Second-line afatinib versus methotrexate in patients with recurrent and/or metastatic head and neck squamous cell carcinoma: subgroup/biomarker analysis of LUX-Head and Neck 1

Makoto Tahara, Ezra E. W. Cohen, Robert I. Haddad, Jérôme Fayette, Lisa F. Licitra, Paul M. Clement, Jan B. Vermorken, Thomas Gauler, Didier Cupissol, Juan José Grau, Joël Guigay, Joseph M. del Campo, Kenji Okami, Shunji Takahashi, Barbara Burtness, Xiuyu Julie Cong, Neil Gibson, Flavio Solca, Eva Ehrnrooth, and Jean-Pascal H. Machiels on behalf of the LUX-H&N 1 investigators



Disclosures

- Grants and contracts: Eisai, Merck Sharp & Dome
- Honoraria and consultation fees: Merck Serono, Bristol-Myers Squibb, Eisai, Otsuka and Bayer



Background

- R/M HNSCC patients progressing on/after first-line platinum therapy have poor prognosis (36 months median survival) and limited treatment options^{1–3}
- While some molecular biomarkers, including p16 (surrogate for HPV infection), EGFR overexpression and PTEN loss, have been associated with prognosis in HNSCC, there are no established biomarkers predictive of treatment response^{4,5}
- In the Phase III LUX-H&N1 trial, afatinib, an oral irreversible ErbB family blocker, significantly improved PFS (median 2.6 vs 1.7 months; HR=0.80; p=0.030) versus methotrexate in second-line R/M HNSCC patients⁶
 - Complete efficacy and safety findings from the overall population are published⁶
- This report focuses on efficacy outcomes in selected prespecified subgroups and biomarker-defined populations^{6,7}



LUX-Head & Neck 1: study design

Patients (N=474)

- With R/M HNSCC not amenable for curative treatment
- Progression on/after first-line platinum-based therapy
- No more than one previous systemic regimen was allowed
- EGFR TKI-treatment naïve





ECOG PS, Eastern Cooperative Oncology Group performance status; iv, intravenous; mAb, monoclonal antibody; MTX, methotrexate; TKI, tyrosine kinase inhibitor; ORR, objective response rate; OS, overall survival; PFS, progression-free survival; PRO, patient-reported outcome; RECIST v1.1, Response Evaluation Criteria In Solid Tumors version 1.1

Patient characteristics*

		Afatinib	MTX
		(n=322)	(n=161)
Gender, %	Male/female	85/15	85/15
Median age, years (range)		60 (32–82)	59 (32–88)
Ago oubgroup %	<65 years	74	72
Age subgroup, %	≥65 years	26	28
ECOG PS, %	0/1	28/72	26/74
	Asia	8	11
Decies 0/1	Europe	77	74
Region, %	North/Latin America	12	13
	Other	2	2
	<10 pack years	17	19
Smoking history, % [†]	≥10 pack years	79	78
	Unknown	3	3
	Oral cavity	29	26
Primary tumor site % [†]	Oropharynx	31	34
Filmary lumor sile, %	Hypopharynx	20	19
	Larynx	20	22
First-line anti-EGFR mAb, % [‡]		59	61
	Positive	7	7
p16 status, % ^{†,§}	Negative	42	40
	Not performed	51	53



*Biomarker analyses were conducted using tumor samples from a subset of patient volunteers, resulting in a smaller sample size (n=234) than the overall population [†]Percentages may not total 100% due to rounding [‡]One patient received panitumumab and all other patients received cetuximab; [§]Assessed in a central laboratory

PFS in the overall population



SINGAPORE

2015

PFS subgroup analysis*

SINGAPORE

SINGAPORE

2015

Factors		No. of patients		HR (95% CI)		
Total		483	⊢ ,	0.80 (0.65–0.98)		
Baseline ECOG PS	0 1	131 352		0.73 (0.49–1.10) 0.77 (0.60–0.98)		
Prior use of EGFR-targeted antibody for R/M HNSCC	No Yes	196 287		0.63 (0.45–0.88) 0.91 (0.70–1.19)		
Gender	Male Female	412 71		0.74 (0.59–0.92) 0.95 (0.55–1.64)		
Age	<65 years ≥65 years	355 128	F- ♦ -1 F 1	0.79 (0.62–1.01) 0.68 (0.45–1.03)		
Region	Asia Europe North/Latin American	43 369 60		0.62 (0.32–1.20) 0.82 (0.64–1.04) 0.41 (0.21–0.79)		
Smoking pack-years	<10 pack years ≥10 pack years	87 381		1.05 (0.66–1.70) 0.71 (0.56–0.90)		
Alcohol consumption	≤7 units/week >7 units/week	374 91		0.79 (0.62–1.00) 0.73 (0.46–1.14)		
Primary tumor site	Oral cavity Oropharynx Hypopharynx Larynx	136 153 93 101		0.69 (0.46–1.04) 0.99 (0.68–1.44) 0.78 (0.48–1.25) 0.59 (0.38–0.92)		
Recurrence or metastases	Recurrent Metastatic Both	167 64 241		0.59 (0.42–0.84) 1.18 (0.65–2.14) 0.81 (0.60–1.10)		
Response to prior platinum therapy for R/M HNSCC	CR/PR/SD PD	261 146		0.82 (0.62–1.09) 0.66 (0.45–0.96)		
	1/	16 Fa v	1/4 1 vors afatinib ◀─── 1	4 16 Favors MTX		
SINGAPORE SALA 18-21 DECEMBER						

*p16 status evaluated in the biomarker analysis (based on central laboratory assessment); CR, complete response; PD, progressive disease; PR, partial response; SD, stable disease

PFS by age subgroup

SINGAPORE

SINGAPORE

2015



PFS by prior EGFR-mAb therapy



SINGAPORE

2015

Tumor response



[†]OR: 1.5 (1.03–2.26); p=0.035

SINGAPOR

2015

ORR: <65 OR: 2.0 (0.81–5.15); ≥65 OR: 1.7 (0.44–6.64) DCR: <65 OR: 1.4 (0.92–2.26); ≥65 OR: 1.9 (0.89–3.90) ORR: prior OR: 0.90 (0.26–3.17); *No prior OR: 2.8 (1.03–7.73) DCR: prior OR: 1.2 (0.70–1.92); [†]No prior OR: 2.3 (1.24–4.21)



Prespecified biomarkers associated with ErbB pathway dysregulation

Patient disposition					
Biomarker status (cut-offs)*	No. of patients (afatinib vs MTX)	Percentage of total, n/N (%) [†]			
p16-positive (H-score ≥210)	23 vs 12	35/234 (15%)			
p16-negative (H-score <210)	135 vs 64	199/234 (85%)			
EGFR-amplified [‡]	50 vs 16	66/146 (45%)			
EGFR non-amplified	53 vs 27	80/146 (55%)			
HER3-high (H-score >50)	64 vs 26	90/156 (58%)			
HER3-low (H-score ≤50)	49 vs 17	66/156 (42%)			
PTEN-high (H-score >150)	30 vs 12	42/157 (27%)			
PTEN-low (H-score ≤150)	82 vs 33	115/157 (73%)			



*Cut-offs were not prespecified in the protocol and are exploratory; [†]Percentage based on total patients with specific biomarker available; [‡]Amplification defined as ≥50% of cells with ≥4 copies, or ≥1 cell with ≥8 copies H-score, histology-score

PFS according to biomarker status

Factors								HR	(95% CI)	Interaction p value	
Total study pop	oulation (N=483)				⊢_			0.80	(0.65–0.98)		
p16	Positive (n=35)		F		+			0.81	(0.39–1.69)	0.654	
μιο	Negative (n=199	9)			-			0.70	(0.50–0.97)	0.054	
ECER	Amplified (n=66))	⊢		•			0.66	(0.35–1.24)	0 1 9 0	
EGFR	Non-amplified (r	n=80)				•		1.13	(0.68–1.86)	0.180	
	Low (n=66)			•				0.47	(0.25–0.86)	0.012	
HENJ	High (n=90)					•		1.33	(0.79–2.24)	0.013	
	Low (n=115)					•		1.01	(0.65–1.58)	0.006	
FICN	High (n=42)		+					0.36	(0.16–0.81)	0.000	
		1/8	1/4	1/2	1	L	2		4	8	
Favors afatinib ← ── → Favors MTX											



Tumor shrinkage in biomarker populations deriving more pronounced PFS benefit with afatinib



Biological Hypothesis





Conclusions

- The proportion of patients achieving clinical benefit with afatinib over methotrexate was 4 x greater in the EGFR-mAb therapy-naïve subgroup (37% reduction in risk of progression/death) compared with EGFR-mAb pretreated patients (9% reduction)
- The efficacy benefit with afatinib over methotrexate was similar between older (≥65 years) and younger (<65 years) patients
- Afatinib showed more pronounced antitumor effects in patients with p16-negative disease and dysregulation of ErbB pathway-related biomarkers (*EGFR*-amplification, HER3-low, PTEN-high expression)
- Additional samples are being evaluated to provide a more robust readout of clinical outcomes based on these biomarkers



Acknowledgments

 Thank you to all of the patients and their families, and the LUX-Head & Neck study investigators and their teams for participating in this study





Back-up



Methodology for tumor biomarker assessments

Biomarker	Method	Manufacturer: assay	Cut-offs*
p16	IHC	Ventana: CINtec [®] p16	p16-positive= H-score ≥210
EGFR amplification	FISH	Abbott: Vysis™	Amplification= ≥50% of cells with ≥4 copies, or ≥1 cell with ≥8 copies
HER3	IHC	Dako: DAK-H3-IC	H-score ≤50 (low expression)
PTEN	IHC	Cell Signaling: 138G6	H-score >150 (high expression)



Tumor response in biomarker-defined populations

	Afatinib vs MTX		
Biomarker status (cut-offs)*	No. of patients	ORR, %	
p16-positive (H-score ≥210)	23 vs 12	0 vs 8.3	
p16-negative (H-score <210)	135 vs 64	14.1 vs 1.6	
EGFR-amplified [†]	50 vs 16	14.0 vs 0	
EGFR non-amplified	53 vs 27	3.8 vs 0	
HER3-high (H-score >50)	64 vs 26	9.4 vs 0	
HER3-low (H-score ≤50)	49 vs 17	12.2 vs 0	
PTEN-high (H-score >150)	30 vs 12	6.7 vs 0	
PTEN-low (H-score ≤150)	82 vs 33	12.2 vs 0	



Note: This slide shows tumor shrinkage in patients who received prior cetuximab (green) or did <u>not</u> receive prior cetuximab (red) – according to biomarker status



Note: This slide shows tumor shrinkage in all study patients who received prior cetuximab (green) or did <u>not</u> receive prior cetuximab (red)

Tumor shrinkage by treatment and prior treatment with cetuximab

