www.esmo.org



European Society for Medical Oncology

# ESMO Hamilton Fairley Award lecture From empirical to rational treatment of human cancers cells and their stroma

JY Blay Lyon, France FSG, EORTC

# Major successes in clinical oncology came from an in- depth understanding of the biology of the tumor



# Target the histotype?

# The histotype, the driver mutation, the drug

- Leukemia
- Sarcoma
- Melanoma
- NSCLC
- BCC, Medulloblastoma
- Breast Carcinoma
- Gastric adenocarcinoma
- Renal cell carcinoma

CML,CMML, HES GIST, DFSP, PVNS, IMT, WPLPS KIT or BRAF mutations HER1 or Alk or DDR2 mutations Hh pathway alterations HER2, BRCA1 HER2 amplification VHL loss..



. . .

# **Target the primary mutation?**

# The histotype, the driver mutation, the drug

- KIT
- PDGFR
- Alk
- HER1
- HER2
- Hh
- VHL/HIF1A/VEGF
- mTOR (TSC/PI3K/Akt)
- BRAF

GIST, Melanoma, ALL, Mast. **CMML, HES, DFSP NSCLC, IMT, Neuroblastoma? NSCLC, HN?** Breast Ca, Gastric Ca **BCC**, Medullo, chondroS RCC, NET **RCC, NET, Breast Ca** MMM, other BRAF mut?





# **Target the primary mutation?**

## The disease, the driver mutation, the drug

Imatinib	KIT, PDGFR,CSF1R,+	GIST, MMM, ALL, Mast. CMML, HES, DFSP, others?
Crizotinib	Alk, Met,+	NSCLC, IMT, GC, others?
Trastuzumab	HER2	BC,GC, others?
Erlotinib	HER1	NSCLC, others?
Sunitinib	KIT,PDGFR,VEGFR,RET,+	RCC, NET, others?
Vemurafenib	BRAF	MMM, others?
Everolimus	mTOR	RCC,NET,Br.Ca, others?







# Major successes in clinical oncology came from an in-depth understanding of the biology of the tumor

Vol 463 18 February 2010 doi:10.1038/nature08822

nature

#### ARTICLES

# The landscape of somatic copy-number alteration across human cancers

Rameen Beroukhim<sup>1,3,4,5</sup>\*, Craig H. Mermel<sup>1,3</sup>\*, Dale Porter<sup>8</sup>, Guo Wei<sup>1</sup>, Soumya Raychaudhuri<sup>1,4</sup>, Jerry Donovan<sup>8</sup>, Jordi Barretina<sup>1,3</sup>, Jesse S. Boehm<sup>1</sup>, Jennifer Dobson<sup>1,3</sup>, Mitsuyoshi Urashima<sup>9</sup>, Kevin T. Mc Henry<sup>8</sup>, Reid M. Pinchback<sup>1</sup>, Azra H. Ligon<sup>4</sup>, Yoon-Jae Cho<sup>6</sup>, Leila Haery<sup>1,3</sup>, Heidi Greulich<sup>1,3,4,5</sup>, Michael Reich<sup>1</sup>, Wendy Winckler<sup>1</sup>, Michael S. Lawrence<sup>1</sup>, Barbara A. Wei<sup>1,1</sup>, Kumiko E. Tanaka<sup>1,3</sup>, Derek Y. Chiang<sup>1,3,1,3</sup>, Adam J. Bass<sup>1,3,4</sup>, Alice Loo<sup>8</sup>, Carter Hoffman<sup>1,3</sup>, John Prensner<sup>1,3</sup>, Ted Liefeld<sup>1</sup>, Qing Gao<sup>1</sup>, Derek Yecies<sup>3</sup>, Sabina Signoretti<sup>3,4</sup>, Elizabeth Maher<sup>10</sup>, Frederic J. Kaye<sup>11</sup>, Hidefumi Sasaki<sup>12</sup>, Joel E. Tepper<sup>13</sup>, Jonathan A. Fletcher<sup>4</sup>, Josep Tabernero<sup>14</sup>, José Baselga<sup>14</sup>, Ming-Sound Tsao<sup>15</sup>, Francesca Demichelis<sup>16</sup>, Mark A. Rubin<sup>16</sup>, Pasi A. Janne<sup>3,4</sup>, Mark J. Daly<sup>1,17</sup>, Carmelo Nucera<sup>7</sup>, Ross L. Levine<sup>18</sup>, Benjamin L. Ebert<sup>1,4,5</sup>, Stacey Gabriel<sup>1</sup>, Anił K. Rustgi<sup>19</sup>, Cristina R. Antonescu<sup>18</sup>, Marc Ladanyi<sup>18</sup>, Anthony Letai<sup>3</sup>, Levi A. Garraway<sup>1,3</sup>, Massimo Loda<sup>3,4</sup>, David G. Beer<sup>20</sup>, Lawrence D. True<sup>21</sup>, Aikou Okamoto<sup>22</sup>, Scott L. Pomeroy<sup>6</sup>, Samuel Singer<sup>18</sup>, Todd R. Golub<sup>1,3,23</sup>, Eric S. Lander<sup>1,2,5</sup>, Gad Getz<sup>1</sup>, William R. Sellers<sup>8</sup> & Matthew Meyerson<sup>1,3,5</sup>

A powerful way to discover key genes with causal roles in oncogenesis is to identify genomic regions that undergo frequent alteration in human cancers. Here we present high-resolution analyses of somatic copy-number alterations (SCNAs) from 3,131 cancer specimens, belonging largely to 26 histological types. We identify 158 regions of focal SCNA that are altered at significant frequency across several cancer types, of which 122 cannot be explained by the presence of a known cancer target gene located within these regions. Several gene families are enriched among these regions of focal SCNA, including the *BCL2* family of apoptosis regulators and the NF- $\kappa$ B pathway. We show that cancer cells containing amplifications surrounding the *MCL1* and *BCL2L1* anti-apoptotic genes depend on the expression of these genes for survival. Finally, we demonstrate that a large majority of SCNAs identified in individual cancer types are present in several cancer types.

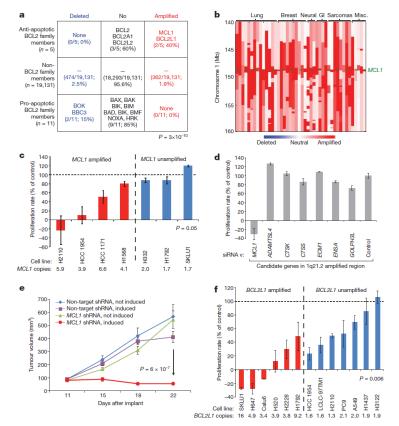


Figure 3 | Dependency of cancer cell lines on the amplified *BCL2* family members, *MCL1* and *BCL2L1*. a, Enrichment of pro- and anti-apoptotic



### **Translational research in oncology research**

From empiric to cosmetic to integrated translational research

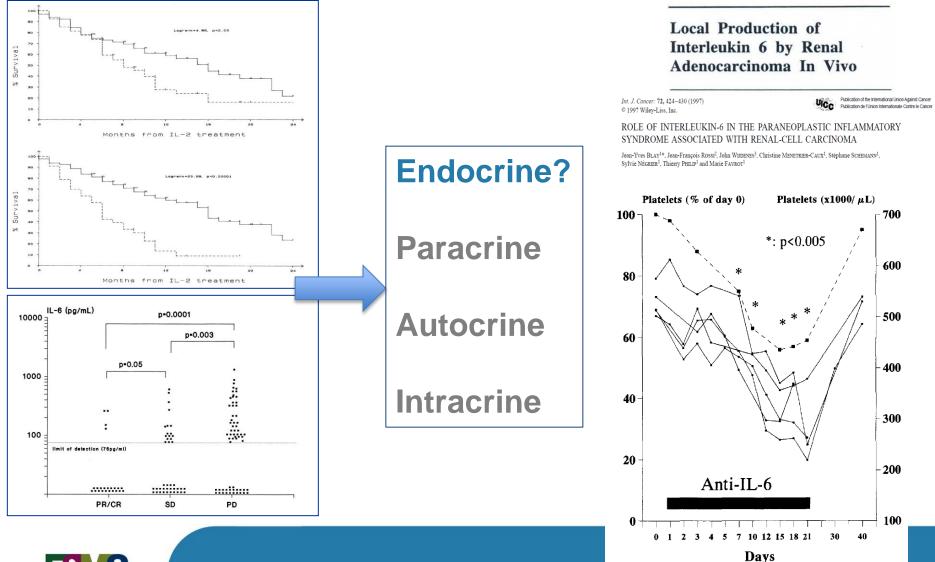
The tumor cell

# The surrounding cells

The patient

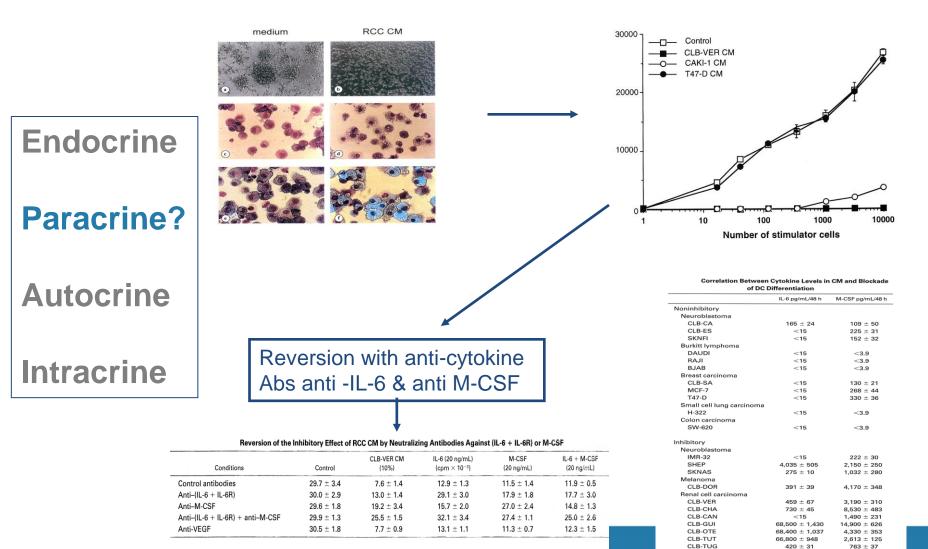


Intracrine, paracrine, endocrine roles of IL-6



ESMU

#### Intracrine, paracrine, endocrine roles of IL-6



CAKI-1

CAKI-2

HT-29

Colon carcinoma

 $2,010 \pm 370$ 

 $3.560 \pm 426$ 

439 + 59

9,780 ± 512

6,500 ± 731

<3.9



#### All IL-6+tumors

#### Intracrine, paracrine, endocrine roles of IL-6

*Int. J. Cancer:* **111,** 653–661 (2004) © 2004 Wiley-Liss, Inc.

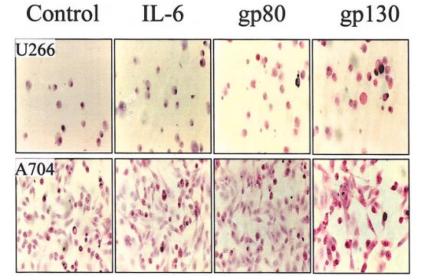


Publication of the International Union Against Cancer

#### IL-6 AS AN INTRACRINE GROWTH FACTOR FOR RENAL CARCINOMA CELL LINES

Laurent Alberti<sup>1</sup>, Marie Cécile THOMACHOT<sup>1</sup>, Thomas BACHELOT<sup>1</sup>, Christine MENETRIER-CAUX<sup>1</sup>, Isabelle PUISIEUX<sup>1</sup> and Jean Yves BLAY<sup>1,2\*</sup>

<sup>1</sup>Equipe Cytokine et Cancer, Unité INSERM 590, Lyon, France <sup>2</sup>Hôpital Édouard Herriot. Lyon. France



#### TABLE VII - ANTIPROLIFERATIVE EFFECT OF IL-6 ANTISENSE OLIGONUCLEOTIDES AND/OR IL13

3H TdR uptake ( $\times 10^3$ cpm) (% of control) Culture conditions									
Cell lines	Medium	IL-6 antisense ON (20 µM)		IL-13 (100 n	$g \cdot mL^{-1}$ )	IL-6 antisense ON (20 $\mu$ M) 13 (100 ng $\cdot$ mL <sup>-1</sup> )			
A704	$59.7 \pm 7.1$	$23.6 \pm 1.7$	(40%)	$39.0 \pm 7.5$	(65%)	$15.7 \pm 1.0$	(26%)		
ACHN	$62.1 \pm 2.4$	$27.6 \pm 2.4$	(44%)	$32.1 \pm 3.5$	(52%)	$16.2 \pm 3.9$	(26%)		
CAKI1	$23.0 \pm 0.5$	$8.7 \pm 0.0$	(37 %)	$13.9 \pm 1.8$	(61%)	$6.3 \pm 0.1$	(27 %)		
CAKI2	$13.9 \pm 3.3$	$6.4 \pm 1.2$	(46 %)	$16.3 \pm 1.3$	(117%)	$7.3 \pm 1.1$	(53 %)		

#### Endocrine

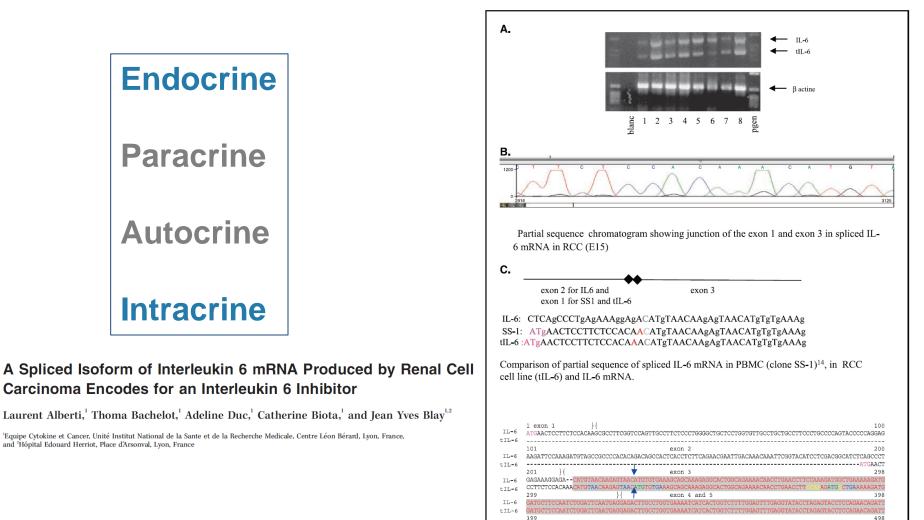
Paracrine

#### Autocrine

#### Intracrine



Intracrine, paracrine, endocrine roles of IL-6



IL-6 tIL-6

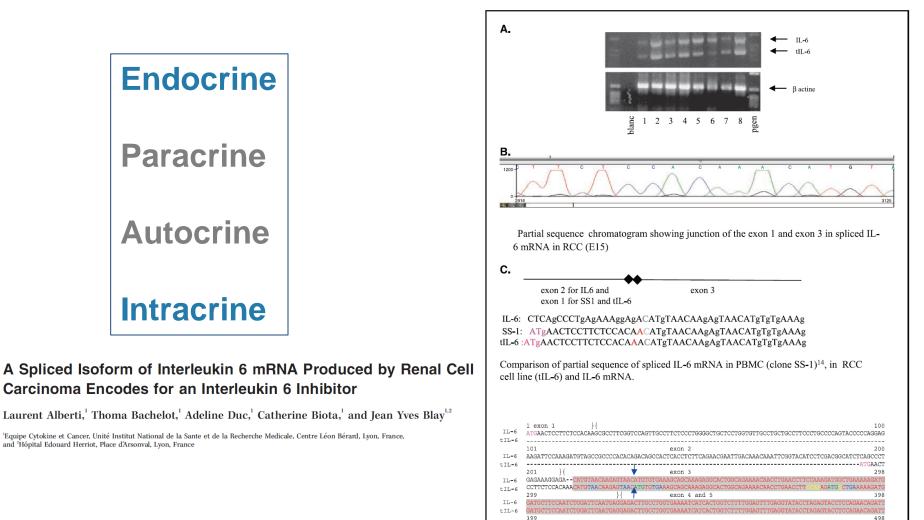
IL-6 tIL-6 tIL-6



Alignment sequence of spliced IL-6 mRNA in RCC cell line (tIL-6) and IL-6 mRNA.

598

Intracrine, paracrine, endocrine roles of IL-6



IL-6 tIL-6

IL-6 tIL-6 tIL-6



Alignment sequence of spliced IL-6 mRNA in RCC cell line (tIL-6) and IL-6 mRNA.

598

#### Intracrine, paracrine, endocrine roles of IL-6 and VEGF

VOLUME 22 · NUMBER 12 · JUNE 15 2004

JOURNAL OF CLINICAL ONCOLOGY

ORIGINAL REPORT

Interleukin-6, Interleukin-10, and Vascular Endothelial Growth Factor in Metastatic Renal Cell Carcinoma: Prognostic Value of Interleukin-6—From the Groupe Français d'Immunothérapie

Sylvie Negrier, David Perol, Christine Menetrier-Caux, Bernard Escudier, Michel Pallardy, Alain Ravaud, Jean-Yves Douillard, Christine Chevreau, Christine Lasset, and Jean-Yves Blay

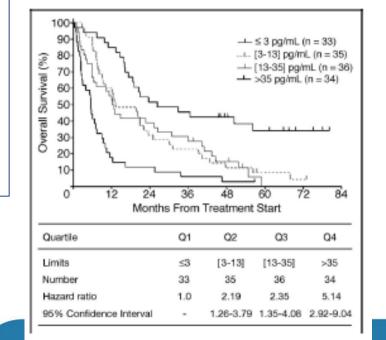


Fig 1. Results of the quartile analysis of interleukin-6 serum levels (pg/mL), with Kaplan-Meier overall survival analysis and univariate Cox proportional hazard regression model (n = 138).

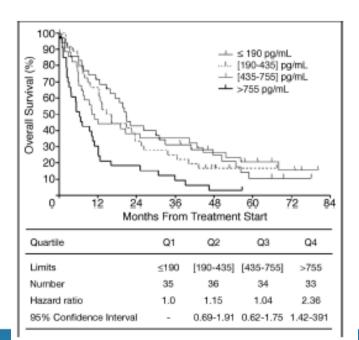


Fig 2 Results of the quartile analysis of vascular endothelial growth factor serum levels (pg/mL), with Kaplan-Meier overall survival analysis and univariate Cox proportional hazard regression model (n = 138).

#### Endocrine

#### Paracrine

#### Autocrine

#### Intracrine



### **Translational research in oncology research**

From empiric to cosmetic to integrated translational research

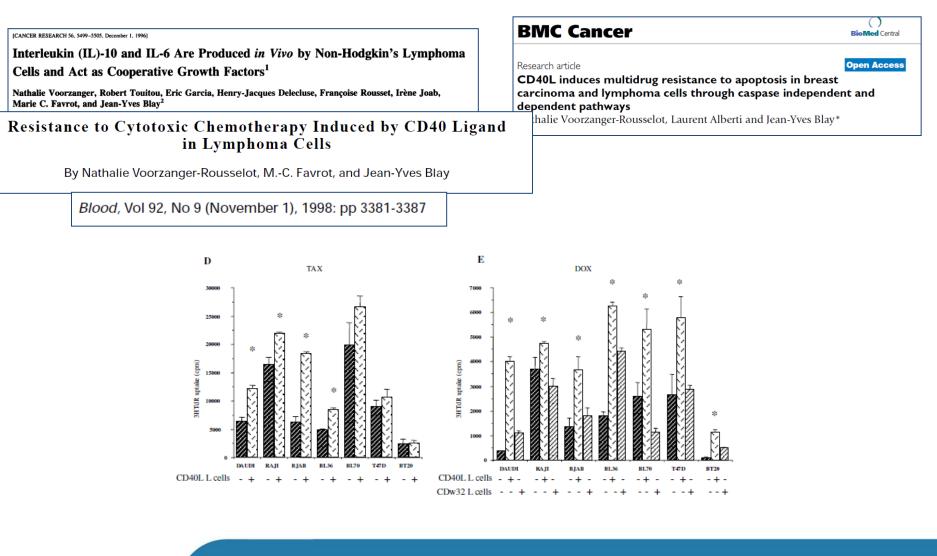
The tumor cell

# The surrounding cells

The patient

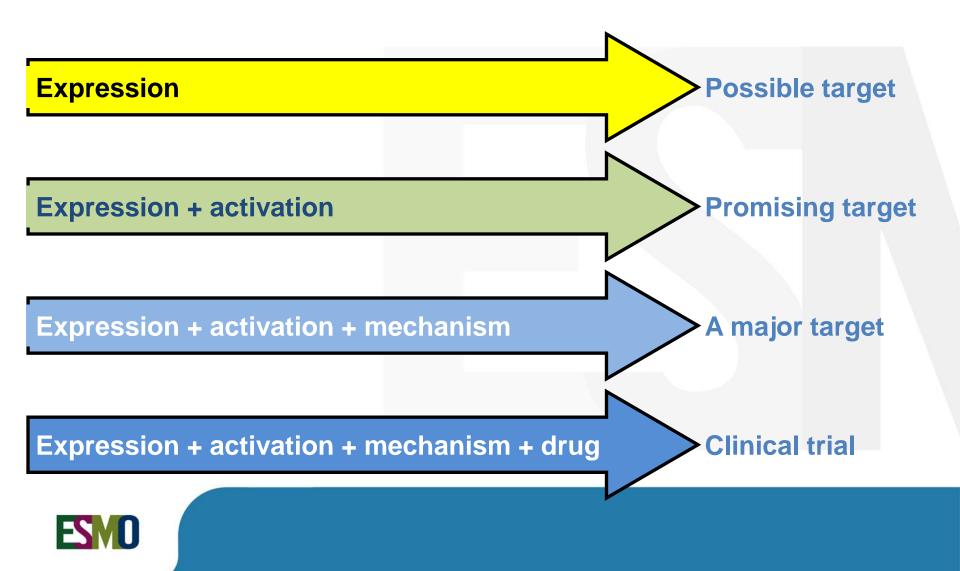


## Cytokine as growth factors in NHL and breast Ca



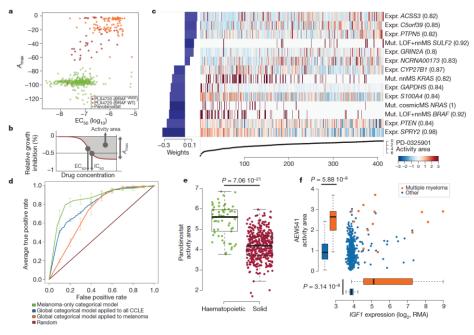


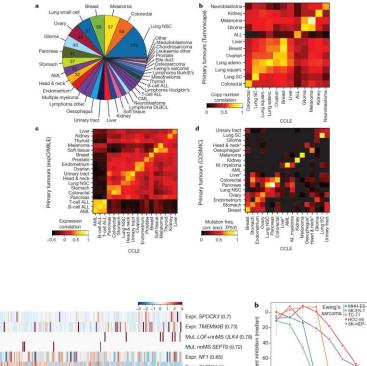
# What is a good target?

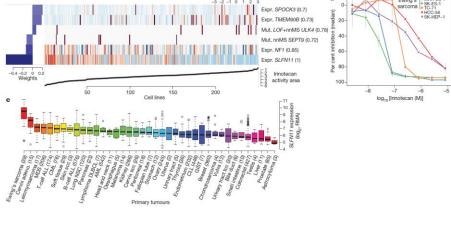


# The Cancer Cell Line Encyclopedia enables predictive modelling of anticancer drug sensitivity

Jordi Barretina<sup>1,2,3</sup>†\*, Giordano Caponigro<sup>4</sup>\*, Nicolas Stransky<sup>1</sup>\*, Kavitha Venkatesan<sup>4</sup>\*, Adam A. Margolin<sup>1</sup>†\*, Sungjoon Kim<sup>5</sup>, Christopher J. Wilson<sup>4</sup>, Joseph Lehár<sup>4</sup>, Gregory V. Kryukov<sup>1</sup>, Dmitriy Sonkin<sup>4</sup>, Anupama Reddy<sup>4</sup>, Manway Liu<sup>4</sup>, Lauren Murray<sup>1</sup>, Michael F. Berger<sup>1</sup>†, John E. Monahan<sup>4</sup>, Paula Morais<sup>1</sup>, Jodi Meltzer<sup>4</sup>, Adam Korejwa<sup>1</sup>, Judit Jané–Valbuena<sup>1,2</sup>, Felipa A. Mapa<sup>4</sup>, Joseph Thibault<sup>5</sup>, Eva Bric-Furlong<sup>4</sup>, Pichai Raman<sup>4</sup>, Aaron Shipway<sup>5</sup>, Ingo H. Engels<sup>5</sup>, Jill Cheng<sup>6</sup>, Guoying K. Yu<sup>6</sup>, Jianjun Yu<sup>6</sup>, Peter Aspesi Jr<sup>4</sup>, Melanie de Silva<sup>4</sup>, Kalpana Jagtap<sup>4</sup>, Michael D. Jones<sup>4</sup>, Li Wang<sup>4</sup>, Charles Hatton<sup>3</sup>, Emanuele Palescandolo<sup>3</sup>, Supriya Gupta<sup>1</sup>, Scott Mahan<sup>1</sup>, Carrie Sougnez<sup>1</sup>, Robert C. Onofrio<sup>1</sup>, Ted Liefeld<sup>1</sup>, Laura MacConaill<sup>3</sup>, Wendy Winckler<sup>1</sup>, Michael Reich<sup>1</sup>, Nanxin Li<sup>5</sup>, Jill P. Mesirov<sup>1</sup>, Stacey B. Gabriel<sup>1</sup>, Gad Getz<sup>1</sup>, Kristin Ardlie<sup>1</sup>, Vivien Chan<sup>6</sup>, Vic E. Myer<sup>4</sup>, Barbara L. Weber<sup>4</sup>, Jeff Porter<sup>4</sup>, Markus Warmuth<sup>4</sup>, Peter Finan<sup>4</sup>, Jennifer L. Harris<sup>5</sup>, Matthew Meyerson<sup>1,2,3</sup>, Todd R. Golub<sup>1,3,7,8</sup>, Michael P. Morrissey<sup>4</sup>\*, William R. Sellers<sup>4</sup>\*, Robert Schlegel<sup>4</sup>\* & Levi A. Garraway<sup>1,2,3\*</sup>



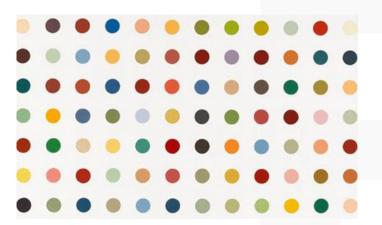




# Lineage Gene expression

# Biomarkers shared across lineages

# **Towards a major fragmentation** of nosological entities





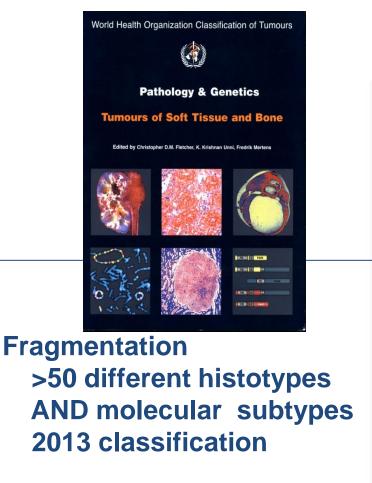


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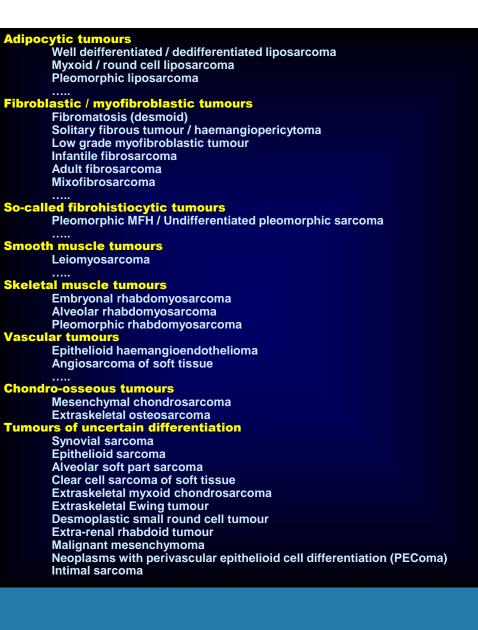
- « Acivicin »
- « Arginosuccinic acid »

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# **Even for rare tumors...**

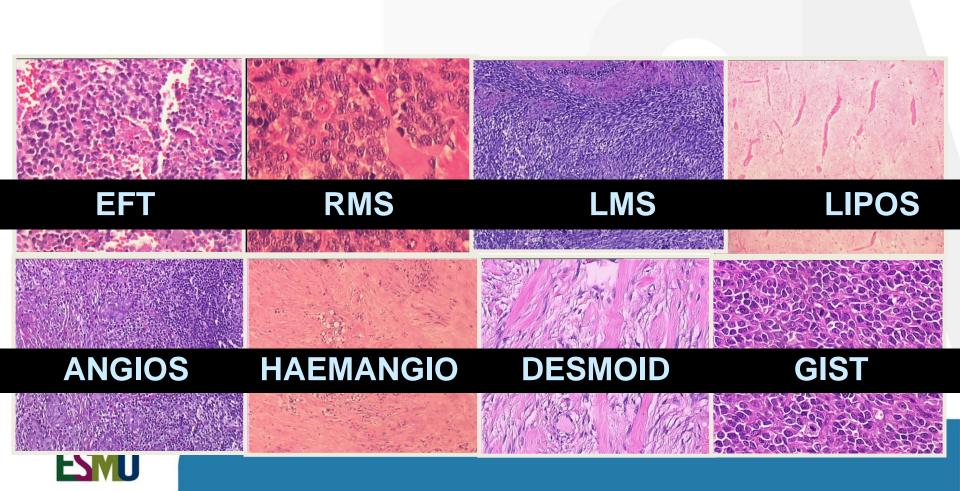


ESM





## Soft Tissue Sarcomas



# **Connective tissue tumours**

<ul> <li>Sarcoma with translocations</li> <li>Ewing, DFSP, Synovial sarcomas,</li> </ul>	~15%
<ul> <li>Sarcoma with kinase mutations</li> </ul>	~15%
GIST, few Angiosarcomas	
<ul> <li>Sarcoma with tumor suppressor gene inactivation</li> <li>MPNST NF1, Rhabdoid tumors- INI1, PEComas TSC</li> </ul>	~10%
<ul> <li>Sarcomas with chromosome 12q14-15 amplification</li> <li>WD/DDLPS, intimal sarcomas, LG OS</li> </ul>	~15%
<ul> <li>Sarcomas with complex genetic alterations</li> <li>Pleomorphic sarcomas, LMS,</li> </ul>	~50%
Low grade or locally agressive	
• Desmaid tumora	Constation

- Desmoid tumors
- Giant cell tumor of the bone ? (RANK involved)

beta catenin or APC mutation 2 (RANK involved)

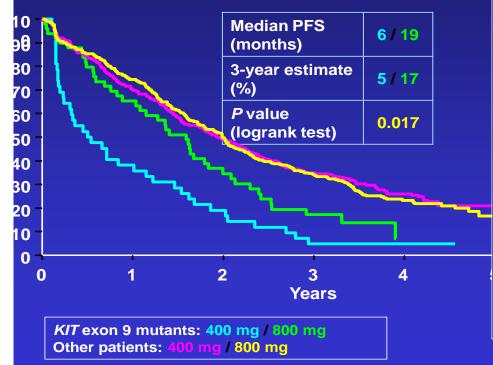
• Giant cell tumor of the soft part (PVNS) translocation



# **Fragmentation even in rare diseases**

# **GIST are at least 10 diseases ESMO 2012**

#### *KIT* exon 9 mutants (10% of patients)



	Dose A	djuvant
KIT Exon 11	lm 400	+
KIT exon 9	lm 800	+
PDGFRA		
Non D842V	lm 400	+
D842V:	0	0
KIT/PDGFR WT	lm 400	+/?
NF1	?/Im 400	+/?
SDHB	?/lm 400	+/?
Raf	?	?
Pediatric	?	?

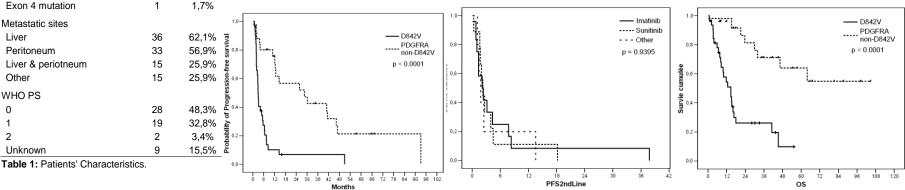


# **PDGFRA GIST in advanced phase**

Characteristic	Ν	%	_
Total	58	100	
Gender			
Male	34	58,6%	
Female	24	41,4%	
Primary tumor location			
Stomach	40	69,0%	
Small bowel	7	12,1%	
Peritoneum/Mesentery	2	3,4%	
Rectum/Anus	1	1,7%	
Other	4	6,9%	
Unknown	4	6,9%	
KIT/CD117 expression			
Positive	38	65,5%	
Negative	7	12,1%	
Unknown	13	22,4%	
Type of mutation			
Exon 18 D842V substitution	32	55,2%	
Other exon 18 mutation	17	29,3%	
Exon 12 mutation	8	13,8%	
Exon 4 mutation	1	1,7%	
Metastatic sites			1
Liver	36	62,1%	val
Peritoneum	33	56,9%	o survi
Liver & periotneum	15	25,9%	ree s
Other	15	25,9%	oion-1
WHO PS			Probability of Progression-free survival
0	28	48,3%	6 Pro
1	19	32,8%	lity o
2	2	3,4%	babi
Unknown	9	15,5%	Pro
Table 1. Patiente' Characteristic	·c		0

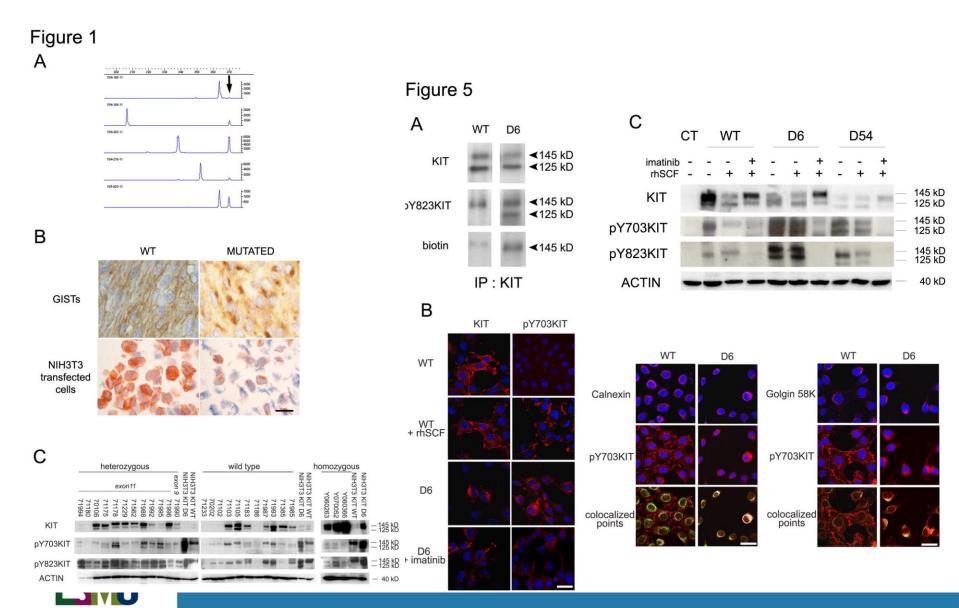
Response	D842V*		Non-D842V D842V* Exon 18		E	con 12	Exc	on 4	Overall*		
	Ν	%	Ν	%	N	%	N	%	N	%	
CR	0	0%	1	6%	1	13%	0	-	2	4%	
PR	0	0%	4	24%	3	38%	1	-	8	14%	
SD	10	32%	10	59%	3	38%	0	-	23	40%	
PD	21	68%	2	12%	1	13%	0	-	24	42%	

**Table 2:** response rate to imatinib per group of PDGFRA mutation and overall. (\*): one patient with a D842V-mutant GIST died of gastrointestinal hemorrage before his first assessment and was therefore not evaluable for response.



P. Cassier, E Fumagalli, P Rutkowski, P Schoffski, M Van Glabbeke, M Debiec Rychter, JF Emile, F Duffaud, J Martin, B Landi, A Adenis, F Bertucci, E Bompas, S Leyvraz, I Judson, J Verweij, P Hohenberger, P Casali, JY Blay (unpublished data)

# Intracellular localization of mutated & activated KIT receptors



#### Tabone-Eglinger et al 2008

# Timely proof of require international collaborations



# **MCSFR** inhibitors in PVNS with t(1,2)

Figure: Response to imatinib in PVNS

- Case report in 2008
  - (Ann Oncol 2008)

- Retrospective study 2011
  - (Cancer 2011)

18/09/06 08/11/06 28/02/07 20 10 -10 -20 -30 -40 -50 -60 -70 -80 % tumor growth -90 -100 Figure 1. The best tumor shrinkage is illustrated according to Response Evaluation Criteria in Solid Tumors (RECIST) 1st interim analysis 2nd interim analysis 3 3rd interim analysis 0 0.1 0.2 0.3 0.4 0.5 0.6 0.7 0.8 0.9

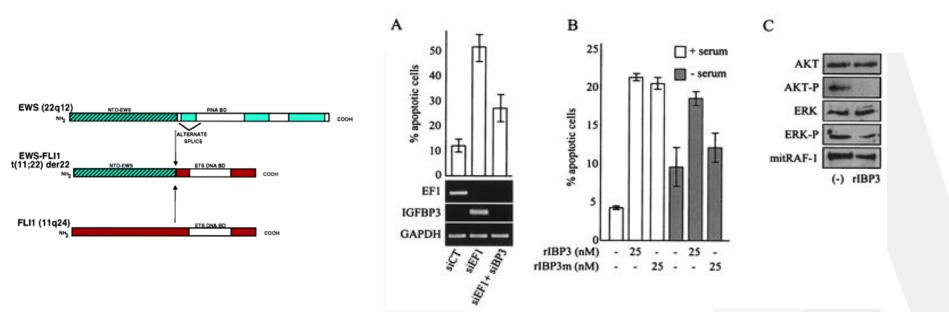
- Prospective study 2012
  - (Proc ASCO 2012)



# Genomic characterisation and cellular models are required



# Ewing cells depend on the IGF1 pathway

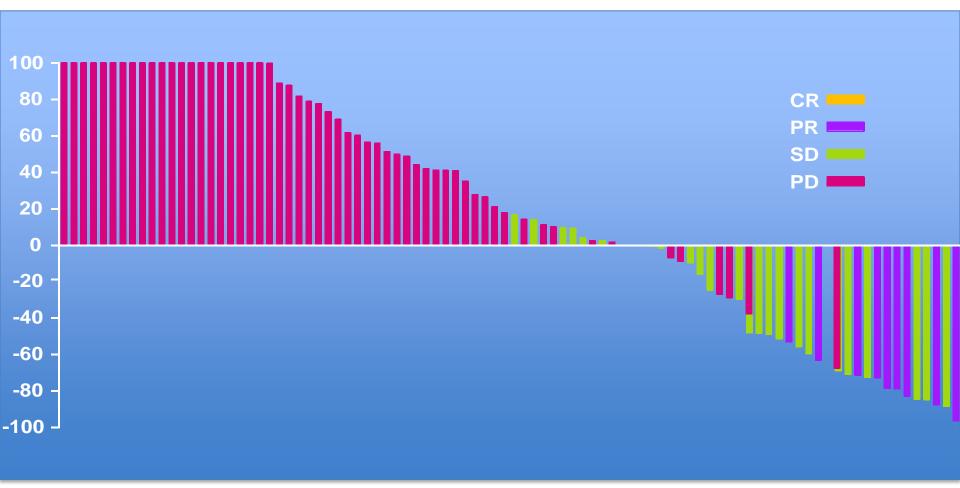


IGF1 inhibitors as potential targeted therapy in ES?



Delattre 2003, Prieur 2004

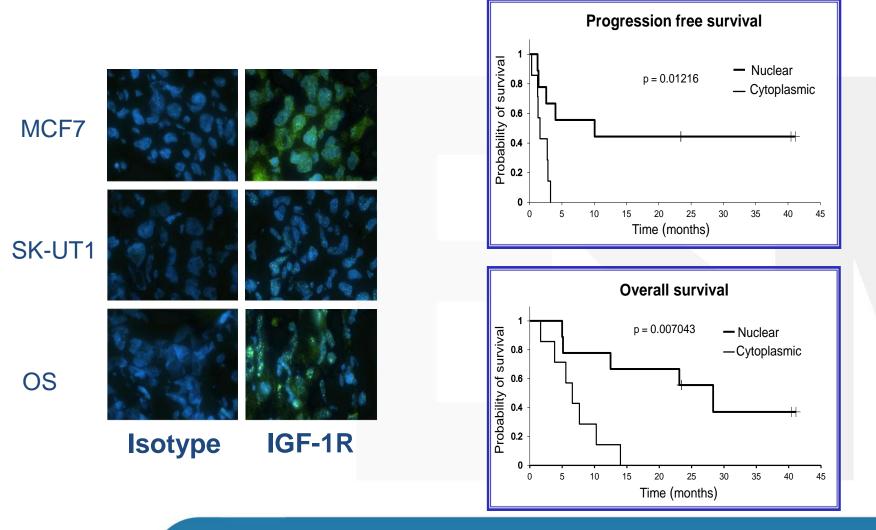
### Ewing sarcoma and IGF1R Ab





Patel 2012

# Nuclear staining for IGF1R: a biomarker for response in sarcoma?



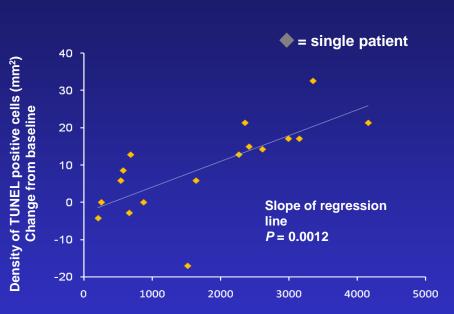


Asmane I, Blay JY, AACR 201

ESMO 2012

#### Nutlin 3a (RG7112) in Liposarcoma with MDM2 amplification

- MDM2 inhibition in human tumors activates p53, arrests cell proliferation, and induces apoptosis
- This proof of mechanism study in patients with LPS demonstrates:
  - Pharmacological p53 activation by an inhibitor of the p53-mdm2 interaction
  - Post-treatment Increases in p53, p21, and mdm2 levels
  - Exposure-related increases in MIC-1 levels
  - Post-treatment decreases proliferation as measured by change in Ki-67
  - Exposure-related induction of apoptotic signals
  - While not designed as an efficacy study, early signs of clinical activity included:
    - 1 PR after a single cycle
    - 13 SD
- This study also supports the feasibility of multiple biopsies in patients with liposarcomas eligible for surgery



MIC-1 C1D8 Mean change from baseline



### **Translational research in oncology research**

From empiric to cosmetic to integrated translational research

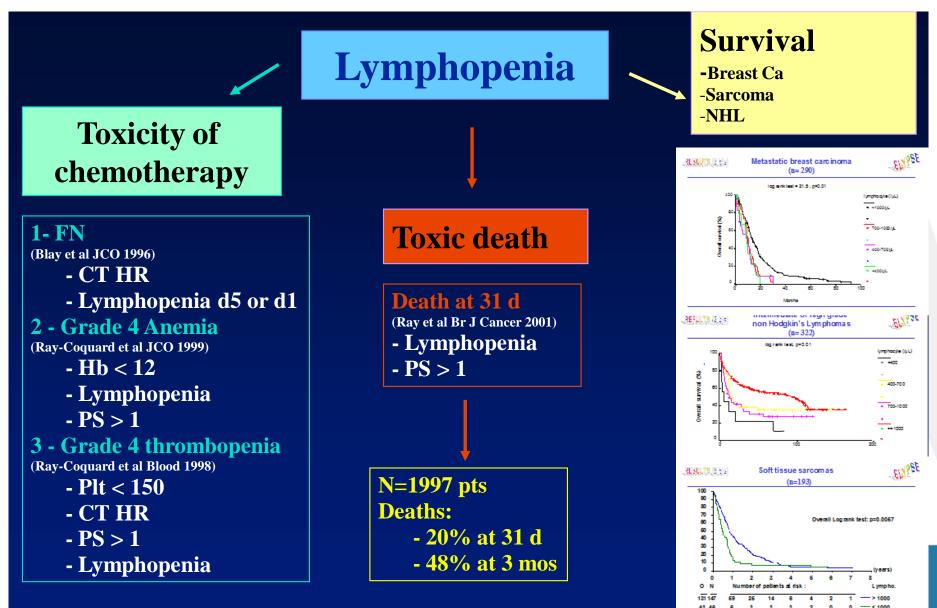
The tumor cell

# The surrounding cells

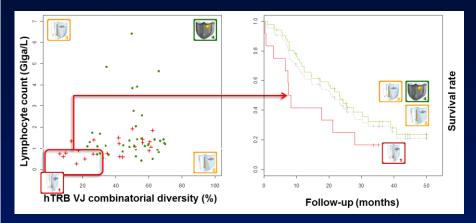
The patient



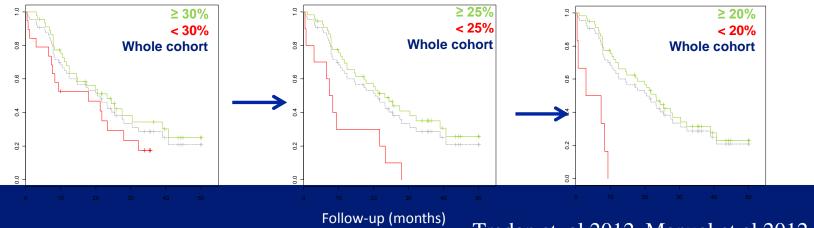
# Lymphopenia and cancer



#### Lympho-Divpenia predicts overall survival



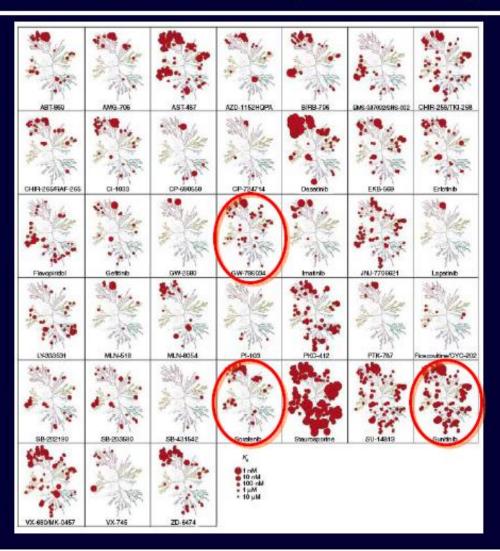
OS according to baseline diversity



Tredan et al 2012, Manuel et al 2012



### Kinase inhibitory selectivity



Karaman MW, et al. Nature Biotech 2008

## Kinase inhibitory selectivity



Karaman MW, et al. Nature Biotech 2008





# From empirical to rational treatment of human cancers cells and their stroma

•Genomic characterisation : opportunities and challenges

•Functional assays / in vivo models

•Fragmentation of nosological entities: lineage matters!

•The fragmented small groups of tumors are challenging for clinical research

•Novel dimensions of complexity:

- International legal requirements
- Health economics



# Thank you

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Maud Brunat	Pierre Meeus	JM Coindre	D. Lacombe
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	Simon Baconnier		Many others
	 Alain Puisieux, Sylvie	EuroSarc partners	
	Negrier, Patrick		
	Mehlen, M. Rousseau	Netsarc partners	
	Many others in the CLB & CRCL & UCC,	Many others	
	,		

The World Sarcoma Networks: G. Demetri, P. Casali, A Gronchi, AP Dei Tos, P. Hohenberger, I Judson, V. van der Graaf, R. Maki, M. von Mehren, S.Patel, R. Benjamin, T. Nishida, D. Thomas, J. Martin, J Garcia... and many others

# Thank you

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# Understanding the biology is as important as molecular characterisation

The examples of cytokines and growth factors



## **Challenges of targeted therapeutics**

Ineluctable emergence of resistance?

Endless fragmentation of nosological entities.



# Cells of the stroma are guilty by association and need to be treated accordingly

A contrasted role of the immune system

Promoting tolerance

Quantitativeluy and qualitatively altered.



# Rare tumors of 2012 are models for the future rare tumors.

The fragmented small groups of tumors are challenging for clinical research



