

LUX-Lung 7

## ■1 ■LUX-LUNG 7 STUDY DESIGN

Patients(N=319)

- Stage IIIB/IV adenocarcinoma of the lung
- EGFR mutation(Del 19 and/or L858R) in the tumor tissue\*
- No prior treatment for advanced/metastatic disease
- ECOG PS0/1

Afatinib 40mg QD<sup>†</sup>

Stratified by

- 1 : 1
- Mutation type(Del 19/L858R)
  - Brain metastases(present/absent)

Gefitinib 250mg QD

Co-primary endpoints:

- PFS(independent review)
- TTF
- OS

Secondary endpoints:

- ORR
- Time to response
- Duration of response
- Tumor shrinkage
- HRQoL

- Treatment beyond progression allowed if deemed beneficial by investigator
- RECIST assessment performed at Week4, 8 and every 8 weeks thereafter until Week 64, and every 12 weeks thereafter
- Primary PFS analysis conducted after ~ 250 events ; primary OS analysis conducted after ~ 213 events and ≥ 32-mo follow-up
- All statistical testing at two-sided 5%alpha level with no adjustment for multiplicity

\*Central or local test :<sup>†</sup>Dose modification to 50, 30 or 20mg was permitted in line with prescribing information

ECOG PS, Eastern Cooperative Oncology Group performance status ; HRQoL, health-related quality of life ; QD, once daily ; RECIST, Response Evaluation Criteria in Solid Tumors :

# LUX-Lung 7 (EGFR-TKI-65)

2011年12月から2013年8月までに北米や欧州、アジア  
(日本は不参加) など13カ国の64施設から患者が登録

Primary Endpoint: PFS, TTF, OS

Secondary Endpoint: ORR, DCR, safety, QOL

## Phase IIb

Adeno

化学療法未治療  
非小細胞肺癌患者

StageIIIB/IV

(1次治療)

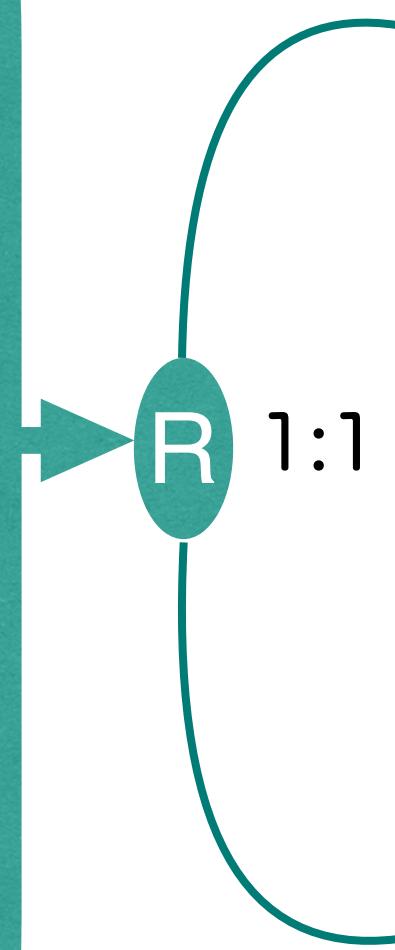
- EGFR遺伝子変異

Exon19 deletion

Exon21 L858R

- Asymptomatic

- No Brain meta



n=319

afatinib (40mg/day)

n=160

Exon 19 del :58%  
Exon 21 L858R:42%

n=159

Gefitinib (250mg/day)

mPFS

11ヶ月

HR=0.74  
p=0.0178

OS

27.9ヶ月

HR=0.86  
p=0.258

ORR

73%

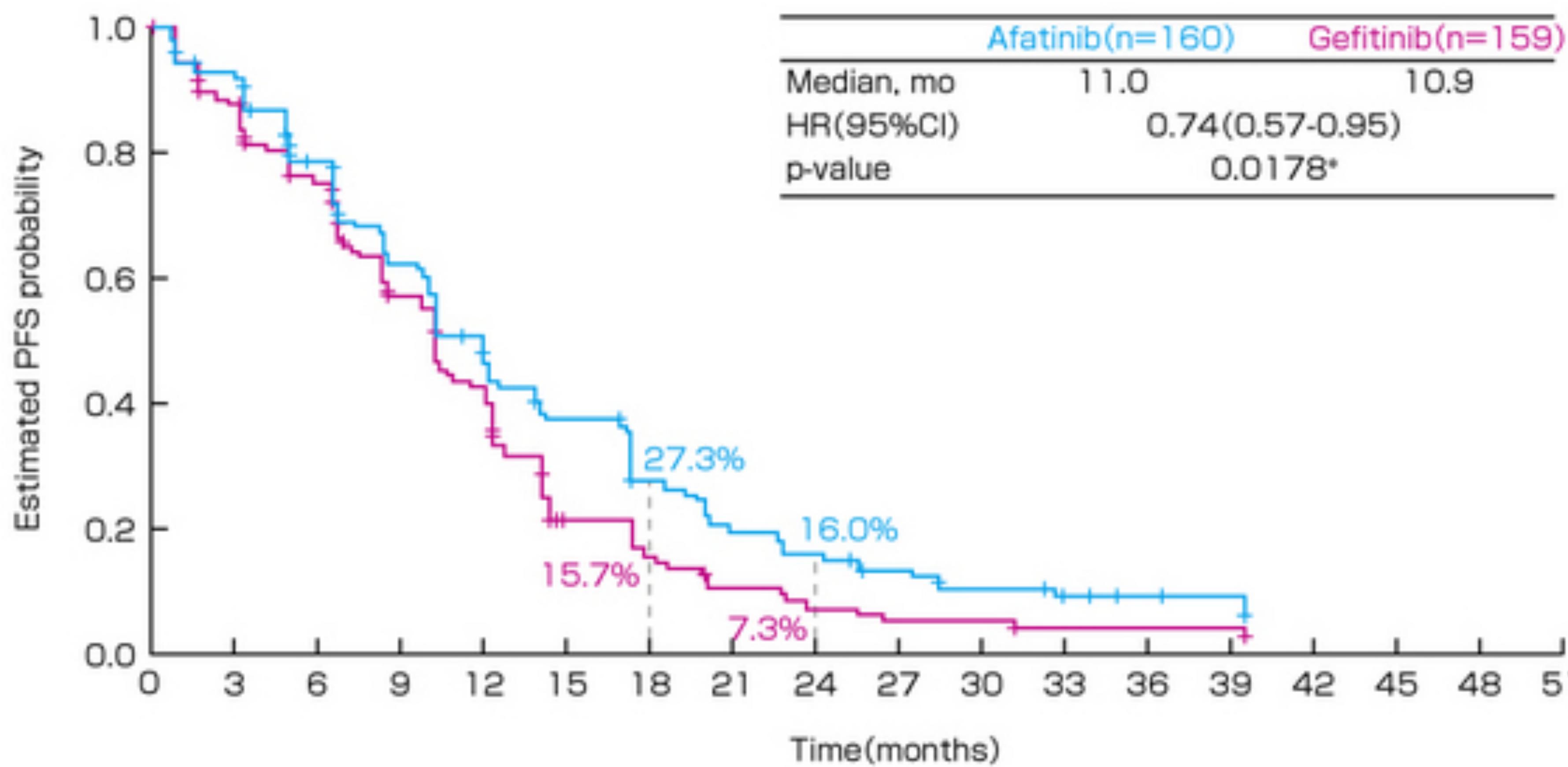
p=0.002

TTF

13.7ヶ月

11.5ヶ月

## ■2 ■ UPDATED PFS (INDEPENDENT REVIEW)



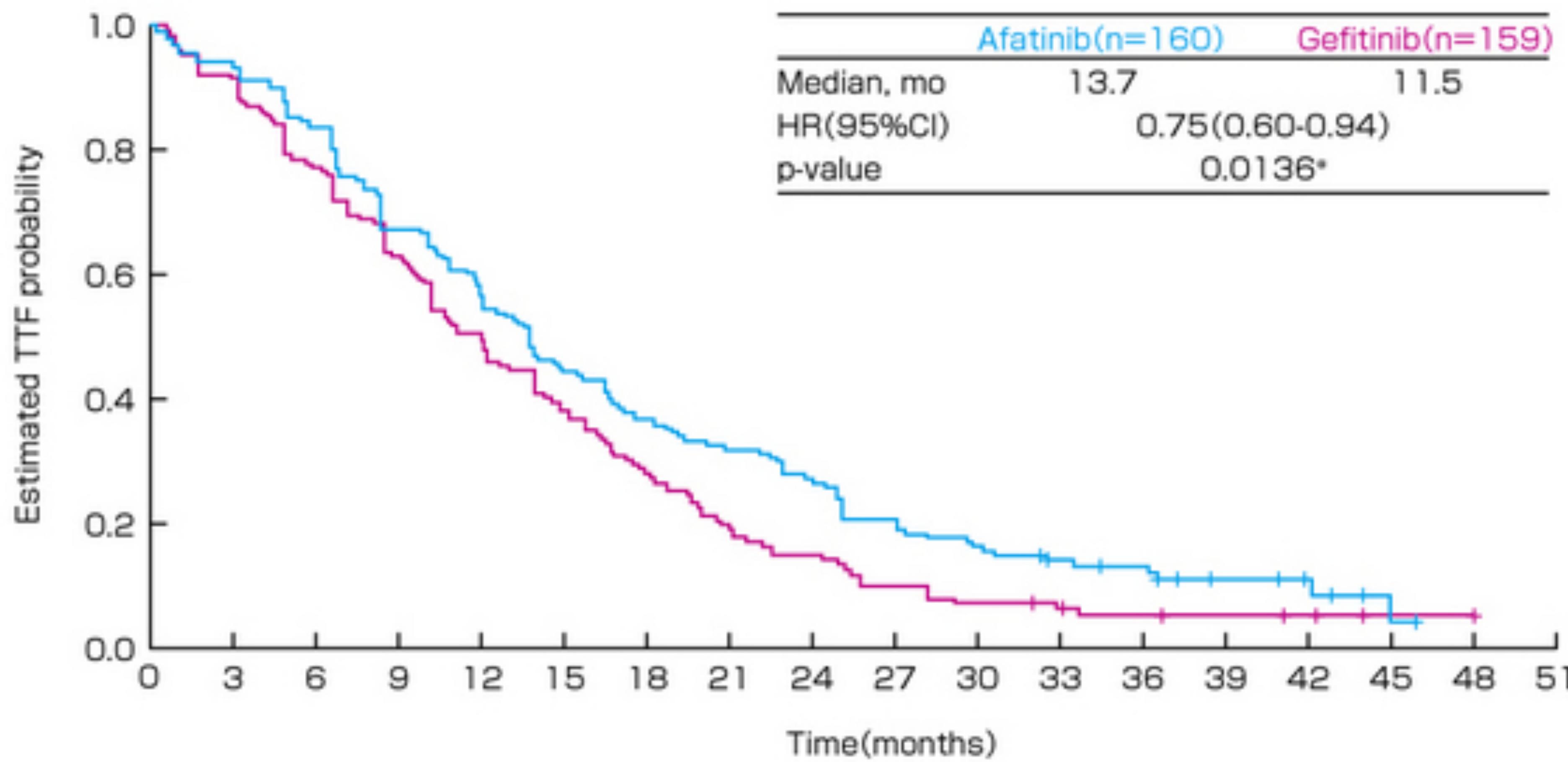
No. at risk

Afatinib	160	142	113	94	67	47	34	26	20	13	10	8	4	3	0	0	0	0
Gefitinib	159	132	105	82	51	21	15	10	7	5	5	3	3	0	0	0	0	0

\*unadjusted

(Luis Paz-Ares, et al. ESMO2016 Abstract No. LBA43)

### ■3 ■ UPDATED TTF



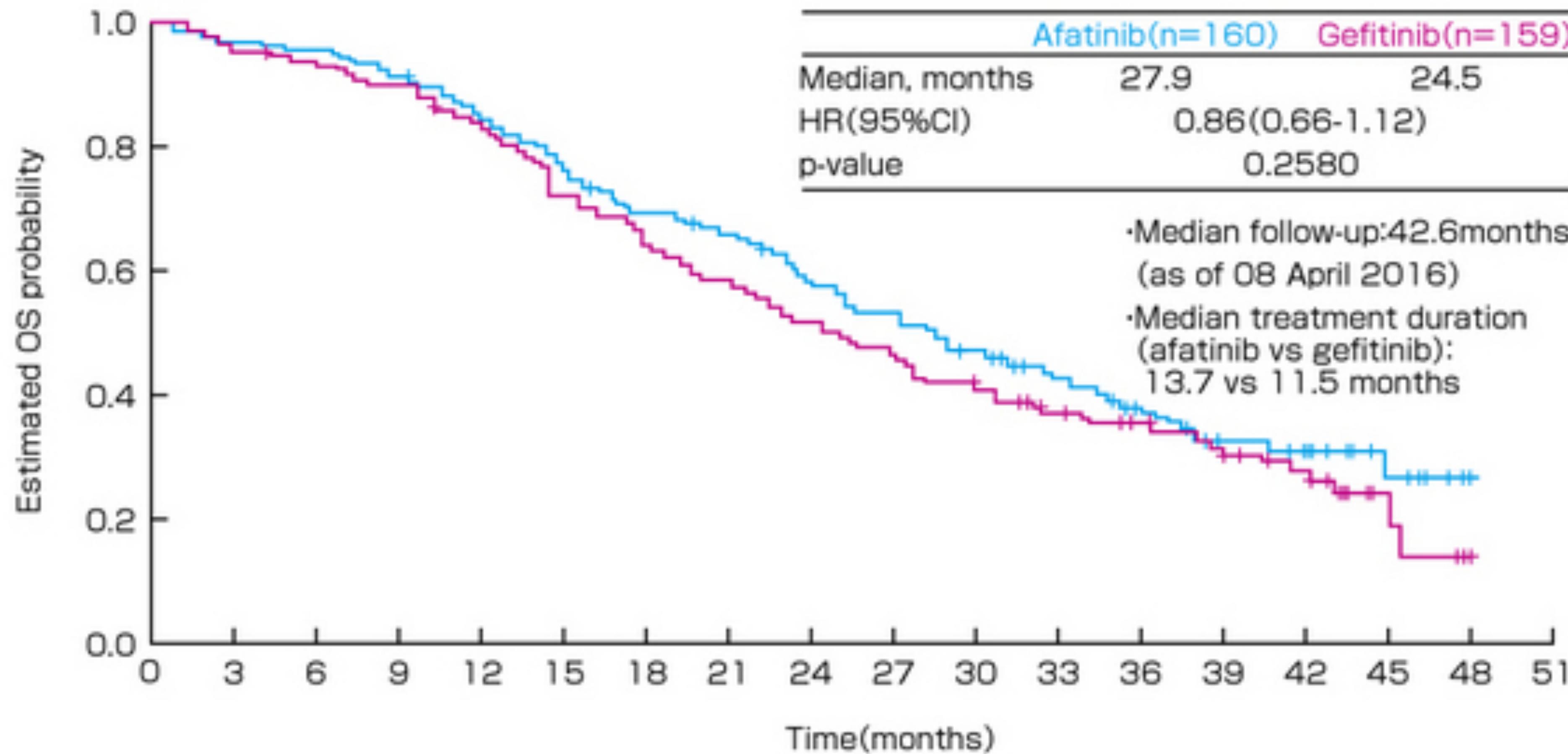
No. at risk

Afatinib	160	148	133	113	91	68	56	48	40	29	25	19	16	7	6	1	0	0
Gefitinib	159	144	120	103	74	59	43	30	21	14	10	9	6	5	4	2	0	0

\*unadjusted

(Luis Paz-Ares, et al. ESMO2016 Abstract No. LBA43)

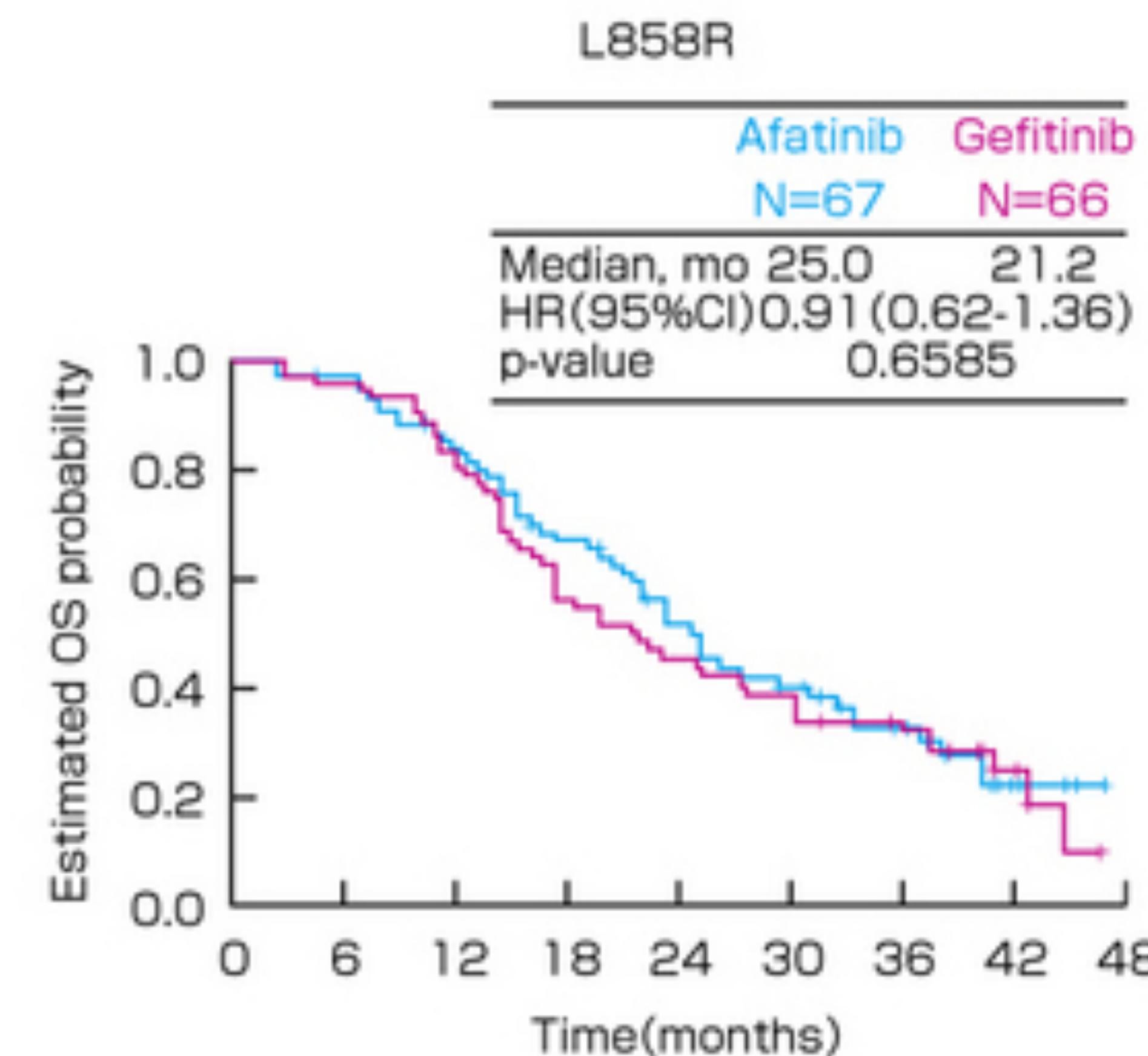
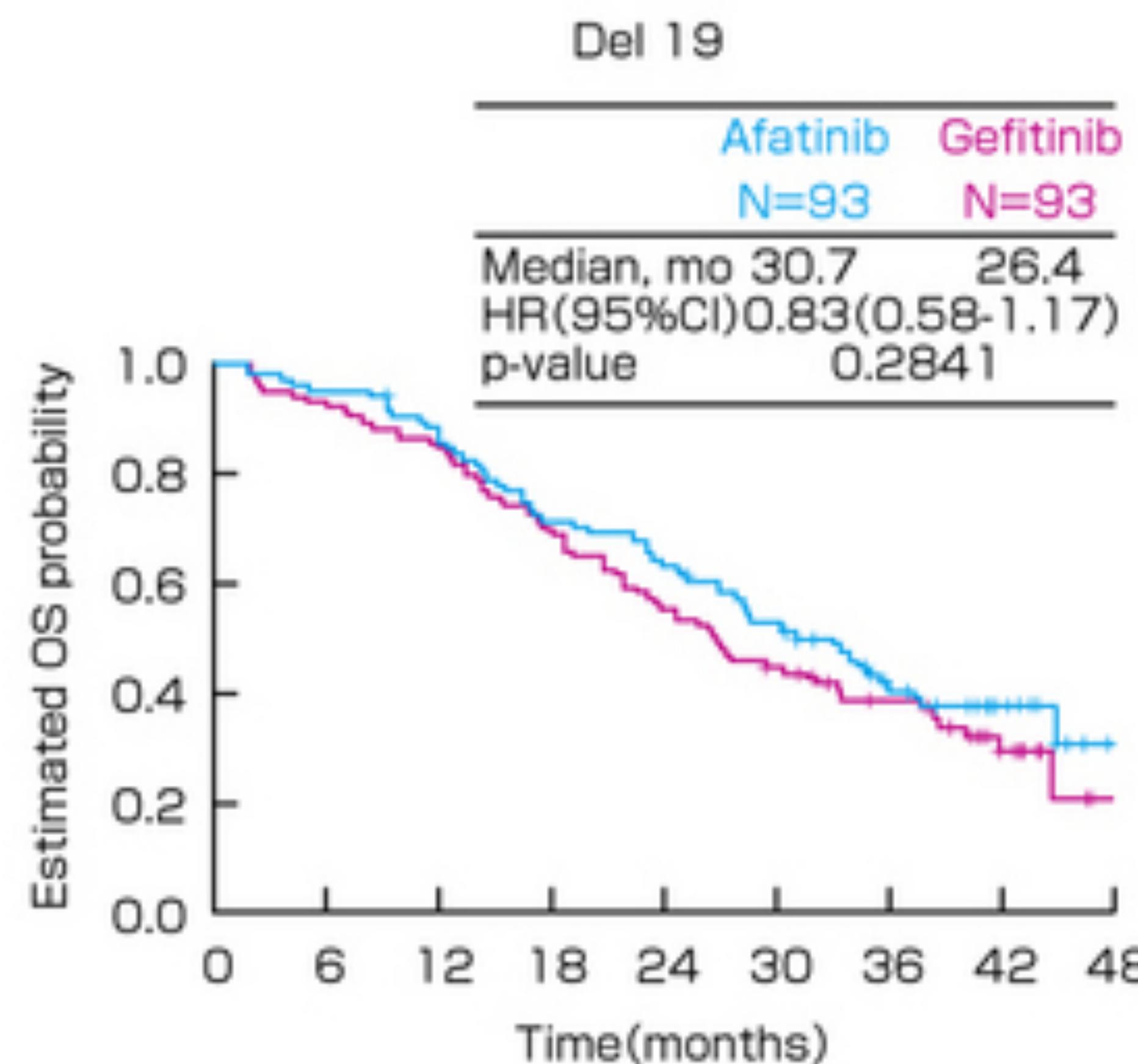
#### ■ OS(OVERALL POPULATION)



No. at risk

Afatinib	160	156	153	148	139	125	111	104	94	81	74	61	50	36	30	12	2	0
Gefitinib	159	153	148	142	133	119	105	90	80	71	62	56	48	44	27	7	0	0

## FIGURE 5 OS BY EGFR MUTATION SUBTYPE



No. at risk									
Afatinib	93	88	82	68	61	50	35	20	1
Gefitinib	93	86	79	66	52	39	29	17	0

No. at risk									
Afatinib	67	65	57	43	33	24	15	10	1
Gefitinib	66	62	54	39	28	23	19	10	0

Afatinib (n=160) Gefitinib (n=159)

	Afatinib (n=160)	Gefitinib (n=159)
Sex		
Men	69 (43%)	53 (33%)
Women	91 (57%)	106 (67%)
Age	63 (30–86)	63 (32–89)
Ethnic origin		
Asian	94 (59%)	88 (55%)
Black/African American	1 (1%)	0
White	48 (30%)	54 (34%)
Missing*	17 (11%)	17 (11%)
Smoking status		
Never smoked	106 (66%)	106 (67%)
Light ex-smoker†	21 (13%)	19 (12%)
Other current or ex-smokers	33 (21%)	34 (21%)
Baseline ECOG PS		
0	51 (32%)	47 (30%)
1	109 (68%)	112 (70%)
Histological classification		
Adenocarcinoma	159 (99%)	158 (99%)
Mixed	1 (1%)	1 (1%)
Clinical stage at screening		
IIIB	8 (5%)	3 (2%)
IV	152 (95%)	156 (98%)
EGFR mutation category		
Leu858Arg	67 (42%)	66 (42%)
Leu858Arg alone	67 (42%)	65 (41%)
Leu858Arg+exon 19 deletion	0	1 (1%)
Exon 19 deletion‡	93 (58%)	93 (58%)
Metastases at screening		
Adrenal glands	12 (8%)	16 (10%)
Bone	80 (50%)	73 (46%)
Brain	26 (16%)	24 (15%)
Liver	16 (10%)	24 (15%)
Lung ipsilateral	86 (54%)	88 (55%)
Lung contralateral	65 (41%)	73 (46%)
Other	100 (63%)	104 (65%)

Data are n (%) or median (range). ECOG PS=Eastern Cooperative Oncology Group performance status. \*Patients recruited in French sites did not have their ethnic origin recorded. †Less than 15 pack-years and stopped more than 1 year before diagnosis. ‡One patient in the afatinib group with wild-type EGFR was erroneously included in the trial and was reported as exon 19 deletion at the time of randomisation by the investigator.

Table 1: Baseline demographics and disease characteristics

