

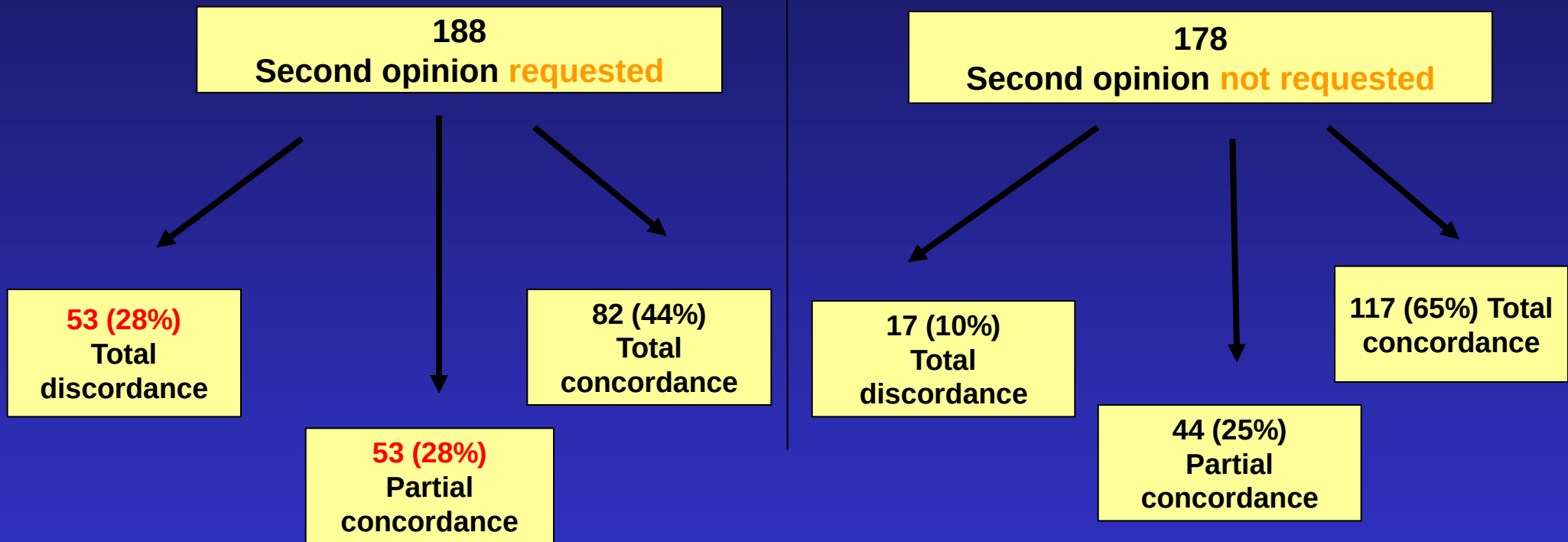
Improving clinical research on rare cancers

JY Blay
Lyon, France
FSG, EORTC

Issues

- Recognition
- Fragmentation
- Solutions

Rate of concordance by patient sub-group



→ For **56 %** the diagnosis is not totally correct

→ For **35 %** the diagnosis is not totally correct

Clinical practice guidelines

Soft tissue sarcomas: ESMO Clinical Recommendations for diagnosis, treatment and follow-up

P. G. Casali¹, L. Jost², S. Sleijfer³, J. Verweij⁴ & J.-Y. Blay⁵

On behalf of the ESMO Guidelines Working Group*



Bone sarcomas: ESMO Clinical Recommendations for diagnosis, treatment and follow-up

P. G. W. Hogendoorn[†]

On behalf of the ESMO/EUROBONET Working Group*

University Medical Center, Leiden, Netherlands

clinical recommendations

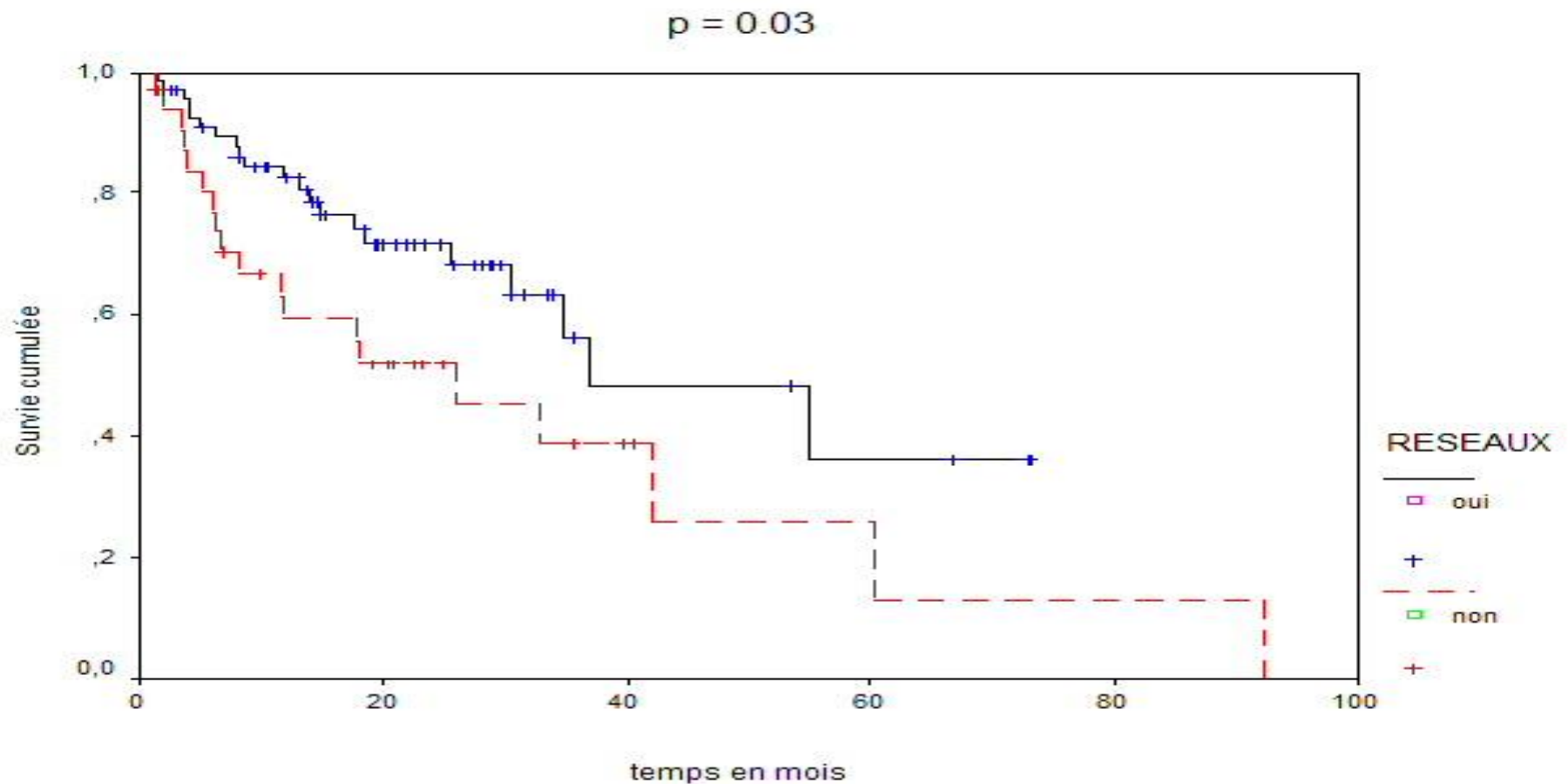
*Annals of Oncology 21 (Supplement 5):
doi:10.1093/annonc*

Gastrointestinal stromal tumours: ESMO Clinical Recommendations for diagnosis, treatment and follow-up

P. G. Casali¹ & J.-Y. Blay²

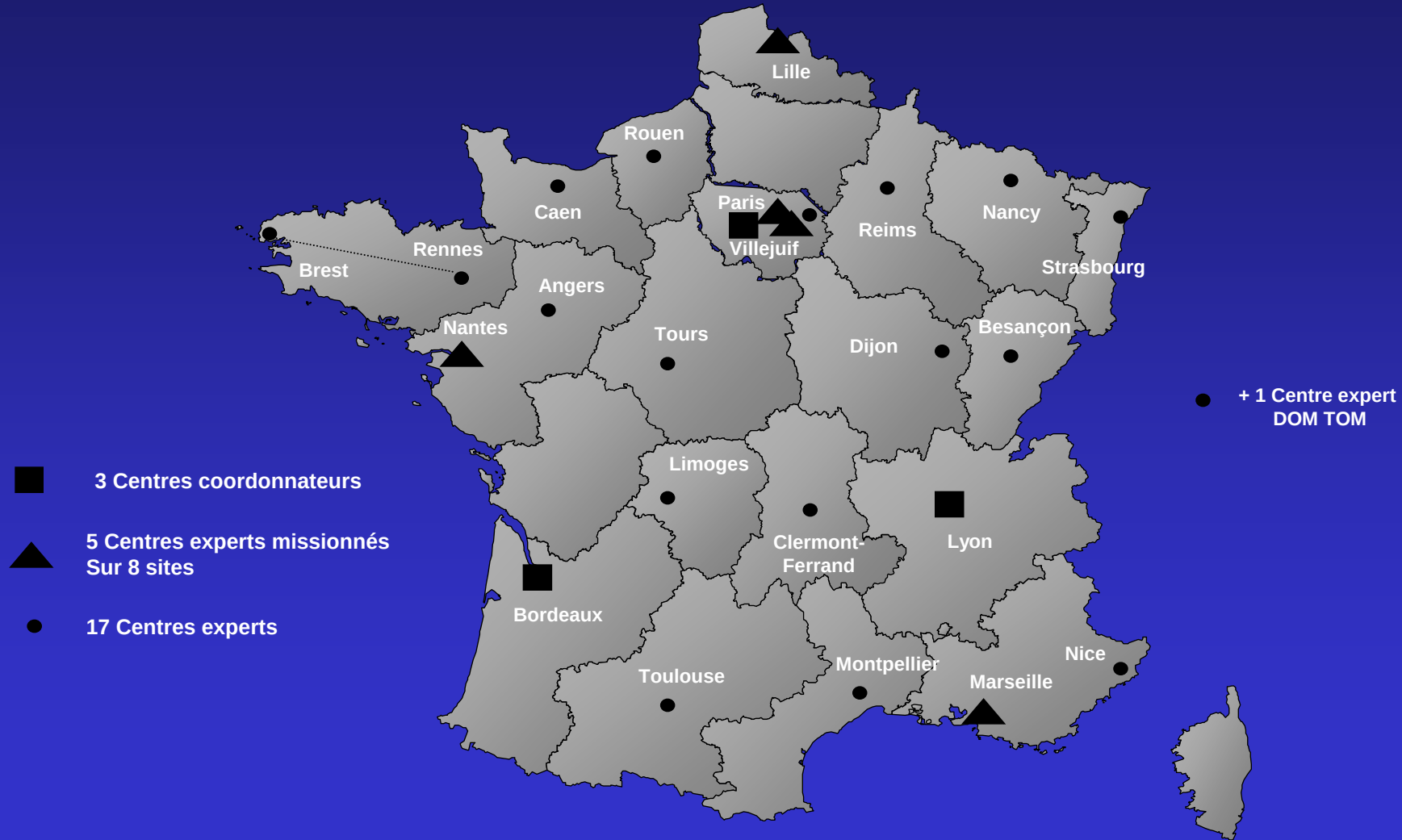
On behalf of the ESMO/CONTICANET/EUROBONET Consensus Panel of Experts*

Distant metastasis and multidisciplinary assessment



French Clinical network (NetSarc)

Brussels 25 5 2012



Netsarc 2010-2012

- N=8954 patients with sarcomas
- Primary surgery in
 - Reference center
 - R0 : 60%, R1: 36%, R2: 4%
 - Primary care or non reference center
 - R0 : 30%, R1: 49%, R2: 21%

($p < 0.000$)

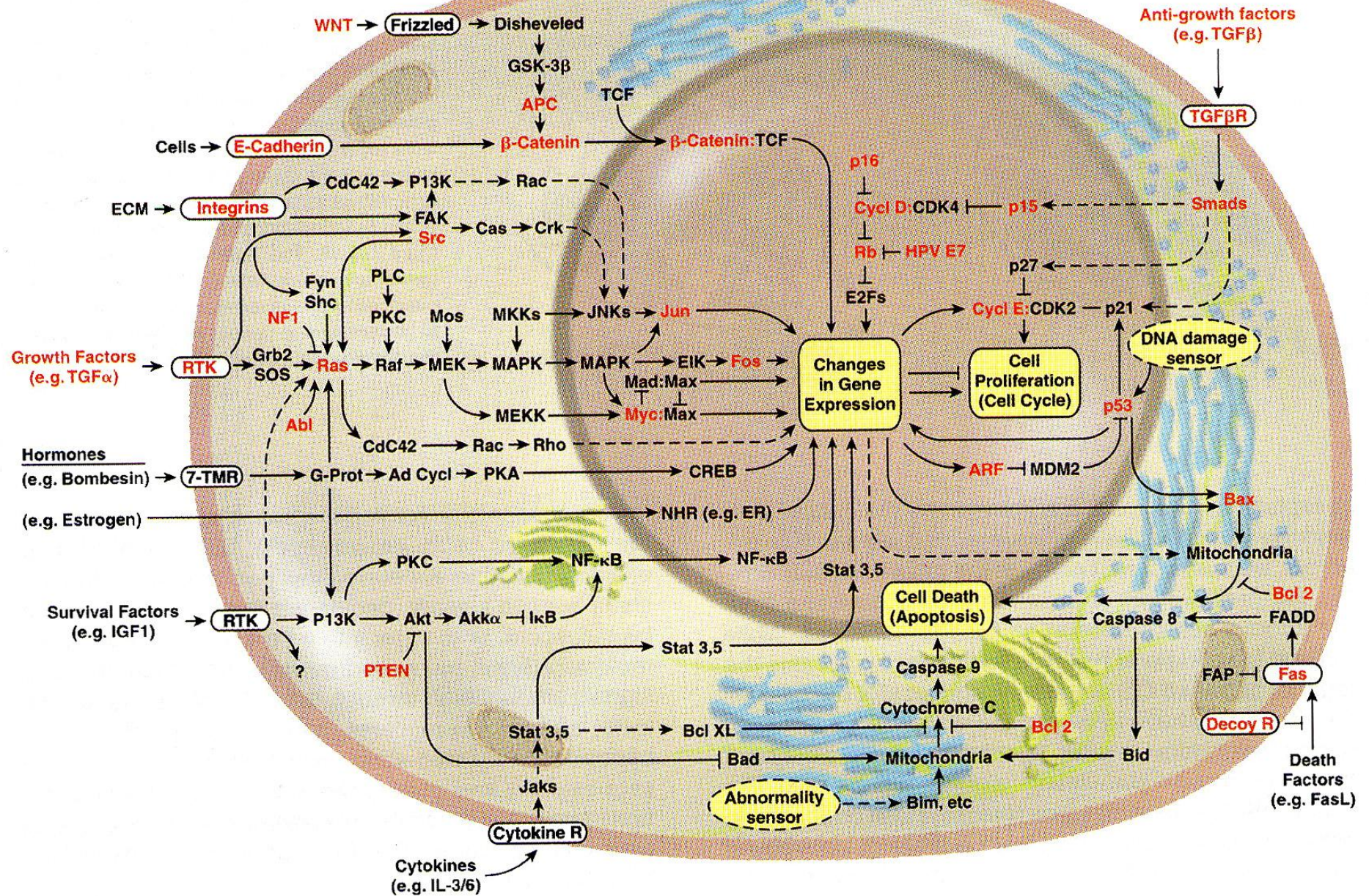
Issues

- Recognition
- Fragmentation
- Solutions

The Consensus Coding Sequences of Human Breast and Colorectal Cancers

Tobias Sjöblom,^{1*} Siân Jones,^{1*} Laura D. Wood,^{1*} D. Williams Parsons,^{1*} Jimmy Lin,¹ Thomas Barber,¹ Diana Mandelker,¹ Rebecca J. Leary,¹ Janine Ptak,¹ Natalie Silliman,¹ Steve Szabo,¹ Phillip Buckhaults,² Christopher Farrell,² Paul Meeh,² Sanford D. Markowitz,³ Joseph Willis,⁴ Dawn Dawson,⁴ James K. V. Willson,⁵ Adi F. Gazdar,⁶ James Hartigan,⁷ Leo Wu,⁸ Changsheng Liu,⁸ Giovanni Parmigiani,⁹ Ben Ho Park,¹⁰ Kurtis E. Bachman,¹¹ Nickolas Papadopoulos,¹ Bert Vogelstein,^{1†} Kenneth W. Kinzler,^{1†} Victor E. Velculescu^{1†}

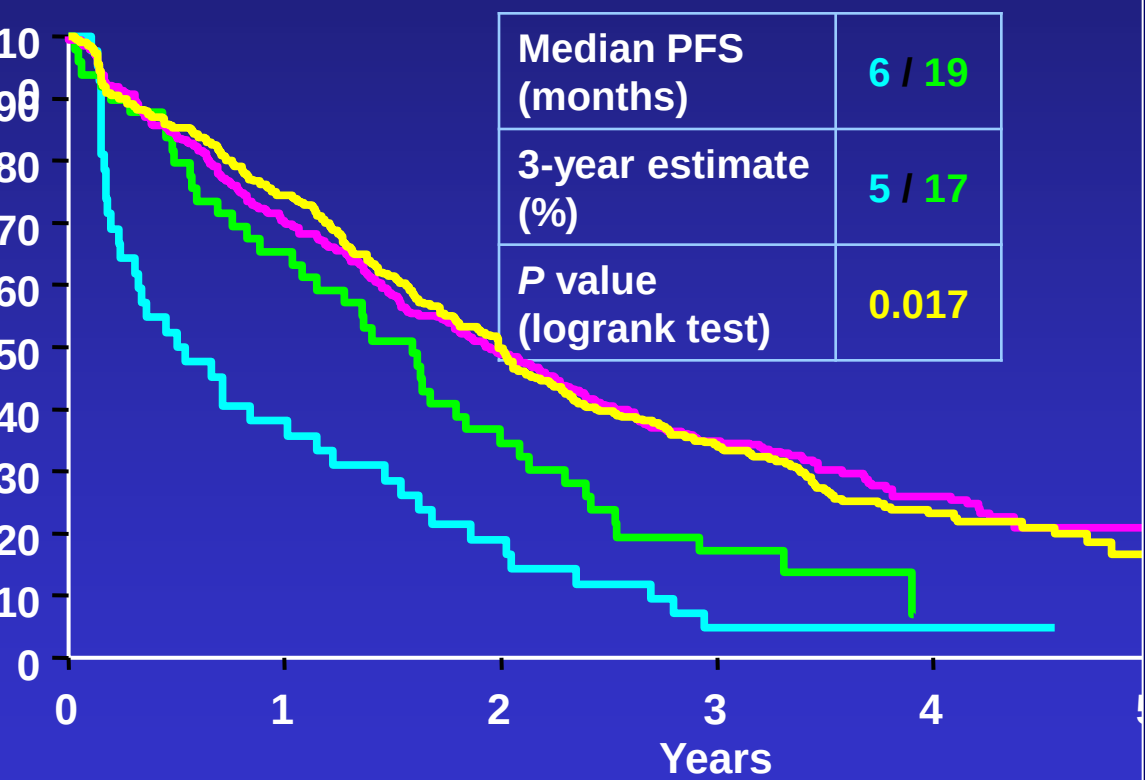
- Breast or colon carcinoma: cell lines or xenografts (n=22)
- 13023 genes sequenced, 3 millions PCR, 452 MB
- total : n=90 mutated genes per cell lines
- Identification of **n=189 significant genes**
- Mean: n=11 (total genome n=14 à 20)
- Transcription, adhesion, invasion
- **CANdicates CANcer (CAN) genes**
 - « Expected »: p53, KRAS, APC, MRE11...
 - Oncogenes for other tissues : MLL3, EPHB6..
 - Unexpected: PKHD1, tubuline tyr ligase TTLL3
- CAN genes C. du sein ≠ C. colorectal



Treatment of cancer needs genomics

GIST are at least 10 diseases

KIT exon 9 mutants (10% of patients)



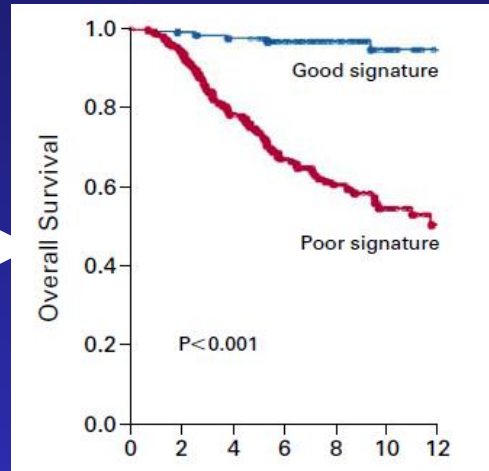
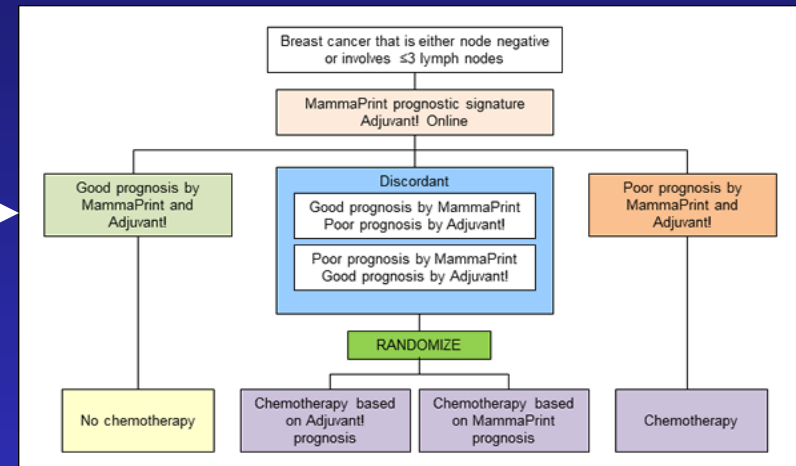
KIT exon 9 mutants: 400 mg / 800 mg
Other patients: 400 mg / 800 mg

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

Rapid complexification of the molecular informations

- Gene expression profile

Mindact EORTC 10041



VOLUME 347 DECEMBER 19, 2002 NUMBER 25

A GENE-EXPRESSION SIGNATURE AS A PREDICTOR OF SURVIVAL IN BREAST CANCER

MARC J. VAN DE VLIJVER, M.D., PH.D., YUDONG D. HE, PH.D., LAURA J. VAN 'T VEER, PH.D., HONGYUE DAI, PH.D., AUGUSTINUS A.M. HART, M.Sc., DORREN W. VOSKUIL, PH.D., GEORGE J. SCHREIBER, M.Sc., JOHANNES L. PETERSE, M.D., CHRIS ROBERTS, PH.D., MATTHEW J. MARTON, PH.D., MARK PARRISH, DOUGIE ATZMA, ANKE WITTEVEEN, ANNUSKA GLAS, PH.D., LEONIE DELAHAYE, TONY VAN DER VELDE, HARRY BARTELINK, M.D., PH.D., SJOERD RODENHUIS, M.D., PH.D., EMIEL T. RUTGERS, M.D., PH.D., STEPHEN H. FRIEND, M.D., PH.D., AND RENÉ BERNARDS, PH.D.

Gene expression profiling predicts clinical outcome of breast cancer

Laura J. van 't Veer^{*,†}, Hongyue Dai^{*,†}, Marc J. van de Vijver^{*,†}, Yudong D. He[‡], Augustinus A. M. Hart^{*}, Mao Mao[‡], Hans L. Peterse^{*}, Karin van der Kooy^{*}, Matthew J. Marton[‡], Anke T. Witteveen^{*}, George J. Schreiber[‡], Ron M. Kerkhoven^{*}, Chris Roberts[‡], Peter S. Linsley[‡], René Bernards^{*} & Stephen H. Friend[‡]

NATURE | VOL 415 | 31 JANUARY 2002

Prediction of cancer outcome with microarrays: a multiple random validation strategy

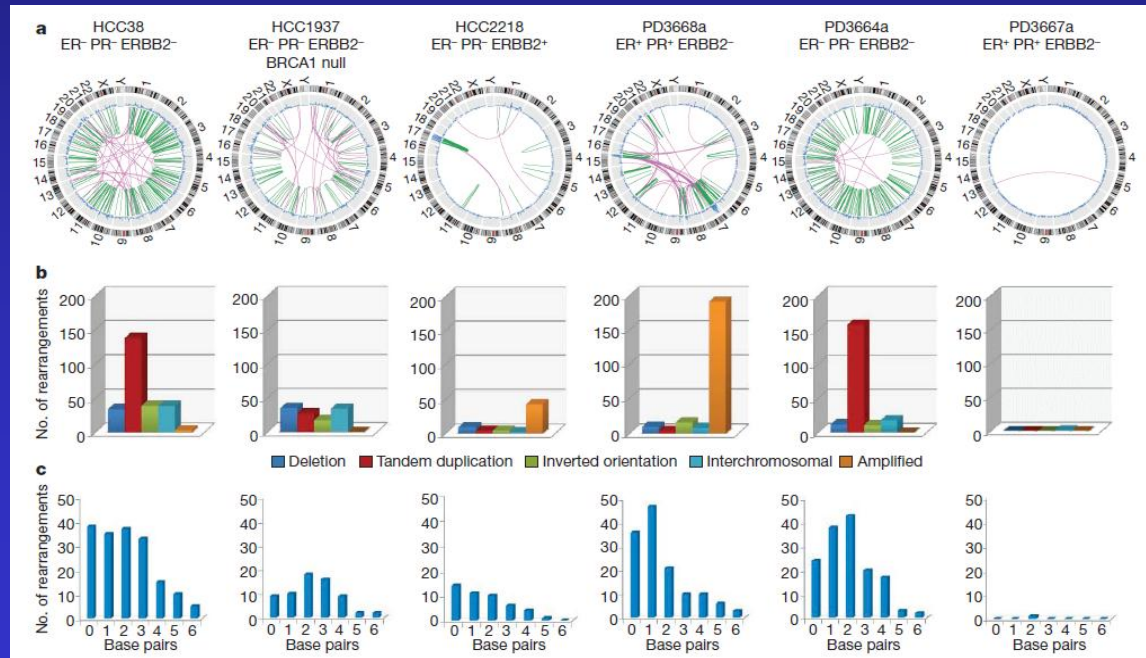
Stefan Michiels, Serge Koscielny, Catherine Hill

Rapid complexification of the molecular informations

Complex landscapes of somatic rearrangement in human breast cancer genomes

Vol 462 | 24/31 December 2009 | doi:10.1038/nature08645

Philip J. Stephens¹, David J. McBride¹, Meng-Lay Lin¹, Ignacio Varella¹, Erin D. Pleasance¹, Jared T. Simpson¹, Lucy A. Stebbings¹, Catherine Leroy¹, Sarah Edkins¹, Laura J. Mudie¹, Chris D. Greenman¹, Mingming Jia¹, Calli Latimer¹, Jon W. Teague¹, King Wai Lau¹, John Burton¹, Michael A. Quail¹, Harold Swerdlow¹, Carol Churcher¹, Rachael Natrajan², Anieta M. Sieuwerts³, John W. M. Martens³, Daniel P. Silver⁴, Anita Langerød⁵, Hege E. G. Russnes⁵, John A. Foekens⁵, Jorge S. Reis-Filho⁶, Laura van 't Veer⁶, Andrea L. Richardson^{4,7}, Anne-Lise Børresen-Dale^{5,8}, Peter J. Campbell¹, P. Andrew Futreal¹ & Michael R. Stratton^{1,9}



Which subset? Which target? Which agents?

The environment of the cancer cell

- uPA/PAI1



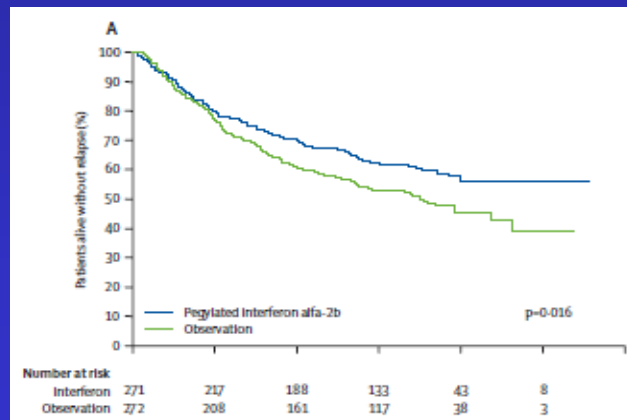
**TUMOR-ASSOCIATED PROTEOLYTIC FACTORS uPA AND PAI-1:
Critical Appraisal of Their Clinical Relevance in Breast Cancer and
Their Integration into Decision-Support Algorithms**

Nadia Harbeck, Manfred Schmitt, and Stefan Paepke

Critical Reviews in Clinical Laboratory Sciences, 44(2):179–201 (2007)

- Immune effectors

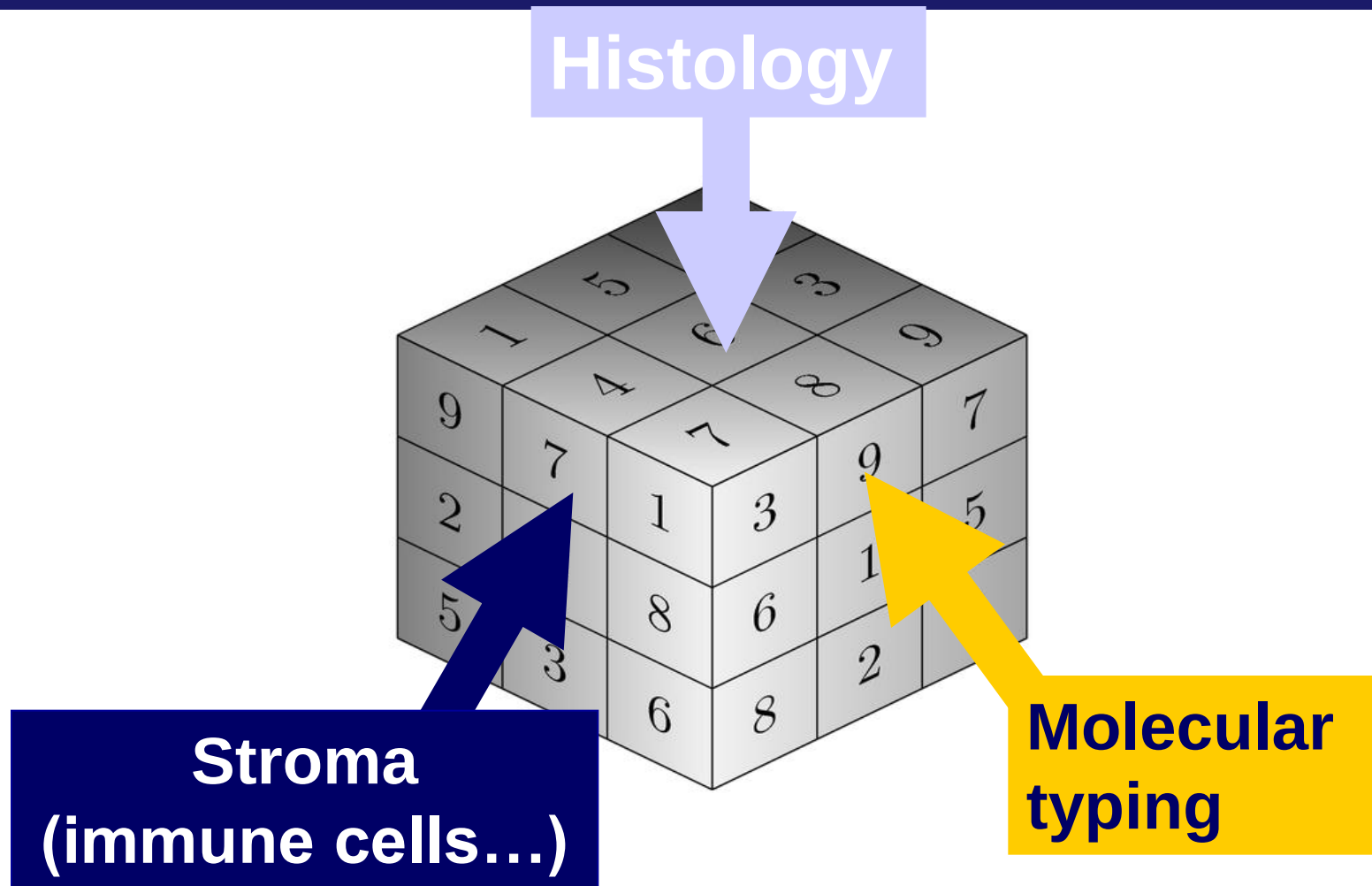
18991: Interferon



**18071
Adjuvant
ipilimumab**

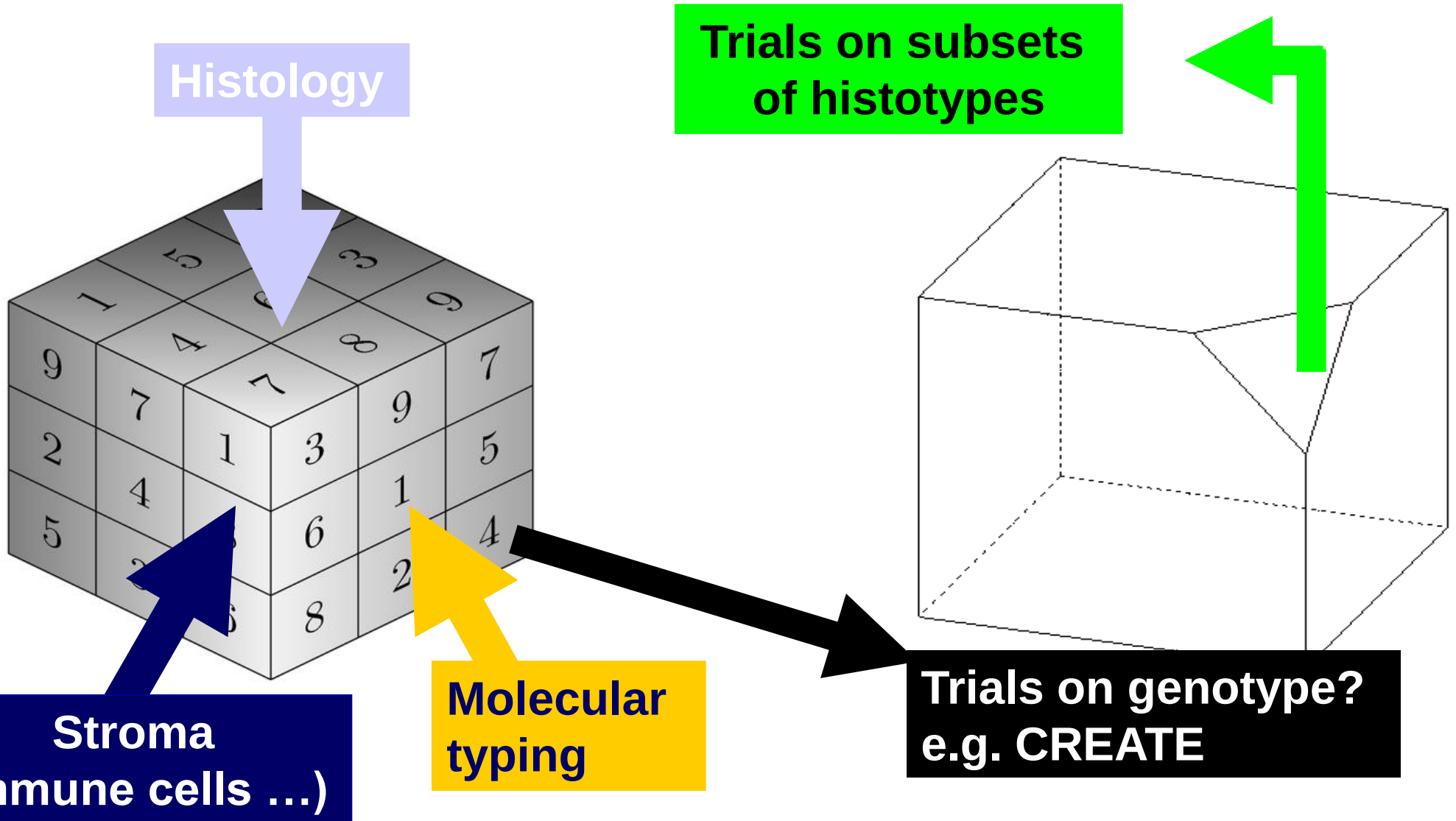
A new vision of the disease

Brussels 25 5 2012



A new vision of the disease

Brussels 25 5 2012



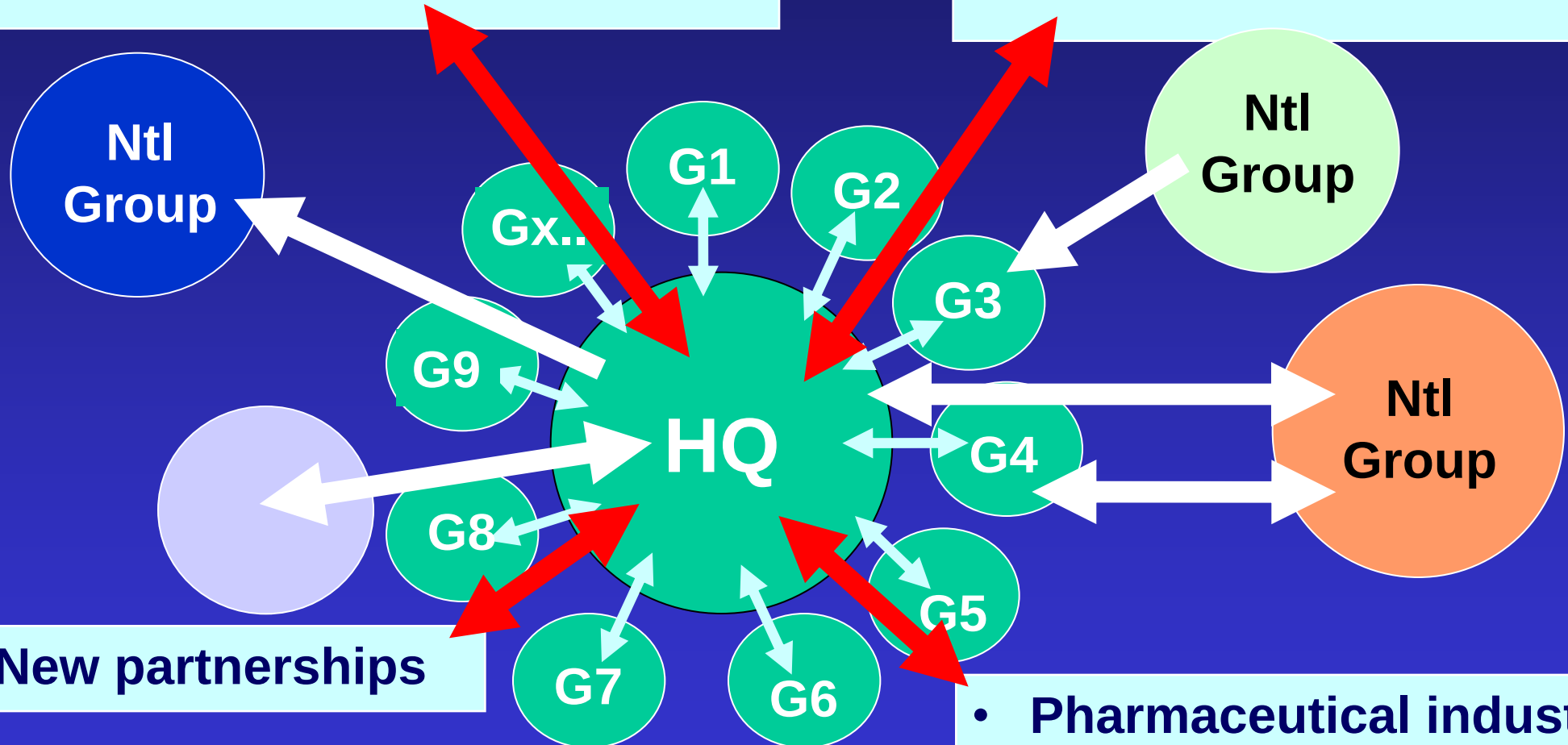
Issues

- Recognition
- Fragmentation
- Solutions

Reinforcing interactions

- New academic partners
- Novel models of collaboration

- Cancer Leagues
- Patient advocacy groups



- New partnerships

- Pharmaceutical industry

Tumor banking and registries

- Tumor banks
- Facilitating the **donation of tumor tissues** across countries
 - Less specific informed consent
 - useful for advancing science
 - Economic value (competition with Far east)
- **Prospective clinical data bases and registries**
- **Collaborative networks** focusing on health care
 - properly funded for quality of care reasons,
- **Observational clinical studies** on selected patient subgroups
 - tailored to answer specific open questions
 - value of retrospective and observational research
 - research resource allocation decisions.

EORTC CRC Screening platform

Tumor blocks
screened for
mutations /
expression

	ras/raf wt, high ligands	1 st line Phase II trial: EGFR monotherapy	treatment outside EORTC trials	treatment outside EORTC trials
	MSI	treatment outside EORTC trials	Trial X and chemotherapy	Phase I with X and Y
	b-raf mut	b-raf inhibitor and chemotherapy	treatment outside EORTC trials	treatment outside EORTC trials
	k-ras mut	treatment outside EORTC trials	MEK-inhibitor (?) + chemotherapy	treatment outside EORTC trials
	k-ras mut	treatment outside EORTC trials		treatment outside EORTC trials
	PI3KCAmut	Chemotherapy and antibody W	treatment outside EORTC trials	Phase I with
	xxx	treatment outside EORTC trials	Phase II with new chemotherapy	A and B
	xxx	No Chemotherapy Inhibitor X and imaging	treatment outside EORTC trials	treatment outside EORTC trials
	xxx	treatment outside EORTC trials	treatment outside EORTC trials	Phase I with Drug F
	xxx	treatment outside EORTC trials	treatment outside EORTC trials	treatment outside EORTC trials

EORTC Clinical Trials

“Rational treatments to right patients”



Menu

[Home](#)[Charter](#)[Survey](#) →[Export](#) →[Patient](#) →[Audit](#) →[Online Help](#) →

Welcome to conticabase

The CONTICANET database and tumour bank

This database contains anonymised information describing the tumour, treatment and follow-up as well as tumour sample availability and molecular biology analyses for mesenchymal tumours except GIST and bone tumours.

The tool can be used as a local center database thanks to its **rules for access to patient data and material**. It will be maintained and updated centrally. Please follow this **this link** to fill the account application form

The query tool allows users to ask questions about the overall content of the database in order to evaluate the feasibility of specific collaborative studies.

We hope this database will become an important tool for increasing our knowledge on these rare tumours and for developing joint research programmes.

Website requirements

This website has been designed for both **Firefox 3** (advised) and **Internet Explorer 7** (or older versions). Please note that some features may not work correctly with other web browsers.

Content overview

conticabase currently contains the following data from **26** out of the **43** registered centres :

- **4804** Patients
- **4826** Tumours
- **5699** Samples (**5493** Paraffins and **2600** Frozen)

Networks and PAGs

- Health care Networks
- **Reference centers**, within quality control programs. They improve health care and they improve accrual in trials as well as clinical quality within clinical trials.
- **Patient information about trials** (what they are, where they are available). There is even a greater added value for the today patient in entering a trial.

How can
YOU help?

Sign
our petition
NOW...

...and ask FRIENDS,
COLLEAGUES and
CONTACTS...

...to support
this initiative
by SIGNING UP
too!

The European Action Against Rare Cancers is a joint initiative based on a partnership between the European Society for Medical Oncology (ESMO), the European Organisation for Rare Diseases (Eurordis), the European Cancer Patient Coalition (ECPC), Conticanet, the Association of European Cancer Leagues (ECL), the Chronic Myeloid Leukaemia Support Group, the International Brain Tumour Alliance (IBTA), Orphanet, the Chronic Myeloid Leukaemia Advocates Network, the European Institute of Oncology (EIO) and the Fondazione IRCCS Istituto Nazionale dei Tumori, as well as Novartis Oncology as the founding sponsor.

The organisations collaborate as equal partners and all decisions are made on the basis of consensus. The initiative is moreover supported by eight corporate organisations.

For more information about this European initiative, please visit our Web site:
www.rarecancers.eu

or contact us:
European Action Against Rare Cancers
c/o Robert Schaefer

Brunnenstrasse 178/179
10119 Berlin, Germany
T +49 (0)30 288 797 55
F +49 (0)30 288 797 66

schaefer.robert@esmo.org
www.rarecancers.eu



Collaborating Partners:



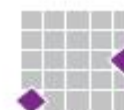
Initiating Sponsors:



**RARE CANCERS:
MORE COMMON
THAN
YOU THINK!**

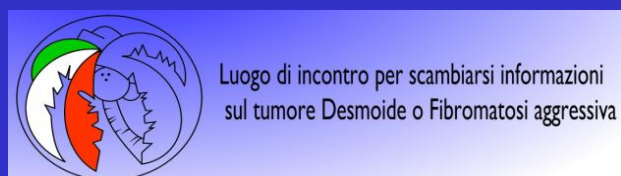
☒ Sign the call to
ACTION AGAINST RARE CANCERS

www.rarecancers.eu



European Action Against Rare Cancers

18 SPAEN Full Members



Pharma and academic trials

- Share the results across trials including with industry sponsored trials.
- Collaboration across companies
 - Comparing drugs in tumors with unmet needs
 - Combining drugs
- Large multisources Databases, Registries
 - Funding issues

How to improve collaboration ?

Centres

National
European
Worldwide

Phase II, not so rare subtypes

Phase III - not so rare subtypes

Phase II super rare subtypes

Incentives for clinical trials

- Incentives for orphan drugs devt for pharma companies
- Drug supply by pharma to academics before any approval?
- Screen rare tumors in Phase 1 setting
 - based on molecular screening ?
- A need **framework study protocols** on specific rare cancers liable to be sequentially exploited for different drugs ?

Enrich clinical trials

- Enrich the informations collected from Rare cancer trials need to be richer in information in order to maximize their efficiency, e.g. a long follow up for each patient

Collaborations

- In rare cancers, national, international, even global collaborations should be pursued to make investigator-driven studies possible.
- Clinical Trial Directive is currently under revision. It is recommended that it is improved in some crucial aspects.
- Sponsorship of international trials for investigator-driven studies are sponsored by academic institutions, collaborative groups,
 - difficult to comply with regulations which differ from country to country,

A World Sarcoma Network is needed



The World Sarcoma Network (2009)

Studies pipeline

- nilotinib in PVNS with t(1,2) M-CSF-col6A3 fusion gene
- mTOR inhibitors in PEComas, and in tumours of the TSC complex
- Aplidin in Dedifferentiated Liposarcomas with JUNK overexpression
- Alk inhibitors in inflammatory myofibroblastic tumours with Alk amplification and over expression
- IGF1R inhibitors in GIST with IGF1R over-expression and amplification
- MDM2 inhibitors (nutlin3a) in WDLPS with MDM2 amplification
- MET inhibitors in sarcomas with translocation involving fusion genes encoding for abnormal transcription factors (ASPS, CCS)
- VEGFR2 inhibitors in ASPS

How to improve collaboration ?

Perimeter

National
European
Worldwide

Structure

Centre
Group
Intergroup
EU (EORTC)

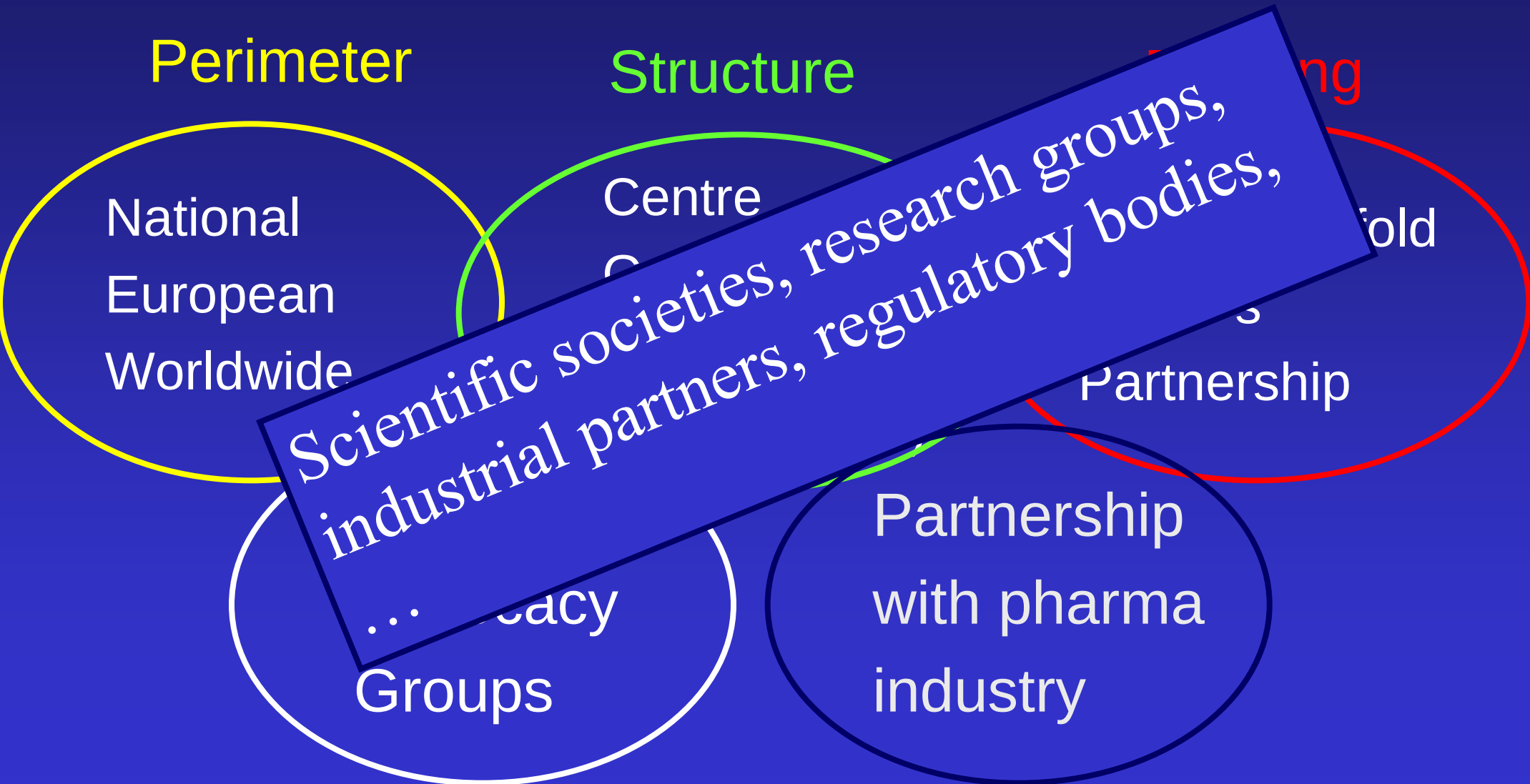
Funding

National- x-fold
EU FPs
Partnership

Patient
Advocacy
Groups

Partnership
with pharma
industry

How to improve collaboration ?



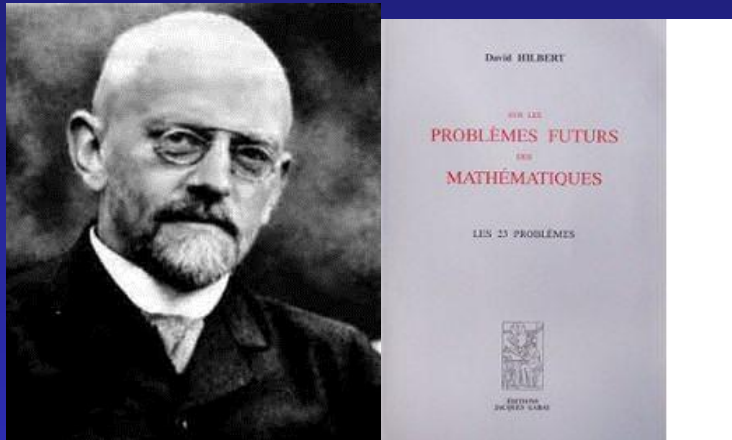
Regulatory aspects

- Regular consultations between regulatory bodies, pharma, academia, scientific societies-ESMO, and PAGs
- Simplify and streamline the access to compassionate use programs

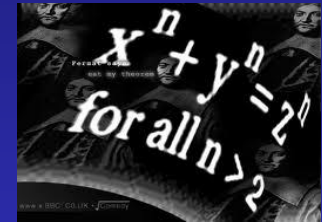
To address major scientific questions

Oncology research ... a « hard science »
with multidimensional complexity

1900



17/23 solved



2000



1/7 solved

Questions for the next 20 years

Some are organisational

1. Is it possible to organize the health care systems to ensure optimal local treatments, surgery and radiotherapy, for all patients?
2. How to build simple academic clinical trials?
3. Is it possible to organize annotated multinational tumor collection and storage to enable clinical research on molecular subtypes?
4. How can we integrate the volume of molecular information generated to the routine care of the patient?
5. Can health care systems absorb the cost of targeted treatments?

Questions for the next 20 years

Brussels 25 5 2012

Some are scientific

6. How to recognize the driving mutations in individual patients to guide the treatment?
7. How to generate algorithms with tumors+ constitutional genomics+patients info to guide treatment decisions?
8. Does adjuvant treatment with targeted therapies prevent or postpone relapse?
9. Can we define better surrogate markers for overall survival?

Questions for the next 20 years

10. Is resistance to targeted treatment unavoidable in advanced phase?
11. What are the paradigms of combination or sequential treatment to prevent resistance?
12. Can surgical removal of the metastatic burden reduce the risk of emergence of clonal and clinical resistance?
13. Can immunotherapy cure minimal residual disease in man?
14. Is it possible to generate models on when to stop specific treatment?

Conclusions

Improving clinical research on rare cancers

- Recognition : improved through education and networking
- Management : poor to be improved with reference center
- Fragmentation : ineluctable- an opportunity
- Solutions : collaboration, interactions