

Francois Pognan, PhD, Discovery Investigative Safety Basel, Switzerland BioMed 21: A Human Pathways Approach to Disease Research 26-27 June 2017 – Bethesda, MD, USA

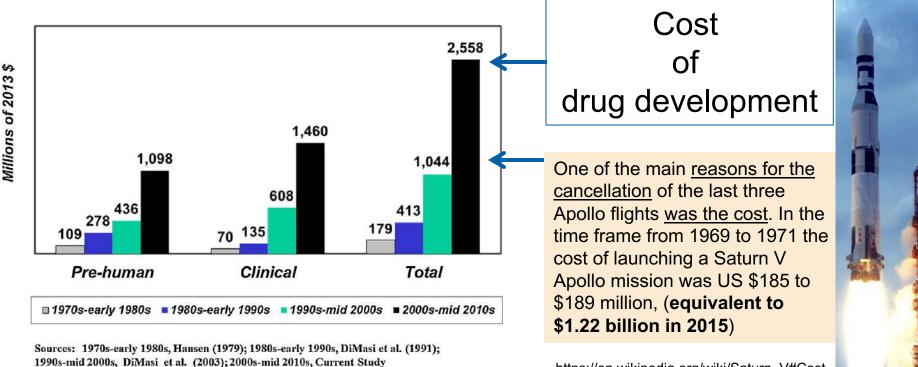


Thoughts for today's debate

- Drug development cost and challenges
- Data waste
- Data mess with order
- New order needed for data recycling
- What is being done and will be done
- Examples of new knowledge out of old data
- Discussion: precompetitive data sharing

"Give, and you will receive"



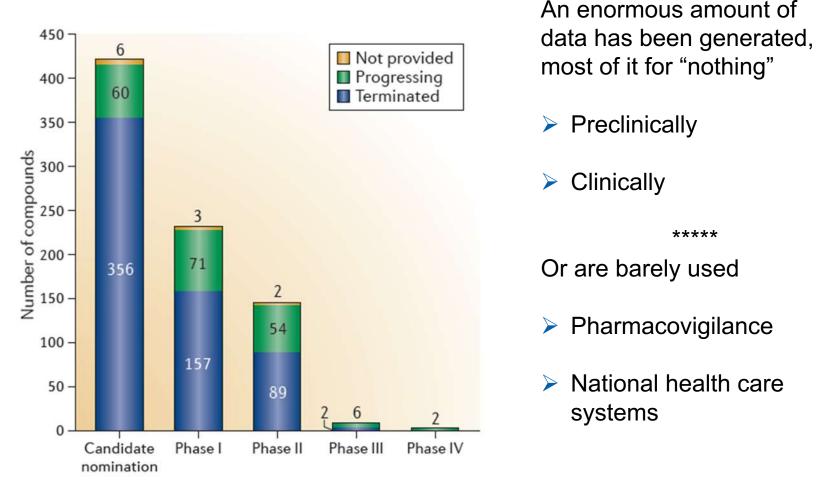


DiMasi, J. A., et Al., J Health Econ 47, 20-3

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https://en.wikipedia.org/wiki/Saturn V#Cost

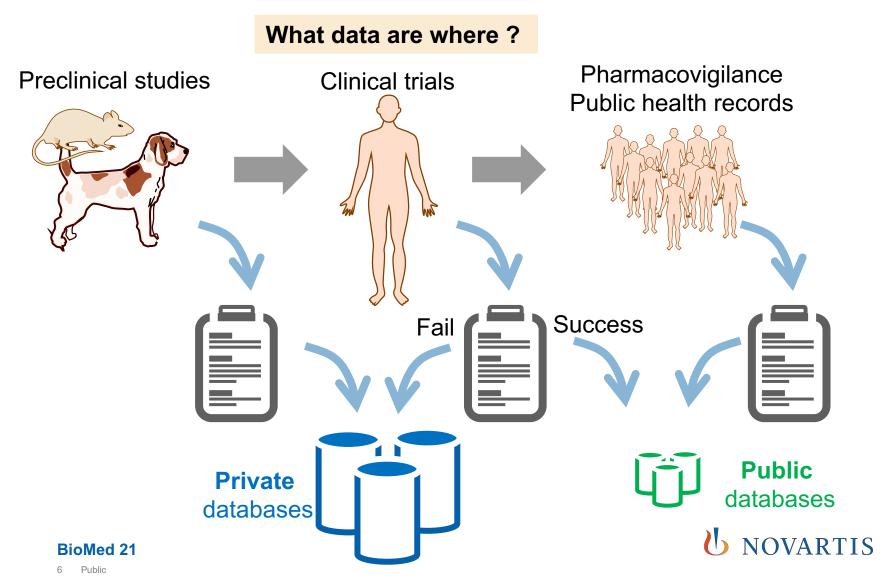
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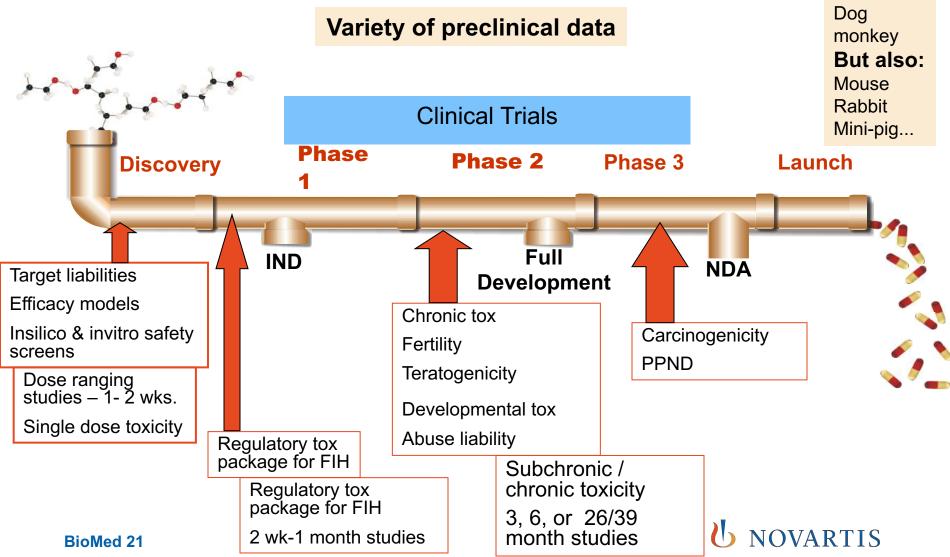
Waring, M. J. et al.. Nature Reviews Drug Discovery 14, 475-486

- Collectively, the pharma industry has generated a gigantic amount of preclinical and clinical data
- These data are conscientiously kept in archives for retrieval when required by HAs
- Data are archives in a way that allows almost immediate retrieval, but do not permit cross or meta-analysis
- Pharma DBs contain data for compounds successfully developed to drug, and many logs more of unsuccessful molecules
- Data therein are mostly considered as competitive advantage
- These data are mostly sleeping despite containing an unmined wealth of data that could be turned into knowledge



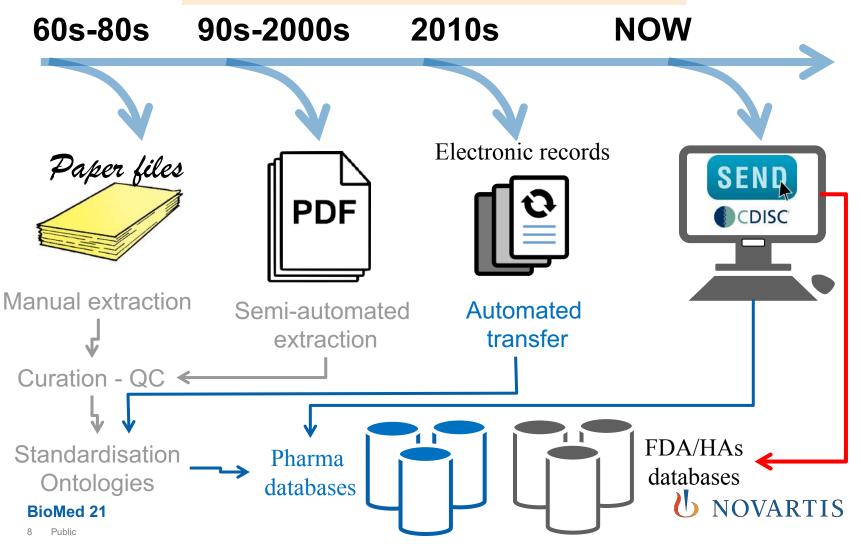
Mostly:

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Heterogeneity of preclinical data supports



 The eTOX initiative under the aegis of the European Union (IMI – Innovative Medicine Initiative)



Project vision: to develop innovative strategies and novel software tools to better predict the potential side-effects of new drug candidates on the basis of integrative approaches.







Challenges and needs to use existing data in drug development From data to knowledge to wisdom

EFPIA data



Organised eTOX db







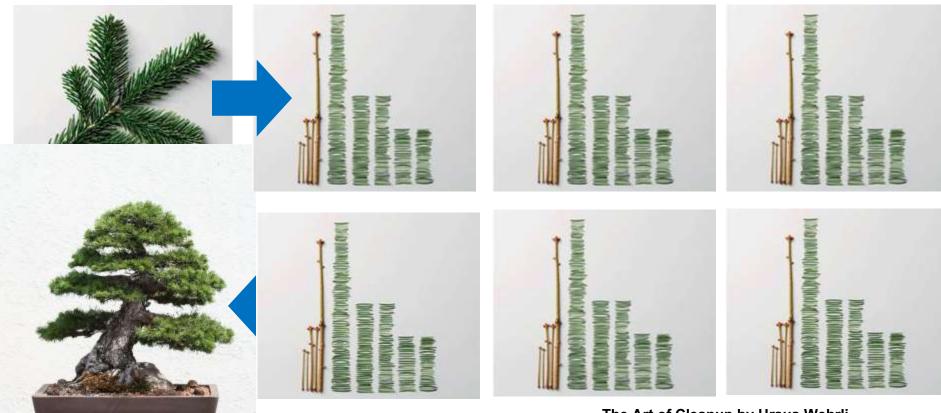


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Challenges and needs to use existing data in drug development From data to knowledge to wisdom

Organised eTOX db

Enabled ontology classifications - knowledge



Meta-analysis, extra knowledge : wisdom

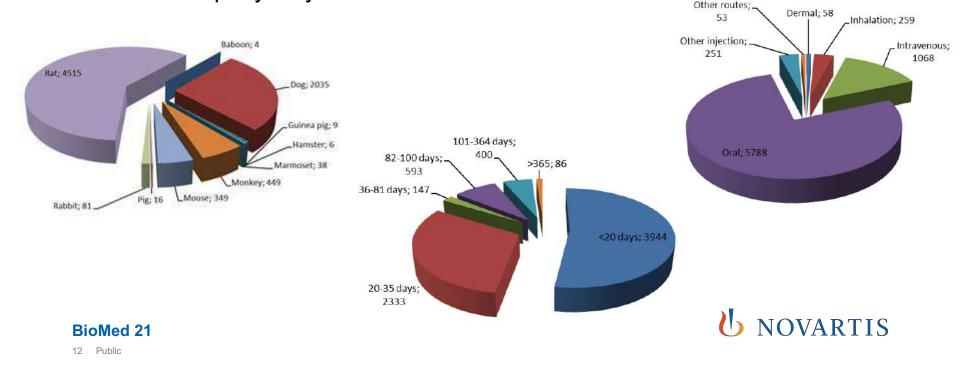
The Art of Cleanup by Ursus Wehrli https://www.brainpickings.org/2013/03/28/the-art-of-cleanup-ursus-wehrli/



As of Jan 2017, the eTOX database is

Total number of structures: 2112 (476 confidential compounds) **Total number of studies:** 8238 This represent about 20 millions entry lines

Private: company only and not shared



ontologies | dealing with 80 000 terms

Current workflow allows efficient ontology maintenance and term mapping More than **80 000 finding verbatim terms** out of **20 millions entry lines** mapped to the appropriate locations.

- 1. Ontobrowser collects verbatim terms from Vitic-Nexus when they are loaded (lines of data in scope).
- Unknown verbatim terms are manually assigned as synonyms to > 7 000 preferred terms.
- 3. Pending synonyms are approved (or rejected) by a second curator.
- 4. The Vitic-Nexus database is loaded with preferred terms and synonyms during build.

OntoBrowser <u>http://opensource.nibr.com/projects/ontobrowser/</u>

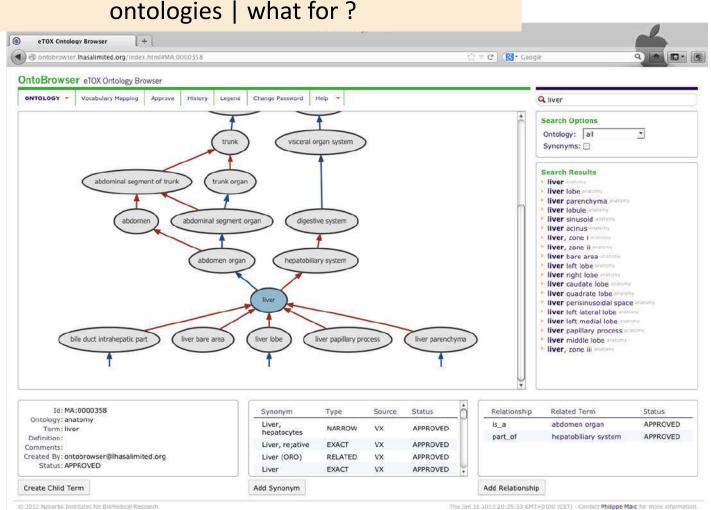
Ontology/Codelist	Preferred Terms	Synonyms	Cross Refs*
anatomy	3141	18873	13%
effect level	8	49	0%
histopathology	1008	33310	38%
in-life observation	698	12294	0%
laboratory test name	1324	11301	99%
moa	465	0	0%
pk parameters	309	1868	95%
route of administration	123	411	100%
sex of participants	3	123	100%
species	27	32	93%
strain/substrain	84	1121	98%

*Percentage of preferred terms with cross references public nomenclatures such as SEND/NCI, INHAND, MedDRA etc...

https://ontobrowser.lhasalimited.org/index.html#MC:0000301

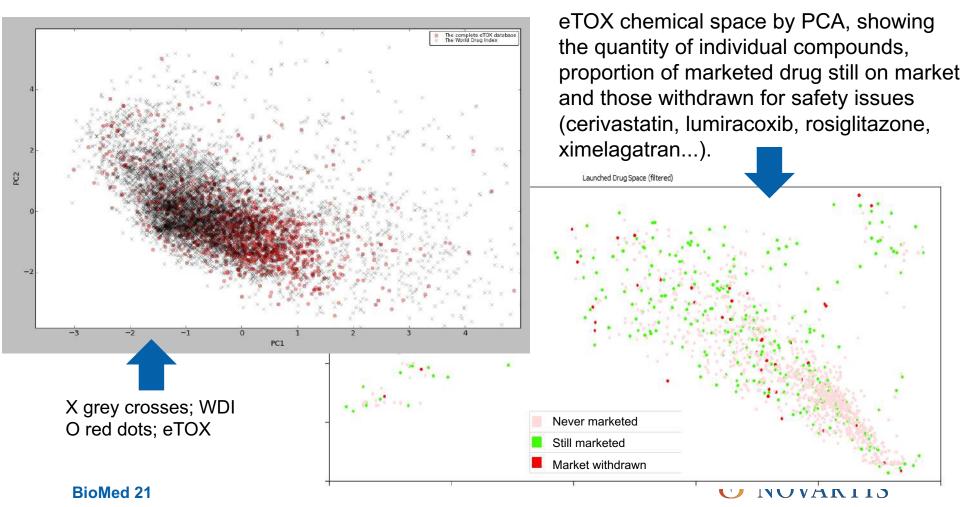


Collecting data is fine, but is a specific finding related to "liver", hepato*, hepatic Kupffer, cholangiocyte, bile duct ...???

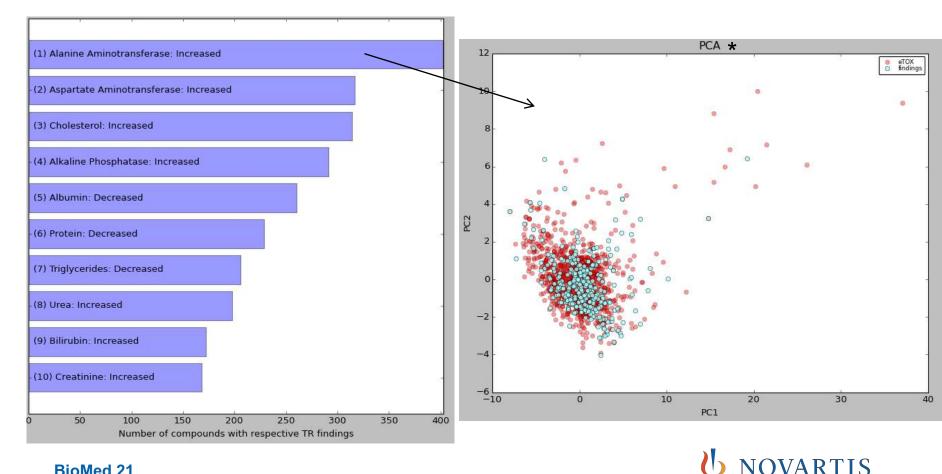


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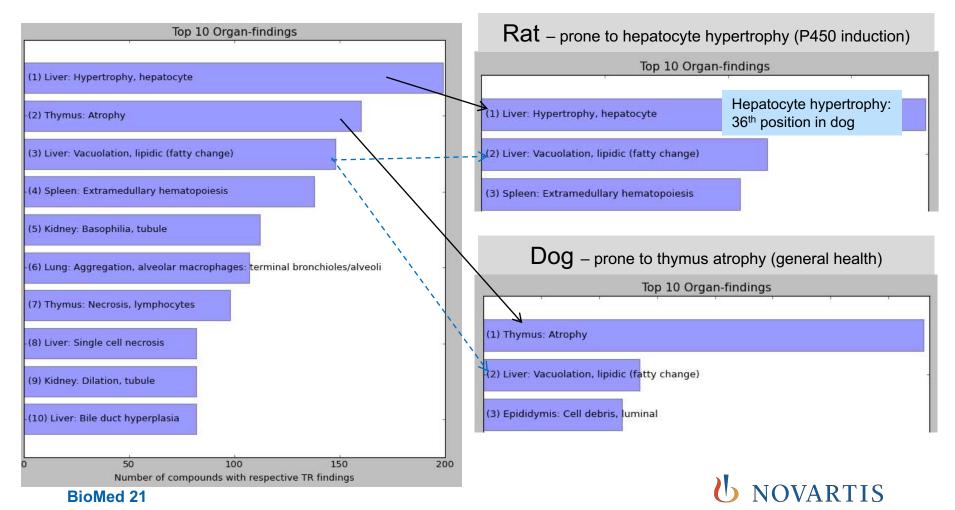
eTOX chemical space and content



- Meta-analyses allow extraction of larger knowledge, not otherwise possible, eg:
 - Clin chem: Most common findings all species



Histopathological findings – All species

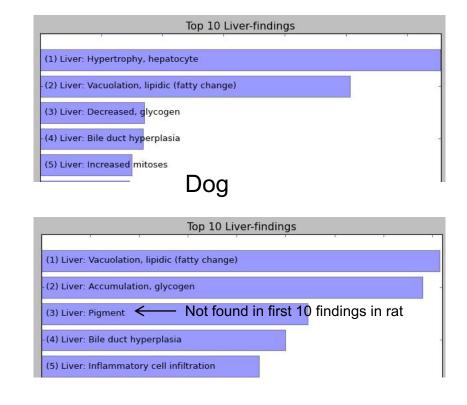


Histopathological findings – liver only

All species

Top 10 Liver-findings	
(1) Liver: Hypertrophy, hepatocyte	
(2) Liver: Vacuolation, lipidic (fatty change)	1
(3) Liver: Bile duct hyperplasia	
(4) Liver: Decreased, glycogen	-
(5) Liver: Accumulation, glycogen	
(6) Liver: Single cell necrosis	15
(7) Liver: Inflammatory cell infiltration	
(8) Liver: Inflammation	
(9) Liver: Increased mitoses	
(10) Liver: Lymphohistioplasmacytic inflammatory cell infiltration	15
	10
200 400 600 800 1000 1200 1400 Total number of TR findings	160

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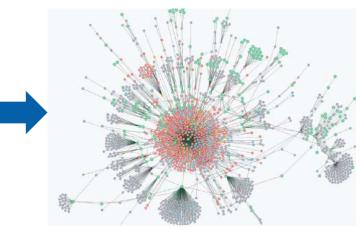


New structural alerts from the eTOX database (chemistry + toxicology)

Dataset

(structures flagged has positive or negative for a given tox. end-point)



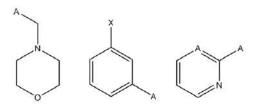


Fragmentation

(generation of unique fragments from every dataset structures, using several fragmentation algorithms)

Matrix analysis

(Retrieval of structural alerts and scoring coming from the fragments that are the most frequently encounter in toxic structures) e.g: fragment_combination_51263:



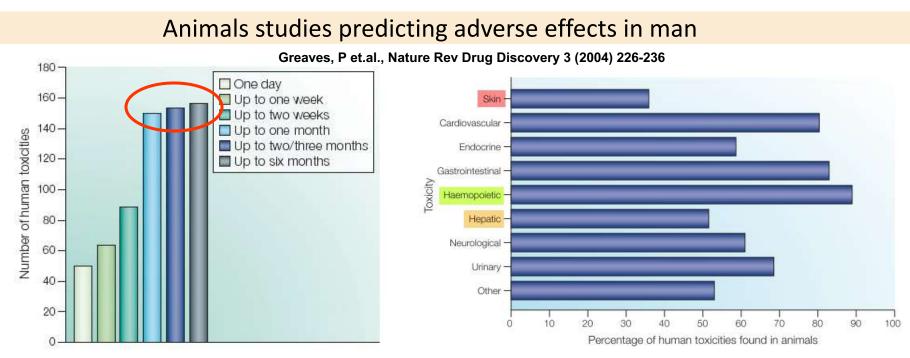
42 substructure-relationships with the dataset (40 with toxic and 2 with non toxic structures)

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	structure_1	structure_2	structure_3	structure_	4 structure_5	structure_6	structure_7	structure_8	structure_9	structure_10	structure_11	structure_12	 structure_1500
ination_1		Contra Separativ			a av seres er terres			b men mensen e					
ination_2													1. Sec.
ination_3													
ination_4													
ination_5													
ination_6													
ination_7													
ination_8													
ination_9													
ination_10													
ination_11													
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ination 60000													

Matrix of substructure-relationship

(example with 1.500 structures X 600.000 fragments where red shows substructure relationships with toxic structures and green shows substructure relationships with non-toxic structures)



With limited data:

94% of animal toxicities that correlate with human toxicities were detected in preclinical studies of ≤ 1 month duration.

If some parameters or studies are irrelevant to human, why running them, need better studies, better design, different approach → Impact on guidelines ?

What is next after eTOX - eTRANSAFE (eTS)

• Three objectives:



- exhaustive analysis of correspondence and validity of animal data for human safety
- discovery of translational and reverse-translational biomarkers
- predict animal toxicities

• Four deliverables:

- preclinical data base with retrospective and prospective data (SEND) from multiple companies
- mining and visualisation tools for cross-analysis with human data
- in silico predictive platform (algorithms)
- new translation and reverse-translation biomarkers

• Impacts

- preclinical studies adapted to human outcome = increased safety, reduced attrition
- 3Rs
- preclinical knowledge management



Outlook

- New knowledge extractable from old data
- Need to very large amount of data \rightarrow consortia
- Need for quality and standard format (e.g., SEND)
- Need for large, comprehensive systematic and objective assessment of preclinical data for their validity of human safety prediction
- Impact on
 - 3Rs
 - Future design of preclinical studies
 - Better understanding of toxicology
 - Translational aspects of safety assessment
 - Perhaps on duration of development
 - Perhaps on overall cost of development



Outlook

- Emergence and understanding of:
 - Concept of data sharing
 - "Pre-competitive" data
 - International consortia
 - The pharma industry has to collaborate in order to compete
 - Health Authorities, and academics have to join or even to push for initiation of collaborations



Data sharing is a gain, not a loss

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