

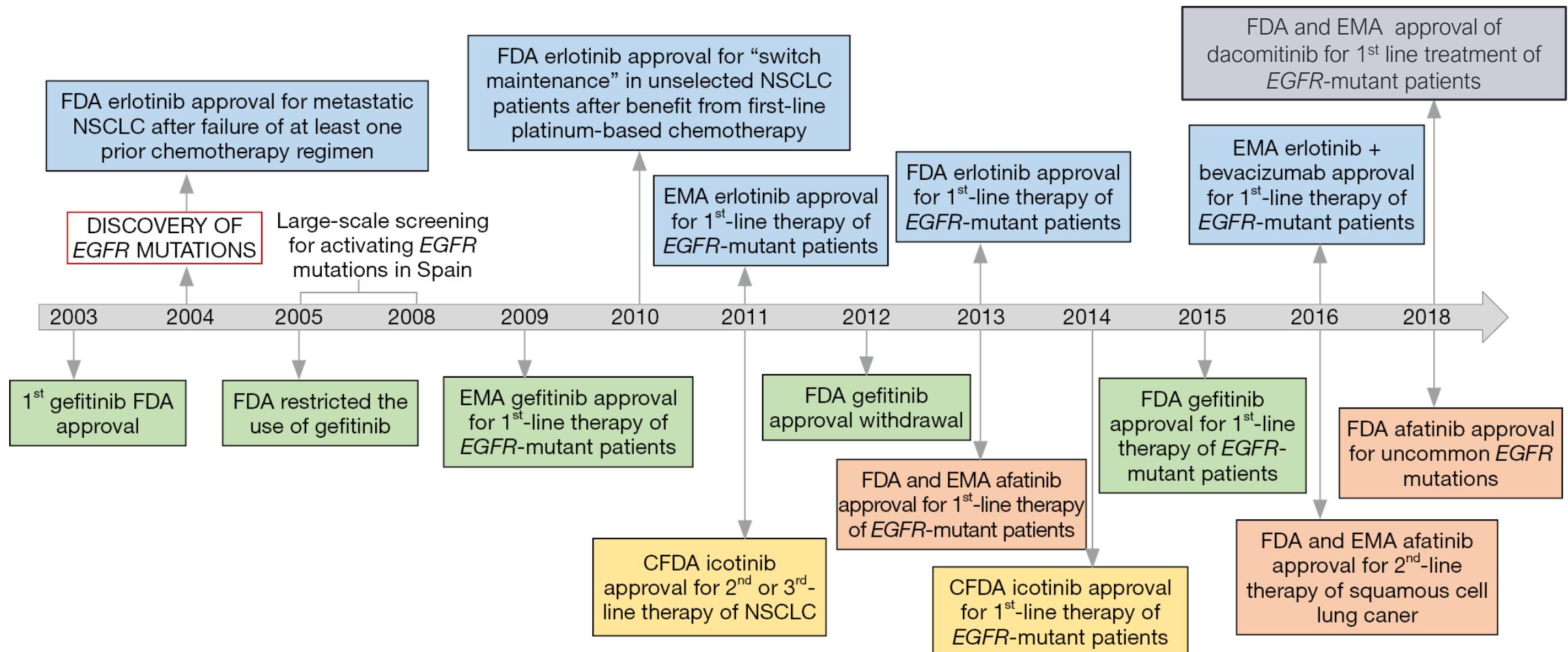
Bevacizumab plus erlotinib versus erlotinib alone in Japanese patients with advanced, metastatic, EGFR-mutantnon-small-cell lung cancer (NEJ026): overall survival analysis of an open-label, randomized, multicenter, phase 3 trial

Dr Arvind Kumar

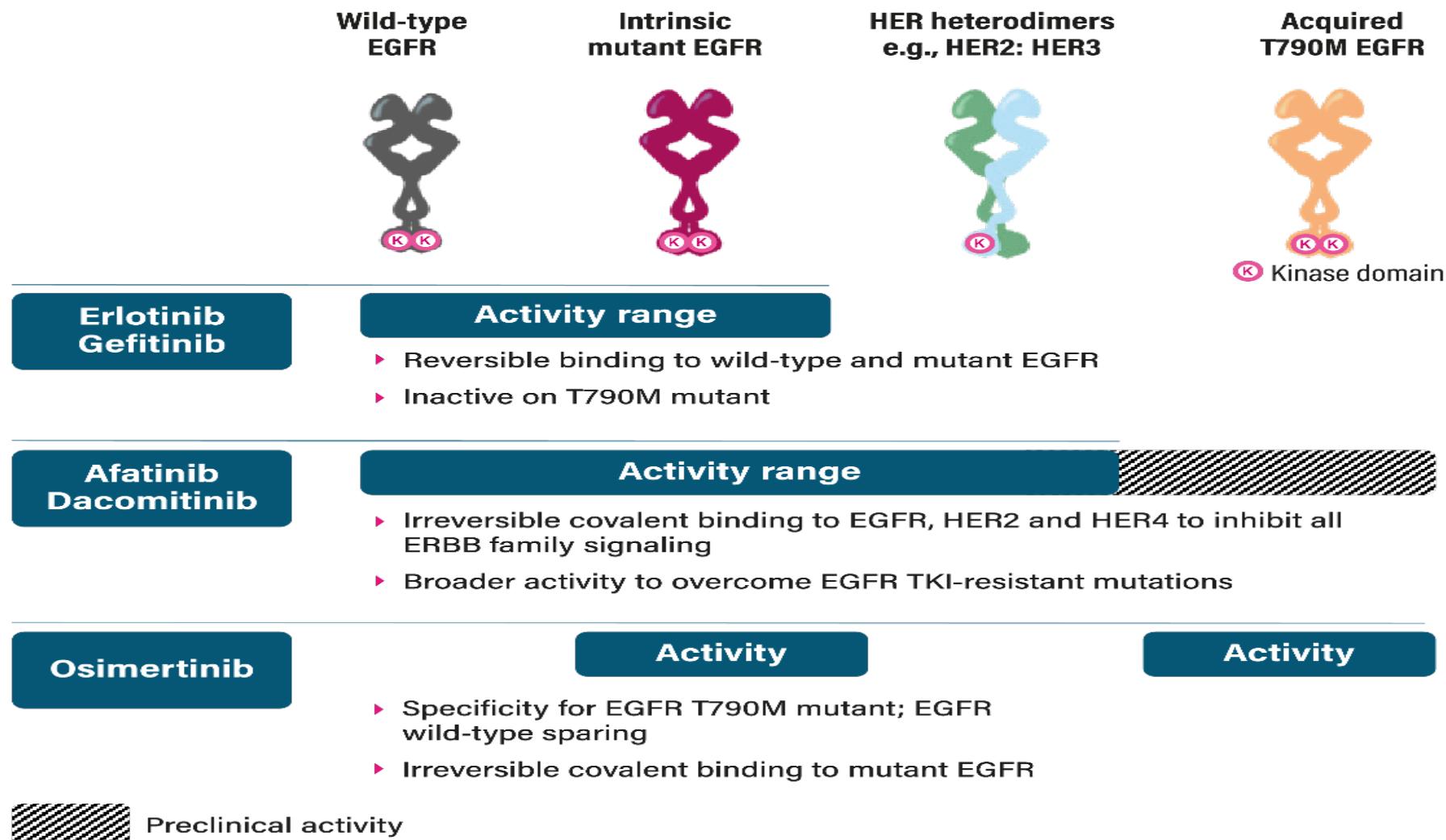
MD, DM (Medical Oncology AIIMS Delhi)

Director & Consultant Medical Oncologist at Buddha Cancer Centre, Patna.

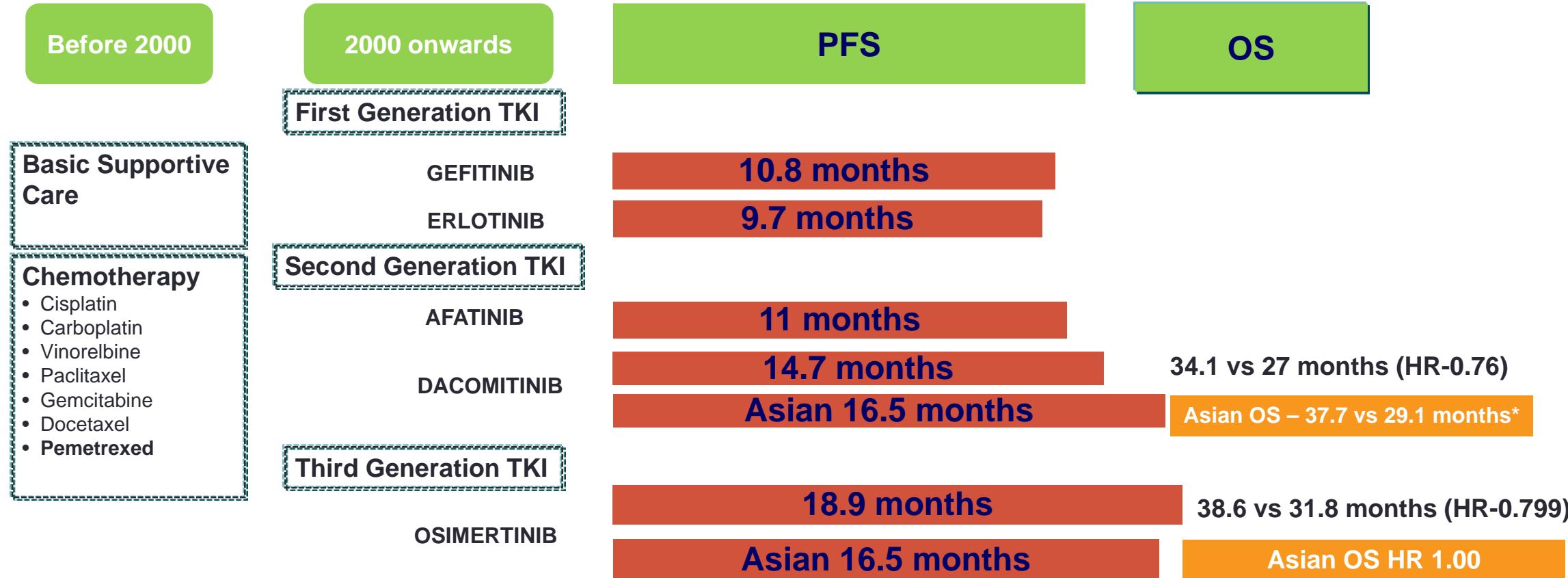
Personalized Therapy with EGFR TKIs: Timeline of Development



Overview of activity range of first-, second- and third-generation EGFR tyrosine kinase inhibitors



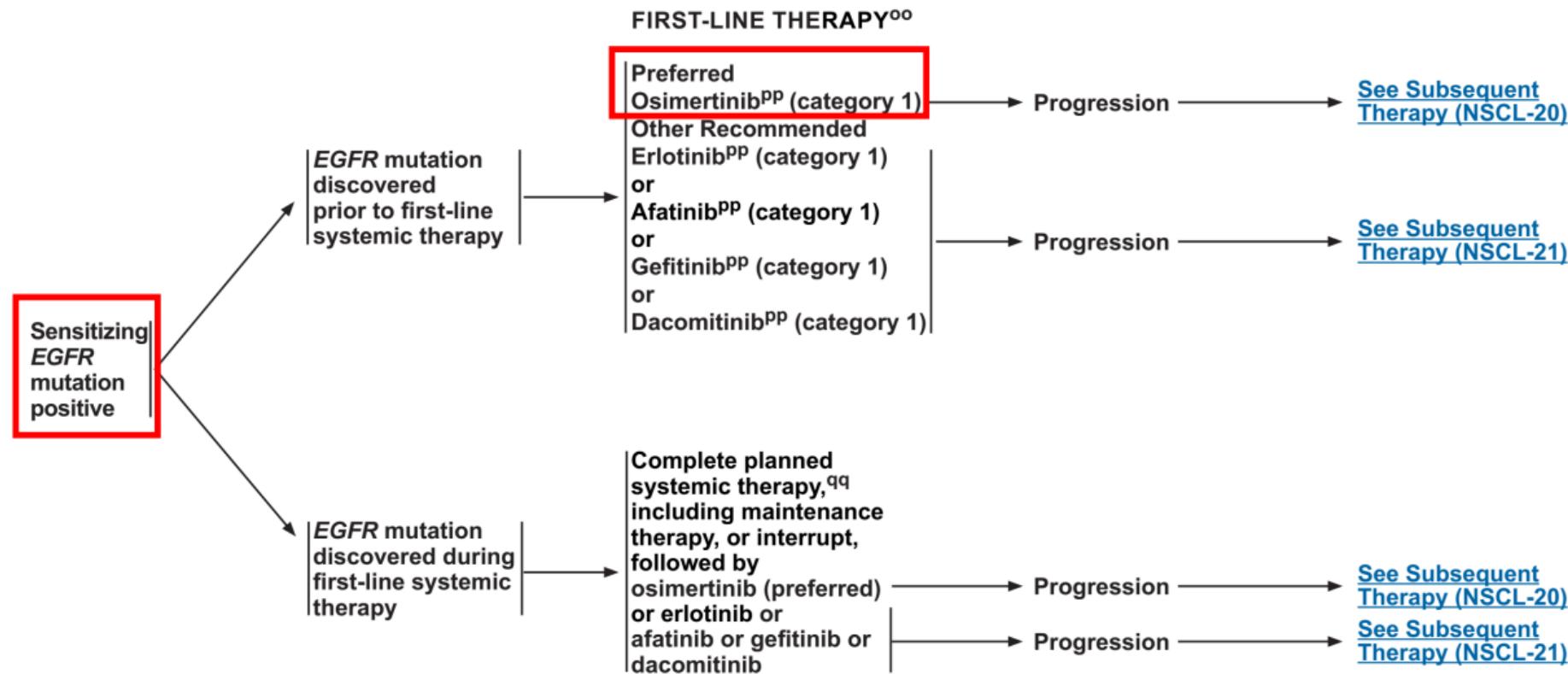
Treatment evolution in EGFR positive NSCLC



Data for understanding only, no cross-trial comparison

NSCLC with EGFR mutation treatment

SENSITIZING EGFR MUTATION POSITIVE^{jj}



EGFR-TKI Plus anti-VEGF/VEGFR

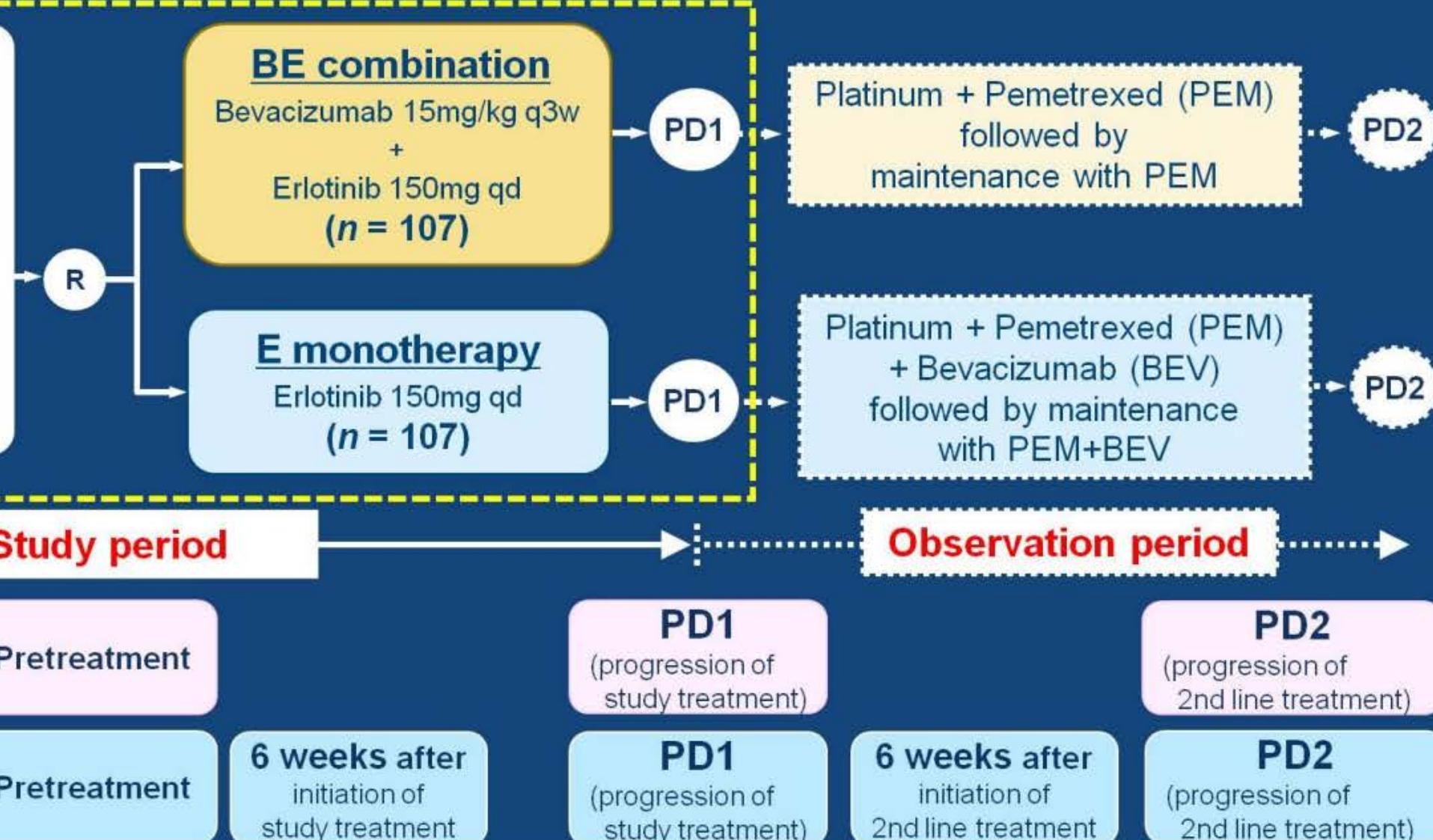
Phase III study comparing bevacizumab plus erlotinib to erlotinib in patients with untreated NSCLC harboring activating EGFR mutations: NEJ026

Furuya N, Fukuhara T, Saito H, Watanabe K, Sugawara S, Iwasawa S, Tsuneyzuka Y, Yamaguchi O, Okada M, Yoshimori K, Nakachi I, Gemma A, Azuma K, Hagiwara K, Nukiwa T, Morita S, Kobayashi K, Maemondo M

Journal of clinical oncology, 2018, 36(15)

Study Design : NEJ 026 (Phase III study)

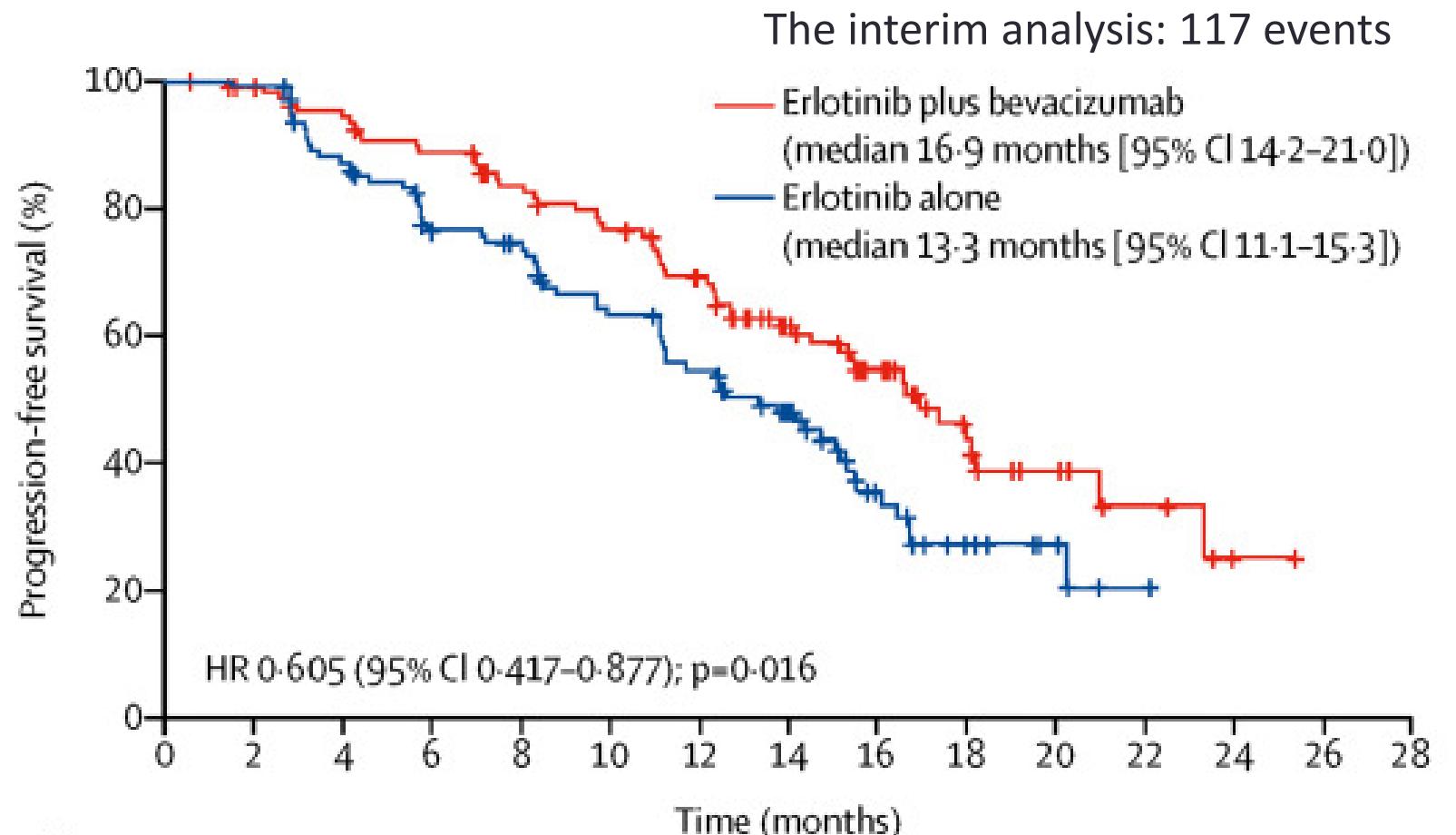
- Chemotherapy-naïve
 - Non-Sq NSCLC
 - Stage IIIB/IV or postoperative recurrence
 - Activating *EGFR* mutations
Ex19 del, Ex21 L858R
 - Asymptomatic CNS metastases allowed
- UMIN 000017069



Baseline Characteristics

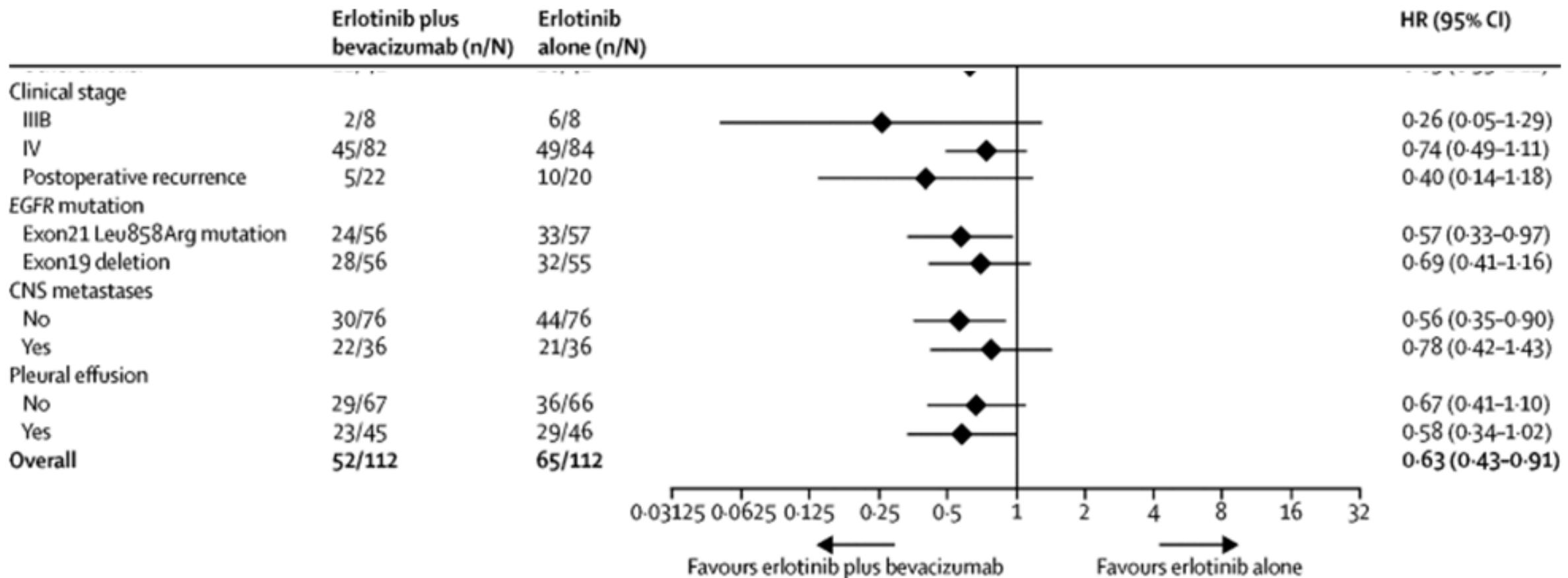
| | Erlotinib plus bevacizumab (n=112) | Erlotinib (n=112) |
|---|---------------------------------------|-------------------|
| Histopathological classification | | |
| Adenocarcinoma | 110 (98%) | 112 (100%) |
| Large cell carcinoma | 1 (1%) | 0 |
| Other | 1 (1%) | 0 |
| EGFR genomic aberration | | |
| Exon 19 deletion | 56 (50%) | 55 (49%) |
| Exon 21 Leu858Arg mutation | 56 (50%) | 57 (51%) |
| Clinical stage at screening | | |
| IIIB | 8 (7%) | 8 (7%) |
| IV | 82 (73%) | 84 (75%) |
| Postoperative recurrence | 22 (20%) | 20 (18%) |
| CNS metastases | | |
| Yes | 36 (32%) | 36 (32%) |
| No | 76 (68%) | 76 (68%) |

Primary endpoint: PFS by independent review

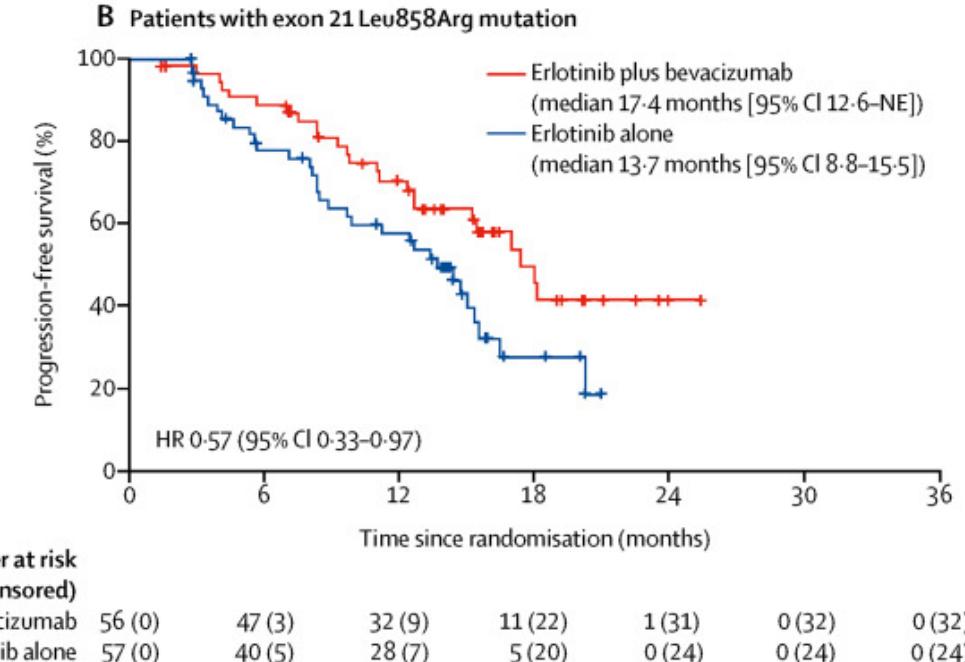
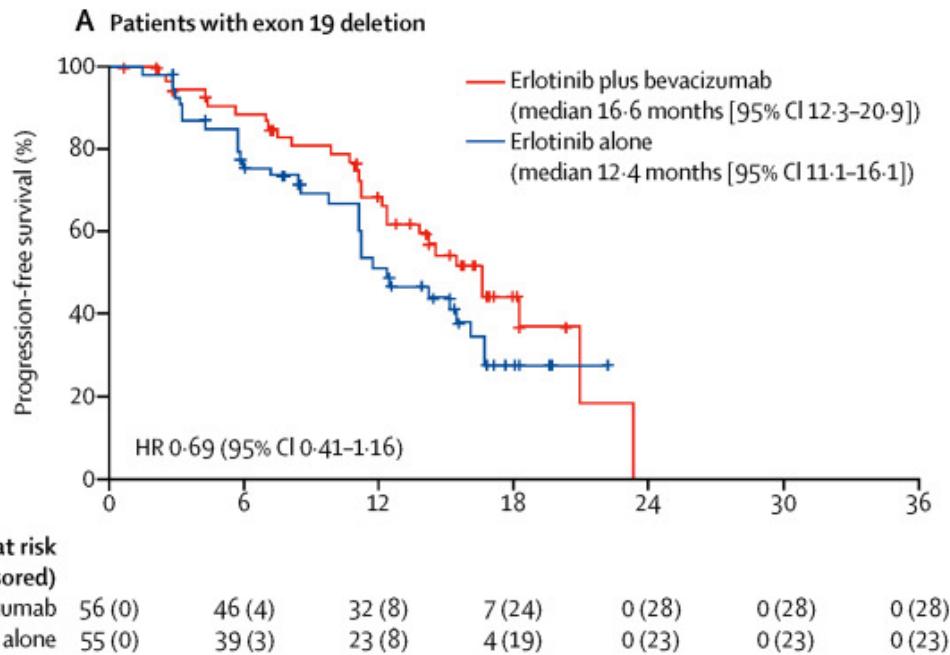


| | Number at risk (number censored) | | | | | | | | |
|--|-------------------------------------|---------|---------|---------|---------|---------|---------|--------|--------|
| | Erlotinib plus bevacizumab | 112 (0) | 100 (6) | 83 (7) | 64 (17) | 34 (35) | 10 (52) | 1 (59) | 0 (60) |
| | Erlotinib alone | 112 (0) | 94 (4) | 73 (12) | 51 (15) | 18 (34) | 5 (43) | 0 (47) | 0 (47) |

Subgroup analysis



PFS by EGFR mutation subtype

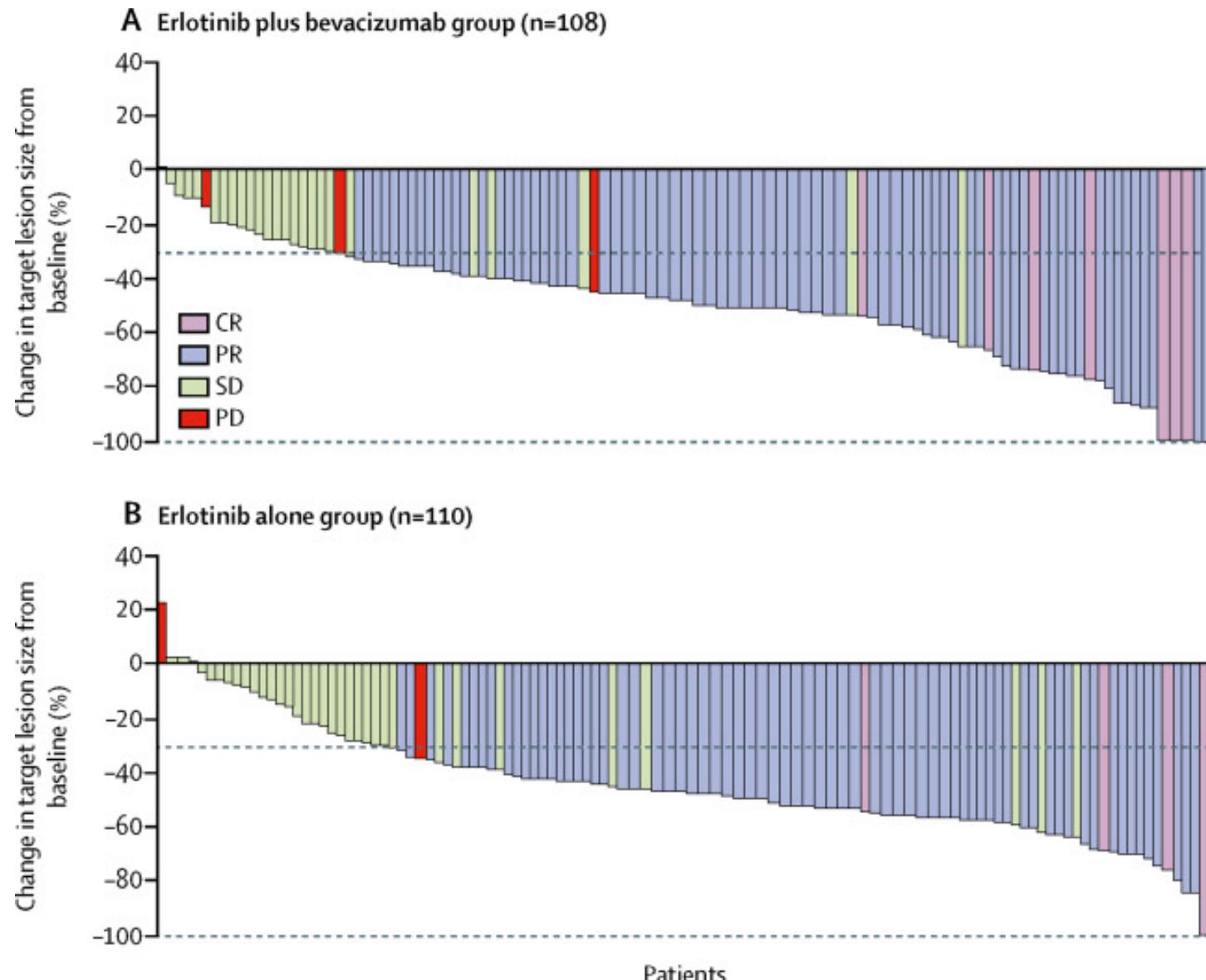


Response as per independent review committee's assessment

| | Erlotinib plus bevacizumab group (n=112) | Erlotinib alone group (n=112) | p value* |
|---------------------|--|-------------------------------|----------|
| Complete response | 8 (7%) | 4 (4%) | .. |
| Partial response | 73 (65%) | 70 (63%) | .. |
| Stable disease | 25 (22%) | 34 (30%) | .. |
| Progressive disease | 4 (4%) | 2 (2%) | .. |
| Not evaluable | 2 (2%) | 2 (2%) | .. |
| Objective response | 81 (72%; 63·1-80·4) | 74 (66%; 56·5-74·7) | 0·31 |
| Disease control | 106 (95%; 88·7-98·0) | 108 (96%; 91·1-99·0) | 0·52 |

Data are n (%). *p value for χ^2 test.

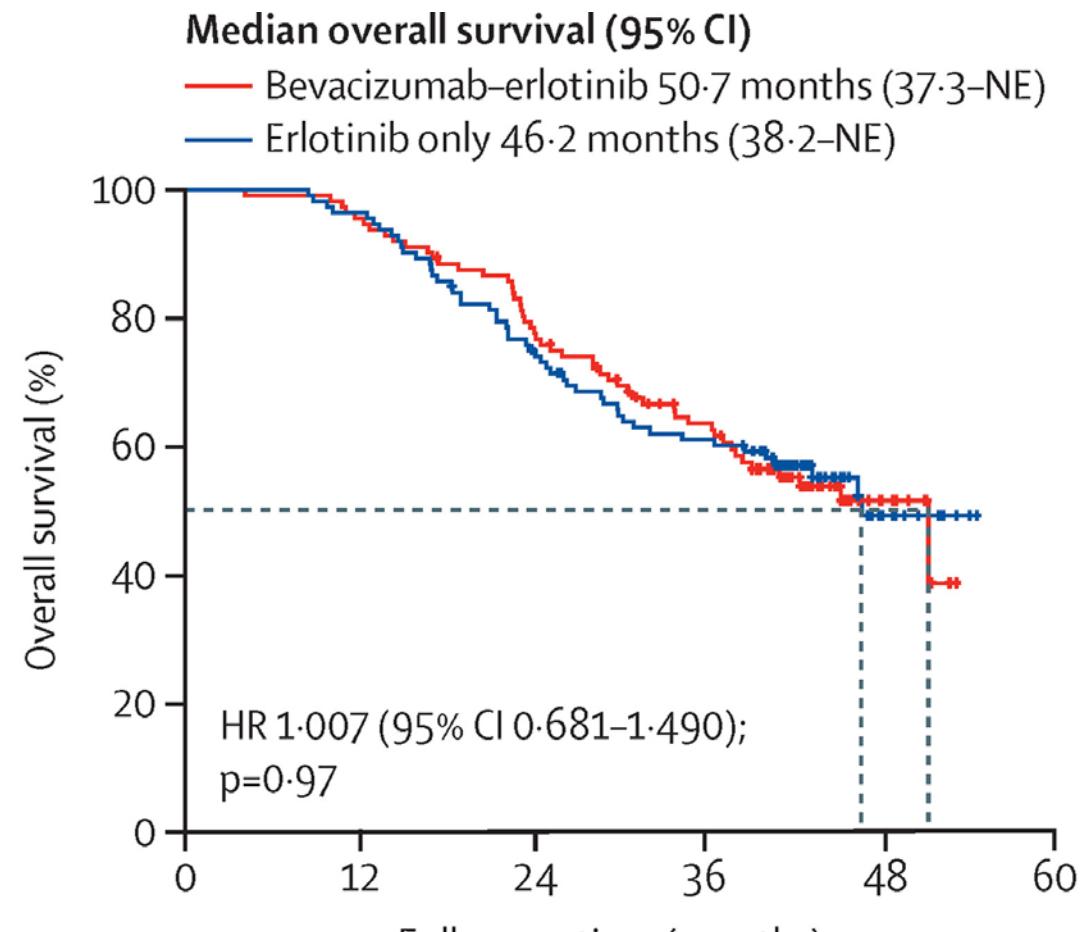
Best percentage change from baseline in target lesion size



Adverse events in the safety population

| | Erlotinib plus bevacizumab group (n=112) | | | | Erlotinib alone group (n=114) | | | |
|----------------------------|--|-----------|----------|---------|-------------------------------|-----------|----------|---------|
| | All | Grade 1–2 | Grade 3 | Grade 4 | All | Grade 1–2 | Grade 3 | Grade 4 |
| Rash | 98 (88%) | 75 (67%) | 23 (21%) | 0 | 99 (87%) | 75 (66%) | 24 (21%) | 0 |
| Diarrhoea | 53 (47%) | 47 (42%) | 6 (5%) | 0 | 47 (41%) | 45 (39%) | 2 (2%) | 0 |
| Proteinuria | 36 (32%) | 28 (25%) | 8 (7%) | 0 | 6 (5%) | 5 (4%) | 1 (1%) | 0 |
| Pulmonary haemorrhage | 2 (2%) | 2 (2%) | 0 | 0 | 0 | 0 | 0 | 0 |
| Non-pulmonary haemorrhage | 29 (26%) | 27 (24%) | 2 (2%) | 0 | 3 (3%) | 2 (2%) | 1 (1%) | 0 |
| Hypertension | 52 (46%) | 26 (23%) | 26 (23%) | 0 | 11 (10%) | 10 (9%) | 1 (1%) | 0 |
| Increased aminotransferase | 30 (27%) | 21 (20%) | 6 (5%) | 3 (3%) | 34 (30%) | 28 (25%) | 5 (4%) | 1 (1%) |
| Stomatitis | 23 (21%) | 22 (20%) | 1 (1%) | 0 | 12 (11%) | 11 (10%) | 1 (1%) | 0 |
| Paronychia | 17 (15%) | 15 (13%) | 2 (2%) | 0 | 18 (16%) | 15 (13%) | 3 (3%) | 0 |

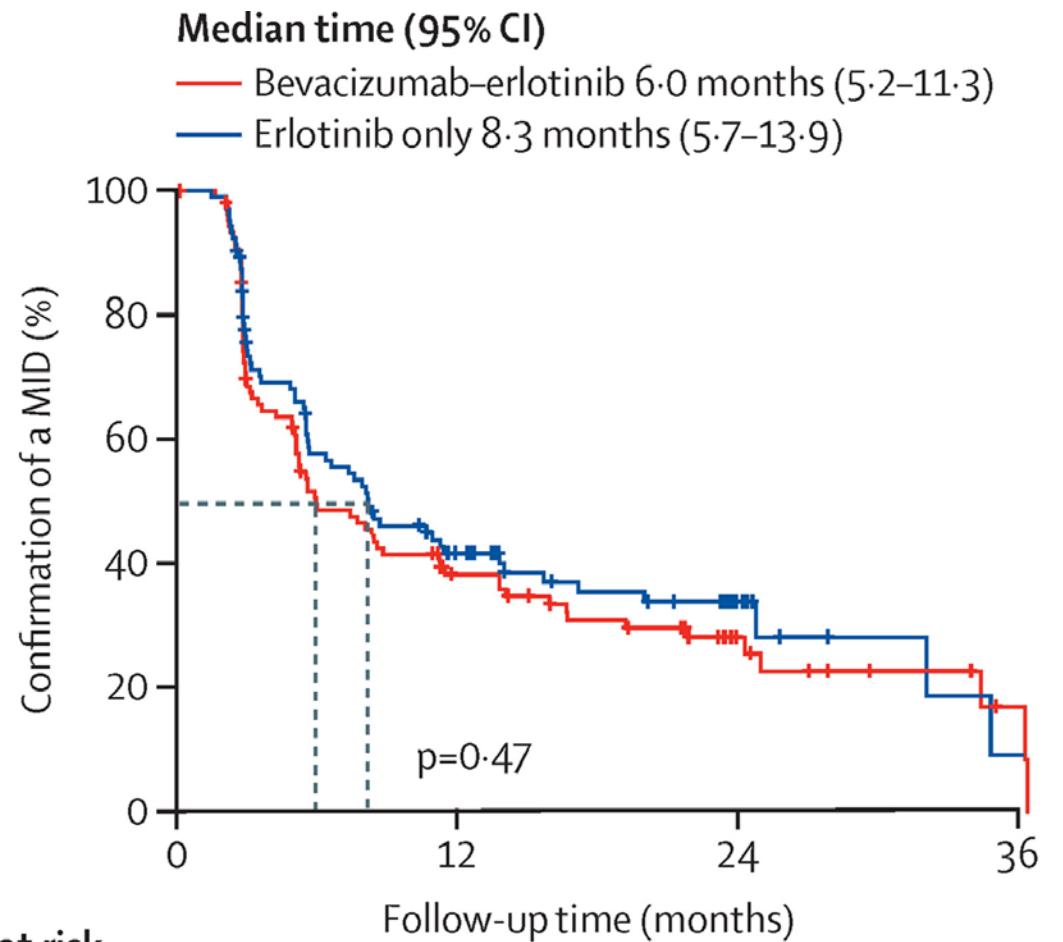
Overall survival



**Number at risk
(number censored)**

| | | | | | | |
|-----------------------|---------|---------|--------|--------|---------|--------|
| Bevacizumab–erlotinib | 112 (0) | 107 (0) | 85 (1) | 62 (9) | 12 (50) | 0 (61) |
| Erlotinib only | 112 (0) | 108 (0) | 81 (2) | 65 (4) | 10 (52) | 0 (62) |

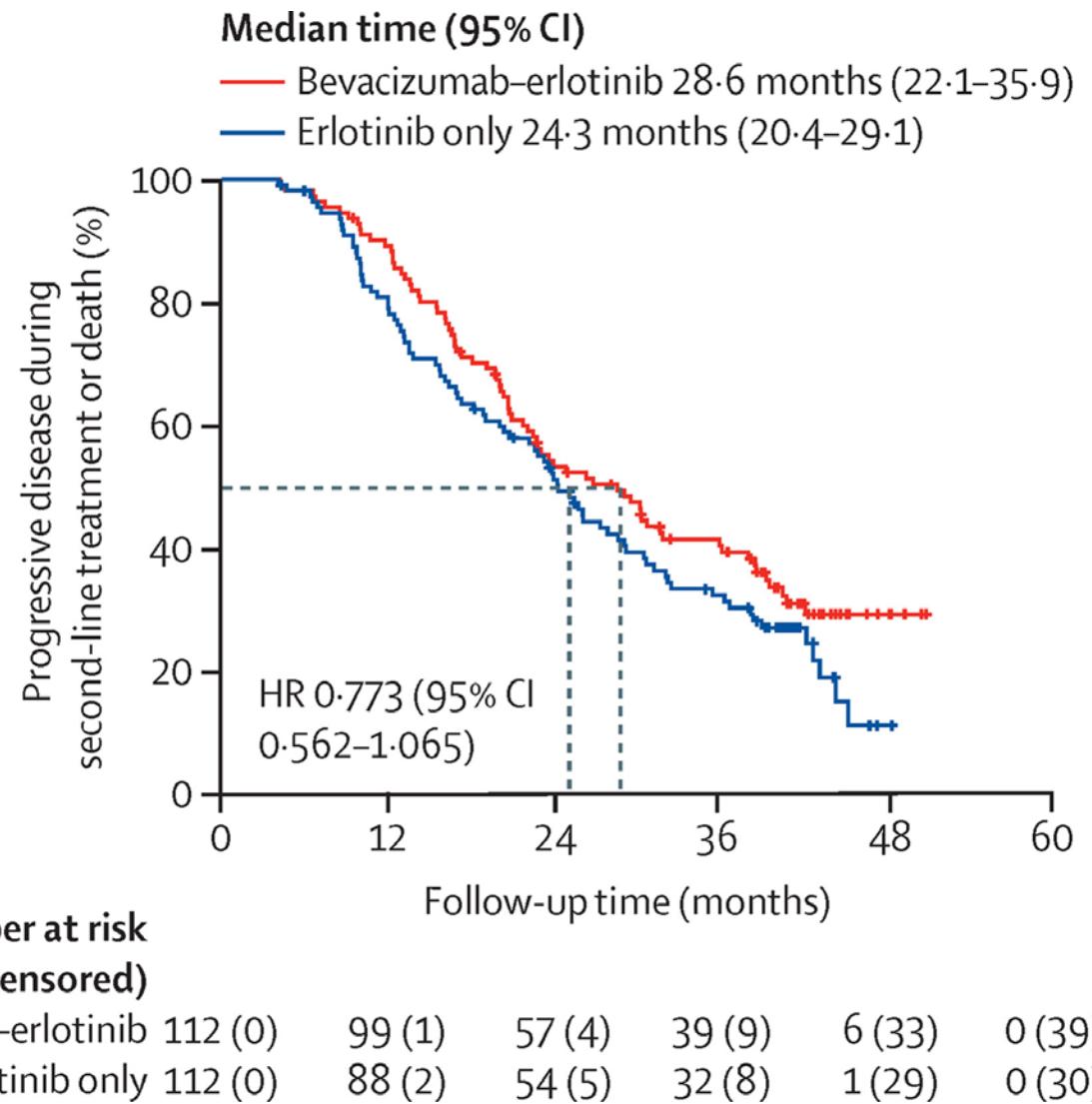
Quality of life with completed questionnaires



Number at risk
(number censored)

| | | | | |
|-----------------------|---------|---------|---------|--------|
| Bevacizumab-erlotinib | 108 (0) | 33 (12) | 12 (25) | 2 (32) |
| Erlotinib only | 104 (0) | 36 (11) | 13 (29) | 1 (38) |

Time from enrolment to progressive disease during second-line treatment or death



Conclusion

- Unmet Need for EGFR mutation NSCLC patients
- Despite PFS benefit this trial didn't show Significant OS benefit.
- The probable reasons for this might be second line treatment.
- Further Randomized trial like combination of first Generation TKI + CT+/- Bev may be planned in L858R subset only.

Thank You