



## ESMO 2012 NSCLC - IMMUNOTHERAPY, SCLC AND MESOTHELIOMA

# Poster Discussion - 31180PD - #1181PD- #1521PD- #1192PD

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VIENNA  
2012

**ESMO** congress



Institut de cancérologie  
**GUSTAVE ROUSSY**  
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## → Disclosures

- No personal financial disclosures
- Institutional grants for clinical and translational research
  - Abbott, Amgen, AstraZeneca, Boehringer-Ingelheim, Lilly, Pfizer, Roche-Genentech, Sanofi-Aventis, Clovis

## → Early NSCLC - Abstracts overview

- **Early diagnosis**

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# → Classical Hodgkin's lymphoma

## Standard treatments in adults patients - EORTC

5 yrs OS

Stages I – II*	ABVD x 3 +	
W/O risk factor	Radiotherapy 30 Gy involved-fields	94-100%

Stages I – II*	ABVD x 4 +	
With Risk factors	Radiotherapy 30 Gy involved-fields	92-95%

\* supra-diaphragmatic stages

### EORTC - Risk factors

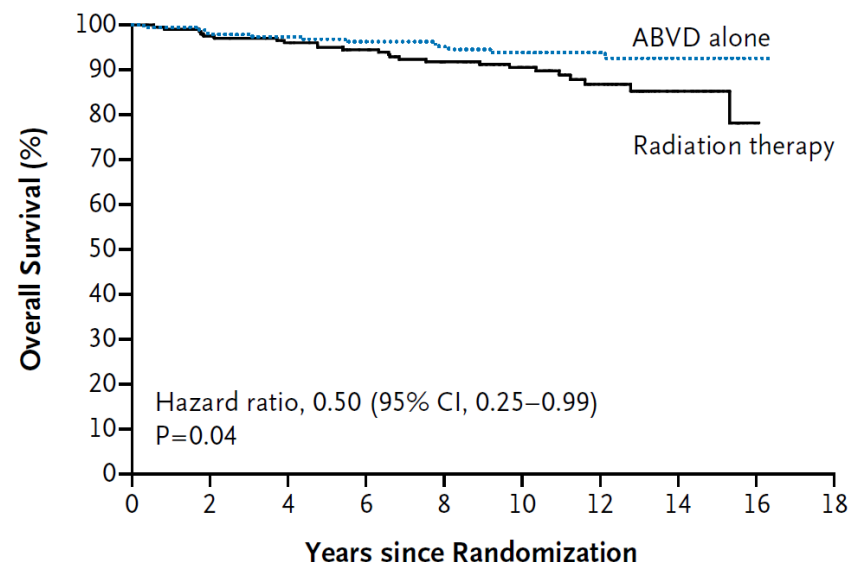
A - Large mediastinal mass\*

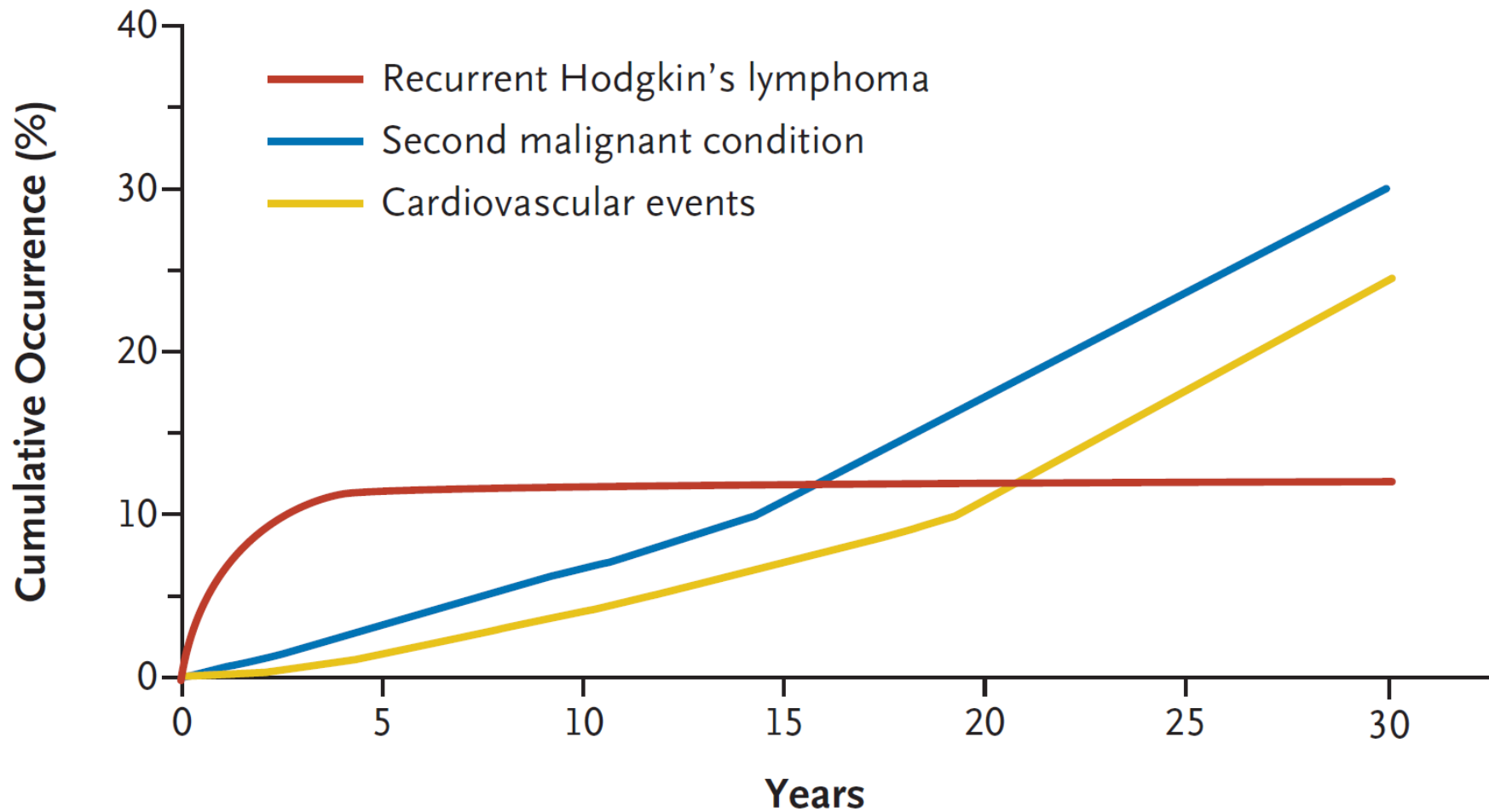
B - Age  $\geq 50$  ans

C - A and ESR  $\geq 50$  mm

D -  $\geq 4$  nodal sites

E - B + ESR  $> 30$





## → Hodgkin's lymphoma

### Second malignancies in 32591 pts

	Relative risk (95% CI)	Absolute excess risk
All solid tumours	2.0 (1.9–2.4)	33.1
Oesophagus	2.8 (1.8–4.0)	0.7
Stomach	1.9 (1.5–2.4)	1.5
Small intestine	1.5 (0.3–3.5)	0.1
Colon	1.6 (1.4–1.9)	2.0
Lung	2.9 (2.6–3.2)	9.7
Female breast	2.0 (1.8–2.3)	10.5
Leukaemia	9.9 (8.7–11.2)	8.8

Data from Dores and colleagues.<sup>6</sup> All  $p < 0.05$ , except for small intestine.

## → Hodgkin's lymphoma Second lung cancer in 5519 pts

### BY TIME SINCE TREATMENT (YEARS)

	Number of lung cancers	Relative risk (95% CI)	Absolute excess risk
0-4	18	1.9 (1.1-2.9)	3.9
5-9	25	3.8 (2.5-5.5)	13.8
10-14	21	4.9 (3.1-7.4)	21.5
≥15	14	5.2 (3.0-8.5)	28.3



## → Hodgkin's lymphoma

### Second lung cancer in 222 pts

#### • RELATION WITH PREVIOUS TREATMENT AND CIGARETTE SMOKING

Treatment		Other*		Moderate to heavy smoker†	
Radiation $\geq 5$ Gy	Alkylating agents	Relative risk (95% CI)	p	Relative risk (95% CI)	p
No	No	1.0‡		6.0 (1.9–20.4)	0.002
Yes	No	7.2 (2.9–21.2)	<0.001	20.2 (6.8–68.0)	<0.001
No	Yes	4.3 (1.8–11.7)	<0.001	16.8 (6.2–53.0)	<0.001
Yes	Yes	7.2 (2.8–21.6)	<0.001	49.1 (15.1–187)	<0.001

\*Includes non-smokers, light cigarette smokers (less than one pack per day), former cigarette smokers, smokers of cigars and pipes only, and patients for whom tobacco smoking habit was not stated. †At least one pack-per-day cigarette smokers.

‡Reference group.

# LUNG MALIGNANCIES AND SECOND NEOPLASIAS IN PATIENTS WITH HODGKIN'S LYMPHOMA

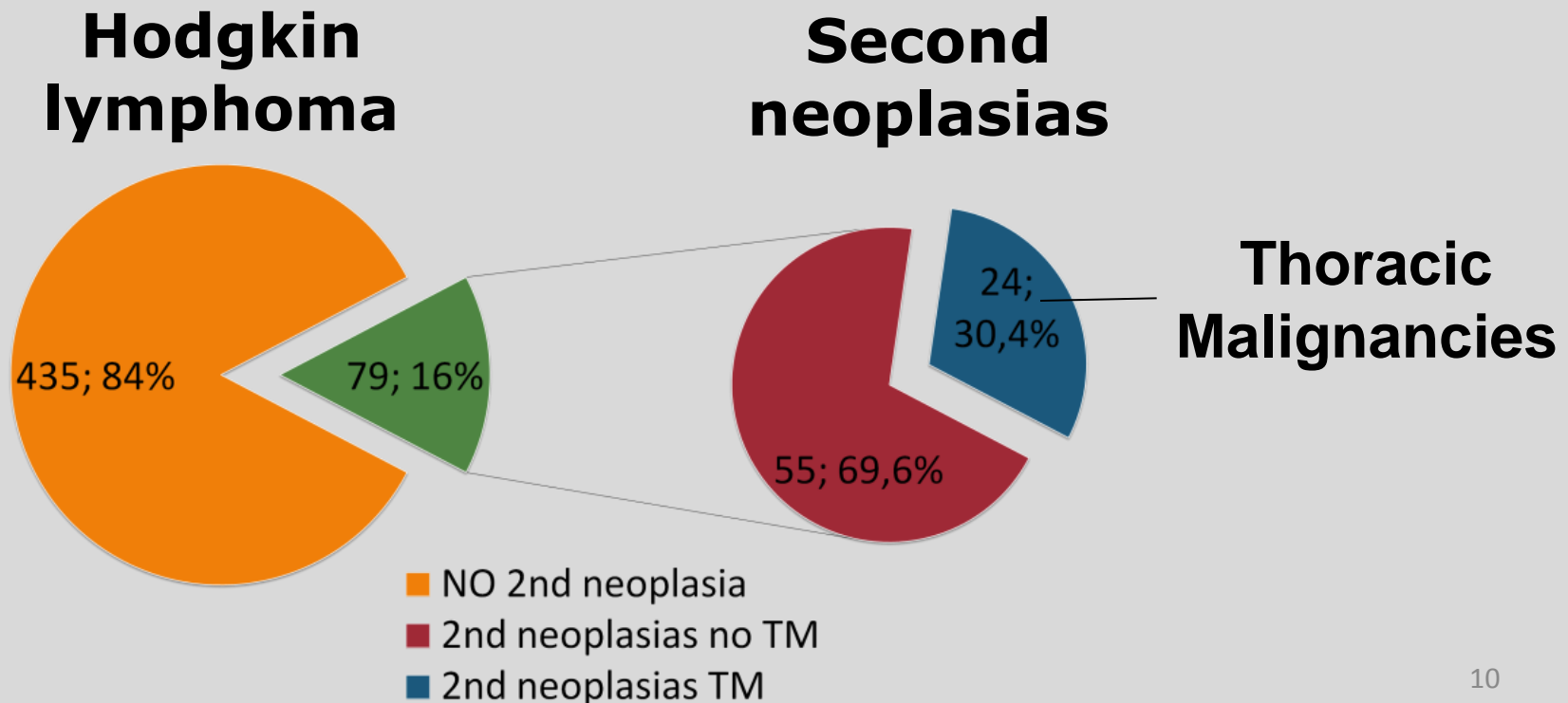
E. Almagro-Casado<sup>1</sup>, D. Pérez-Callejo<sup>1</sup>, A. López-González<sup>3</sup>, P. Ibeas<sup>1</sup>, A. Ruiz-Valdepeñas<sup>1</sup>, M. Palka<sup>1</sup>, C. Maximiano<sup>1</sup>, M. Méndez García<sup>1</sup>, S. Mellor<sup>2</sup>, M. Provencio Pulla<sup>1</sup>;

<sup>1</sup>Medical Oncology ,Hospital Puerta de Hierro, Majadahonda (Spain), <sup>2</sup>Internal Medicine (idem) <sup>3</sup> Medical Oncology ,Hospital de León (Spain). Corresponding e-mail:

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## BACKGROUND AND METHODS

### Patients distribution (n=514)



# LUNG MALIGNANCIES AND SECOND NEOPLASIAS IN PATIENTS WITH HODGKIN'S LYMPHOMA

## RESULTS

- Thoracic malignancies were more prevalent in men (88%  $P=0,001$ )

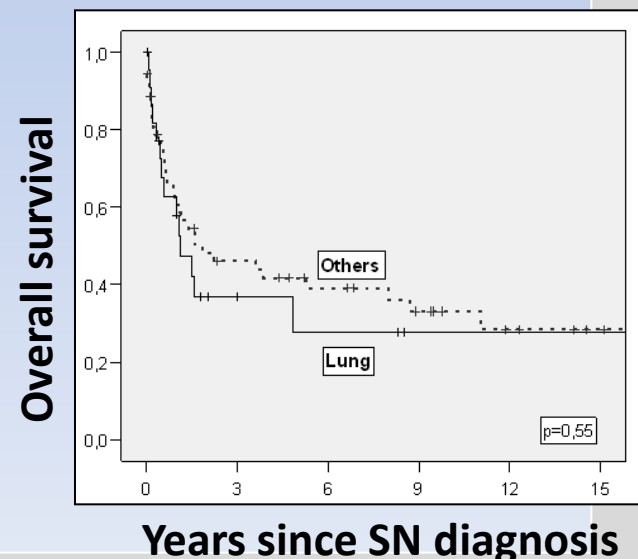
What is the smoking status ?

- Treatment with radiotherapy was associated with increased risk of thoracic malignancies (OR 2,9 IC 95% 1,1- 7,7  $P=0,033$ )

Confirmation of previous studies

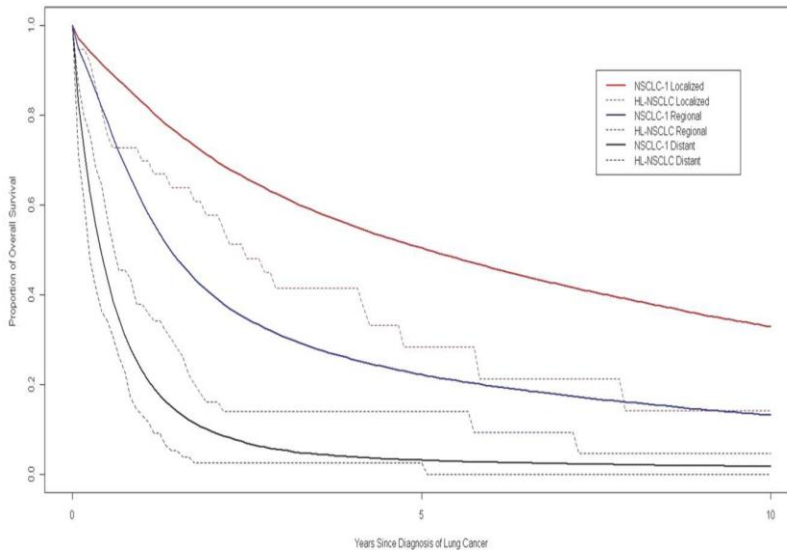
- Elapsed time to diagnosis of second neoplasia was longer in TM group  
(median of 16,4 years in TM and 9,7 in HL  $P=0,03$ )

Late screening ?

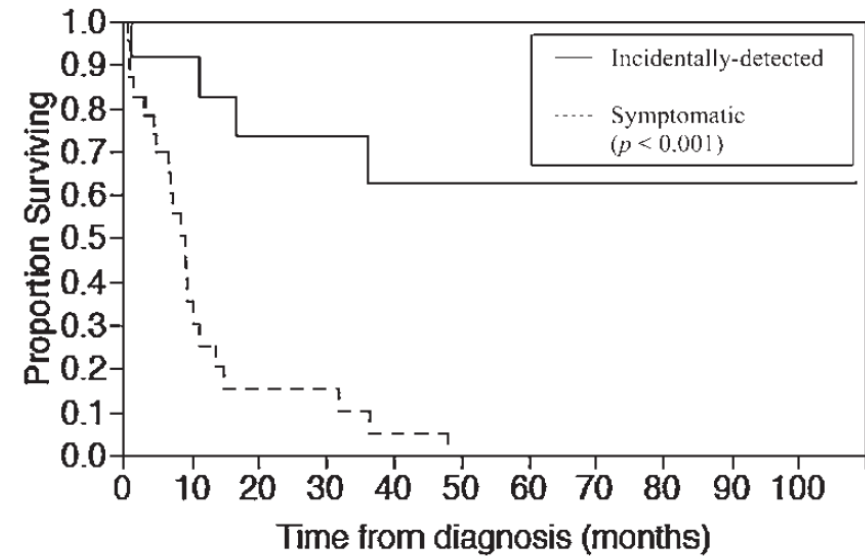


## → Lung cancer after Hodgkin's lymphoma

Poorer prognosis  
(SEER database)



Better survival  
if incidentally-detected



# Population to screen ?

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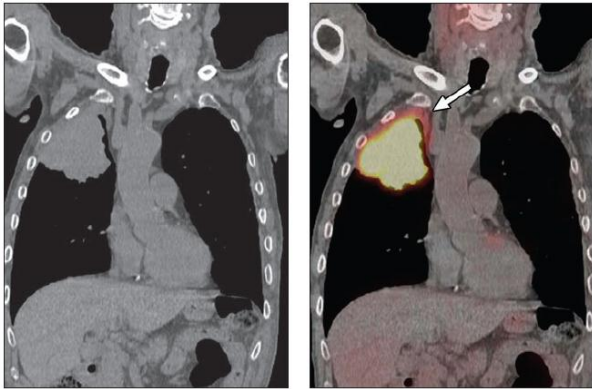
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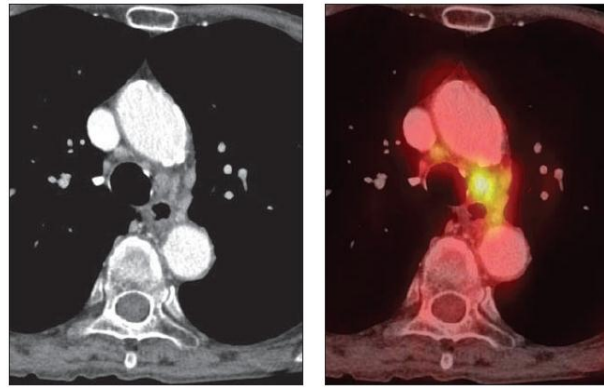
## → PET/CT in NSCLC staging

# T



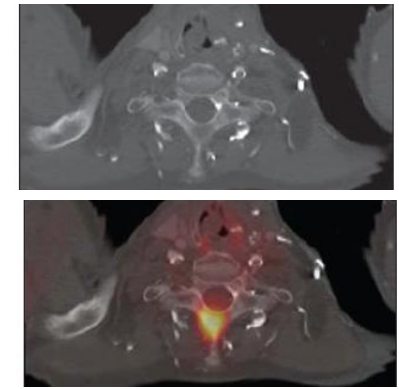
PET/CT accurately predicted the T stage in 82% of cases compared with 68% with CT alone

# N



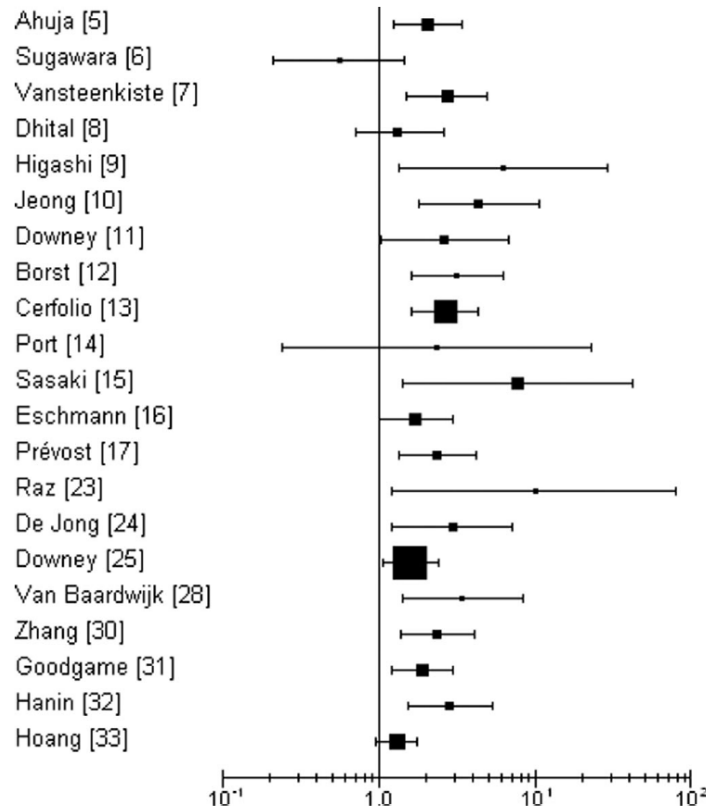
The accuracy of lymph node staging by PET and PET/CT is up to 56% and 78% when compared with surgical staging

# M



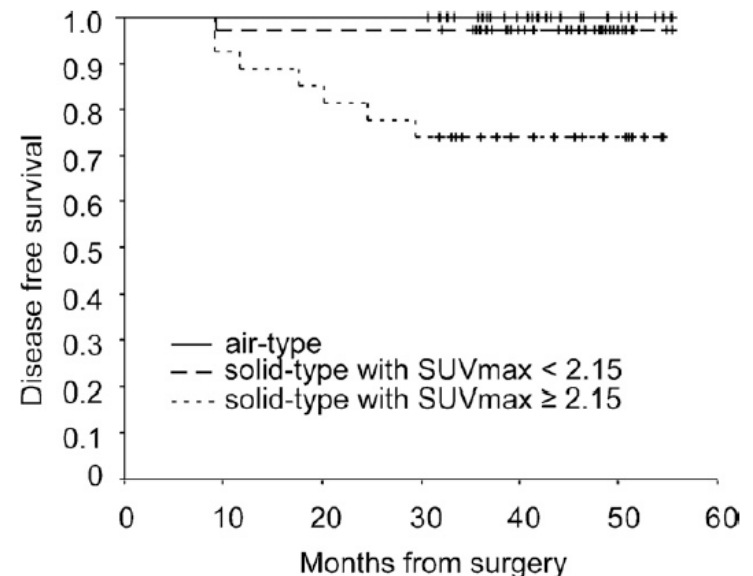
PET/CT discovers occult metastatic disease in up to 29% of patients.

# → Tumor (T) FDG uptake is of prognostic value



**HR (95% CI) on a logarithmic scale.**

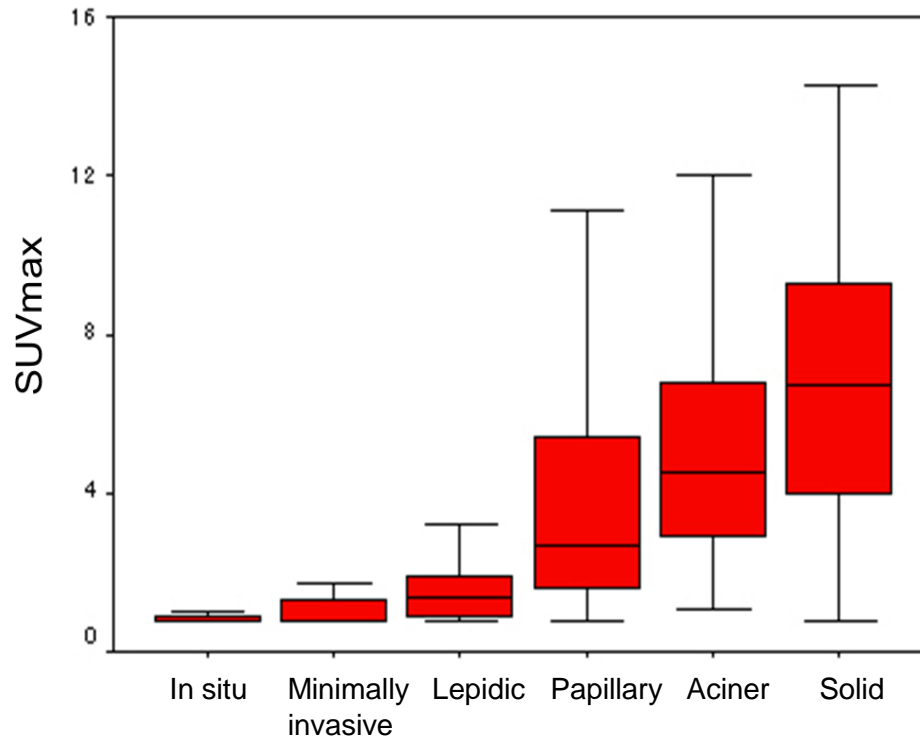
*By convention, HR 1 means that patients with a higher standard uptake value (SUV) on the primary tumor have a worse prognosis.*



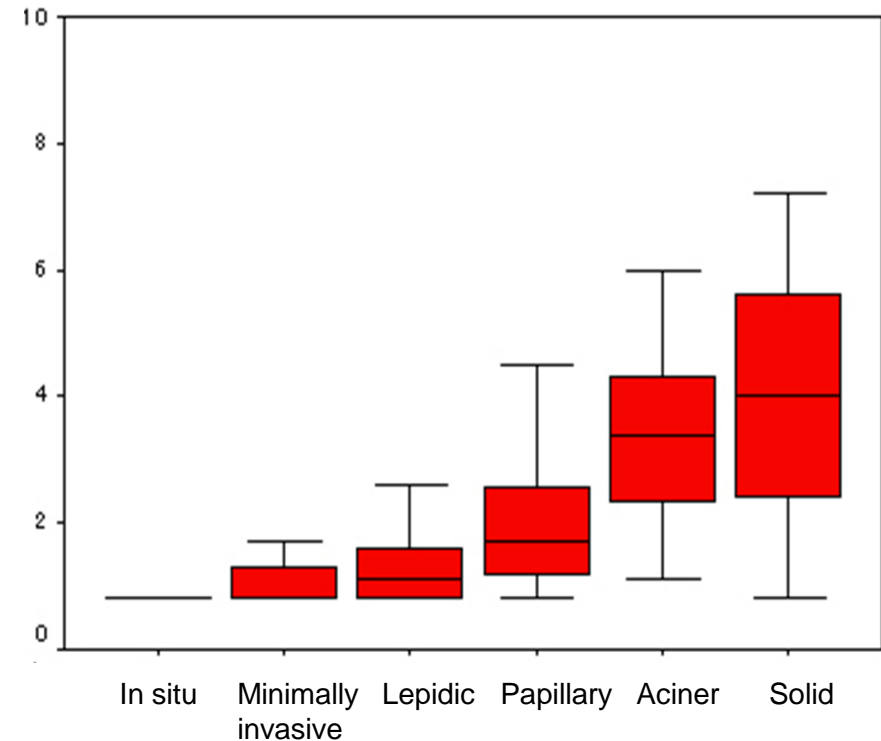
**Stage Ia disease, n=100**

# Correlation of different histologic subtypes with SUVmax

【All case (n=287)】

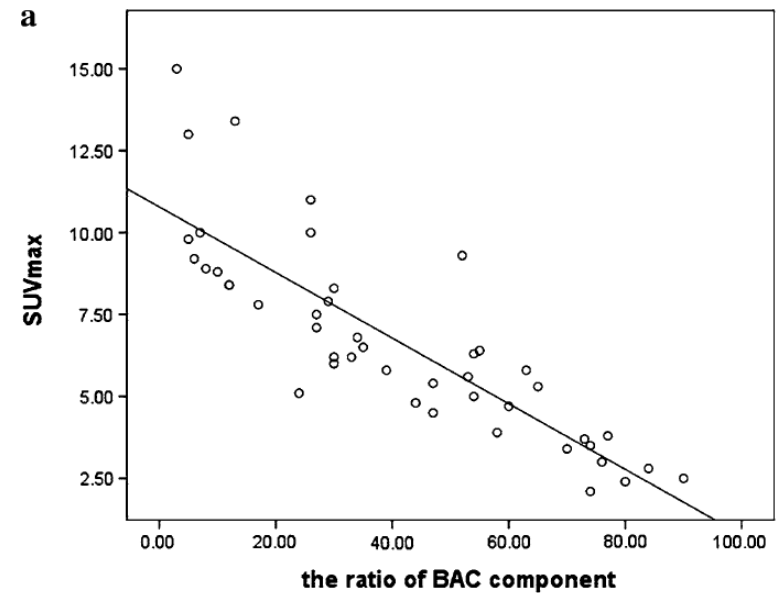
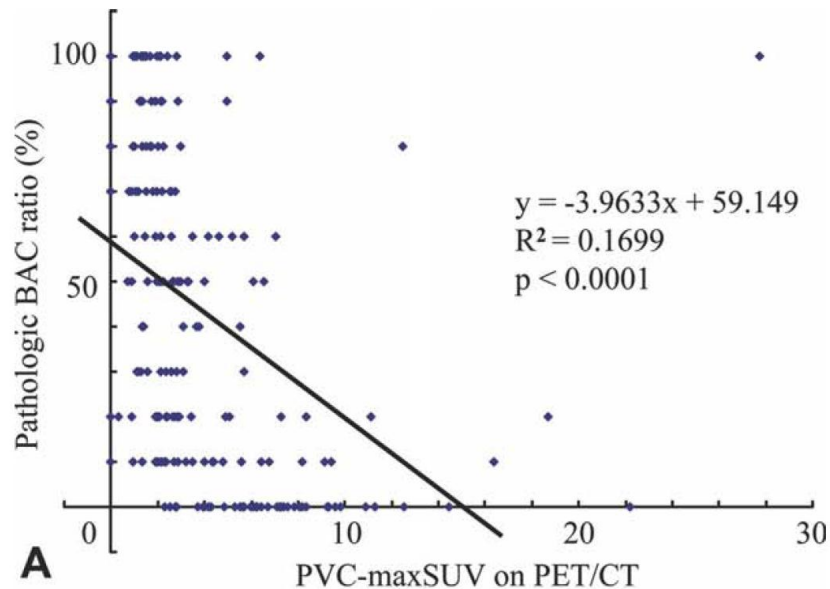


【Stage IA (n=166)】





## → Lepidic (ex-BAC) influences FDG uptake



The SUVmax and the ratio of lepidic component had significant inverse correlation

# Correlation of different histologic subtypes with GLUT1 and HIF1 $\alpha$

## 【GLUT1】

	in situ:13	Lepdic:36	Papillary: 52	Acinar:34	Micropapillary: 9	Solid:3
<b>0</b>	11	34	20	14	1	0
<b>1</b>	2	1	17	5	3	0
<b>2</b>	0	0	10	9	4	0
<b>3</b>	0	1	5	6	1	3
mean $\pm$ SD	0.15 $\pm$ 0.36	0.11 $\pm$ 0.52	1.0 $\pm$ 0.98	1.2 $\pm$ 1.2	1.6 $\pm$ 0.83	3

P=0.019

## 【HIF1 $\alpha$ 】

	in situ:13	Lepdic:36	Papillary: 52	Acinar:34	Micropapillary: 9	Solid:3
<b>0</b>	13	35	45	26	8	0
<b>1</b>	0	1	5	3	1	1
<b>2</b>	0	0	1	2	0	0
<b>3</b>	0	0	1	3	0	2
mean $\pm$ SD	0	0.028 $\pm$ 0.16	0.19 $\pm$ 0.56	0.47 $\pm$ 0.95	0.11 $\pm$ 0.31	2.3 $\pm$ 0.94

P=0.010



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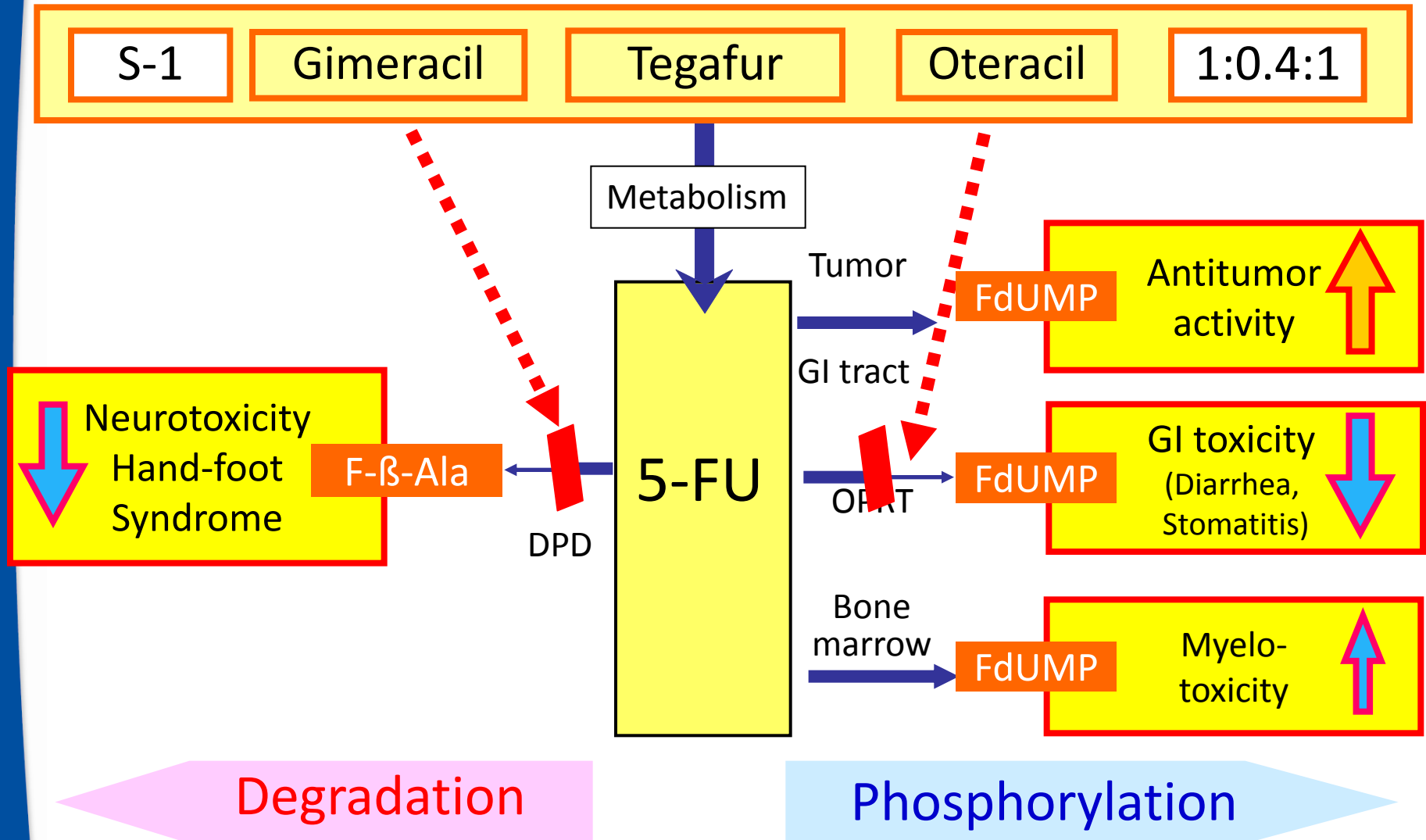
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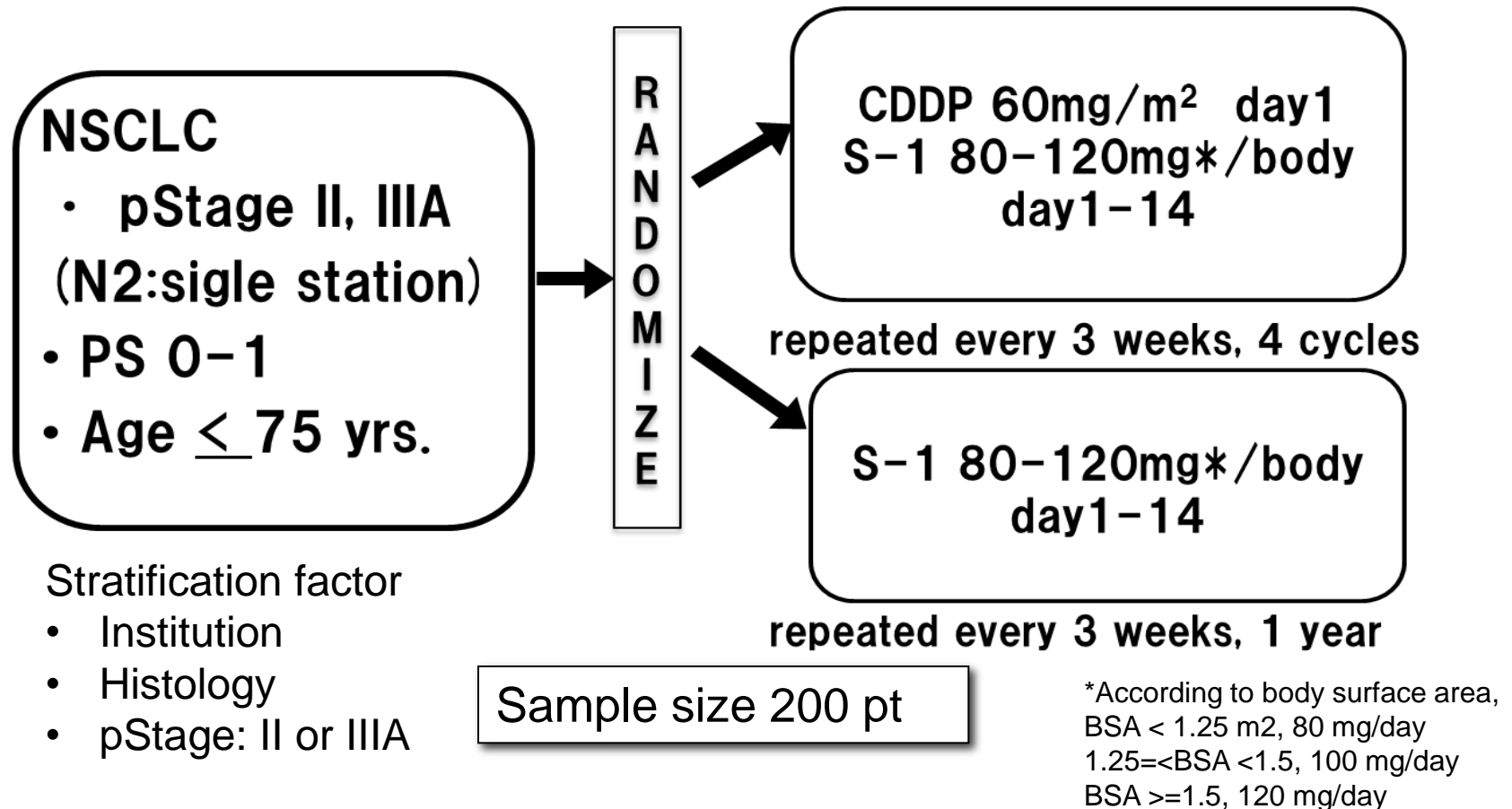
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## → Biochemical Action of S-1



## → Design



**Primary endpoint: Two year relapse free survival**

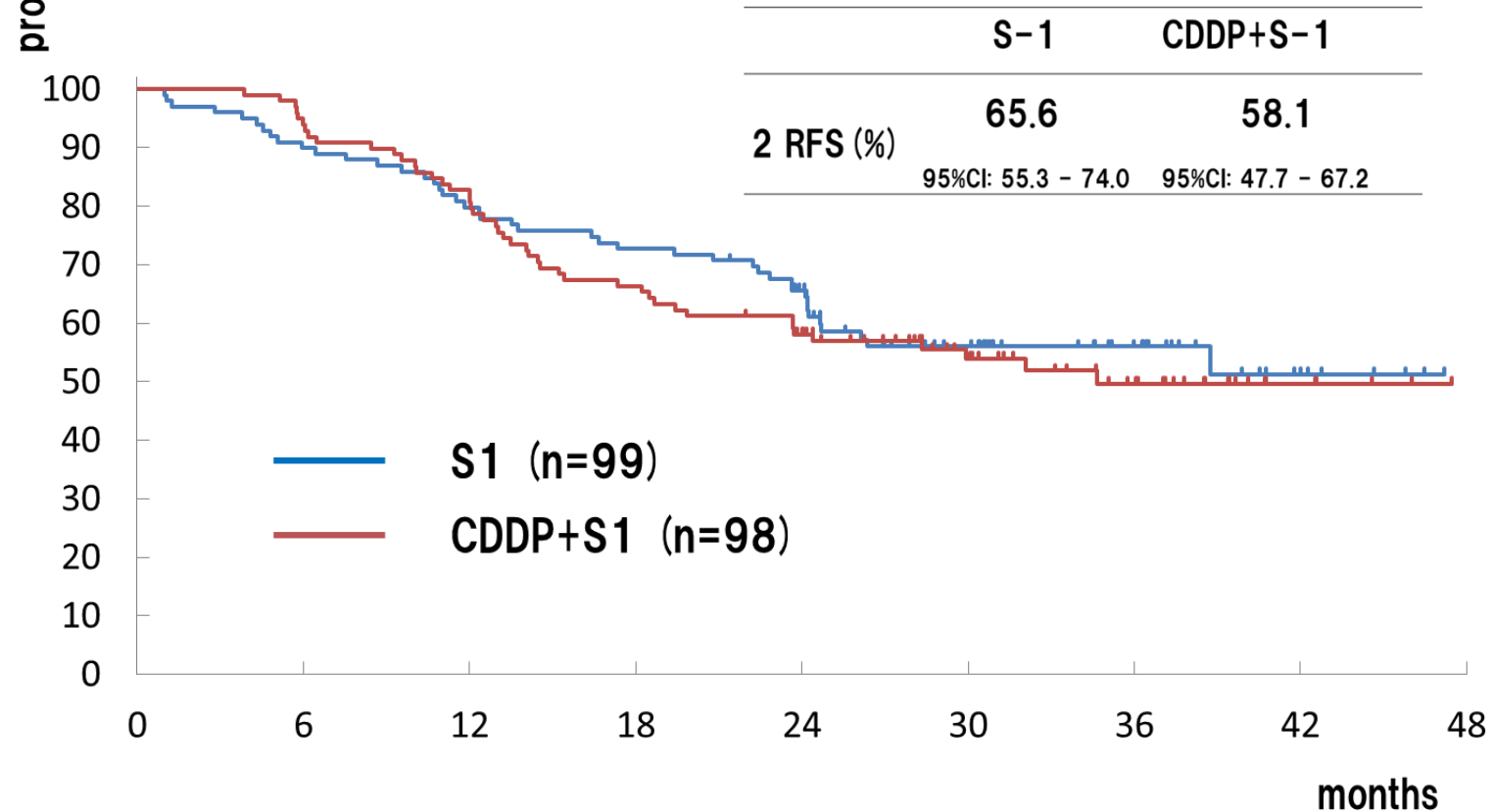
# Adverse events

	S-1 (n=97)	CDDP+S-1 (n=95)	P
	%	%	
	Grade 3 or 4	Grade 3 or 4	
<b>General</b>			
Fatigue	1	4.2	0.209
Anorexia	2.1	9.5	0.032
Nausea	0	6.3	0.014
Vomiting	0	2.1	0.244
Eruption	0	0	
Febrile neutropenia	0	5.3	0.028
<b>Hematological</b>			
Anemia	1	8.4	0.018
Neutropenia	13.4	27.4	0.020
Thrombocytopenia	0	2.1	0.244
<b>Biochemical</b>			
ALT elevation	1	0	1.000
Creatinin elevation	0	1.1	0.495

## Completion of protocol treatment

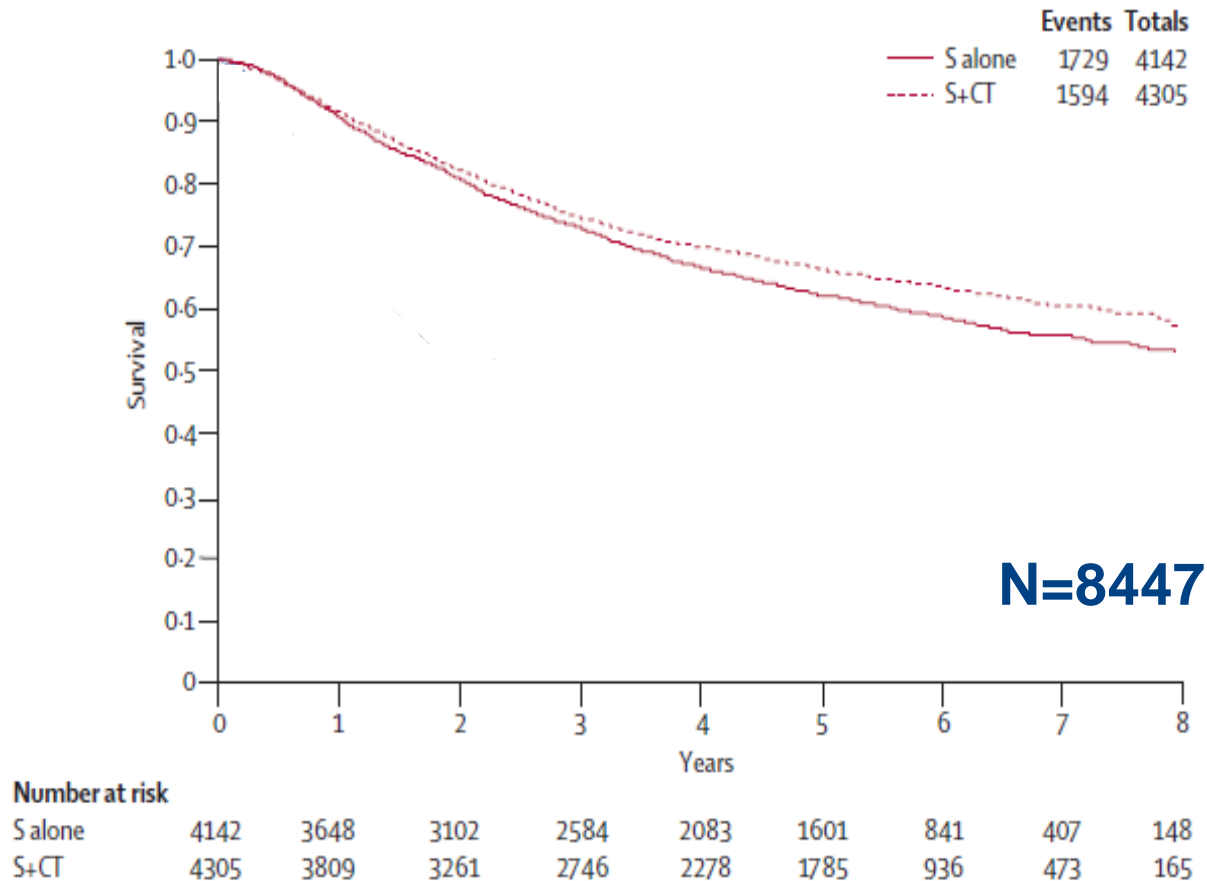
S-1	CDDP+S-1
52.6%	74.7%

# Relapse free survival



Conclusion: S-1 and CDDP+S-1 demonstrated a promising activity with a favorable 2-year relapse free survival and safety as an adjuvant chemotherapy in patients with completely resected NSCLC.

## → NSCLC Meta-analyses Collaborative Group

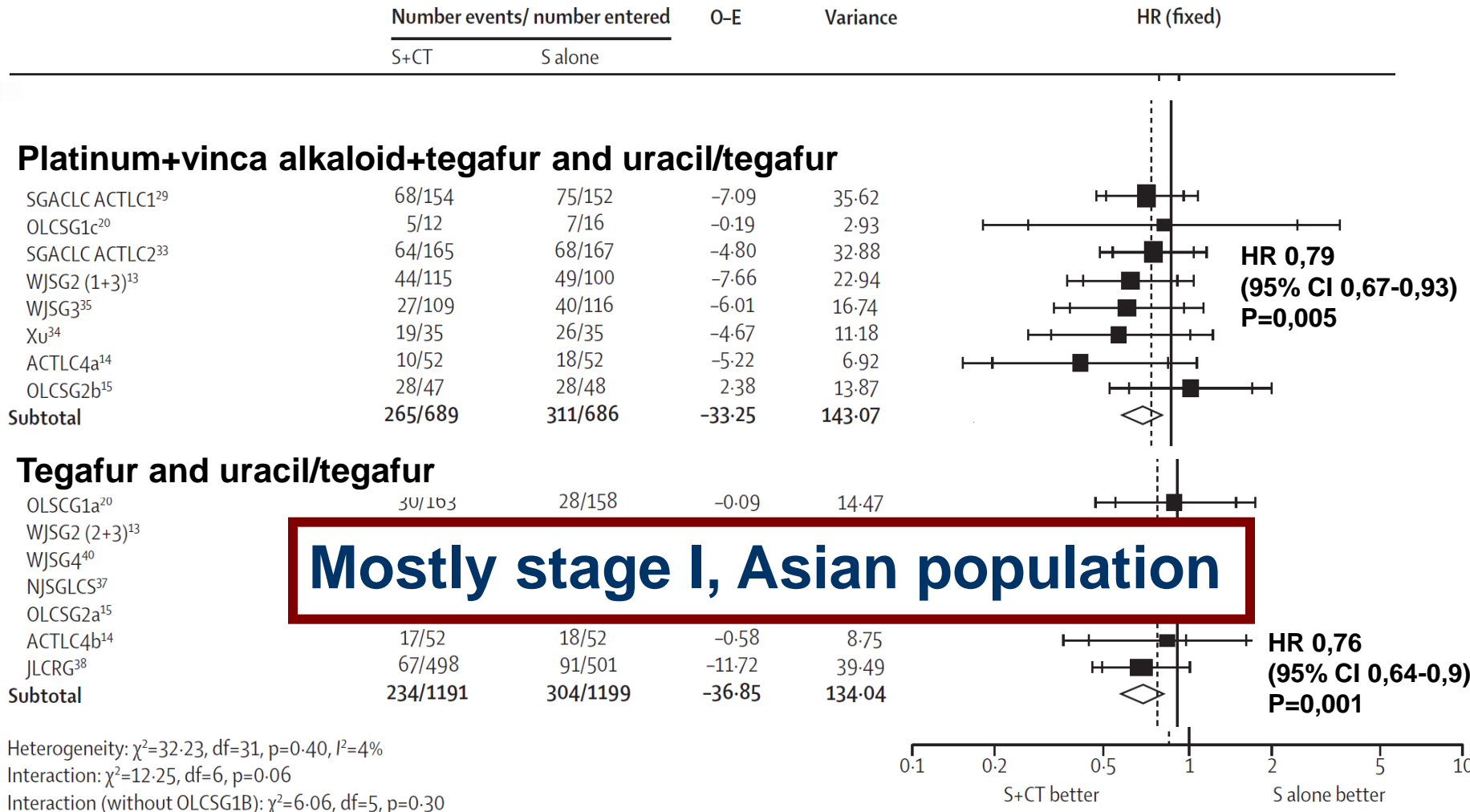


**HR = 0.87 (0.81-0.93)  $p < 0.000001$**

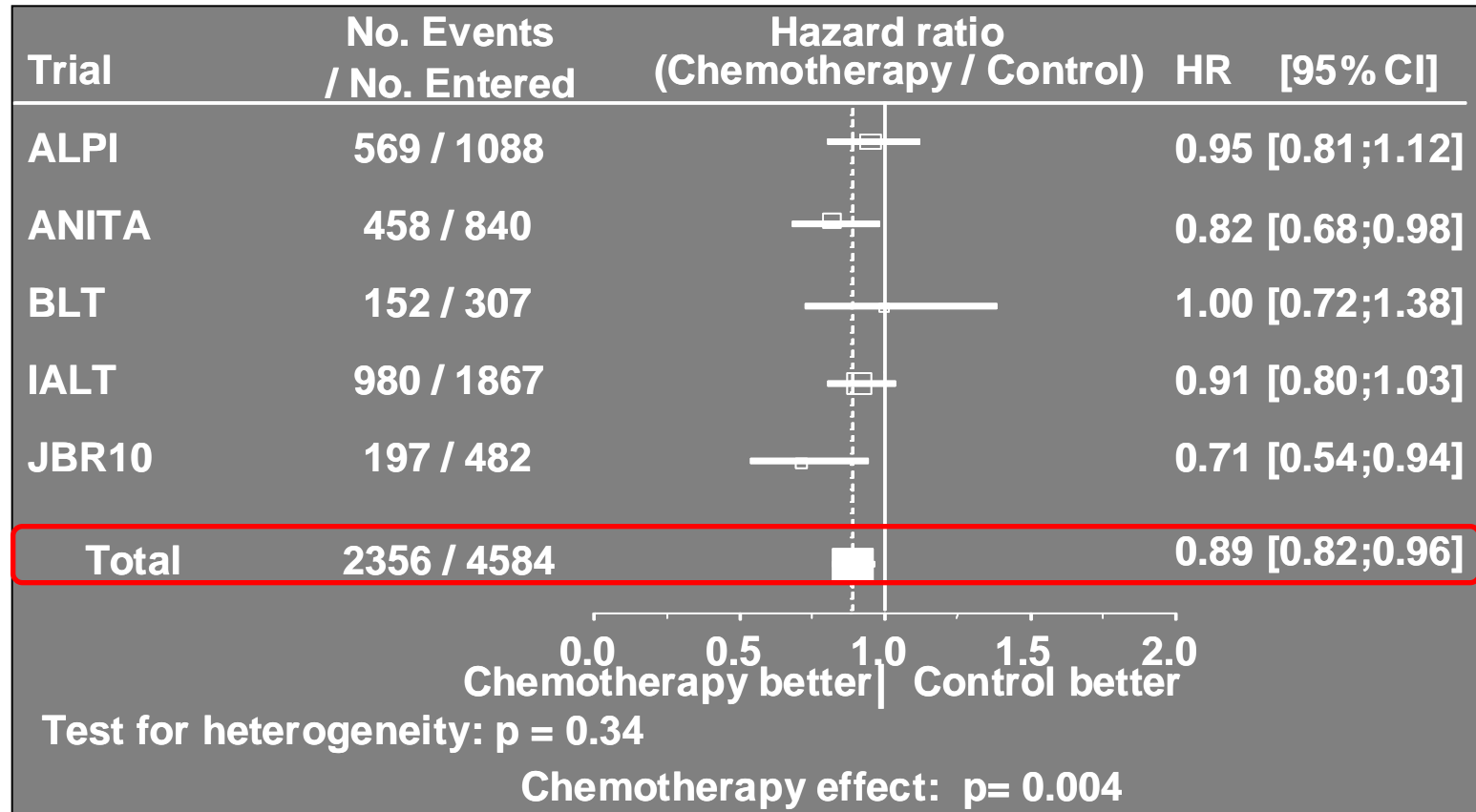
**Absolute benefit of 4% at 5 years**



## → NSCLC Meta-analyses Collaborative Group



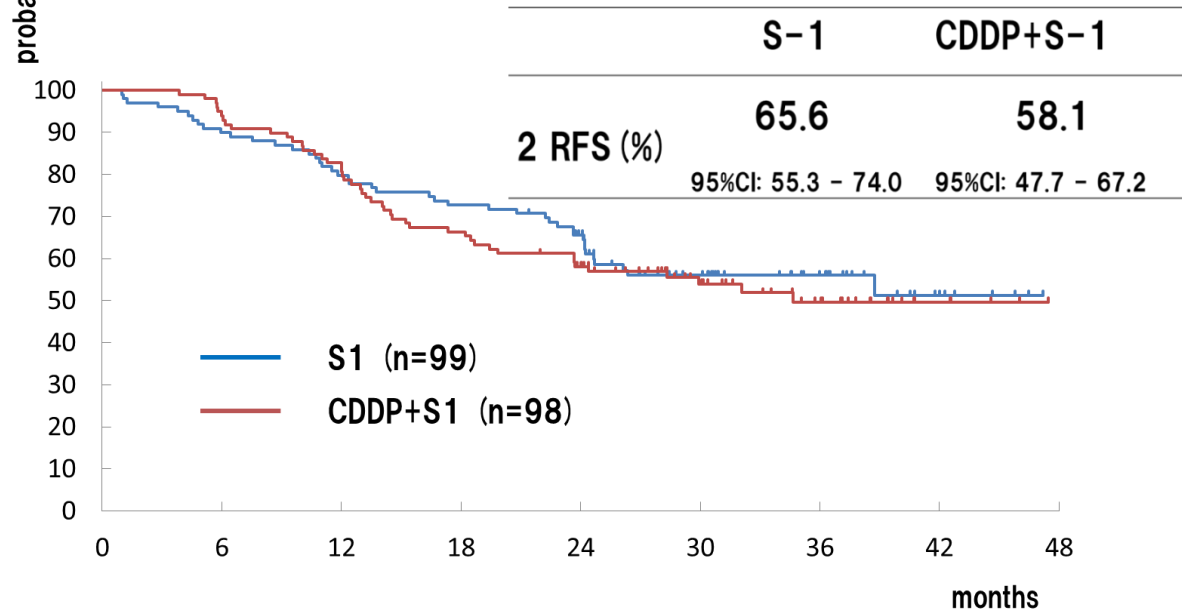
## → LACE : adjuvant cisplatin-based regimens



**Absolute OS benefit at 5 years = 5.3%  $\pm$  1.6%**

**Toxic death = 0.8 to 2 %**

## Relapse free survival



- No implication for clinical practice
- Both arms may be evaluated vs. standard treatment
  - in larger population
  - in western population

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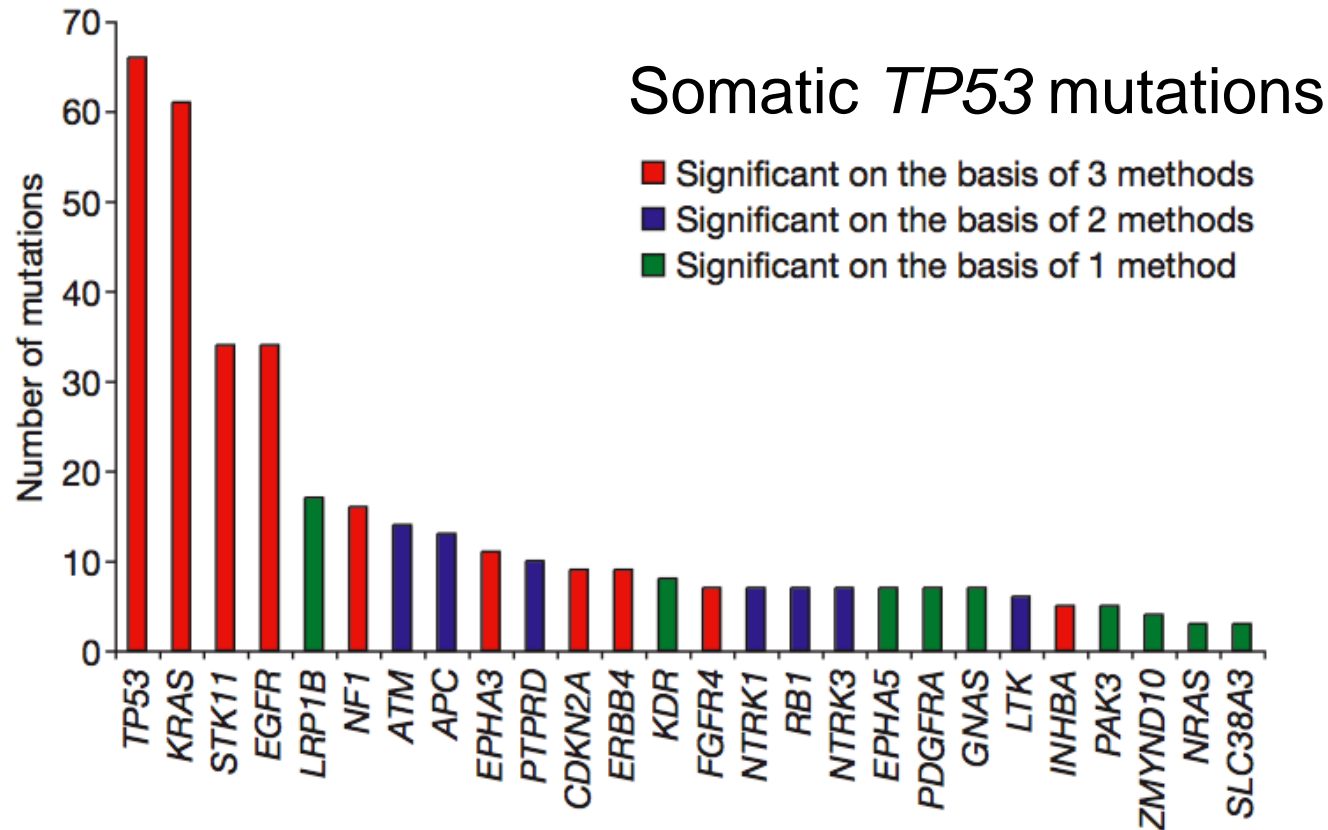
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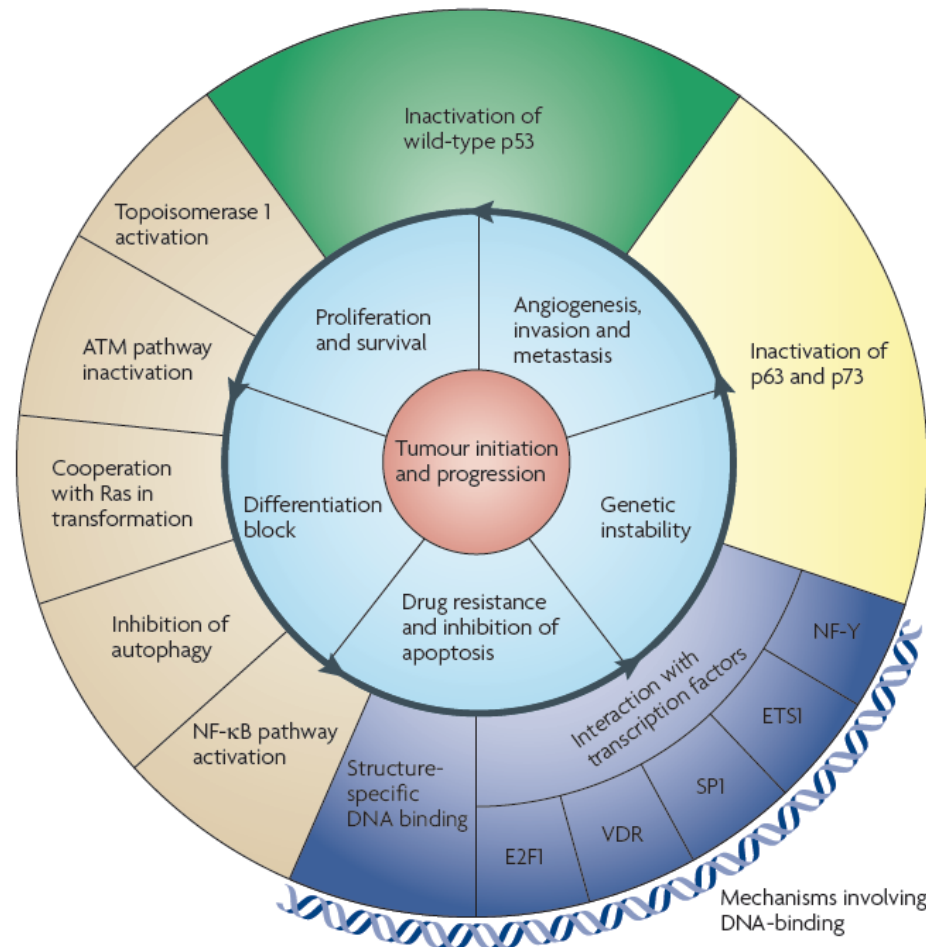
## → *TP53* mutation in NSCLC : the most frequent



Germline *TP53* mutations : Li-Fraumeni Syndrome

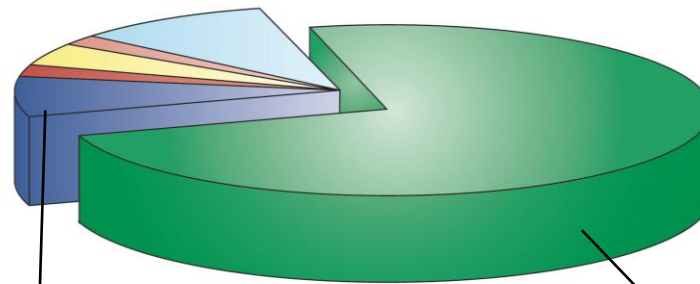
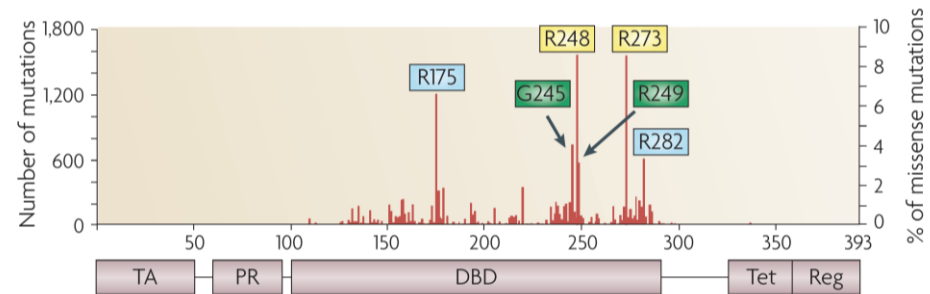
## → p53 role

### Decision making by p53: life, death and cancer



## → *TP53* vs. p53

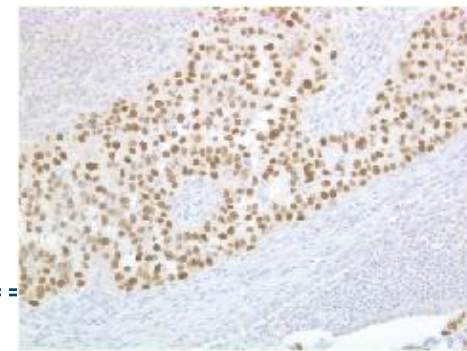
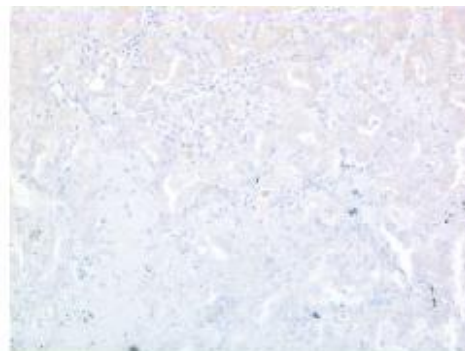
### *TP53* gene (mutations)



Truncated protein

Accumulation of abnormal p53

### P53 protein (IHC)



# *TP53* Mutation Description

Four Mutation Classification systems were tested:

Classification 1: WT/MT

Classification 2: WT/3 classes of MT based on measured residual p53 transactivation activity in a standard functional assay

Classification 3: WT/ 3 classes of MT based on effects of mutation on p53 protein structure

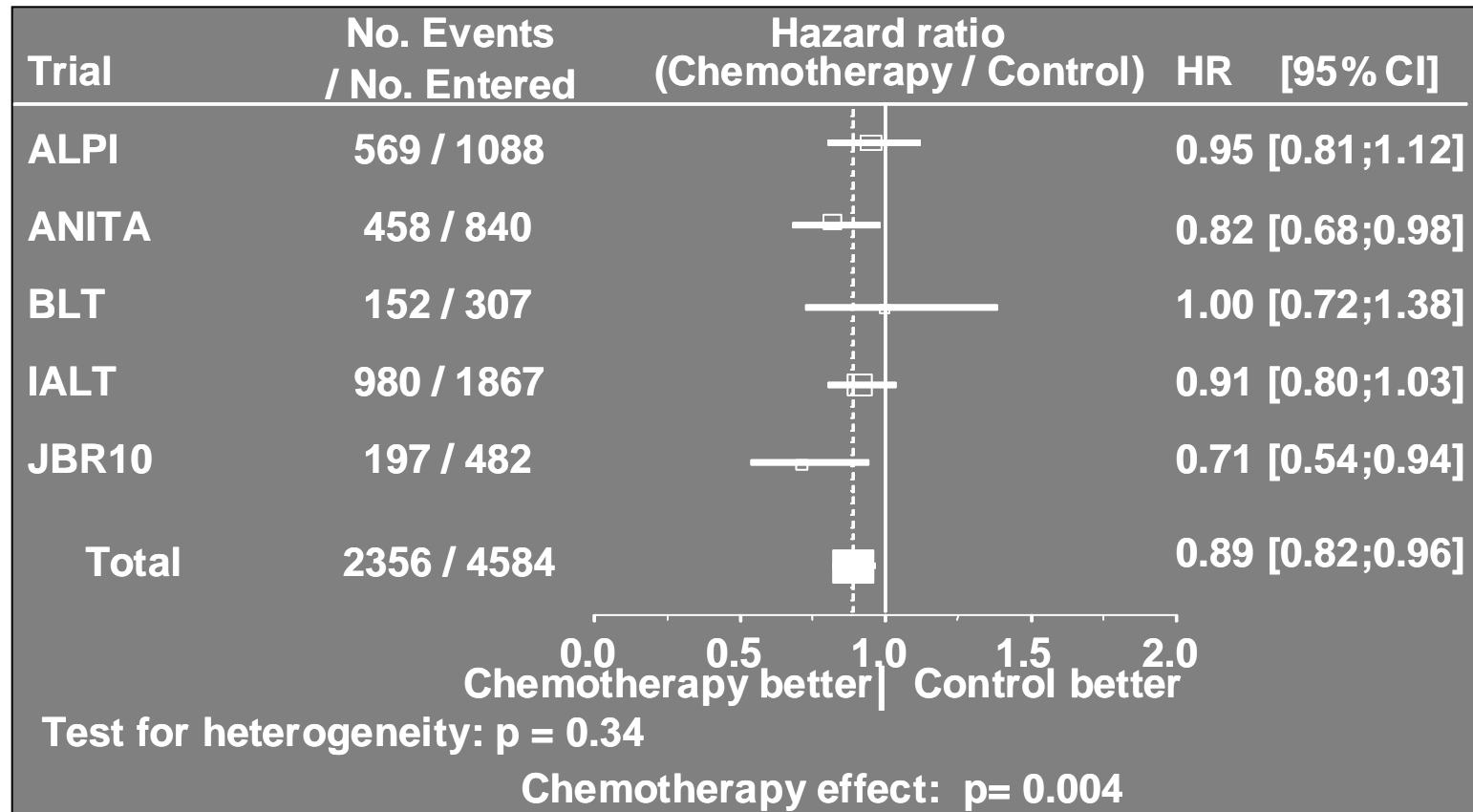
Classification 4: WT/ 3 classes of MT based on interpolating classifications 2 and 3 above.

Data are presented for classifications 1 and 3 only. Classifications 2 and 4 do not add to the message.

Analysis of classification 1 was presented before. However, in present analysis, all datasets were reviewed using the same criteria. Silent mutations were reclassified as WT, as well as mutations not previously found in TP53 database and predicting no functional/structural effects



## → LACE : adjuvant cisplatin-based regimens



**Absolute OS benefit at 5 years = 5.3%  $\pm$  1.6%**

**Toxic death = 0.8 to 2 %**

# Dataset/Classification 1

TRIAL		WT	MT	TOTAL
IALT	Nb	303	221	524
	%	58	42	
JBR10	Nb	286	111	397
	%	72	28	
ANITA	Nb	74	31	105
	%	70	30	
CALGB	Nb	112	71	183
	%	61	339	
TOTAL		775	434	1209

# Analysis

Cox model used in LACE-BIO

**stratified on trial**

**adjusted on :**

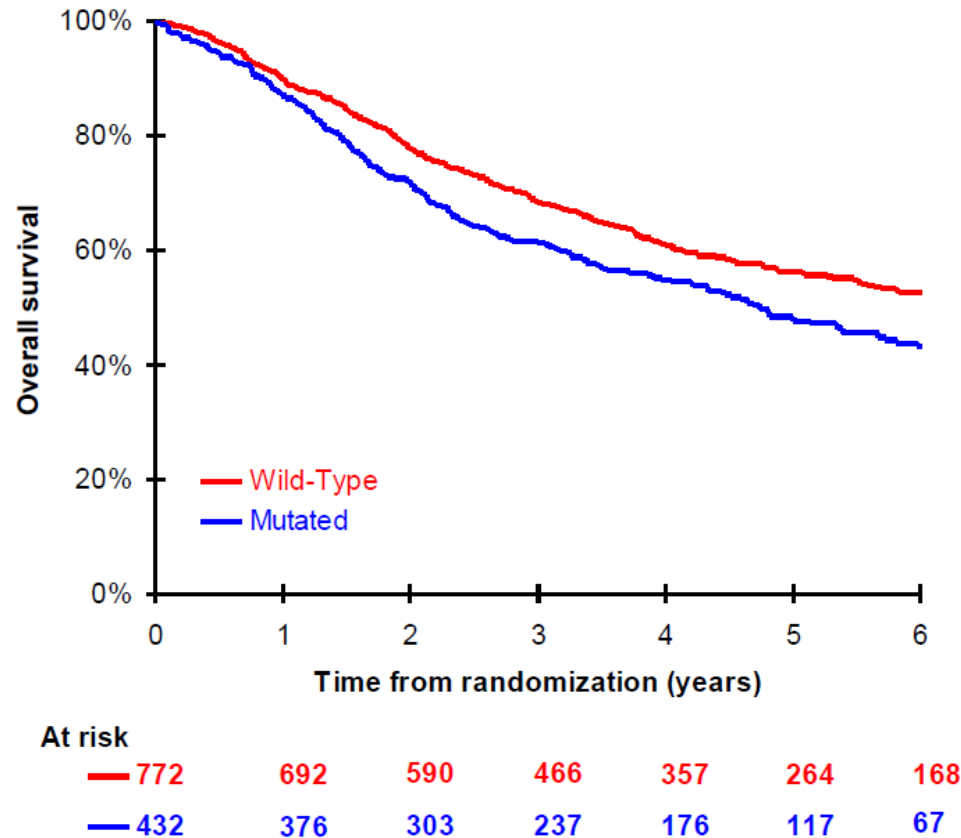
- **Attributed treatment (no chemotherapy, chemotherapy)**
- **Sex (male, female)**
- **Age (<55, 55-64, ≥65)**
- **Histology (squamous cell carcinoma, adenocarcinoma, other)**
- **T (T1, T2, T3-4)**
- **N (N0, N1, N2)**

**A test was considered significant if p-value  $\leq 0.01$**

Test for heterogeneity will not be presented: They all are non-significant

# → Classification 1

## Prognostic analysis : unadjusted Kaplan-Meier survival curves



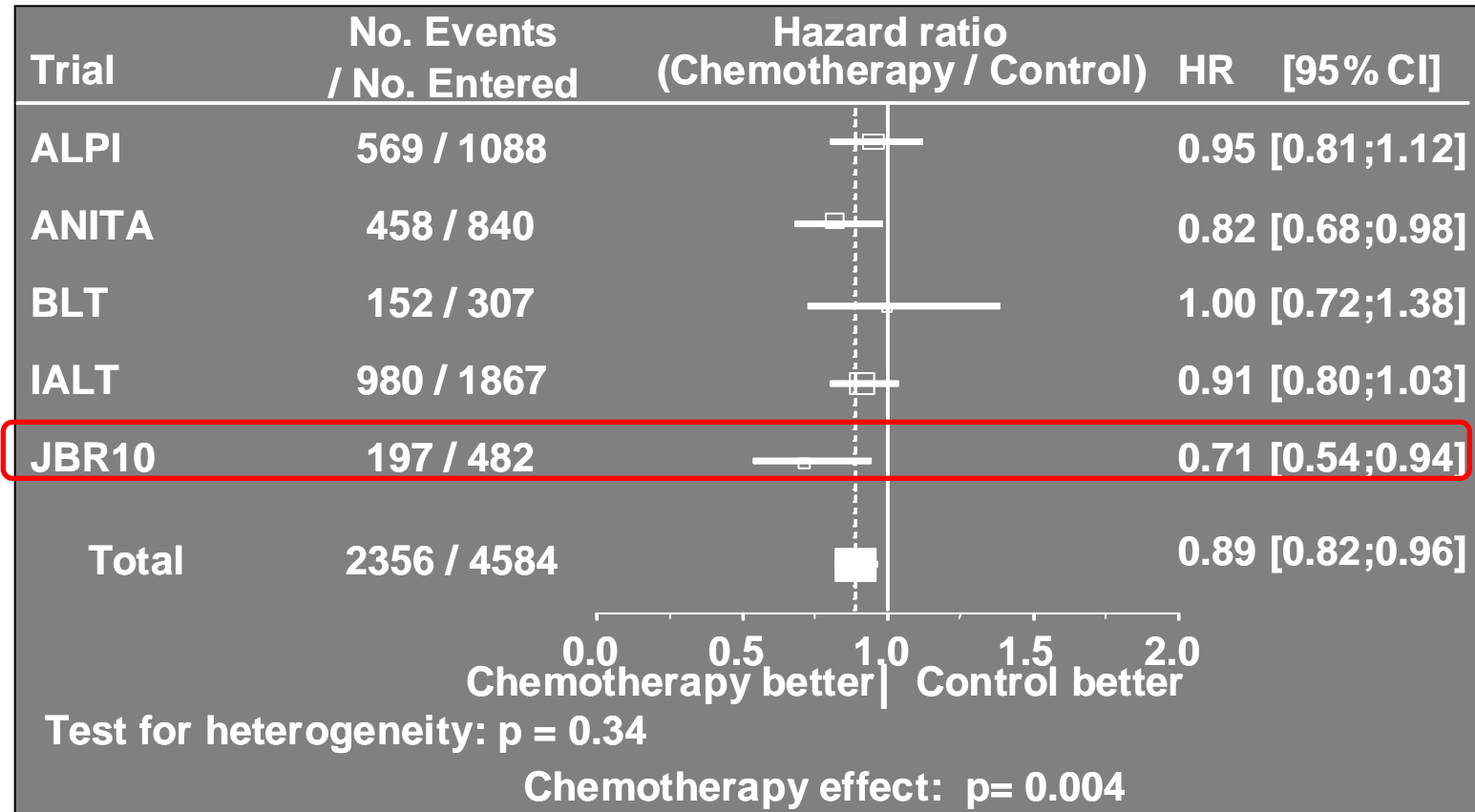
# Classification 1 – Overall Survival

## Predictive analysis

<b>P53 mutation</b>	<b>Chemotherapy group</b> (Nb deaths / Nb patients)	<b>Control group</b> (Nb deaths / Nb patients)	<b>HR for event CT vs. no CT</b> [95% CI]
Wild-Type n=772	153/377	190/395	0.77 [0.62;0.96] p=0.02
Mutant n=432	127/233	100/199	1.05 [0.81;1.37] p=0.71
HR for event Mutant vs. WT [95% CI]	1.39 [1.09;1.77] p=0.008	1.02 [0.79;1.30] p=0.90	Test for interaction p53*treatment p=0.07

<b>TP53 mutation</b> (classification 3)	<b>CT group</b> (Nb events / Nb patients)	<b>No CT group</b> (Nb events / Nb patients)	HR for event CT vs. no CT [95% CI]
Wild-type n=772	179/377	218/395	0.75 [0.62;0.92] <b>p=0.005</b>
Non-missense n=91	34/56	18/35	1.12 [0.63;1.98] p=0.71
HR for event Non-missense vs. WT	1.31 [0.90;1.90] p=0.16	0.88 [0.54;1.44] p=0.62	1.48 [0.81;2.72] p=0.21
Missense-nonDBM n=153	48/78	45/75	1.03 [0.69;1.55] p=0.88
HR for event Missense-nonDBM vs. WT	1.52 [1.10;2.10] <b>p=0.01</b>	1.11 [0.80;1.54] p=0.53	1.37 [0.87;2.16] p=0.18
Missense-DBM n=188	56/99	52/89	0.88 [0.60;1.29] p=0.50
HR for event Missense-DBM vs. WT	1.15 [0.85;1.56] p=0.36	0.99 [0.73;1.34] p=0.94	1.17 [0.76;1.79] p=0.48

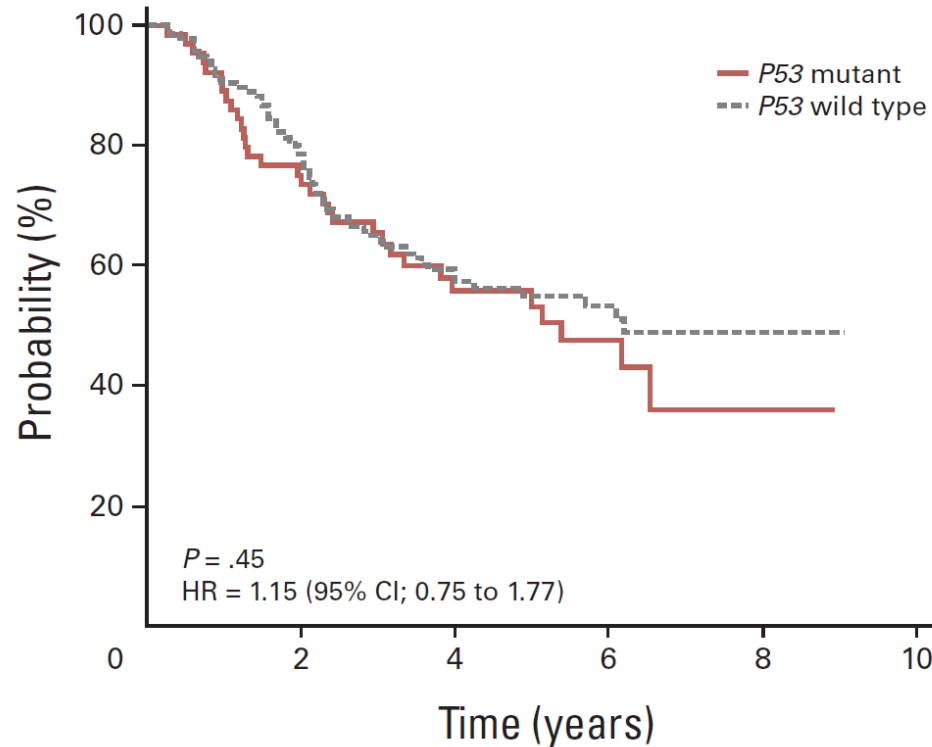
## → LACE : adjuvant cisplatin-based regimens



**Absolute OS benefit at 5 years = 5.3%  $\pm$  1.6%**

**Toxic death = 0.8 to 2 %**

## → P53 in BR10



- Neither prognostic or predictive

at risk						
3 mutant	64	48	26	14	3	0
3 wild type	136	106	55	27	10	0



- Many biomarkers analyzed in LACE-BIO (KRAS, EGFR, ERCC1...)
- Pathway analysis > single biomarker analysis ?
  - Pasi Janne : KRAS & P53 correlation in LACE-BIO - 17:15 Hall D
- Validation in prospective trial
- P53  
Discovered 33 yrs ago, still so much to learn !