







## 27 SEPTIEMBRE - 1 OCTUBRE 2019



Con la colaboración de:









# Terapias dirigidas (II)

Dra. Noemí Reguart

Con la colaboración de:



# **Targeted Therapies (Sunday 29, Monday 30)**



Poster Discussion – NSCLC, metastatic (ID 256)

Proffered Paper 2 – NSCLC, metastatic (ID 257)

#### EGFR+ Disease

- LBA85- Longitudinal circulating tumour DNA (ctDNA) monitoring for early detection of disease progression and resistance in advanced NSCLC in FLAURA (ID 2685). *Presenter Jhanelle E. Gray*
- **14800- CTONG 1509:** Phase 3 study of bevacizumab with or without erlotinib in untreated Chinese patients with advanced EGFR-mutated NSCLC (ID 3200). *Presenter Qing Zhou*

### ROS1/NTRK+ Disease

- **1485PD-** Safety and efficacy of **WX-0593** in ALK-positive or ROS1-positive non-small cell lung cancer (ID 6131). *Presenter Yuan-Kai Shi*
- **1488PD- Entrectinib** in Locally Advanced/Metastatic ROS1 and NTRK Fusion-Positive NSCLC: Updated Integrated Analysis of **STARTRK-2, STARTRK-1 and ALKA-372-001** (ID 4178). *Presenter Filippo Guglielmo M. De Braud*
- **1489PD-** Secondary **ROS1** mutations and lorlatinib sensitivity in crizotinib-refractory ROS1 positive NSCLC: results of the prospective **PFROST trial** (ID 4899). *Presenter Lorenza Landi*



# **Targeted Therapies (Sunday 29, Monday 30)**



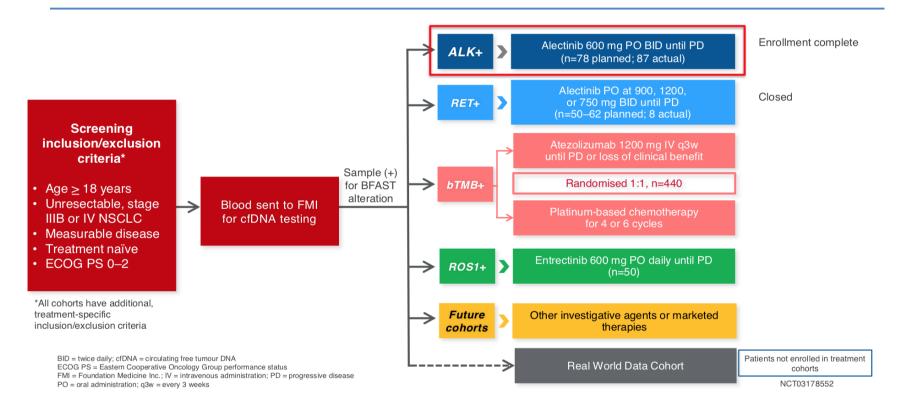
#### ALK+ Disease

- 1484PD- Final PFS, updated OS and safety data from the randomised, phase III ALEX study of alectinib (ALC) versus crizotinib (CRZ) in untreated advanced ALK+ NSCLC (ID 3850). Presenter Tony S.K. Mok
- **1486PD-** Exposure-response analyses of ALK-inhibitors **crizotinib** and **alectinib** in NSCLC patients (ID 1906). *Presenter Steffie L. Groenland*
- 1487PD- Intracranial and extracranial efficacy of lorlatinib in the post second-generation ALK tyrosine kinase inhibitor (TKI) setting (ID 2577). Presenter Steffie L. Groenland
- LBA81\_PR- Phase II/III Blood First Assay Screening Trial (BFAST) in patients (pts) with treatment-naïve NSCLC: initial results from the ALK+ cohort (ID 5026). Presenter Shingo Miyamoto. *Presenter Shirish M. Gadgeel*





## Study design







# Goal

Demonstrate consistency of benefit with alectinib in a population selected by blood-based NGS as opposed to tissue-based assay, using ALEX alectinib data as reference

- Unresectable, stage IIIB or IV NSCLC
- ALK+ by centralised blood screening only
- Treatment naïve
- ECOG PS 0-2
- Primary endpoint
   Confirmed ORR by investigator
- Exploratory endpoint Confirmed ORR by investigator for patients with baseline CNS metastases

Alectinib 600mg BID

Until PD,
toxicity,
withdrawal
or death

Secondary endpoints

By investigator By independent review facility

DOR ORR

PFS DOR

PFS

CNS = central nervous system; DoR = duration of response
ORR = overall response rate; PD = disease progression; PFS = progression-free survival
ECOG PS = Eastern Cooperative Oncology Group performance status; NGS = next generation sequencing

NCT03178552





### **Demographics and baseline characteristics**

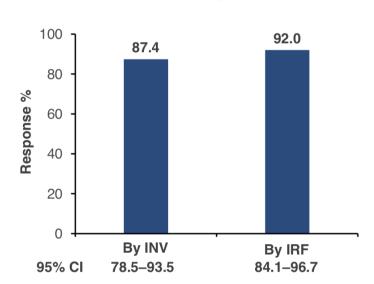
| Characteristic                        | ALK+ Cohort (N=87) | ALEX alectinib arm <sup>1</sup> (N=152) |
|---------------------------------------|--------------------|---|
| Median age, years (range)             | 55.0 (25-82)       | 58.0 (25-88)                            |
| Gender                                |                    |   |
| Male                                  | 35 (40)            | 68 (45)                                 |
| Female                                | 52 (60)            | 84 (55)                                 |
| Race, n (%)                           |                    |   |
| Asian                                 | 29 (33)            | 69 (45)                                 |
| Non-Asian                             | 58 (67)            | 83 (55)                                 |
| ECOG PS, n (%)                        |                    |   |
| 0-1                                   | 82 (94)            | 142 (93)                                |
| 2                                     | 5 (6)              | 10 (7)                                  |
| Smoking status, n (%)                 |                    |   |
| Active smoker                         | 5 (6)              | 12 (8)                                  |
| Past smoker                           | 32 (37)            | 48 (32)                                 |
| Never smoked                          | 50 (58)            | 92 (61)                                 |
| Disease state, n (%)                  |                    |   |
| IIIb                                  | 5 (6)              | 4 (3)                                   |
| IV                                    | 82 (94)            | 148 (97)                                |
| Histology, n (%)                      |                    |   |
| Adenocarcinoma                        | 81 (93)            | 136 (90)                                |
| Other                                 | 4 (5)              | 16 (10)                                 |
| Missing                               | 2 (2)              | 0 (0)                                   |
| CNS metastases by investigator, n (%) |                    |   |
| Yes                                   | 35 (40)            | 60 (40)                                 |
| No                                    | 52 (60)            | 92 (60)                                 |





#### **Results: Confirmed response (INV vs IRF)**

#### **Overall Response Rate**



| Me | dian | duration | of | fol | low-up: | 12.58 | months |
|----|------|----------|----|-----|---------|-------|--------|
|----|------|----------|----|-----|---------|-------|--------|

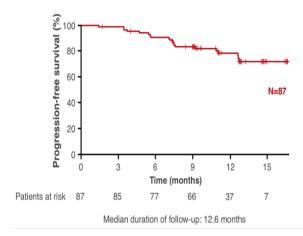
|                                      | INV<br>(N=87)                  | IRF<br>(N=87)                  |
|--------------------------------------|--------------------------------|--------------------------------|
| Complete Response, n (%)<br>95% Cl   | <b>0</b> (0.00–4.15)           | <b>11 (12.6)</b> (6.48–21.50)  |
| Partial Response, n (%)<br>95% Cl    | <b>76 (87.4)</b> (78.50–93.52) | <b>69 (79.3)</b> (69.29–87.25) |
| Progressive Disease, n (%)<br>95% CI | <b>1 (1.1)</b> (0.03–6.24)     | <b>1 (1.1)</b> (0.03–6.24)     |

ALEX confirmed ORR = 71.7% (95% 63.8–78.7)<sup>1</sup>





#### **Results: PFS by investigator**



| Patients with event, n (%) | 20 (23)       |
|----------------------------|---------------|
| Median PFS, months         | NE            |
| (95% CI)                   | (NE)          |
| 12-month PFS, %            | 78.38         |
| (95% CI)                   | (69.07–87.69) |
| ALEX <sup>1</sup>          | N=152         |
| 12-month PFS, %            | 68.4          |
| (95% CI)                   | (61.0–75.9)   |

#### **Conclusions**

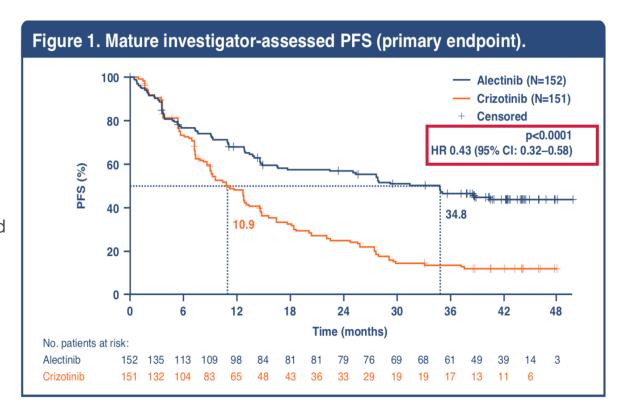
- BFAST is the first prospective trial to use blood-based NGS testing as the sole method
  of identifying actionable genetic alterations and assigning NSCLC patients to targeted or
  immunotherapy
- The primary endpoint of the ALK+ cohort was achieved with an investigator-confirmed ORR of 87.4% and an independent review facility confirmed ORR of 92.0%
- Key secondary endpoints: 6-month DoR 90.4%, 12-month PFS 78.38%
- The safety profile of alectinib was consistent with that established in previous phase III trials and post-marketing experience
- These results demonstrate the clinical utility of blood-based NGS as a method to inform clinical decision-making in ALK+ NSCLC



1484PD- Final PFS, updated OS and safety data from the randomised, phase III ALEX study of alectinib (ALC) versus crizotinib (CRZ) in untreated advanced ALK+ NSCLC (ID 3850). *Presenter Tony S.K. Mok* 



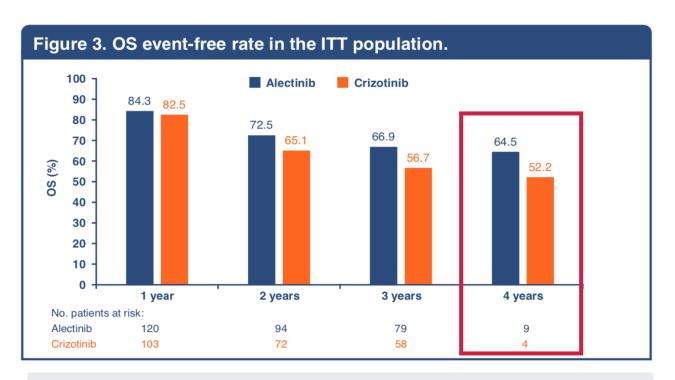
- Last update: PFS alectinib vs crizotinib HR 0.43, (95% CI: 0.32–0.58): Median 34.8 versus 10.9 months. OS immature: stratified HR 0.76, 95% CI: 0.50–1.15.2
- Present updated data median duration of follow-up 4 years (37.8 months with alectinib and 23.0 months with crizotinib)





1484PD- Final PFS, updated OS and safety data from the randomised, phase III ALEX study of alectinib (ALC) versus crizotinib (CRZ) in untreated advanced ALK+ NSCLC (ID 3850). *Presenter Tony S.K. Mok* 





OS data remain immature, with a 4-year OS rate of 64.5% (95% CI: 55.6–73.4) with alectinib versus 52.2% (95% CI: 42.6–61.8) with crizotinib





Survival

follow-up

## STUDY DESIGN

- Locally advanced, metastatic or recurrent non-squamous NSCLC
- Chemo-naïve
- EGFR mutation positive(exon 19 deletion or exon 21 L858R)
- ECOG PS 0-1
- Bevacizumab-eligible N=311
- Primary endpoint: PFS (Independent Review Committee, IRC)
- Secondary endpoints:
  - PFS(Investigator, INV), ORR, DCR, DOR, OS, TTF, safety
- · Exploratory endpoints:
  - ✓ To identify biomarkers in tissue and plasma that are associated with acquired resistance to bevacizumab combined with erlotinib or erlotinib alone in NSCLC.

1:1

- Stratified by
  - ✓ Sex (female vs. male)

first

Until PD.

unacceptable

toxicity, study ends

or withdrawal of

patient's consent,

whichever occurs

- ✓ Disease stage(stage IIIb vs. stage IV vs. recurrence)
- ✓ EGFR gene mutation (exon 19 del vs. exon 21 L858R)



PFS: Progression-free Survival, ORR: Objective Response Rate; DCR: Disease Control Rate; DOR: Duration of Response; OS: Overall Survival, TTF: Time to Treatment Failure

Erlotinib 150mg qd

(N=154)

Bevacizumab 15mg/kg q3w

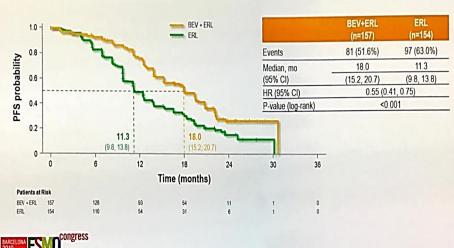
Erlotinib 150mg qd

(N=157)





# PRIMARY ENDPOINT: PFS BY IRC (ITT POPULATION)



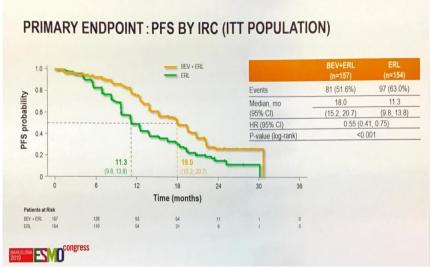
#### SUBGROUP ANALYSIS OF PFS BY INV (ITT POPULATION)

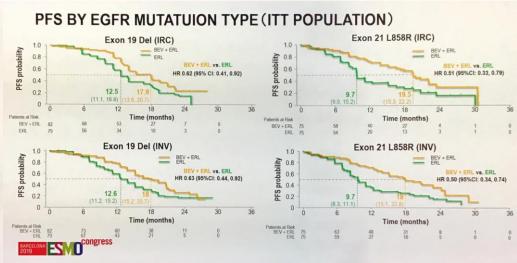
| Subgroup                          | Total number of<br>patients/events (n/N) | BEV + ERL (n/N) | ERL (n/N) | Hazard Ratio and 95% CI | HR with its 95%CI |
|-----------------------------------|--|-----------------|-----------|-------------------------|-------------------|
| All                               | 311/221                                  | 157/100         | 154/121   | Her                     | 0.57 (0.44, 0.75) |
| Gender: Male                      | 118/81                                   | 60/37           | 58/44     | H                       | 0.54 (0.35, 0.85) |
| Gender: Female                    | 193/140                                  | 97/63           | 96/77     | H                       | 0.59 (0.42, 0.82) |
| Stage: IIIb                       | 10/6                                     | 4/4             | 6/2       | 1                       | 2.09 (0.36, 12.1) |
| Stage: IV                         | 275/199                                  | 141/89          | 134/110   | Hell                    | 0.52 (0.39, 0.69) |
| Stage: recurrence                 | 26/16                                    | 12/7            | 14/9      | H .                     | 0.77 (0.28, 2.12) |
| EGFR mutation type: Exon 19 Del   | 161/115                                  | 82/56           | 79/59     | Hel                     | 0.63 (0.44, 0.92) |
| EGFR mutation type: Exon 21 L858R | 150/106                                  | 75/44           | 75/62     | HH                      | 0.50 (0.34, 0.74) |
| Age: ≥75yr                        | 6/2                                      | 3/0             | 3/2       |                         | NA                |
| Age: <75yr                        | 305/219                                  | 154/100         | 151/119   | Hel                     | 0.59 (0.45, 0.77) |
| Pathological type; adenocarcinoma | 311/221                                  | 157/100         | 154/121   | Het I                   | 0.57 (0.44, 0.75) |
| Baseline ECOG score: 0            | 42/31                                    | 25/16           | 17/15     | 1                       | 0.41 (0.20, 0.85) |
| Baseline ECOG score: 1            | 269/190                                  | 132/84          | 137/106   | in                      | 0.60 (0.45, 0.80) |
| Baseline brain metastasis: Yes    | 91/75                                    | 44/32           | 47/43     | HH                      | 0.42 (0.26, 0.67) |
|                                   | 220/146                                  | 113/68          | 107/78    | Hel                     | 0.63 (0.45, 0.87) |















### RESISTANCE BIOMARKER ANALYSIS

22 baseline-PD paired frozen tissue samples in ERL group and 10 paired tissue samples in BEV + ERL group underwent Next-generation sequencing (NGS) of a 448-gene panel and transcriptome sequencing (RNA-Seq) was used for resistance biomarker analysis. Relatively less T790M was found in BEV + ERL group and more mutations and amplifications were found in ERL group at the time point of progression disease.

