

EGFR-targeted antibodies: clinical and molecular sensitivity predictors

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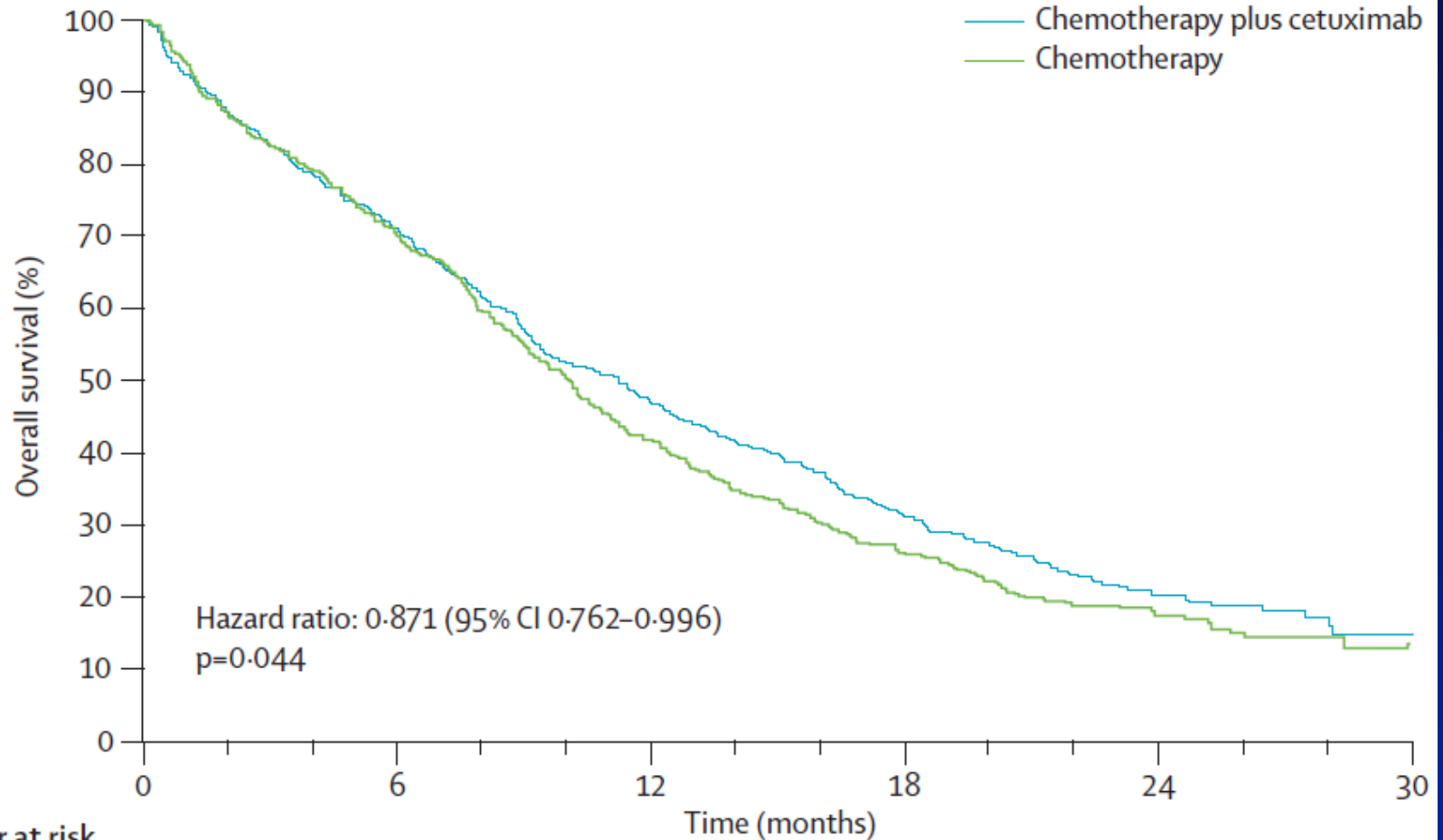
Conflict of Interest

- Honoraria for Advisory Board/Consulting
 - Merck KGaA
 - Roche
 - Eli Lilly
 - AstraZeneca
- Speaker's fee
 - Merck KGaA
 - Roche
 - Eli Lilly
 - Pierre Fabre
 - AstraZeneca
 - Sanofi-Aventis

1st-line chemotherapy plus Cetuximab in advanced NSCLC

- Single-arm phase II studies
- Randomized phase II trials
- Phase III studies
 - FLEX
Pirker R et al. J Clin Oncol 2008, 18 S (abstract 3)
Pirker R et al. Lancet 2009, 373, 1525
 - BMS 099
Lynch T et al. J Thorac Oncol 2007, 2 (Suppl. 4), S340
- Meta-analysis
Thatcher N et al. WCLC 2009

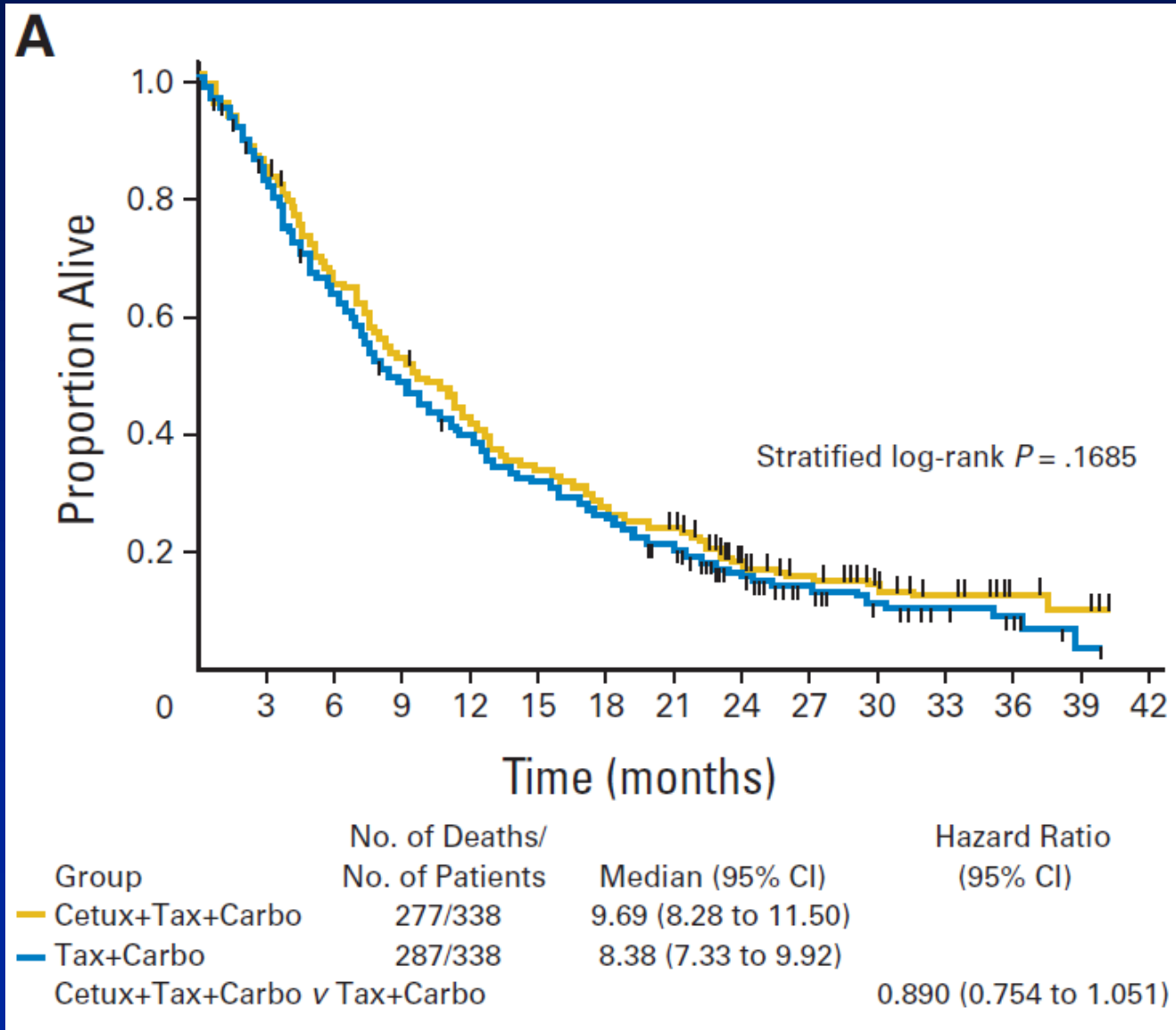
FLEX: Overall Survival



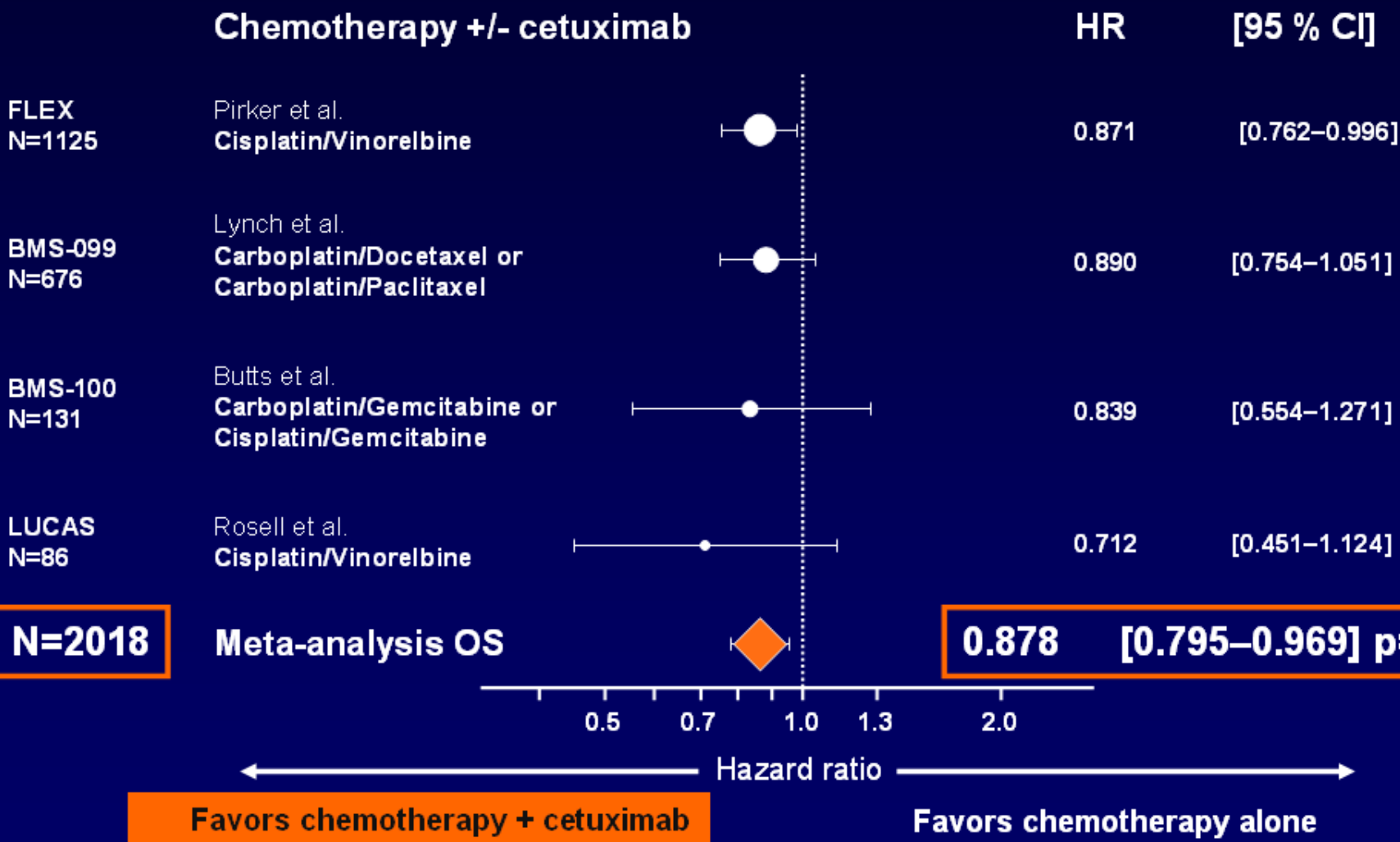
Number at risk		0	6	12	18	24	30
Chemotherapy plus cetuximab	557	383	251	155	53	3	
Chemotherapy	568	383	225	134	48	0	

OS

Lynch TL et al. JCO 2010, 28



Meta-analysis of Overall Survival

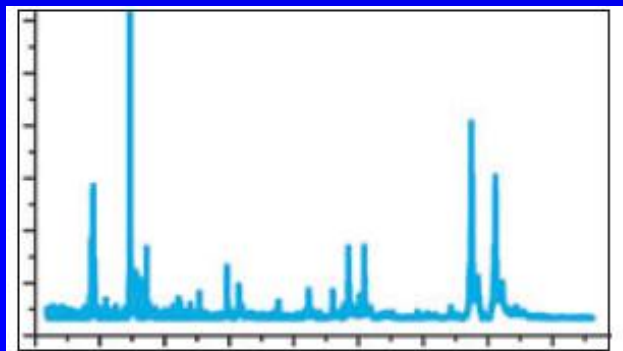
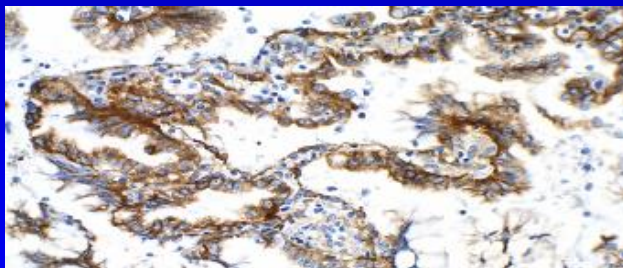
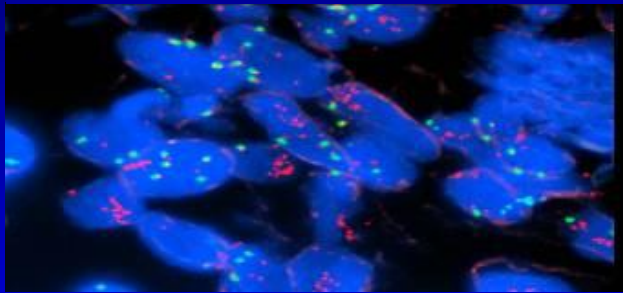
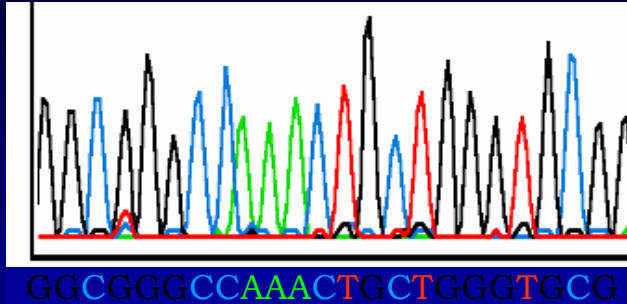


Cetuximab

Predictive markers

- Molecular markers
 - EGFR status
 - KRAS mutations
 - E-cadherin expression
 - HER-2, PI3 kinase mutations
 - Epithelial membrane protein 1 overexpression
 - others
- Immunological markers
- Clinical parameters

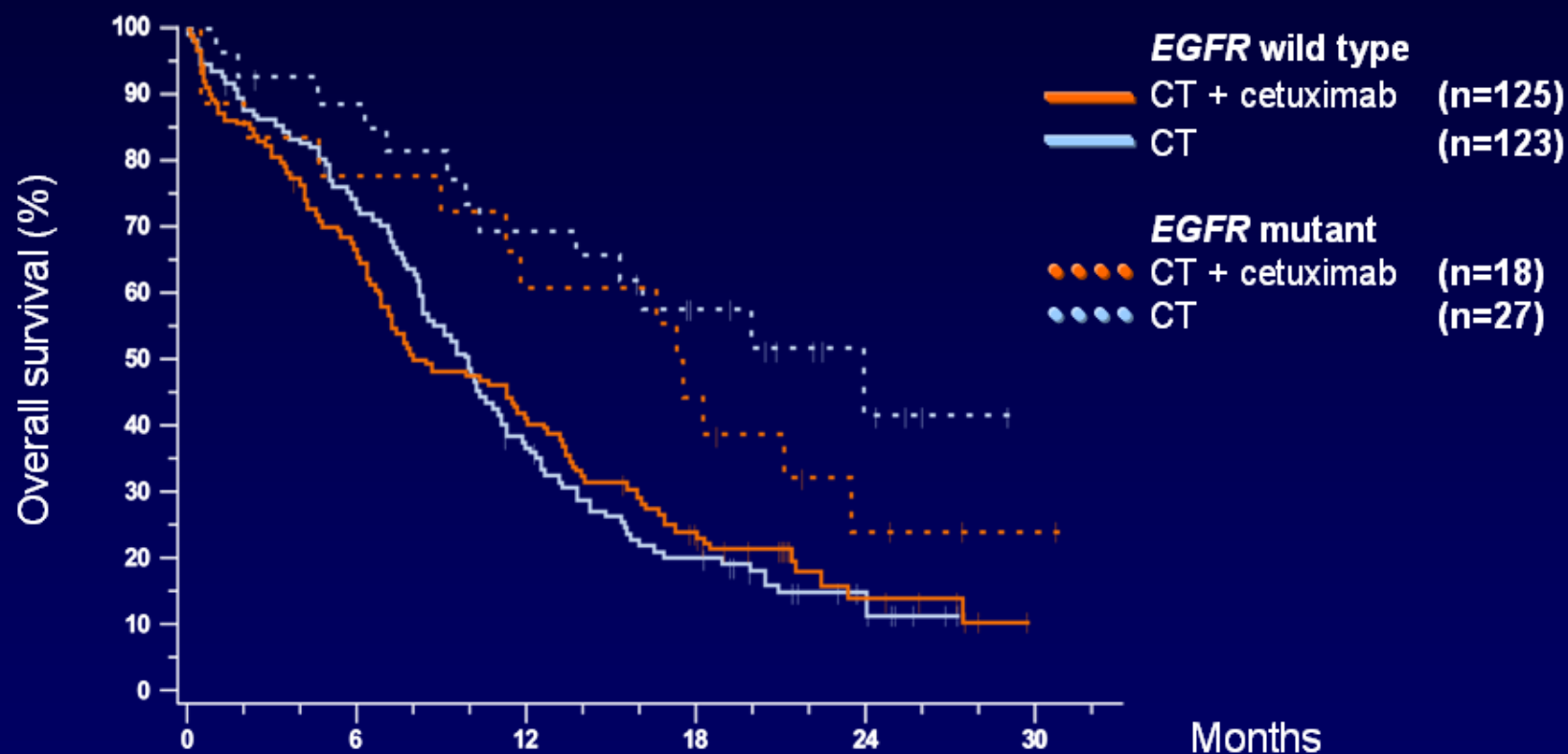
Assessment of EGFR Status



- EGFR mutation status by gene sequencing
- EGFR gene copy number by fluorescence in situ hybridization (FISH)
- EGFR protein expression by immunohistochemistry:
 - 85% positive in FLEX
- Serum Proteomics by MALDI MS

Modified from Courtesy D Gandara

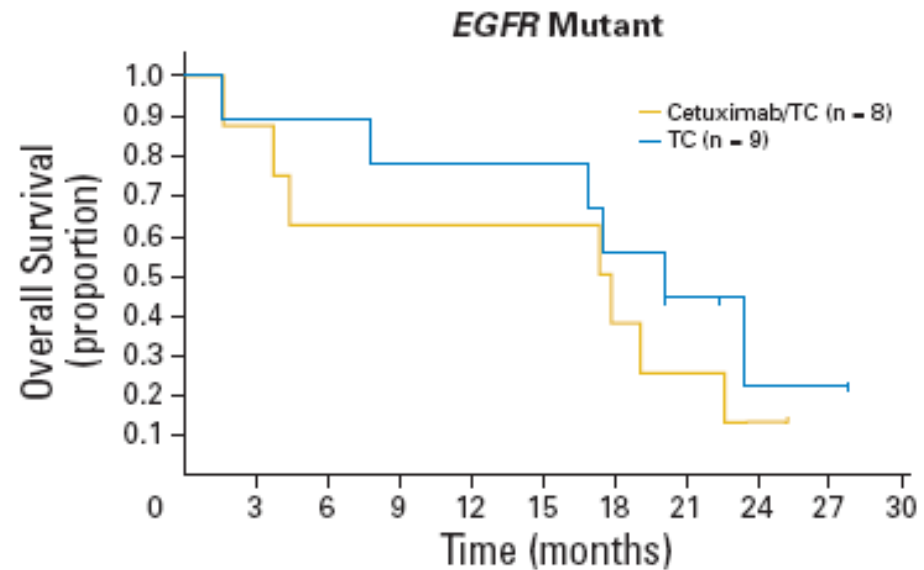
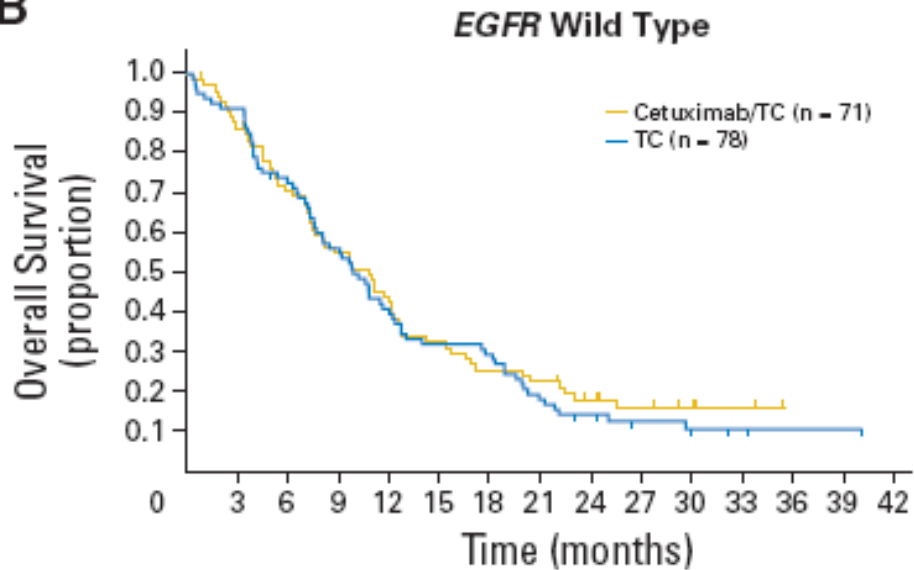
EGFR mutation analysis: OS



Biomarkers: BMS 099

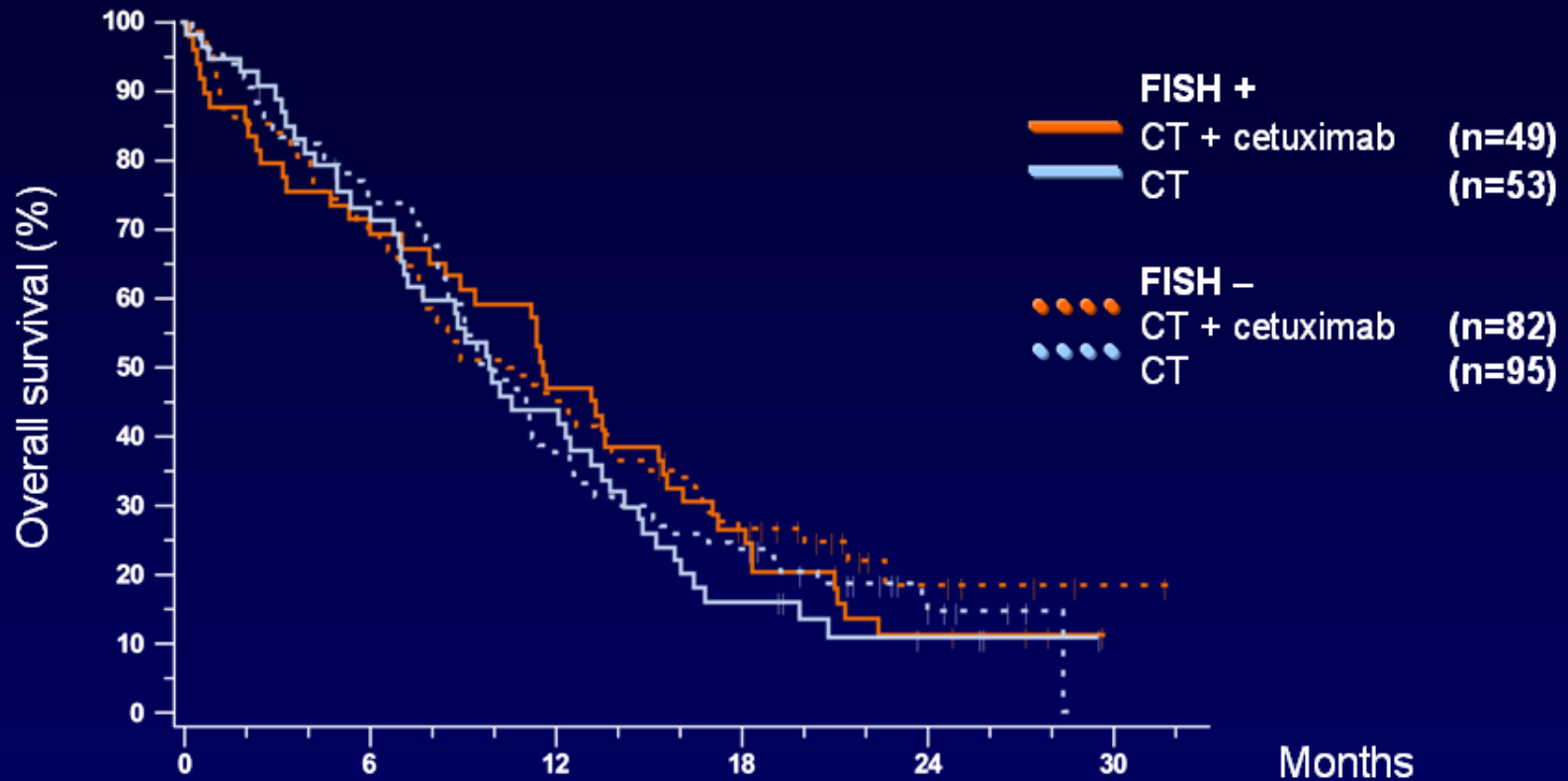
Khambata-Ford S et al. JCO 2010, 28

B



	EGFR Status	Cet + T/C	T/C	HR (95% CI)	P
OS, median	Wild type	9.8 mo	9.8 mo	0.91 (0.64 to 1.29)	.61
	Mutant	17.6 mo	20.0 mo	1.62 (0.54 to 4.88)	.38

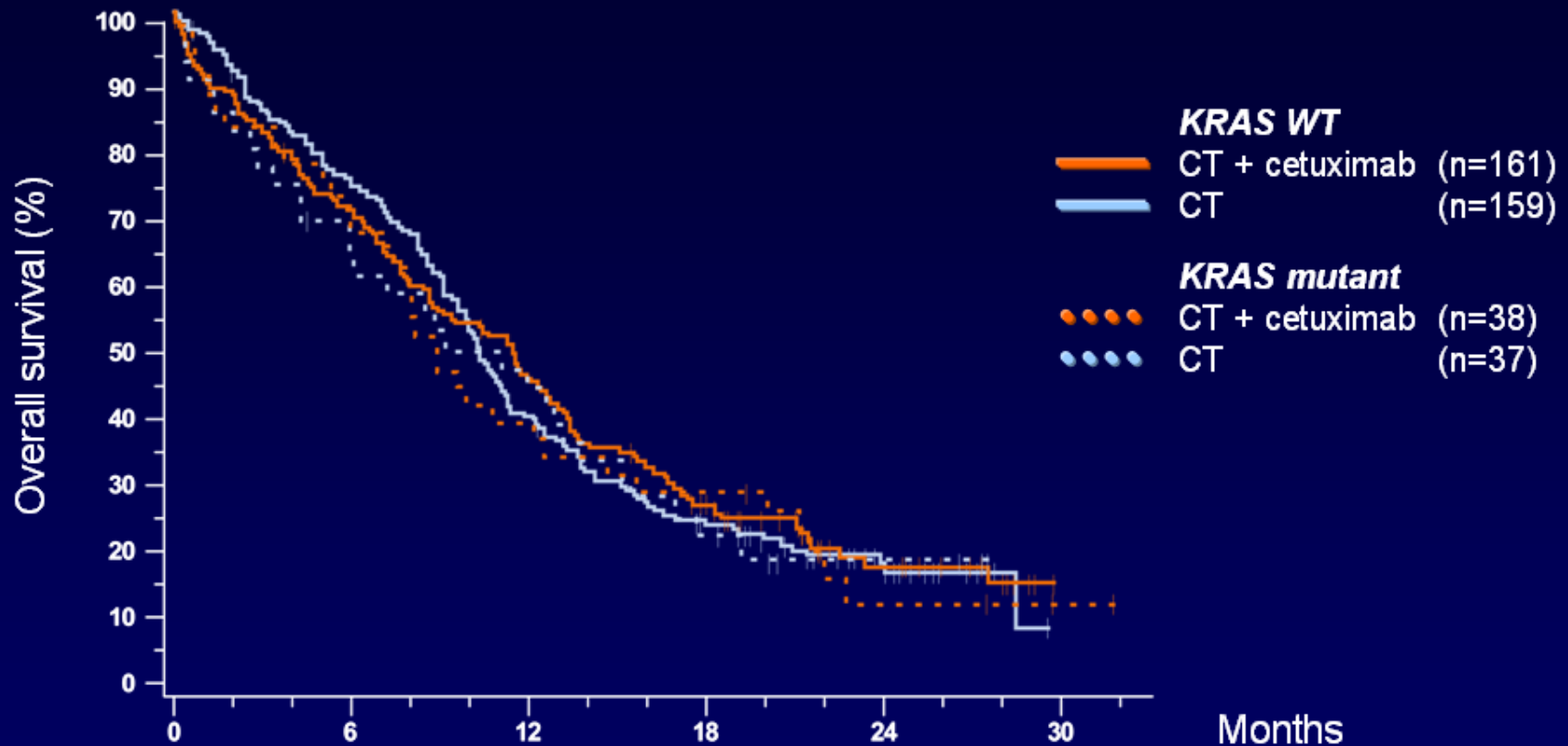
EGFR FISH analysis: OS



	FISH status	CT + cetuximab	CT	HR (95% CI)	p-value
Median OS	FISH -	10.6 mo	10.0 mo	0.91 (0.65–1.26)	0.56
	FISH +	11.6 mo	9.9 mo	0.85 (0.56–1.29)	0.44

CI, confidence interval; CT, chemotherapy; HR, hazard ratio; OS, overall survival

KRAS mutation analysis: OS

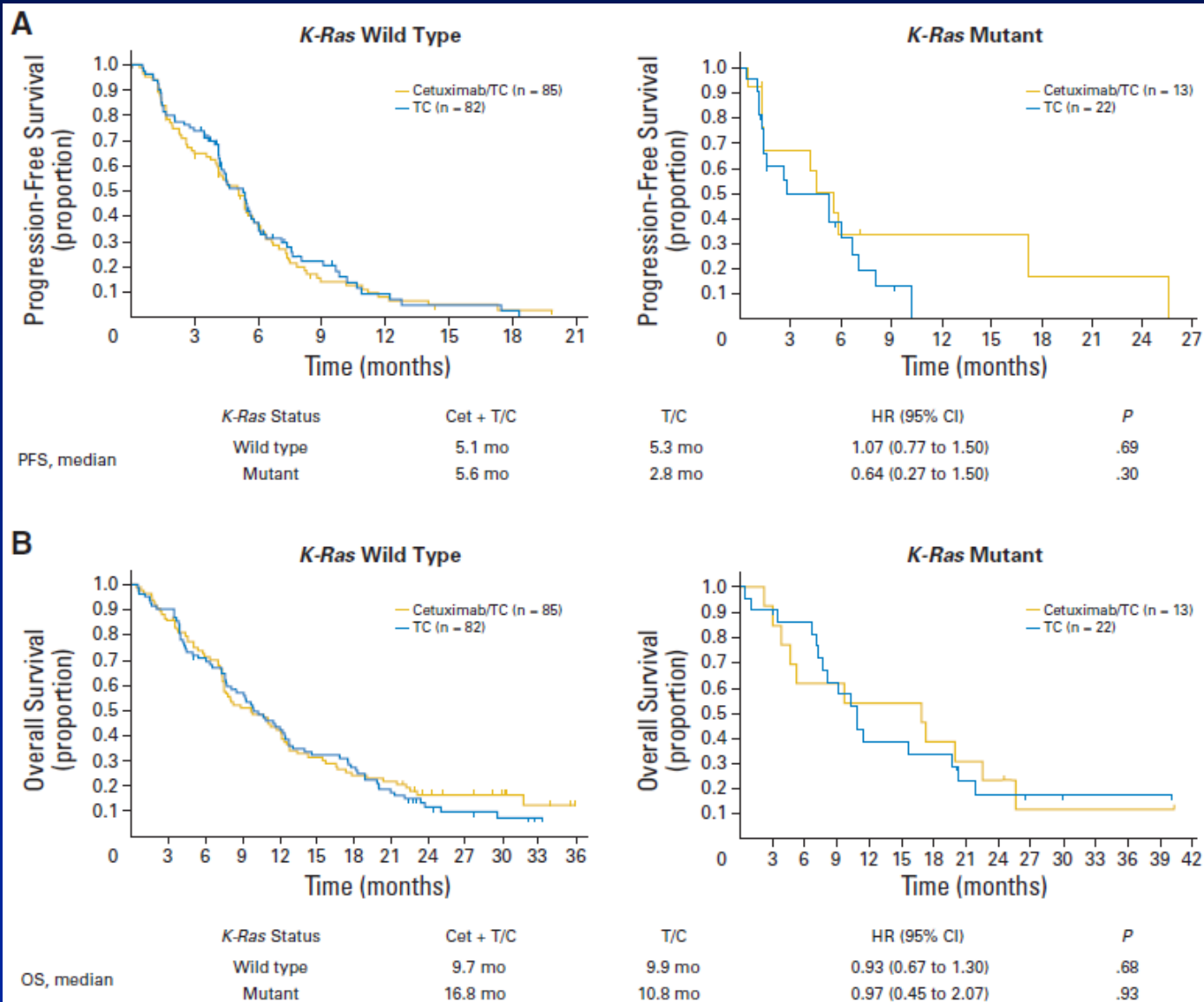


	<i>KRAS</i> status	CT + cetuximab	CT	HR (95% CI)	p-value
Median OS	Wild type	11.4 mo	10.3 mo	0.96 (0.75–1.23)	0.75
	Mutant	8.9 mo	11.1 mo	1.00 (0.60–1.66)	1.00

CI, confidence interval; CT, chemotherapy; HR, hazard ratio; OS, overall survival

Biomarkers: BMS 099

Khambata-Ford S et al. JCO 2010, 28



Cetuximab trials (FLEX, BMS 099)

Predictive factors

- No predictive role of
 - EGFR mutation status
 - EGFR copy numbers (FISH)
 - K-ras mutation status
- Number of samples do not allow to exclude small differences
- Assays reliable enough ?
- Results raise several issues on predictive factors in advanced NSCLC

Predictive factors in NSCLC

Issues

- Heterogeneous disease (molecularly & clinically)
- Complex molecular mechanisms/pathways
- Clinical heterogeneity
 - Many organs as metastatic sites
 - Brain metastasis
 - Co-morbidity
 - Performance status
- Are tumor specimens representative of a complex disease ?
- Assays: reliable, fast, available for routine practice
- Molecular imaging in the future
 - Sensitivity might be too low borderline cases

1st-cycle rash: CT + cetuximab incidence and grade

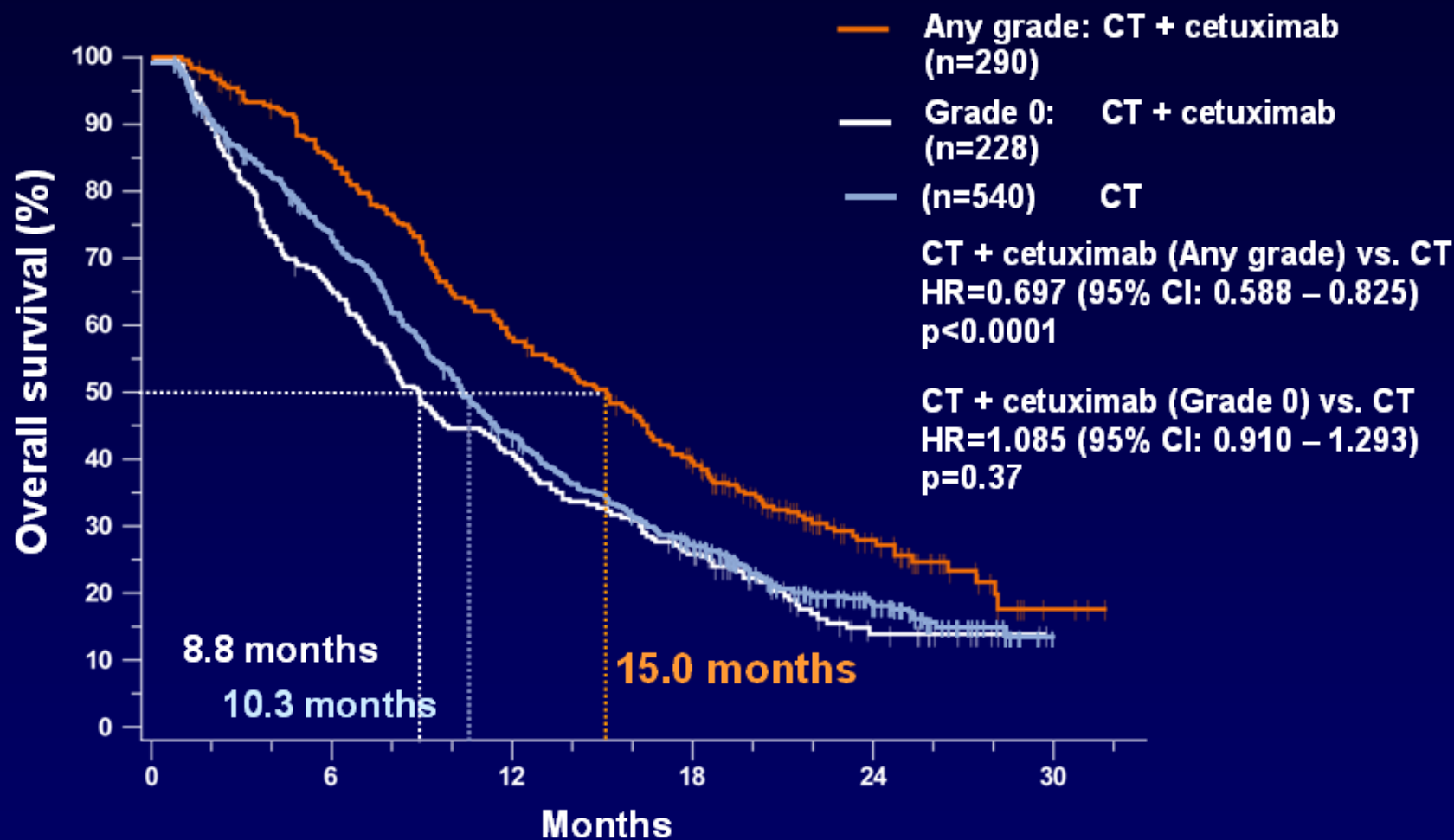
Rash	vs	No rash
Any grade (1-3)		Grade 0
n=290 (56%)		n=228 (44%)

518/557 patients were included in analysis

1st-cycle rash	No. of patients (%)	
Grade 0	228	(44%)
Grade 1	170	(33%)
Grade 2	92	(18%)
Grade 3	28	(5%)
Grade 4	0	—

CT, chemotherapy

1st-cycle rash and OS



CI, confidence interval; CT, chemotherapy; HR, hazard ratio; OS, overall survival
O'Byrne KJ, et al. J Clin Oncol 2009; 27(15s):suppl; abstr 8007
Gatzemeier et al, WCLC 2009, oral presentation (Abstract B2.3)

Summary

- Cetuximab added to first-line chemotherapy increases survival of patients with advanced NSCLC
- EGFR-activating mutations, EGFR copy numbers and KRAS mutation status do not predict benefit
- First-cycle rash associated with better prognosis