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# **Association between Tumor EGFR and KRAS Mutation Status and Clinical Outcomes in NSCLC Patients Randomized to Sorafenib plus Best Supportive Care (BSC) or BSC Alone: Subanalysis of the Phase III MISSION Trial**

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# Conflict of interest disclosure

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Tony S. Mok

- Honoraria

- |                        |                             |
|------------------------|-----------------------------|
| – AstraZeneca          | Hoffmann-La Roche           |
| – Eli Lilly            | Merck Serono                |
| – Eisai                | Bristol-Myers Squibb        |
| – BeiGene              | AVEO                        |
| – Pfizer               | Taiho                       |
| – Boehringer Ingelheim | GlaxoSmithKline Biologicals |

- Speaker

- |                        |                   |
|------------------------|-------------------|
| – AstraZeneca          | Hoffmann-La Roche |
| – Eli Lilly            | Merck Serono      |
| – Boehringer Ingelheim |                   |

- Research Funding

- AstraZeneca

# Background

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- Personalized treatment with targeted agents has become standard in lung cancer patients with specific driver oncogenes<sup>1-3</sup>
- Sorafenib is a multi-kinase inhibitor that has been investigated in phase II and III trials as second/third-line monotherapy in patients with advanced NSCLC<sup>4-6</sup>
- Prior studies suggested that KRAS is a potential predictive biomarker for sorafenib efficacy in NSCLC patients<sup>7,8</sup>
- MISSION is a multi-nation multi-center randomized phase III study comparing sorafenib plus BSC with BSC alone as third/fourth line therapy in an unselected population with advanced NSCLC
- Current study explores the predictive value of EGFR and KRAS mutations in patients from MISSION with available tumor/plasma samples

<sup>1</sup>Mok TS, et al. *N Engl J Med* 2009;361:947-957.

<sup>2</sup>Rosell R, et al. *Lancet Oncol* 2012;13:239-246.

<sup>3</sup>Gandhi L, Janne PA, *Clin Cancer Res* 2012;18:3737-3742.

<sup>4</sup>Blumenschein GR Jr, et al. *J Clin Oncol* 2009;27:4274-4280.

<sup>5</sup>Schiller JH, et al. *J Clin Oncol* 2008;26 Suppl:Abstract 8014.

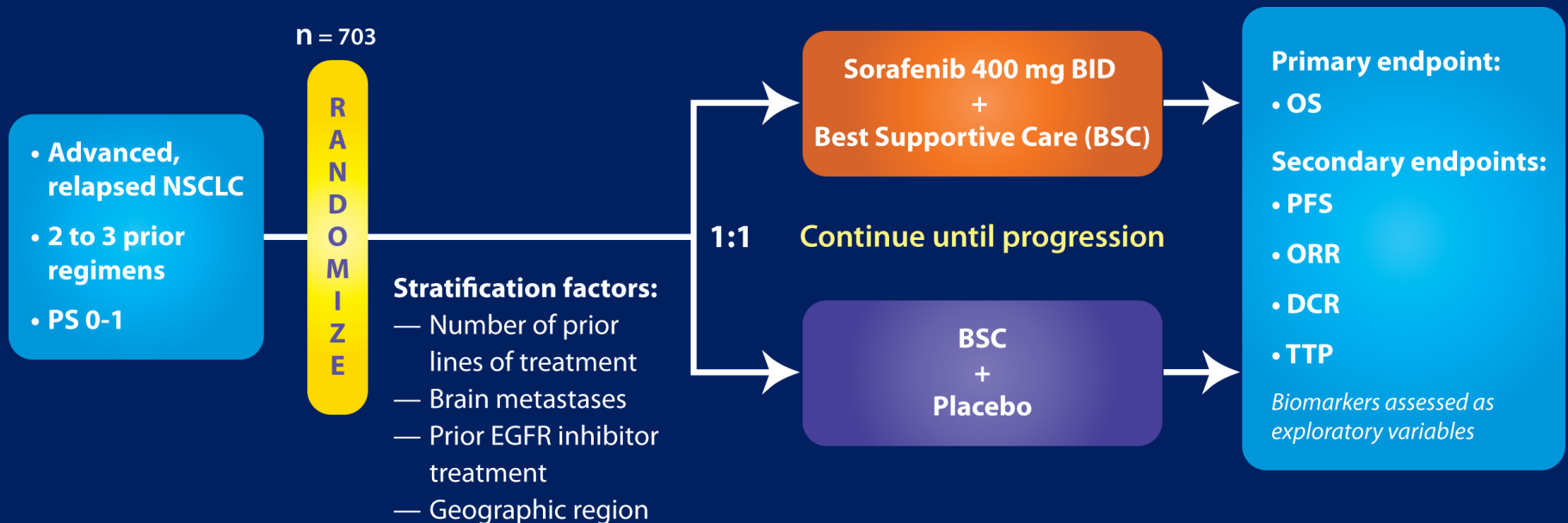
<sup>6</sup>Paz-Ares L, et al. *ESMO* 2012, Abstract LBA 33.

<sup>7</sup>Dingemans AM, et al. *2011 European Multidisciplinary Cancer Congress*, LBA27.

<sup>8</sup>Blumenschein GR Jr, et al. *Cancer Biomark* 2011;10:287-298.

# MISSION: Study objective and design

- Objective
  - To compare the efficacy and safety of sorafenib plus BSC with BSC alone in patients with relapsed or refractory, advanced, predominantly non-squamous NSCLC, with disease progression after two or three prior treatment regimens
- Design
  - Randomized, double-blind, placebo-controlled phase III trial conducted in 33 countries in Europe, North and South America, and Asia Pacific



# MISSION: Overall study results to be presented\*

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**Berlin, May 22, 2012** – Bayer HealthCare and Onyx Pharmaceuticals, Inc. today announced that a Phase III trial evaluating Nexavar® (sorafenib) tablets in patients with advanced relapsed or refractory non-squamous non-small cell lung cancer (NSCLC) whose disease progressed after two or three previous treatments, **did not meet its primary endpoint of improving overall survival**. An improvement in the secondary endpoint of progression-free survival (PFS) was observed... (From May 22 press release)

\* Full results to be presented on Monday, 1 October

Abstract: “Monotherapy Administration of Sorafenib in Patients with Non-Small Cell Lung Cancer: Phase III, Randomized, Double-Blind, Placebo-Controlled MISSION Trial” (LBA33\_PR)

Session: Proffered Paper - NSCLC, metastatic II

Date/time/place: Monday, 1 October 2012, 12:00 PM, Hall A

Presenter: L. Paz-Ares

# Methods: Biomarker analysis

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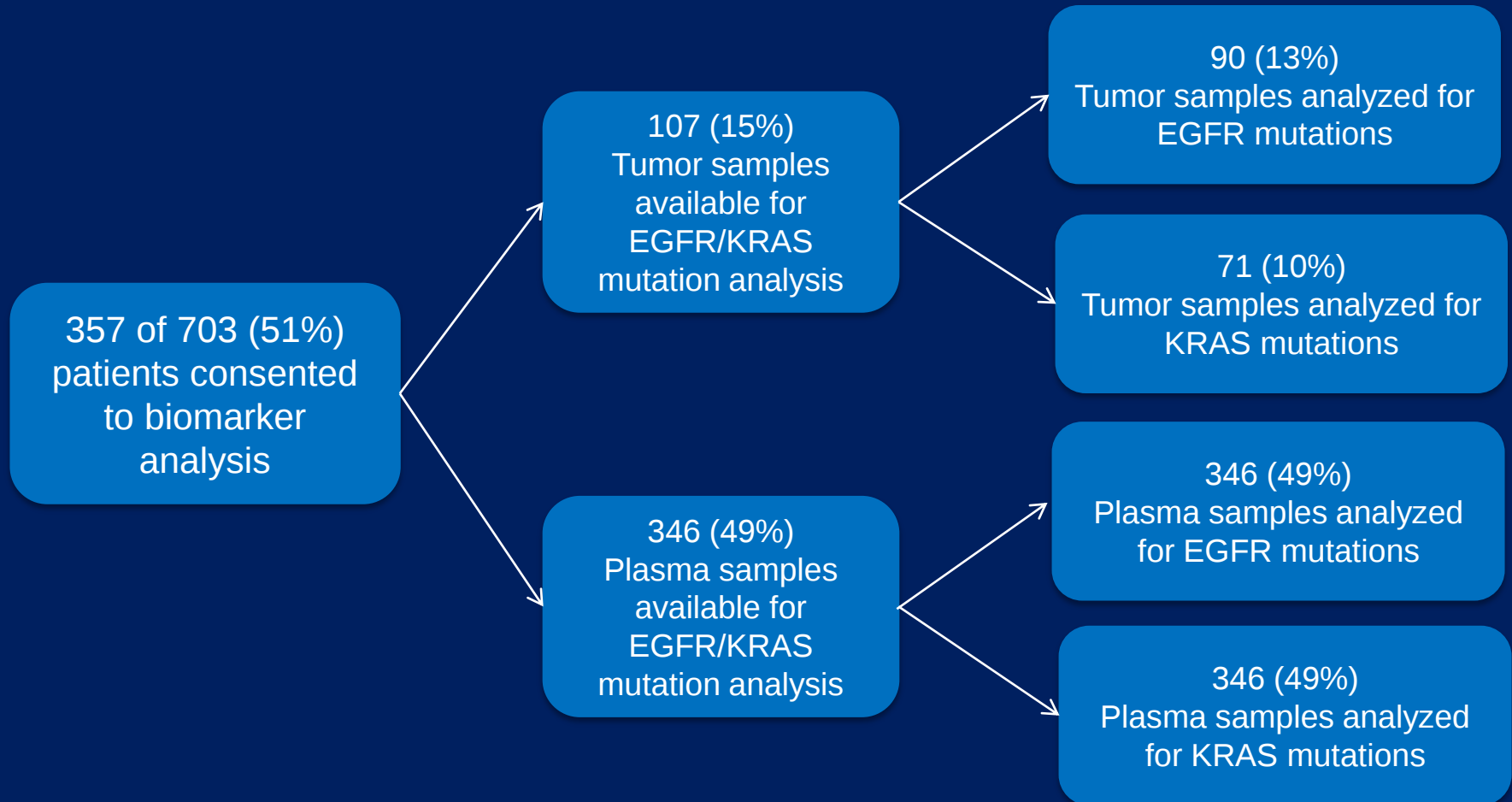
- Archival tumor samples and/or fresh blood samples (during screening) were collected from randomized patients who consented to biomarker study
- EGFR and KRAS mutation status were analyzed in tumor samples and/or in circulating tumor DNA from plasma using BEAMing (Beads, Emulsions, Amplification, and Magnetics; Inostics, Hamburg, Germany).<sup>1,2</sup> Assay sensitivity defining mutation positivity was 0.02% for plasma and 1% for tumor tissue
- Statistical plan
  - The relationships between baseline biomarker concentrations and the effect of sorafenib treatment on OS and PFS were evaluated using Cox proportional hazards models with an interaction term
  - Due to the exploratory nature of the biomarker analyses, p-values must be interpreted with caution because they were not corrected for multiple testing, subset not balanced by randomization, sample size not powered for the analyses

<sup>1</sup>Dressman D, et al. *Proc Natl Acad Sci USA* 2003;100:8817-8822.

<sup>2</sup>Diehl F, et al. *Proc Natl Acad Sci USA* 2005;102:16368-16273.

# Biomarker analysis

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# Results: biomarker analysis

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|                                 | EGFR mutation   | KRAS mutation   |
|---------------------------------|-----------------|-----------------|
| Tumor positive                  | 12 / 90 (13%)   | 20 / 71 (28%)   |
| Tumor negative                  | 78 / 90 (87%)   | 51 / 71 (72%)   |
| Plasma positive                 | 85 / 346 (25%)  | 62 / 346 (18%)  |
| Plasma negative                 | 261 / 346 (75%) | 284 / 346 (82%) |
| Either tumor or plasma positive | 89 / 347 (26%)  | 68 / 347 (20%)  |

# Concordance of tumor and plasma mutations

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| EGFR            |           | Tumor     |        |
|-----------------|-----------|-----------|--------|
|                 |           | Wild-type | Mutant |
| Plasma          | Wild-type | 75        | 2      |
|                 | Mutant    | 4         | 8      |
| Total number    |           | 89        |        |
| Concordance (%) |           | 93.3*     |        |

| KRAS            |           | Tumor     |        |
|-----------------|-----------|-----------|--------|
|                 |           | Wild-type | Mutant |
| Plasma          | Wild-type | 45        | 5      |
|                 | Mut       | 6         | 14     |
| Total number    |           | 70        |        |
| Concordance (%) |           | 84.3*     |        |

\*  $P < 0.001$  by chi-square test

# Demographics based on EGFR mutation status

|                             | EGFR mutation         |                     | EGFR wild type         |                      |
|-----------------------------|-----------------------|---------------------|------------------------|----------------------|
|                             | Sorafenib<br>n=44 (%) | Placebo<br>n=45 (%) | Sorafenib<br>n=122 (%) | Placebo<br>n=136 (%) |
| Median age, yr              | 57.5                  | 56                  | 62.5                   | 62.5                 |
| Male                        | 17 (39)               | 24 (53)             | 71 (58)                | 83 (61)              |
| Smoking status              |                       |                     |                        |                      |
| Non-smoker                  | 28 (64)               | 23 (51)             | 40 (33)                | 32 (24)              |
| Past or present smoker      | 15 (34)               | 22 (49)             | 80 (65)                | 104 (76)             |
| ECOG PS                     |                       |                     |                        |                      |
| 0                           | 12 (27)               | 18 (40)             | 45 (37)                | 52 (38)              |
| 1                           | 31 (70)               | 27 (60)             | 77 (63)                | 84 (62)              |
| Brain metastases present    | 9 (20)                | 13 (29)             | 16 (13)                | 20 (15)              |
| Prior EGFR systemic therapy | 37 (84)               | 37 (82)             | 68 (56)                | 72 (53)              |
| Number of prior regimens    |                       |                     |                        |                      |
| 2                           | 21 (48)               | 27 (60)             | 66 (54)                | 73 (54)              |
| 3                           | 22 (50)               | 18 (40)             | 53 (43)                | 63 (46)              |

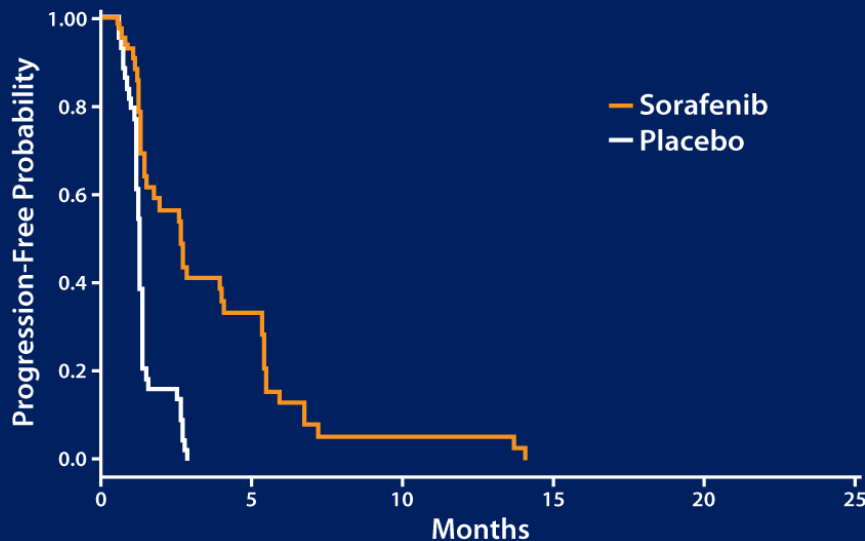
# Demographics based on EGFR mutation status

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# Progression-free survival based on EGFR mutation status

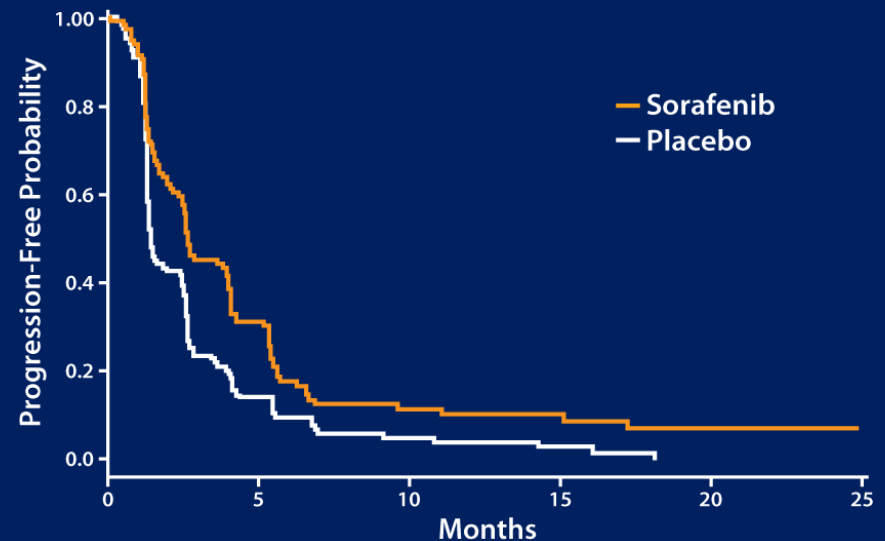
## Pts with EGFR mut (in tumor or plasma)

- Sorafenib N=44; Placebo N=45
- HR=0.27 (95% CI 0.16,0.46)
- P-value<0.001
- Sorafenib median PFS= 2.7 mo (83d)
- Placebo median PFS= 1.4 mo (42d)



## Pts with EGFR wt

- Sorafenib N=122; Placebo N=136
- HR=0.62 (95% CI 0.48,0.82)
- P-value<0.001
- Sorafenib median PFS= 2.7 mo (82d)
- Placebo median PFS= 1.5 mo (46d)

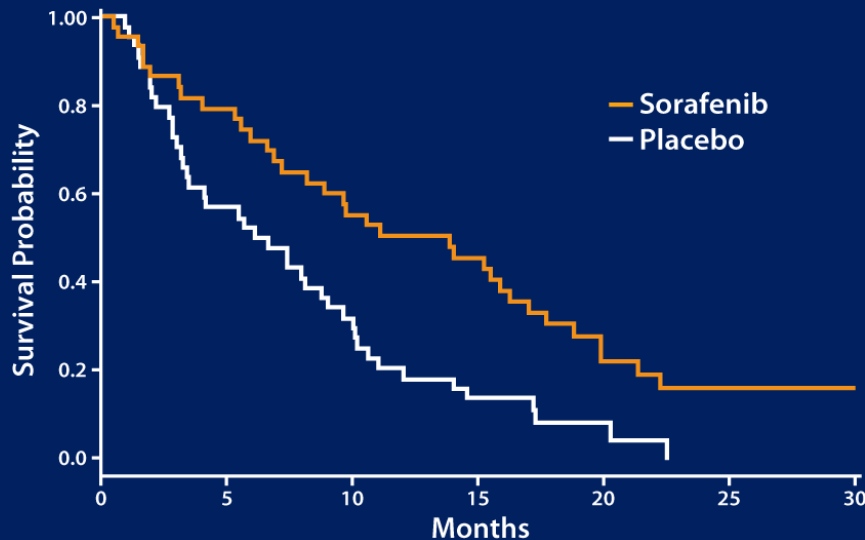


Biomarker\*treatment interaction analysis: p-value=0.015

# Overall survival based on EGFR mutation status

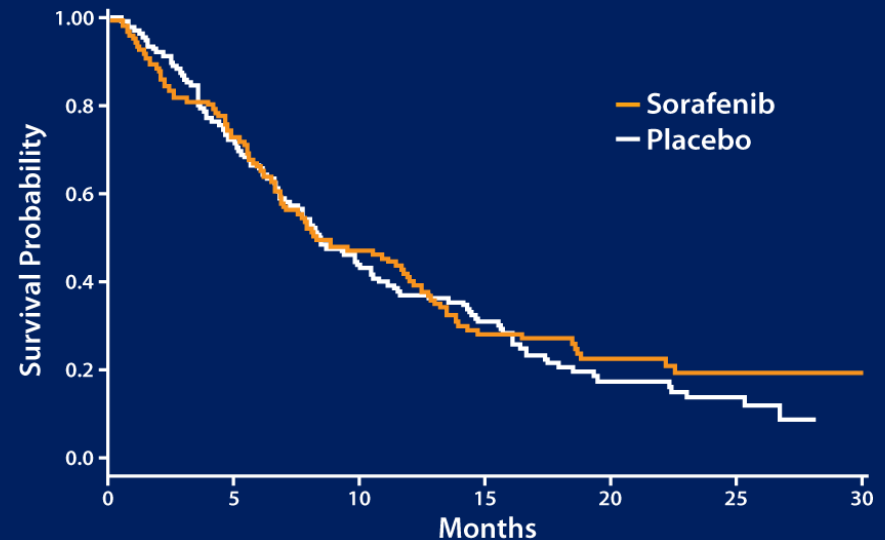
## Pts with EGFR mut (in tumor or plasma)

- Sorafenib N=44; Placebo N=45
- HR=0.48 (95% CI 0.3,0.76)
- P-value=0.002
- Sorafenib median OS= 13.9 mo (423d)
- Placebo median OS= 6.5 mo (197d)



## Pts with EGFR wt

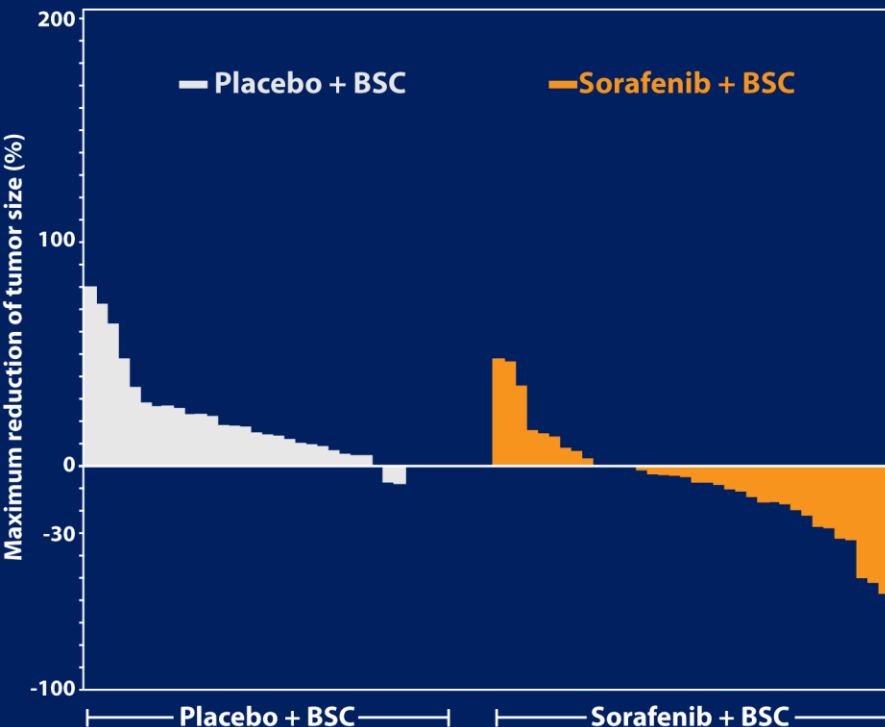
- Sorafenib N=122; Placebo N=136
- HR=0.92 (95% CI 0.7,1.21)
- P-value=0.559
- Sorafenib median OS= 8.3 mo (253d)
- Placebo median OS= 8.4 mo (256d)



Biomarker\*treatment interaction analysis: p-value=0.023

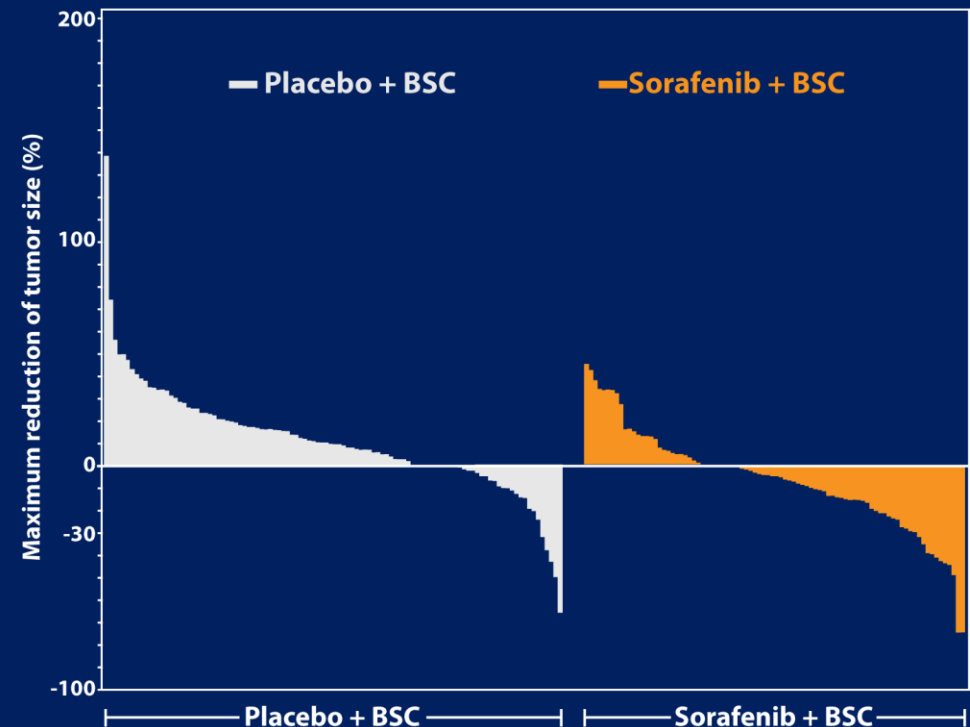
# Tumor response – EGFR status (Investigator assessed)

## EGFR Mutation Positive



ORR – 0% vs. 6.8%  
DCR – 2.2% vs. 40.9%

## EGFR Wild-type



ORR – 1.5% vs. 7.4%  
DCR – 25.8% vs. 46.7%

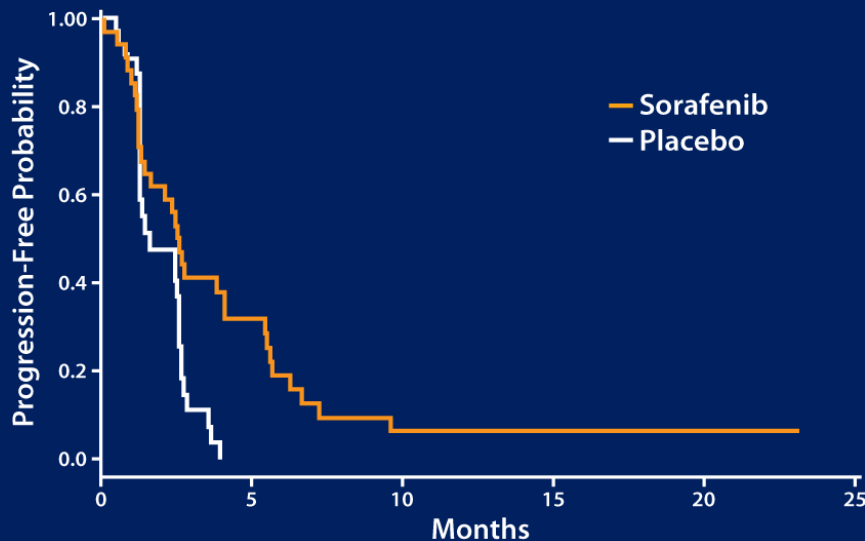
# Treatment summary

|                                 | EGFR mutation         |                     | EGFR wild type         |                      |
|---------------------------------|-----------------------|---------------------|------------------------|----------------------|
|                                 | Sorafenib<br>n=44 (%) | Placebo<br>n=45 (%) | Sorafenib<br>n=122 (%) | Placebo<br>n=136 (%) |
| <b>On-study</b>                 |                       |                     |                        |                      |
| Duration of therapy             |                       |                     |                        |                      |
| Mean (weeks)                    | 16.6                  | 6.1                 | 19.6                   | 12.4                 |
| Dose interruption, n (%)        | 13 (30)               | 6 (13)              | 60 (49)                | 27 (20)              |
| Dose reduction, n (%)           | 8 (18)                | 1 (2)               | 47 (39)                | 8 (6)                |
| <b>Post-progression Therapy</b> |                       |                     |                        |                      |
| Any                             | 26 (59)               | 25 (56)             | 54 (44)                | 84 (62)              |
| 2+                              | 16 (36)               | 9 (20)              | 22 (18)                | 33 (24)              |
| Anti-EGFR                       | 19 (43)               | 8 (18)              | 18 (15)                | 37 (27)              |

# Progression-free survival based on KRAS mutation status

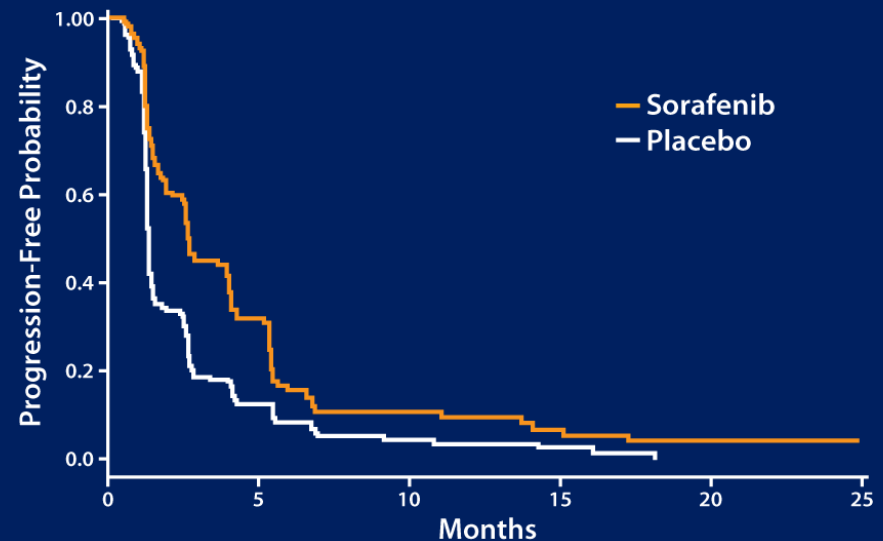
## Pts with KRAS mut (in tumor or plasma)

- Sorafenib N=34; Placebo N=34
- HR=0.46 (95% CI 0.25,0.82)
- P-value=0.007
- Sorafenib median PFS= 2.6 mo (80d)
- Placebo median PFS= 1.7 mo (51d)



## Pts with KRAS wt

- Sorafenib N=132; Placebo N=147
- HR=0.58 (95% CI 0.45,0.75)
- P-value<0.001
- Sorafenib median PFS= 2.7 mo (83d)
- Placebo median PFS= 1.4 mo (43d)

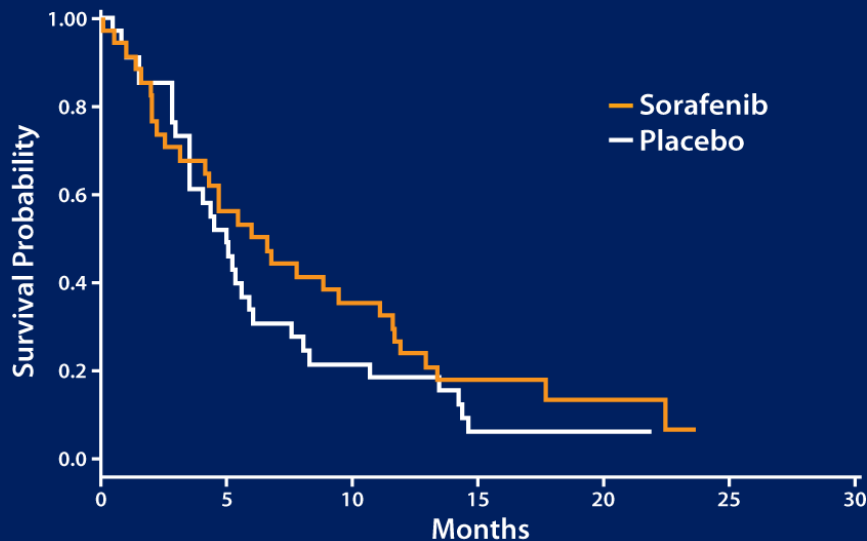


Biomarker\*treatment interaction analysis: p-value=0.696

# Overall survival based on KRAS mutation status

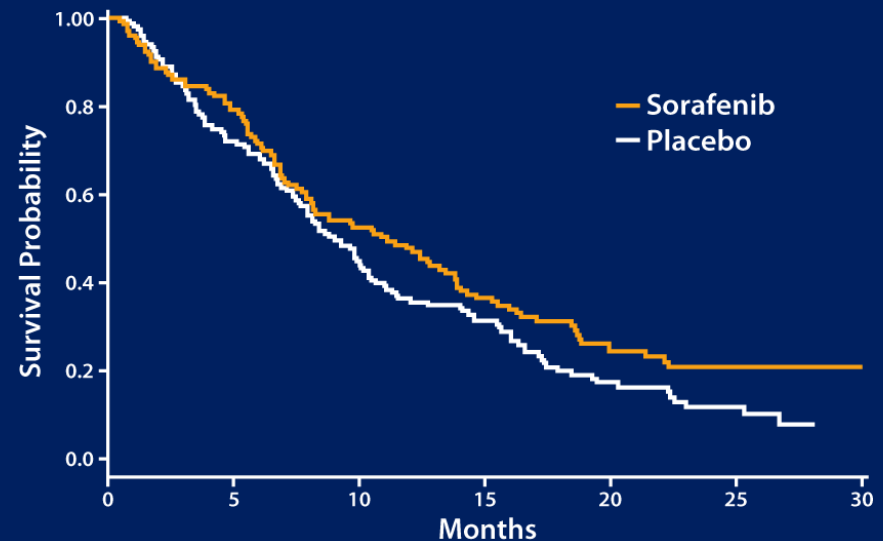
## Pts with KRAS mut (in tumor or plasma)

- Sorafenib N=34; Placebo N=34
- HR=0.76 (95% CI 0.45,1.26)
- P-value=0.279
- Sorafenib median OS= 6.4 mo (195d)
- Placebo median OS= 5.1 mo (156d)



## Pts with KRAS wt

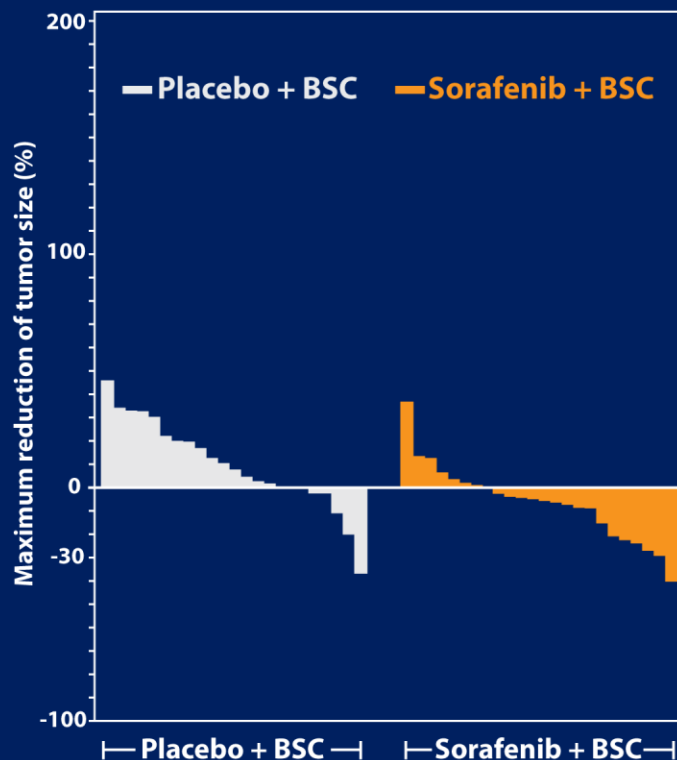
- Sorafenib N=132; Placebo N=147
- HR=0.79 (95% CI 0.6,1.03)
- P-value=0.079
- Sorafenib median OS= 11.0 mo (339d)
- Placebo median OS= 9.1 mo (278d)



Biomarker\*treatment interaction analysis: p-value=0.743

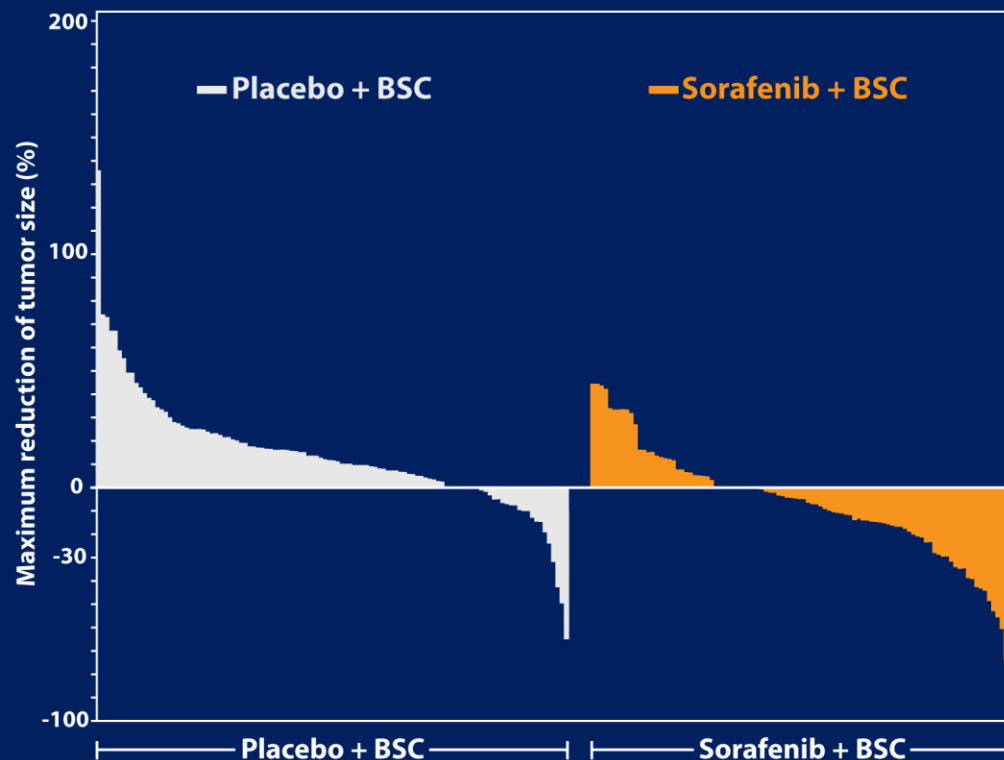
# Tumor response – KRAS status (Investigator assessed)

## KRAS Mutation Positive



ORR – 0% vs. 2.9%  
DCR – 7.6% vs. 44.1%

## KRAS Wild-type



ORR – 1.4% vs. 8.3%  
DCR – 20.4% vs. 45.4%

# Conclusion

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- Based on current data, we hypothesize that EGFR mutation is a predictive biomarker for sorafenib in treatment of patients with advanced NSCLC
  - Positive interaction analysis for both PFS and OS
  - OS outcome may be biased by the unbalanced use of post-study EGFR TKI (sorafenib arm 43%; placebo arm 18%)
- KRAS mutation status did not appear to influence response to sorafenib
  - Negative interaction analysis for both PFS and OS
  - Sorafenib resulted in more favorable PFS outcomes than placebo in patients with both KRAS mutant and wild type tumors
- These results should be interpreted with caution
  - The patient subgroup with available samples for biomarker analysis is not representative of the overall population<sup>1</sup>
  - Limited sample size (47% of overall population)
  - Biomarker analyses in MISSION were retrospective

<sup>1</sup>Paz-Ares L, et al. *ESMO* 2012, Abstract LBA 33.

# MISSION Trial Investigators

## 154 Centers in 33 Countries

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*We thank the patients for their contribution to this study*

**Austria:** Horst Olschewski, Kurt Aigner, Josef Bolitschek

**Belgium:** Philippe Collard, Michiel Thomeer, Benoit Colinet

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**Bulgaria:** Assen Dudov, Danail Damyanov, Vladimir Kanarev, Petar Petrov, Violina Taskova, Dimitar Kalev

**Canada:** Vera Hirsh

**Chile:** Osvaldo Aren, Mónica Ahumada

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**Taiwan:** James Chih-Hsin Yang, Han-Pin Kuo, Gee-Chen Chang, Te-Chun Hsia, Ming-Shyan Huang

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**United States:** J. Beck, Heather Wakelee

*Supported by Bayer HealthCare Pharmaceuticals and Onyx Pharmaceuticals*

# Backup

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# EGFR and KRAS mutations assayed

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| EGFR |                           |                   |
|------|---------------------------|-------------------|
| Exon | Nucleotide change         | Amino acid change |
| 19   | 2235_2249 del15           | E746_A750del      |
| 19   | 2236_2250 del15           | E746_A750del      |
| 19   | 2237_2255>T               | E746_S752>V       |
| 19   | 2239_2248<br>TTAAGAGAAG>C | L747_A750>P       |
| 19   | 2240_2254 del15           | L747_T751del      |
| 19   | 2240_2257 del18           | L747_P753>S       |
| 20   | 2369 C>T                  | 790 T>M           |
| 21   | 2573 T>G                  | 858 L>R           |

| KRAS |                   |                   |
|------|-------------------|-------------------|
| Exon | Nucleotide change | Amino acid change |
| 1    | 34 G>A            | 12 G>S            |
| 1    | 34 G>C            | 12 G>R            |
| 1    | 34 G>T            | 12 G>C            |
| 1    | 35 G>A            | 12 G>D            |
| 1    | 35 G>C            | 12 G>A            |
| 1    | 35 G>T            | 12 G>V            |
| 1    | 38 G>A            | 13 G>D            |