

# EGFR Tyrosine Kinase Inhibitor Sequencing Revisited: From the Revival of Old Tools to the Integration of New Agents



Nicolas Girard, MD, PhD<sup>a,b,\*</sup>

In this issue of the *Journal of Thoracic Oncology*, Piccirillo et al.<sup>1</sup> report the results of the BEVERLY trial, a randomized phase 3 study that reveals the progression-free survival (PFS) benefit of bevacizumab to erlotinib as first-line treatment for Italian patients with metastatic NSCLC with common *EGFR* mutations. In this large study, after a median follow-up of 36.3 months, median investigator-assessed PFS was 15.4 months with erlotinib plus bevacizumab and 9.6 months with erlotinib alone (hazard ratio = 0.66, 95% confidence interval: 0.47–0.92). These results add further evidence to that provided by previously reported randomized studies, mostly conducted in Asian patients, that investigated combinations of first-generation *EGFR* TKIs and anti-angiogenic agents, including J025567, NEJ026, CTONG-1509, and other trials with erlotinib and bevacizumab,<sup>2–5</sup> and the RELAY trial with erlotinib and ramucirumab.<sup>6</sup> Taken together, these studies reported on median PFS ranging from 16.4 to 19.4 months.

This brings back to the fore the historical so-called sequencing strategy consisting of the use of first- or second-generation tyrosine kinase inhibitor (TKI) upfront, keeping a chance to use osimertinib as a subsequent therapy in the setting of *EGFR* T790M-related acquired resistance.<sup>7</sup> First, these studies, including BEVERLY, reveal the opportunity to improve the efficacy of first- or second-generation TKIs in the first-line setting to reach the median PFS reported with first-line osimertinib—18.9 months in the FLAURA trial<sup>8</sup>—considered as the current standard of care wherever available. Second, recent improvements in technologies and access to tissue- and blood-based genomic profiling may allow a higher proportion of patients to be identified with *EGFR* T790M at the time of disease progression, a challenge at the time the AURA-3 trial, revealing the benefit of osimertinib after the failure of first- or second-generation TKIs, was reported.<sup>9</sup> In this study, median PFS was 10.1 months on osimertinib, which may provide additional survival to patients in a sequencing approach. In historical real-world evidence studies, the rate of

patients eligible to osimertinib ranged from 25% to 40%,<sup>10</sup> whereas this was 47% in the control arm of FLAURA.<sup>8</sup> Indeed, in BEVERLY, 57% and 49% of patients in the control and experimental arm, respectively, did receive second-line osimertinib. Third, the use of anti-angiogenic agents may provide protection against central nervous system disease progression, a major challenge for patients with *EGFR*-mutant NSCLC as previously reported in RELAY, although the follow-up of this study remains limited so far.<sup>6</sup> This could actually not be explored in BEVERLY, given the exclusion of patients with brain metastases. Ultimately, the sequencing strategy may offer a prolonged chemotherapy-free time to the patients and more chance of deriving prolonged disease control along with first- and second-line treatments, which may lead to longer median overall survival (OS) as compared with that reported with frontline osimertinib of 38.6 months.<sup>8</sup> From real-world cohorts of patients who received a sequence of first- or second-generation *EGFR* TKI followed by osimertinib, median OS may be similar or even higher: in the GioTag trial, median OS was 37.6 months, rising up to 41.6 and 44.8 months in patients with *EGFR* exon 19 deletion and Asian patients, respectively.<sup>11</sup> Other studies are in line with such finding.<sup>12</sup> The promise of BEVERLY and other trials is

\*Corresponding author.

<sup>a</sup>Thoracic Oncology Service, Institut Curie, Institut du Thorax Curie Montsouris, Paris, France, and <sup>b</sup>UFR Simone Veil, Paris Saclay University, Université de Versailles Saint-Quentin-en-Yvelines (UVSQ), Versailles, France.

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Address for correspondence: Nicolas Girard, MD, PhD, Institut Curie, Institut du Thorax Curie Montsouris, 26 rue d'Ulm, 75248 Paris Cedex 05, France. E-mail: nicolas.girard2@curie.fr

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that OS could be even higher with the optimization of first-line through the combination with antiangiogenic agents. Still in BEVERLY, median OS in the combination arm was 33.3 months, despite the exclusion of patients with brain metastases.

Beyond first- and second-line sequencing, several new options are emerging for patients developing refractory disease, which add subsequent options in a sequencing strategy. Most current developments rely on the targeting of the most frequent molecular mechanisms of resistance to *EGFR* TKIs and/or on a deeper targeting of the *EGFR* signaling pathway. *MET* targeting is a major ongoing focus of clinical research after the failure of first- or second-line osimertinib and is based on *MET* TKIs, such as tepotinib or savolitinib, including on antibodies, such as the dual-specific *EGFR/MET* agent, amivantamab. Amivantamab, when combined with lazertinib, a third-generation *EGFR* TKI, was recently reported to produce response rates as high as 39% in heavily pretreated patients associated with PFS higher than 6 months.<sup>13</sup> Although *MET*- or *EGFR*-expressing tumors may have a better response to amivantamab, the combination is being developed without biomarker-driven selection. Another approach at the time of osimertinib resistance relies on the targeting of *EGFR*-mutant tumor cells with the HER3-specific antibody-drug conjugate patritumab-deruxtecan,<sup>14</sup> providing response rates and PFS in the same range. Interestingly, one striking point was that responses were observed across all the diverse mechanisms of *EGFR* TKI resistance, including those not directly related to HER3, such as *EGFR* C797S, *MET* or *HER2* amplification, and *BRAF* fusion. Meanwhile, new strategies that are currently developed in NSCLC overall are also being tested in *EGFR*-mutant NSCLC after the failure of *EGFR* TKIs, such as anti-TROP2 drugs; response rate to datopotamab deruxtecan in this setting is again in the 30% range, with a median duration on treatment of 13 months.<sup>15</sup> How these new options should be integrated in the treatment sequence remains a major question in the clinic; ultimately, adding new sequences in the patient management strategies is expected to increase the overall outcome. This remains a major concept to keep in mind as (1) those agents are now tested concurrently as first-line treatment, in comparison with osimertinib, and (2) *EGFR* inhibitors are also integrated in the adjuvant, neoadjuvant, or consolidation setting after the curative-intent treatment of nonmetastatic disease.

The BEVERLY study also stresses the question of positioning in the treatment sequence of antiangiogenic agents along with the disease evolution in *EGFR*-mutant NSCLC. Bevacizumab has mostly been used after the failure of TKIs, in combination with chemotherapy. The IMpower150 trial revealed, in a prespecified limited subset of 50 patients, the potential benefit of a

quadruplet with carboplatin, paclitaxel, atezolizumab, and bevacizumab in this setting.<sup>16</sup> Whether and how the early exposure to antiangiogenic agent modifies the natural history of the disease remain unclear. In BEVERLY, no further follow-up is planned, owing to severe acute respiratory syndrome coronavirus 2 pandemic, and sites of recurrence, including OS, will remain unknown. Of note, combination of osimertinib with bevacizumab was disappointing so far.<sup>17</sup> Similarly, the actual sequencing of chemotherapy in the treatment strategy for *EGFR*-mutant NSCLC is a matter of debate. Although it is currently being used in the refractory setting, several trials recently suggested a room for chemotherapy upfront in combination with *EGFR* TKIs.<sup>18,19</sup> FLAURA2 is an ongoing phase 3, randomized trial to evaluate efficacy and safety of first-line osimertinib with platinum-pemetrexed chemotherapy versus osimertinib monotherapy. MARIPOSA2 is a study of amivantamab and lazertinib in combination with platinum-based chemotherapy compared with platinum-based chemotherapy in patients with acquired resistance to osimertinib.

To conclude, whether combination strategies will ultimately provide more prolonged OS to patients, as compared with sequential approaches, is a major question in the clinic. A challenge is that clinical trials, such as BEVERLY, always focus on one line of treatment and may have limited follow-up; even if providing some information on subsequent therapies, these do not address the full sequencing of therapies. The publication of BEVERLY represents another pledge for prospective, adaptive, real-world studies that integrate historical and new options as soon as these become available and analyze treatment sequences along with clinical and molecular evolution of the disease.

## CRedit Authorship Contribution Statement

**Nicolas Girard:** Conceptualization, Validation, Formal analysis, Writing—original draft, Writing—review and editing, Project administration.

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