | 0                    | 0                             | 0.58 (0.43–0.78)                  | 4.4 (2.6–7.3)                 | 2.1 (1.1–4.0)               | 1.12 (0.73–1.72)             |
| LUX-Lung 6 (2013) | Asian         | Aftatinib vs. Cis+G    | 364           | 364 (100)            | 0                    | 0                             | 0.28 (0.20–0.39)                  | 6.8 (4.1–11.2)                | 3.9 (2.1–7.3)               | 0.95 (0.68–1.33)             |

EGFR+ – EGFR mutation positive; EGFR– – EGFR mutation negative; Car – carboplatin; Cis – cisplatin; D – docetaxel; Pa – paclitaxel; Pe – pemetrexed; G – gemcitabine; N.A. – not available; OR – odd ratio; HR – hazard ratio; CI – confidential interval; Pts – patients.

overall survival (OS). In addition, severe drug toxicities (grade III or above adverse effects), including rash, fatigue/asthenia, diarrhea, vomiting/nausea, anemia, neutropenia, thrombocytopenia, and leukocytopenia were extracted for pooled analysis.

Statistical analyses

Cochrane Review Manager (version 5.2, Cochrane Collaboration, Copenhagen, Denmark) was used for statistical analysis. Hazard ratios (HRs) and the associated 95% confidence intervals (CIs) for PFS and OS and odds ratio (OR) and associated 95% CIs for objective response, disease control, and toxicity in original trials were extracted to compared the efficacy of EGFR-TKIs versus chemotherapy in first-line and second-line setting. In addition, subgroup analysis was performed by stratifying EGFR-TKIs within EGFR mut+ and EGFR mut– subgroups. If results of the trials were updated, the most recent OS data was used for analysis. The HR results were pooled by using inverse variance weighted method. A fixed-effects model was applied firstly to test heterogeneity (I² and p values of χ² Cochran Q test were used to detect heterogeneity across the different studies and between subgroups). If I² >50% or p<0.1, a random-effects model would be applied. P<0.05 was considered as significant in Z test of pooled results.

Results

Search results

The literature search identified 17 qualified phase III clinical trials. Among them, 8 studies compared gefitinib, erlotinib, or afatinib versus chemotherapy in first-line treatment and 9 compared gefitinib or erlotinib with chemotherapy in second-line treatment in patients with NSCLC. The search and screening process of qualified trials are described in Figure 1. The 8 first-line trials include IPASS [12], WJTOG3405 [13], NEJ002 [14] and First-SIGNAL [15] which compared gefitinib with chemotherapy,
Table 2. Characteristics of second-line trials included.

| Study        | Country/region | Treatment              | Total no. pts | No. of EGFR+ pts (%) | No. of EGFR- pts (%) | No. of EGFR unknown pts (%) | Progression-free survival HR (95% CI) | Response OR (95% CI) | Disease control OR (95% CI) | Overall survival HR (95% CI) |
|--------------|----------------|------------------------|---------------|----------------------|----------------------|-----------------------------|---------------------------------------|-----------------------|-----------------------------|-----------------------------|
| V-15-32 (2008) | Japan          | Gefitinib vs. D        | 489           | 31 (6)               | 26 (6)               | 432 (88)                    | 0.90 (0.72–1.13)                     | 2.14 (1.21–3.78)       | 1.08 (0.69–1.68)            | 1.12 (0.89–1.41)          |
| KCSG-LU08-01 (2012) | Korean | Gefitinib vs. Pe       | 135           | 33 (24)              | 38 (28)              | 64 (48)                     | 0.54 (0.37–0.79)                     | 1.10 (0.55–2.20)       | 1.02 (0.44–2.34)            | 0.80 (0.50–1.28)          |
| ISTANA (2010) | Korean          | Gefitinib vs. D        | 161           | N.A.                 | N.A.                 | N.A.                        | 0.73 (0.53–1.01)                      | 0.85 (0.44–1.61)       | 0.98 (0.48–2.00)            | 0.87 (0.61–1.24)          |
| Interest (2008) | Global         | Gefitinib vs. D        | 1466          | 44 (3)               | 253 (17)             | 1169 (80)                   | 1.04 (0.93–1.16)                     | 1.22 (0.82–1.84)       | N.A.                        | 1.02 (0.91–1.14)          |
| TITAN (2012)  | Global          | Erlotinib vs. Pe or D   | 424           | 11 (3)               | 149 (35)             | 264 (62)                    | 1.19 (0.97–1.46)                     | 1.27 (0.60–2.66)       | 0.70 (0.47–1.03)            | 0.96 (0.78–1.18)          |
| TAILOR (2012) | Italy           | Erlotinib vs. D        | 219           | 0                    | 219 (100)            | N.A.                        | 1.45 (1.08–1.95)                     | N.A.                  | N.A.                        | N.A.                      |
| PROSE (2014)  | Italy           | Erlotinib vs. Pe        | 263           | 14(5)                | 163 (62)             | 86 (33)                     | 1.27 (0.99–1.63)                     | 0.72 (0.30–1.70)       | 0.47 (0.28–0.77)            | 1.14 (0.88–1.48)          |
| HORG (2013)   | Greece          | Erlotinib vs. Pe        | 332           | 11(3)                | 112(34)              | 209(63)                     | 0.88 (0.73–1.06)                     | 0.77 (0.38–1.57)       | 0.67 (0.42–1.07)            | 0.99 (0.80–1.23)          |
| Delta (2013)  | Japan           | Erlotinib vs. D        | 301           | 56(19)               | 199(66)              | 46(15)                      | 1.22 (0.97–1.53)                     | 0.96 (0.53–1.76)       | N.A.                        | 0.91 (0.68–1.22)          |

Pooled OR (95% CI), p value

|                | Pooled HR (95% CI), p value | Pooled OR (95% CI), p value | Pooled OR (95% CI), p value | Pooled HR (95% CI), p value |
|----------------|-----------------------------|-----------------------------|-----------------------------|-----------------------------|
|                | 1.01 (0.87, 1.17), p=0.92   | 1.13 (0.89, 1.43), p=0.03   | 0.76 (0.59, 0.98), p=0.03   | 1.01 (0.93, 1.08), p=0.86   |

EGFR+ – EGFR mutation positive; EGFR- – EGFR mutation negative; Car – carboplatin; Cis – cisplatin; D – docetaxel; Pa – paclitaxel; Pe – pemetrexed; G – gemcitabine; N.A. – not available; OR – odd ratio; HR – hazard ratio; CI – confidential interval; Pts – patients.

OPTIMAL [16,17] and EURTAC [18] which compared erlotinib with chemotherapy and LUX-lung 3 [19] and LUX-lung 6 [20], which compared afatinib with chemotherapy. The 9 second-line trials include V-15-32 [21], KCSG-LU08-01 [22], ISTANA [23] and Interest [24] that compared gefitinib with chemotherapy and TITAN [25], TAILOR [26], PROSE [27], HORG [28] and Delta [29] that compared erlotinib with chemotherapy. The key information of the 8 first-line and 9 second-line trials are summarized in Tables 1 and 2, respectively. Among the 8 first-line trials, 6 only included patients with EGFR mutation [13,14,16–20]. In second-line trials, EGFR mutation status varied significantly. One study included only patients without mutation [26], 1 study did not report EGFR mutation status [23], while the other 6 had mixed patients with mutation, without mutation, or with unknown mutation status. Table 1 and 2 show the available HR data for PFS, OS, and OR data for objective response and disease control pooled. In the first-line setting, EGFR-TKIs were associated with better effect in prolonging PFS (HR 0.45, 95% CI 0.30–0.67, p<0.0001) and had a high ratio of objective response (OR 4.09, 95% CI 2.35–7.15, p<0.0001) and disease control (OR 1.86, 95% CI 1.01–3.41, p=0.05). But the effect in OS was similar to chemotherapy (HR 0.95, 95% CI 0.86–1.04, p=0.24) (Table 1). In second-line setting, EGFR-TKIs had similar effects in PFS (HR 1.01, 95% CI 0.87–1.17, p=0.02), objective response (OR 1.13, 95% CI 0.89–1.43, p=0.33), and OS (HR 1.01, 95% CI 0.93–1.08, p=0.86) as chemotherapy, but the effect in disease control was significantly worse than chemotherapy (OR 0.76, 95% CI 0.59–0.98, p=0.03) (Table 2).

The effect of EGFR-TKIs vs. chemotherapy on PFS

Eight studies evaluated EGFR-TKIs as first-line agents (4 gefitinib, 2 erlotinib, and 2 afatinib studies) and 9 studies assessed EGFR-TKIs as second-line agent in comparison to chemotherapy. The treatment effect in different settings is given in Figure 2. Generally, EGFR-TKIs were associated with significantly better PFS compared with chemotherapy in first-line setting (HR 0.45, 95% CI 0.30–0.67, p<0.0001) (Figure 2A). No significant heterogeneity was observed in subgroup difference ($I^2=46.3$%). The trend of better PFS was consistent in the other settings.
However, this trend was not observed in second-line setting, and afatinib (HR 0.40, 95% CI 0.20–0.83, p=0.01) (Figure 2A).

Test for subgroup differences: Chi²=1.71, df=2 (P=0.42); I²=0%
Test for overall effect: Z=6.55 (P<0.00001)

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3 EGFR-TKI subgroups: gefitinib (HR 0.62, 95% CI 0.38–1.01, p=0.05), erlotinib (HR 0.25, 95% CI 0.11–0.56, p=0.0008), and afatinib (HR 0.40, 95% CI 0.20–0.83, p=0.01) (Figure 2A). However, this trend was not observed in second-line setting, in which both gefitinib and erlotinib were associated with similar PFS as chemotherapy (gefitinib vs. chemotherapy, HR 0.82, 95% CI: 0.63–1.05, p=0.11; erlotinib vs. chemotherapy, HR 1.17, 95% CI: 0.98–1.38, p=0.08) (Figure 2B).
In patients with EGFR mutations, EGFR-TKIs treatment was associated with significantly lower risk of disease progression as first-line setting (gefitinib: HR 0.43, 95% CI 0.34–0.55, P<0.00001; erlotinib: HR 0.24, 95% CI 0.11–0.55, p=0.0007; and afatinib: HR 0.38, 95% CI 0.17–0.87, p=0.02) (Figure 2C). However, in second-line setting, significantly reduced risk was observed in the gefitinib arm (HR 0.43, 95% CI 0.34–0.55, P<0.00001; erlotinib: HR 0.24, 95% CI 0.11–0.55, p=0.0007; and afatinib: HR 0.38, 95% CI 0.17–0.87, p=0.02) (Figure 2C). Three studies compared erlotinib vs. chemotherapy and the pooled data showed significantly higher risk in the erlotinib arm (HR 1.40, 95% CI 1.17–1.66, p=0.0002) (Figure 2F).

The effect of EGFR-TKIs vs. chemotherapy on OS

Generally, EGFR-TKIs had no significant benefits in OS compared with chemotherapy in both first-line (HR 0.95, 95% CI 0.86–1.04, p=0.24) (Figure 3A) and second-line setting (HR 1.01, 95% CI 0.93–1.08, p=0.86) (Figure 3B). No significant heterogeneity
was observed in subgroup difference in both first-line and second-line settings (P=0.0%). Sub-group analysis showed that all 3 agents had no obvious effects on OS compared with chemotherapy: gefitinib (HR 0.91, 95% CI 0.82–1.02, p=0.11), erlotinib (HR 1.06, 95% CI 0.82–1.37, p=0.65), and afatinib (HR 1.01, 95% CI 0.78–1.32, p=0.93) (Figure 3A). In second-line setting, subgroup analysis also showed gefitinib and erlotinib had similar effects as chemotherapy (gefitinib vs. chemotherapy, HR 1.01, 95% CI 0.92–1.12, p=0.77; erlotinib vs. chemotherapy, HR 1.00, 95% CI 0.88–1.12, p=0.94) (Figure 3B).

**Figure 3A–C.** Meta-analysis of the effect of EGFR-TKIs on OS in first-line and second-line settings. Comparison of the effect on OS between EGFR-TKIs and chemotherapy in first-line setting (A) and in second-line setting (B); Comparison of the effect on OS in mut+ patients in first-line setting (C) and in second-line setting (D); Comparison of the effect on OS in mut– patients in first-line setting (E) and in second-line setting (F).
In patients with EGFR mutations, EGFR-TKIs, including gefitinib, erlotinib, and afatinib, had no significant difference in OS compared with chemotherapy in first-line setting (gefitinib: HR 1.00, 95% CI 0.82–1.22, p=0.97; erlotinib: HR 1.06, 95% CI 0.82–1.37, p=0.65; and afatinib: HR 1.01, 95% CI 0.78–1.32, p=0.93) (Figure 3C). In second-line setting, the difference was also not significant (gefitinib: HR 1.36, 95% CI 0.30–6.29, p=0.69; erlotinib: HR 1.06, 95% CI 0.82–1.31, p=0.49; and afatinib: HR 1.01, 95% CI 0.86–1.52, p=0.36) (Figure 3E) or second-line setting (HR 1.01, 95% CI 0.77–1.31, p=0.97) (Figure 3F). The effect of erlotinib on OS was also not significant in second-line setting (HR 0.92, 95% CI 0.71–1.18, p=0.49) (Figure 3F).

### Drug toxicity of EGFR-TKIs vs. chemotherapy

Severe drug toxicities (grade III or above adverse effects) were extracted for pooled analysis. Compared with chemotherapy, the most common severe adverse effects of EGFR-TKIs were rash in both first-line (OR 24.54, 95% CI 6.81–88.47, p<0.00001) and second-line (OR 7.72, 95% CI 3.70–16.11, p<0.00001) setting (Table 3). However, EGFR-TKIs had significantly lower hematological toxicity compared with chemotherapy. The risks of [Table D](#) and [Table E](#) show the hazard ratios and 95% confidence intervals for various comparisons.

#### Table D

| Study or subgroup | Log[hazard ratio] | SE | Weight | Hazard ratio | Hazard ratio | Hazard ratio |
|------------------|------------------|----|--------|--------------|--------------|--------------|
|                  |                  |    |        | FL random, 95% CI | IV random, 95% CI | IV random, 95% CI |
| 1.1.1.1 (OS/Mut+/Secondline-Gefitinib) | -0.1863 | 0.1398 | 38.1% | 0.83 [0.41, 1.66] | 0.96 [0.80, 1.15] |
| INTEREST (2008) | 1.539 | 1.1814 | 35% | 4.66 [0.46, 47.21] | 4.66 [0.46, 47.21] |
| V-15-32 (2008) | 0.174 | 1.1705 | 36% | 1.19 [0.12, 11.68] | 1.19 [0.12, 11.68] |
| Subtotal (95% CI) | | | | 43.1% | 1.36 [3.36, 6.29] |
| Heterogeneity: Tau²=0.00; Chi²=0.96, df=3 (P=0.81); I²=0% | | | | |
| Test for overall effect: Z=1.10 (P=0.11) | | | | |
| Test for subgroup differences: Not applicable | | | | |

#### Table E

| Study or subgroup | Log[hazard ratio] | SE | Weight | Hazard ratio | Hazard ratio | Hazard ratio |
|------------------|------------------|----|--------|--------------|--------------|--------------|
|                  |                  |    |        | FL random, 95% CI | IV random, 95% CI | IV random, 95% CI |
| 1.1.1 (OS/Mut+/Firstline-Tarceva) | -0.044 | 0.0651 | 11.6% | 0.93 [0.12, 15.54] | 0.93 [0.12, 15.54] |
| INTEREST (2009) | 0.1655 | 0.0184 | 43.1% | 0.66 [0.34, 1.28] | 0.66 [0.34, 1.28] |
| TITAN (2012) | 0.174 | 1.1705 | 36% | 1.19 [0.12, 11.68] | 1.19 [0.12, 11.68] |
| Subtotal (95% CI) | | | | 58.3% | 0.63 [0.36, 1.11] |
| Heterogeneity: Tau²=0.00; Chi²=0.72, df=2 (P=0.49); I²=0% | | | | |
| Test for overall effect: Z=1.10 (P=0.11) | | | | |
| Test for subgroup differences: Not applicable | | | | |

#### Table F

| Study or subgroup | Log[hazard ratio] | SE | Weight | Hazard ratio | Hazard ratio | Hazard ratio |
|------------------|------------------|----|--------|--------------|--------------|--------------|
|                  |                  |    |        | FL random, 95% CI | IV random, 95% CI | IV random, 95% CI |
| 1.1.2 (OS/Mut+/Secondline-Gefitinib) | -0.508 | 0.0212 | 1.3% | 0.60 [0.12, 3.00] | 0.60 [0.12, 3.00] |
| INTEREST (2008) | 0.0384 | 0.0199 | 45.5% | 1.02 [0.78, 1.33] | 1.02 [0.78, 1.33] |
| V-15-32 (2008) | 0.1655 | 0.1664 | 81.0% | 1.19 [0.86, 1.62] | 1.19 [0.86, 1.62] |
| Subtotal (95% CI) | | | | 100% | 1.01 [0.87, 1.17] |
| Heterogeneity: Tau²=0.00; Chi²=0.00, df=1 (P=0.94); I²=0% | | | | |
| Test for overall effect: Z=1.04 (P=0.30) | | | | |
| Test for subgroup differences: Not applicable | | | | |

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**Figure 3D–F.** Meta-analysis of the effect of EGFR-TKIs on OS in first-line and second-line settings. Comparison of the effect on OS between EGFR-TKIs and chemotherapy in first-line setting (A) and in second-line setting (B); Comparison of the effect on OS in mut+ patients in first-line setting (C) and in second-line setting (D); Comparison of the effect on OS in mut− patients in first-line setting (E) and in second-line setting (F).
anemia, neutropenia, thrombocytopenia, and leukocytopenia were all significantly lower than in the chemotherapy group in both first-line and second-line settings (Table 3).

### Discussion

Platinum-based doublet chemotherapy has been considered as the standard first-line treatment for advanced NSCLC cancer patients. However, the efficacy of this therapy is poor in certain groups of patients [30]. Many previous studies have tried to identify predictors of NSCLC development and how they can be used for better disease management [8,31]. The major strengths of this study include comprehensive review of the most recent phase III trials and stratified trials into gefitinib, erlotinib, and afatinib subgroups, rather than assuming these 3 agents have the same efficacy and to pool them into 1 group. Therefore, the exact effects of the 3 agents in EGFR mut+ and mut– patients in both first-line and second-line settings are clearly demonstrated.

This study pooled the most recent clinical results based on 8 first-line phase III trials and 9 second-line phase III trials involving 6761 patients, and addressed the most clinically important molecular factor (EGFR mutation) relevant to the treatment. For EGFR mut+ patients, this meta-analysis found that all of the 3 EGFR-TKIs had a considerable advantage in PFS over chemotherapy in first-line setting. In second-line setting, previous studies only compared gefitinib or erlotinib versus chemotherapy, and subgroup analysis showed that gefitinib was superior to chemotherapy. In EGFR mut- patients, a superior effect of EGFR-TKIs over chemotherapy was not observed. In first-line setting, although only 2 studies [12,15] examined the effect of EGFR-TKI, typically, gefitinib in EGFR mut– NSCLC patients, their findings were consistent and confirmed better efficacy of chemotherapy over gefitinib. In second-line setting for EGFR mut- patients, gefitinib did not present better effect, while erlotinib had worse effect than chemotherapy. Currently, available therapy for patients who failed in first-line treatment includes targeted therapy or further chemotherapy.

### Table 3. Meta-analysis of adverse effects, EGFR-TKIs vs. chemotherapy.

| Grade 3/4 adverse effects | Number of studies | Pooled OR (95%CI) | I² | P-H | P |
|---------------------------|-------------------|-------------------|----|-----|---|
| **Firstline setting**     |                   |                   |    |     |   |
| Rash                      | 5                 | 24.54 [6.81, 88.47] | 0% | 0.65 | <0.00001 |
| Fatigue/Asthenia          | 5                 | 0.23 [0.11, 0.45]  | 11%| 0.34 | <0.0001  |
| Diarrhea                  | 5                 | 13.96 [3.81, 51.14]| 0% | 0.65 | <0.0001  |
| Vomiting/Nausea           | 4                 | 0.18 [0.03, 1.28]  | 80%| 0.02 | 0.09     |
| Aneamia                   | 5                 | 0.07 [0.03, 0.19]  | 33%| 0.2  | <0.00001 |
| Neutropenia               | 5                 | 0.01 [0.00, 0.03]  | 0% | 0.55 | <0.00001 |
| Thrombocytopenia          | 4                 | 0.02 [0.01, 0.09]  | 0% | 0.68 | <0.00001 |
| Leucocytopenia            | 3                 | 0.02 [0.01, 0.06]  | 0% | 0.45 | <0.00001 |
| **Secondline setting**   |                   |                   |    |     |   |
| Rash                      | 7                 | 7.72 [3.70, 16.11] | 3% | 0.41 | <0.00001 |
| Fatigue/Asthenia          | 6                 | 0.42 [0.17, 1.04]  | 56%| 0.04 | 0.06     |
| Diarrhea                  | 6                 | 0.98 [0.57, 1.67]  | 0% | 0.73 | 0.94     |
| Vomiting/Nausea           | 8                 | 0.71 [0.43, 1.18]  | 0% | 0.43 | 0.19     |
| Aneamia                   | 4                 | 0.54 [0.30, 0.96]  | 0% | 0.66 | 0.04     |
| Neutropenia               | 7                 | 0.03 [0.01, 0.06]  | 63%| 0.01 | <0.00001 |
| Thrombocytopenia          | 3                 | 0.10 [0.01, 0.78]  | 0% | 0.77 | 0.03     |
| Leucocytopenia            | 2                 | 0.02 [0.00, 0.78]  | 92%| 0.0006| 0.04     |

CI – confidence interval; OR – odd ratio; P – p value; P-H – P value of Q for heterogeneity test; I² – 0–25%, no heterogeneity; 25–50%, modest heterogeneity; 50% or above, high heterogeneity; fixed effects model was used when I² <50%; random effects model was used when I² ≥50%.
Traditionally, examination of EGFR mutation status is considered unnecessary and time-consuming. However, this meta-analysis demonstrates that in second-line setting, gefitinib had better PFS in EGFR mut+ patients than chemotherapy, while erlotinib had worse PFS in EGFR mut– patients. Even when used as first-line agents, EGFR-TKIs should be recommended to patients with EGFR mutation. Therefore, identifying EGFR mutation status is quite helpful to guide therapeutic regimen development. Based on these findings, it can be concluded that EGFR mutation is a biomarker to predict the benefit of EGFR-TKIs in both first-line and second-line settings.

Even with over 2791 and 3790 patient cases in first-line and second-line settings, respectively, this study failed to demonstrate the OS advantage of EGFR-TKIs over chemotherapy. The most likely reason is the high crossover rate after progression. In both first-line and second-line settings, none of the trials prohibited patients from crossing over to the other treatment arm. For example, in the first-line trial NEJ002, most of the patients randomly assigned to either gefitinib or chemotherapy had subsequent treatment. In the chemotherapy group, 94.6% of patients had second-line gefitinib therapy to control disease progression [32]. In the second-line INTEREST trial, 31% of patients in the gefitinib arm had docetaxel as subsequent therapy, while in the docetaxel arm, 37% of patients had subsequent EGFR-TKI therapy [24]. In addition, in the V-15-32 trial, 36% of patients in the gefitinib arm had docetaxel as subsequent therapy, while 53% of patients in the docetaxel arm had subsequent gefitinib therapy [33]. One recent review indicated that PFS advantage is unlikely to be linked with OS advantage due to the impact of subsequent salvage therapy. Prolongation of survival due to salvage therapy after disease progression limits the use of OS as an indicator of therapeutic efficiency [34].

Regarding severe drug toxicity, this study showed that although EGFR-TKIs produced more rashes, their risks of hematological adverse effects were significantly lower compared with chemotherapy. Rash can be properly managed through supportive care and protocol-defined dose reductions. The rarity of hematological events EGFR-TKIs contributes to substantial improvements in overall health status and quality of life of the patients compared with chemotherapy. This study had several limitations. Firstly, EGFR mutation status was not fully examined in original studies. In second-line trials, 2431 out of 3790 (64%) patients had EGFR mutation status unidentified. Therefore, treatment efficacy in mut+ and mut– patients was only estimated based on a relatively small proportion (36%) of total patients. Secondly, the methods used to detect EGFR mutation were not consistent in the trials involved. Direct sequencing, PCR clamp, and amplification refractory mutation system were used in different trials. Since different methods have different sensitivities in detecting mutations, the possible bias associated with detection methods may directly influence the accuracy of pooled results.

Conclusions

All of the 3 EGFR-TKIs and gefitinib alone regimens had better effect in prolonging progression-free survival in EGFR mut+ patients in first-line and second-line settings, respectively, but chemotherapy seemed more effective in EGFR mut– patients than EGFR-TKIs. Therefore, accurate identification of EGFR mutation status is useful in deciding on an appropriate regimen for NSCLC patients.

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