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[www.imim.es/URECMC/eng](http://www.imim.es/URECMC/eng) Miquel Porta - Mexico City, 7 Sept. 2007  
 PART 2 – 1

**DDT and related compounds and risk of pancreatic cancer**

1.- Garabrant DH et al. *JNCI* 1992; 84: 764-71.  
 2.- Malats N, Real FX, Porta M.  
 + Garabrant DH et al. *JNCI* 1993; 85: 328-9.

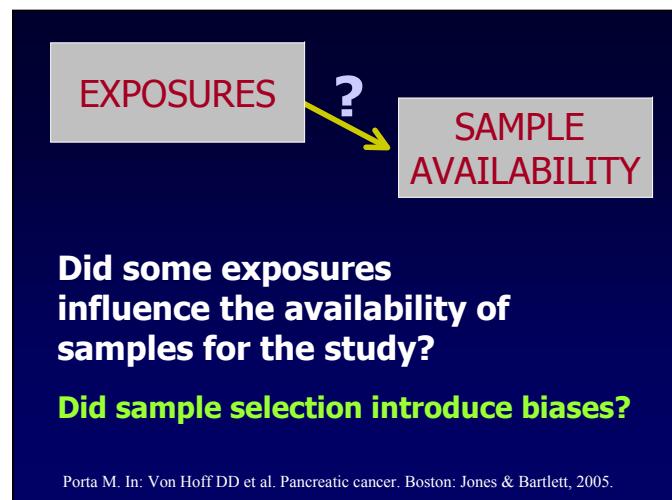
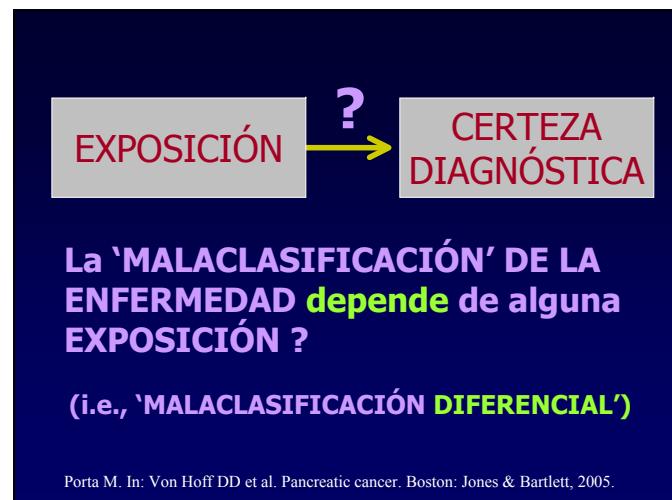
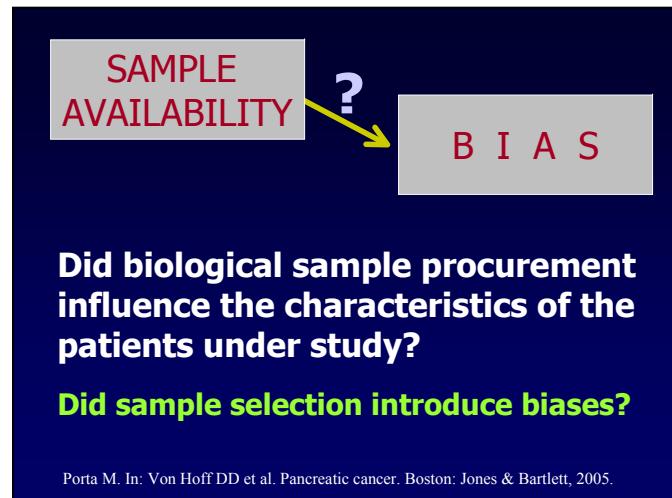
	DDT family	DDT	Ethylan	DDD
All subjects	3.3*	4.8*	5.0*	4.3*
Cytologically confirmed	21.0*	∞*	∞*	15.4*
Death certificate	0.8	1.0	2.6	1.4

\*P ≤ 0.02



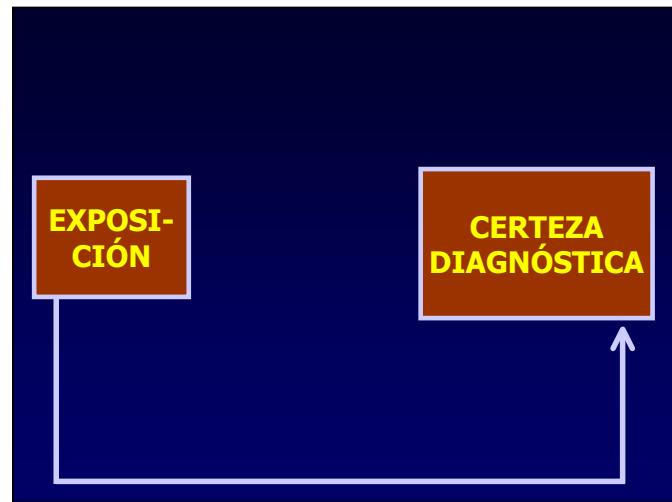
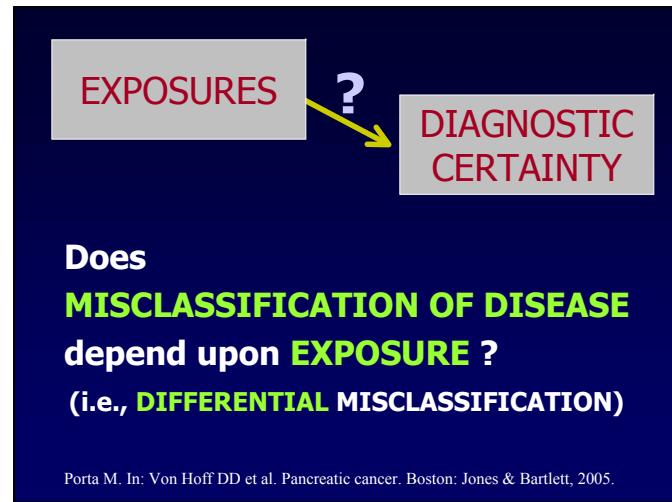
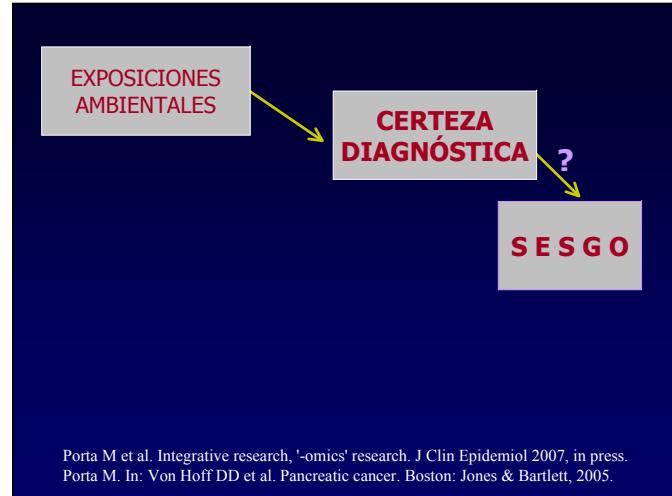
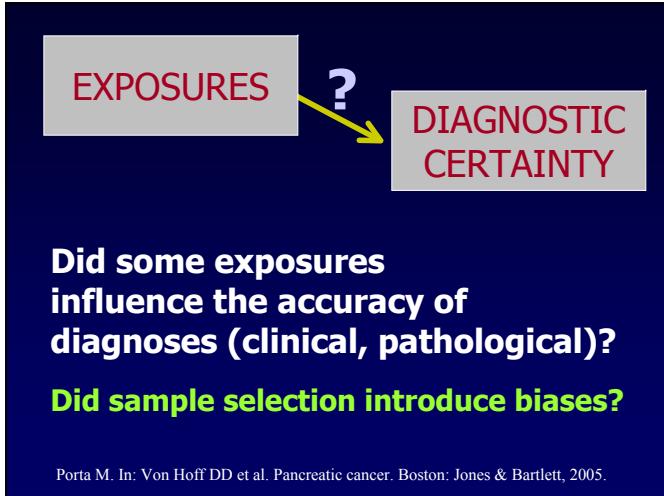
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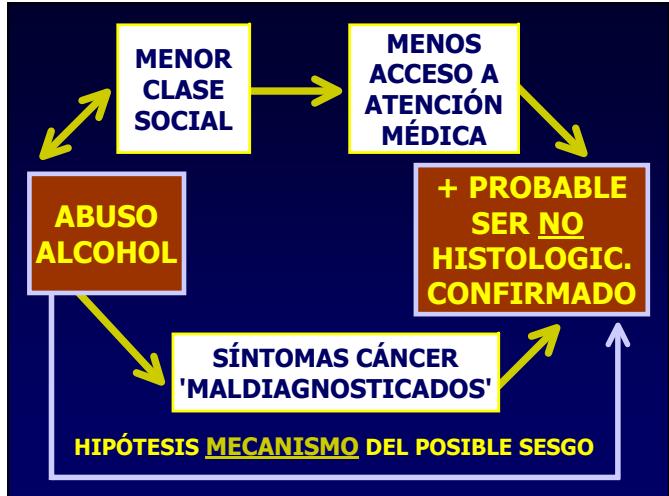
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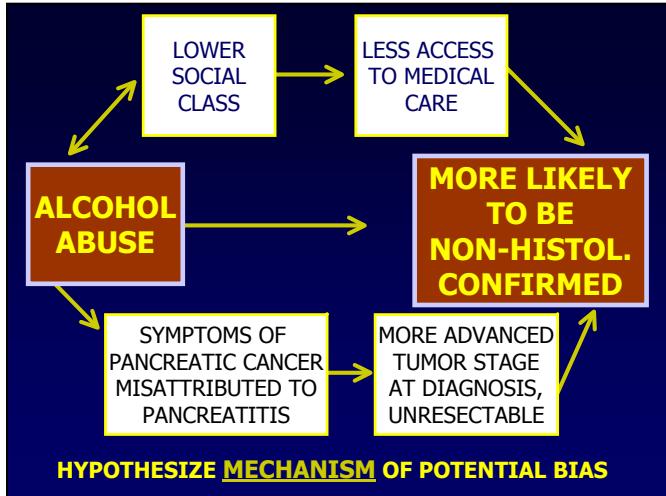


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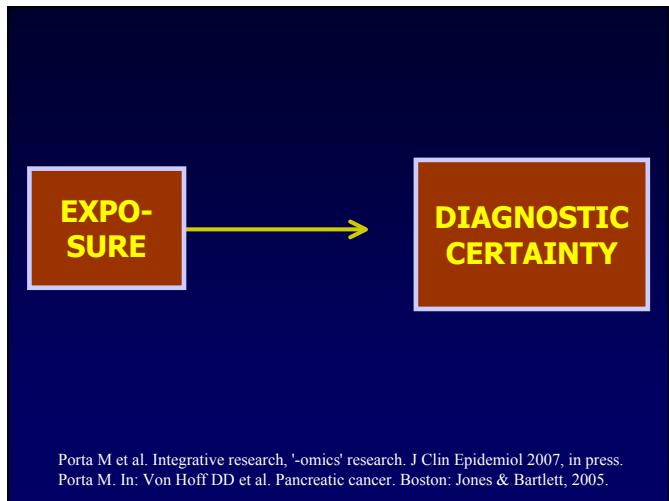
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Porta M et al. Integrative research, '-omics' research. J Clin Epidemiol 2007, in press.  
 Porta M. In: Von Hoff DD et al. Pancreatic cancer. Boston: Jones & Bartlett, 2005.

Silverman DT et al. *Cancer Research* 1995

Case-control studies based exclusively on histologically confirmed cases may preferentially select cases with lower exposure to alcohol. Alcoholics may be more likely to be nonhistologically confirmed than nonalcoholics resulting from less access to medical care or cancer-related symptoms that are misdiagnosed as alcohol related. Because we included all likely cases, regardless of histological confirmation, our study was less prone to this type of selection bias.

**EXCLUDE PATIENTS**  
**>70-80 years, without HV ?**

**PROBLEM:**

Patients without HV  $\approx$  20 - 60% of all cases  
 Patients >70 years  $\approx$  30 - 40% of all cases

**PROBLEM: ↓ EXTERNAL VALIDITY.**

**PROBLEM: ↓ SCIENTIFIC RELEVANCE.**

**PROBLEM: MISCLASSIFICATION  
 may persist even with WITH HV.**

**RESTRICTION  
 "A PRIORI"**



Porta M et al. J Clin Epidemiol 1994 & 1996  
 Silverman DT et al. Cancer Res 1995  
 Skinner HG et al. CEBP 2003  
 Michaud DS et al. Am J Epi 2003

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**IF classification of D was related  
 to E (e.g., through age and sex)**

**THEN → risk of DIFFERENTIAL  
 MISCLASSIFICATION**

**→ ↓ INTERNAL VALIDITY**

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Porta M et al. Integrative research, '-omics' research. J Clin Epidemiol 2007, in press.  
 Porta M. In: Von Hoff DD et al. Pancreatic cancer. Boston: Jones & Bartlett, 2005.



Ann Epidemiol 2002;12:7-14.

**Attention to  
selection biases !**

## Generalizing Molecular Results Arising from Incomplete Biological Samples: Expected Bias and Unexpected Findings

MIQUEL PORTA, MD, MPH, PhD, NÚRIA MALATS, MD, PhD, JOSEP M. COROMINAS, JULI RIFÀ, MD, PhD, JOSEP L. PIÑOL, MD, PhD, AND FRANCISCO X. REAL, MD, PhD, FOR THE PANKRAS I PROJECT INVESTIGATORS\*

**PURPOSE:** In molecular epidemiology, obtaining biological samples for all subjects targeted for study is frequently hampered by ethical, clinical, and logistic factors. The extent to which the incompleteness of biological samples could cause bias is rarely analyzed in depth. Here we report some expected bias and some unexpected findings during a study on mutations in the K-ras gene in exocrine pancreatic cancer

**METHODS:** In this case-case study, all patients registered with EPC between 1980 and 1990 at two general hospitals were retrospectively identified from the hospital tumor registries. Their clinical records were abstracted and paraffin-embedded samples retrieved from pathology records. DNA was amplified, and mutations in codon 12 of the K-ras gene were detected using the artificial RFLP technique.

J Epidemiol Community Health 2002;56:734-738

734

### DEBATE

Incomplete overlapping of biological, clinical, and environmental information in molecular epidemiological studies: a variety of causes and a cascade of consequences

M Porta, N Malats, J Vioque, A Carrato, M Soler,

**Attention to  
selection biases !**

**Out of >600 cases:  
7 wt vs. 17 mutated**

Cancer Epidemiology, Biomarkers & Prevention

Vol. 9, 1223-1232, November 2000

K-ras and p53 in Pancreatic Cancer: Association with Medical History, Histopathology, and Environmental Exposures in a Population-based Study.<sup>1</sup>

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Ann Epidemiol 2002;12:7-14.

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selection biases !**

**RESULTS:** Results on the mutations (RM) were obtained for 51 of the 149 cases of EPC (34.2%). There were no significant differences on the availability of RM by age, gender, and tumor stage at diagnosis, but RM were over five times more likely to be available from one of the hospitals. Subjects with RM were more likely to have received a treatment with curative intent (OR = 11.56, 95% CL: 2.88-46.36). The existence of RM was positively associated with the availability of information on alcohol use and family history of cancer. Subjects with RM tended to belong to higher occupational groups and to smoke less than subjects without RM.

Unexpectedly—given that in EPC K-ras mutations have consistently been found unrelated to age, gender, tumor stage, and other clinical factors—, cases with a K-ras mutation were more likely than wild-type cases to have information on tobacco and alcohol use (OR = 3.29,  $p = .21$ ), medical history (OR = 4.46,  $p = .41$ ), and family history of cancer (OR = 4.80,  $p = .01$ ). The relationship between completeness of clinical records and K-ras mutations among cases with RM could not be accounted by age, gender, and occupational group.

**CONCLUSIONS:** Simple tests of age and gender distributions among subjects with and without available clinical information and molecular results may not rule out selection and information bias. Studies using biologic specimens are even more in need than classic studies to explain clearly the process followed to include and exclude subjects. Additional caution is needed when generalizing molecular results arising from incomplete biological specimens.

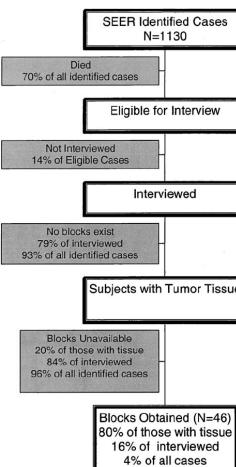
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## Potential for Selection Bias with in Molecular Epidemiology Studi

JANE A. HOPPIN, ScD, PAIGE E. TOLBI

JANE C. SCHROEDER, DVM, PhD, AND

Ann Epidemiol 2002;12:1-6.



Tumor characteristic	Potential biopsy tissue <sup>a</sup>				<i>P</i> -value <sup>b</sup>	
	Yes N = 142		No N = 988			
	Mean	sd	Mean	sd		
Tumor size <sup>c</sup> (mm)	179	343	570	474	<0.0001	
Histologic grade <sup>d</sup>						
G1 (lowest grade)	16	11	49	5	0.001	
G2	59	42	107	11		
G3	41	29	186	19		
G4 (highest grade)	0	0	17	2		
Unknown	26	18	629	64		
Summary stage					0.001	
In situ	2	1	0	0		
Localized	22	15	59	6		
Regional, extension only	49	35	222	23		
Regional, nodes only	7	5	12	1		
Regional, extension and nodes	45	32	58	6		
Remote	13	9	498	51		
Unstaged	4	3	131	13		

## Reporting Participation in Case-Control Studies

Sara H. Olson,<sup>1</sup> Lynda F. Voigt,<sup>2</sup> Colin B. Begg,<sup>1</sup> and Noel S. Weiss<sup>2,3</sup>

EPIDEMIOLOGY 2002;13:123–126

TABLE 4. Reporting Outcomes of Recruiting Respondents in Case-Control Studies; Example 3: Controls Contacted by Telephone from List Obtained from the Health Care Financing Administration

Units Selected from Sampling Frame	No.
Total	128
Unable to determine eligibility, total	8
Respondent not eligible	
Total	63
Prior hysterectomy	63
Respondent screened and eligible	
Total	57
Not interviewed (refused)	8
Interviewed	49

Data from Hill *et al.*<sup>10</sup>

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## Reporting Participation in Case-Control Studies

Sara H. Olson,<sup>1</sup> Lynda F. Voigt,<sup>2</sup> Colin B. Begg,<sup>1</sup> and Noel S. Weiss<sup>2,3</sup>

EPIDEMIOLOGY 2002;13:123–126

TABLE 2. Reporting Outcomes of Recruiting Respondents in Case-Control Studies; Example 1: Controls Contacted by Telephone Using Random-Digit Dialing

Units Selected from Sampling Frame	No.
Total	6,741
Ineligible sampling unit	
Total	3,589
Business, fax, government	1,937
Nonworking numbers	1,436
Institution, group quarters, dateline	216
Unable to determine eligibility	
Total	431
Unknown if residential	274
Residential, unknown if individual eligible	157
Answering machine on all attempts	56
Refusal to answer questions on eligibility	76
Other (language barrier)	25
Respondent not eligible	
Total	1,983
Age	1,749
County	216
Language	18
Respondent screened and eligible, total	738

Data from Rossing *et al.*<sup>8</sup>

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European Journal of Epidemiology 16: 533–541, 2000.

## Validity of the hospital discharge diagnosis in epidemiologic studies of biliopancreatic pathology

M. Porta<sup>1</sup>, S. Costafreda<sup>1</sup>, N. Malats<sup>1</sup>, L. Guarner<sup>2</sup>, M. Soler<sup>1</sup>, J.M. Gubern<sup>3</sup>, E. García-Olivares<sup>4</sup>, M. Andreu<sup>3</sup>, A. Salas<sup>4</sup>, J.M. Coroninas<sup>3</sup>, J. Alguacil<sup>1</sup>, A. Carrato<sup>5</sup>, J. Rifà<sup>6</sup> & F.X. Real<sup>1</sup> for the PANKRAS II Study Group<sup>\*</sup><sup>1</sup>Institut Municipal d'Investigació Médica, Universitat Autònoma de Barcelona and Universitat Pompeu Fabra, Barcelona;<sup>2</sup>Hospital Vall d'Hebron, Barcelona; <sup>3</sup>Hospital del Mar, Barcelona; <sup>4</sup>Hospital de la Mútua de Terrassa, Terrassa, Barcelona;<sup>5</sup>Hospital General de Elche, Alicante; <sup>6</sup>Hospital Son Dureta, Mallorca, Spain

**Table 2.** Changes between the initial hospital discharge diagnosis and the main clinicopathological diagnosis

Initial hospital discharge diagnosis	Main clinicopathological diagnosis (post-review)								Total
	Exocrine pancreatic cancer	Extrahepatic biliary cancer	Acute pancreatitis	Chronic pancreatitis	Syndromic diagnoses	Digestive neoplasm of ill-defined sites	Malignant neoplasm, site unspecified	Other benign diseases	
Exocrine pancreatic cancer	176 (86.3)	8 (3.9)	3 (1.5)	1 (0.5)	4 (2.0)	7 (3.4)	5 (2.5)		204
Extrahepatic biliary cancer	4 (3.1)	114 (88.4)				3 (2.3)	5 (3.9)	3 (2.3)	129
Acute pancreatitis			20 (54.1)	13 (35.1)	1 (2.7)			3 (8.1)	37
Chronic pancreatitis	4 (4.4)	1 (1.1)	2 (2.2)	79 (86.8)	2 (2.2)			1 (1.1)	89
Pancreatic cyst and pseudocyst			13 (34.2)	19 (50.0)			1 (2.6)	5 (13.2)	38
Other specified diseases of pancreas		1 (33.3)	2 (66.7)						3
Syndromic diagnoses				4 (44.4)			1 (11.1)	4 (44.4)	9
Digestive neoplasm of ill-defined sites					1 (50.0)		1 (50.0)		2
Malignant neoplasm, site unspecified						26 (83.9)	2 (6.5)	31	
Other benign diseases	1 (1.7)	2 (3.4)	1 (1.7)	3 (5.2)	2 (3.4)		1 (1.7)	48 (82.8)	58
Total	185	128	40	117	16	13	34	67	600

Figures are number of patients; figures within brackets are the row percent (i.e. the positive predictive value of the hospital discharge diagnosis), taking the main clinicopathological diagnosis as the gold standard. Cells with no patients are left blank.

**Table 3.** Potential predictors of diagnostic change

	Exocrine pancreatic cancer			
	N	n	(%)	p
<b>Diagnostic basis</b>				
Clinical investigation <sup>e</sup>	24	9	(37.5)	
Laparotomy	15	5	(33.3)	
Tumour markers	18	3	(16.7)	
Cytology	66	5	(7.6)	
Histology (primary)	57	5	(8.8)	0.001 <sup>c</sup>
Histology (metastasis)	24	1	(4.2)	
<b>Cyto-histological confirmation</b>				
No	57	17	(29.8)	
Yes	147	11	(7.5)	< 0.001 <sup>a</sup>
Histology <sup>f</sup>	81	6	(7.4)	1.000 <sup>c</sup>
Cytology	66	5	(7.6)	
Missing <sup>d</sup>	9	8	(88.9)	

N, number of subjects with original discharge diagnosis; n, number whose original diagnosis changed; % = (n/N) \* 100

<sup>e</sup> Histology obtained from a primary lesion, a metastasis or in the necropsy.

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**Table 3.** Potential predictors of diagnostic change

	Exocrine pancreatic cancer			
	N	n	(%)	p
Global	204	28	(13.7)	
Gender				
Men	122	18	(14.8)	
Women	82	10	(12.2)	0.603 <sup>a</sup>
Age (tertiles)				
1st	68	5	(7.4)	
2nd	68	11	(16.2)	
3rd	68	12	(17.6)	0.082 <sup>b</sup>
Tumour stage				
Local	43	7	(16.3)	
Regional	55	6	(10.9)	
Distant metastases	97	7	(7.2)	0.103 <sup>b</sup>
Missing <sup>d</sup>	9	8	(88.9)	

N, number of subjects with original discharge diagnosis; n, number whose original diagnosis changed; % = (n/N) \* 100

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**Table 4.** Diagnostic procedures and diagnostic change stratified by presence or absence of cyto-histological confirmation

	Exocrine pancreatic cancer					Exocrine pancreatic cancer			
	N	n	(%)	p	N	n	(%)	p	
<b>Cyto-histologically confirmed<sup>a</sup></b>									
CT	114	8	(7.0)		39	9	(23.1)		
Yes	33	3	(9.1)	0.711 <sup>b</sup>	No	18	8	(44.4)	
ERCP	53	6	(11.3)		33	12	(36.4)		
Yes	94	5	(5.3)	0.205 <sup>b</sup>	No	24	5	(20.8)	
Laparotomy	87	3	(3.4)		22	6	(27.3)		
Yes	60	8	(13.3)	0.051 <sup>b</sup>	No	35	11	(31.4)	
<b>Non cyto-histologically confirmed</b>									
CT	144	8	(7.0)		39	9	(23.1)		
Yes	33	3	(9.1)	0.101 <sup>b</sup>	No	18	8	(44.4)	
ERCP	53	6	(11.3)		33	12	(36.4)		
Yes	94	5	(5.3)	0.206 <sup>c</sup>	No	24	5	(20.8)	
Laparotomy	87	3	(3.4)		22	6	(27.3)		
Yes	60	8	(13.3)	0.051 <sup>b</sup>	No	35	11	(31.4)	

<sup>a</sup> Histology obtained from a primary lesion, a metastasis or in the necropsy.

**The characteristics of the disease  
and the state of knowledge /  
ignorance  
justify (and give meaning to)  
innovative research approaches.**

simple but tough things...  
difficult to achieve, e.g.,  
collaboration + involvement  
of clinical colleagues.

Attention to  
selection biases !

- J Clinical Epidemiology 1998, 2008
- The Lancet 1999
- Digestive Diseases & Sciences 1999, 2007
- Annals of Occupational Hygiene 2000
- International J of Epidemiology 2000
- Occupational & Environmental Med 2000
- European J of Epidemiology 2000, 2006-07
- Gut 2001, 2002
- J Epi Comm Health 1999, 2000, 2002, 2007
- Epidemiology 2001

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## Estudio PANKRAS II

Diseñado en 1990-91, 5 centros,  
prospectivo, unos de sus objetivos  
primarios es valorar interacciones  
entre alteraciones genéticas (K-ras  
y otras) y factores ambientales.

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- Carcinogenesis 2002
- Organohalogen Compounds 2002
- Molecular Carcinogenesis 2003
- International Journal of Cancer 2003
- Clinical & Translational Oncology 2005
- Pancreas 2007
- Environment International 2007
- Gaceta Sanitaria 2002, 2005-2008.
- Int J Occupational & Environ Health 2003



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Table 2

*K-ras mutations and regular coffee consumption among cases of cancer of the pancreas*

Regular coffee drinker	Yes	No	Total
K-ras mutated	73	10	83
K-ras wild type	16	8	24
Total	89	18	107

Source: Porta et al. (1999).

Paul R. Rosenbaum  
 The Case-Only Odds Ratio as a Causal Parameter  
 BIOMETRICS 60, 233–240  
 March 2004

**“In a case-only design (...) the odds ratio in this table is taken as a measure of gene-environment interaction”**

## Indicators

- Time symptom onset → sample extraction
- Site of sample extraction
- Symptoms and clinical status at sample extr.
- Lipid levels                      • Other laboratory findings
- Tumor stage                      • Diagnostic procedures
- Treatment                        • Clinical complications
- Survival                         • Other factors

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- Elucidating the etiologic significance of each OC requires an understanding of its relationships with other OCs, with lipids and with prediagnostic signs & symptoms.

**TABLE 1.** Operational Elements of a Framework to Assess the Etiologic Significance of Associations Between Internal Concentrations of Organochlorine Compounds and Cancer Risk (or Related Biologic Events)

1. Consider which potential indicators may be relevant within the context of knowledge (see 4 below) relevant for your study:
  - Time between symptom onset and sample extraction
  - Site of sample extraction
  - Symptoms and clinical status at sample extraction
  - Lipid levels
  - Other laboratory findings
  - Tumor stage
  - Diagnostic procedures
  - Treatment
  - Clinical complications
  - Survival
  - Other factors

Epidemiology March 2001, Vol. 12 No. 2

**Timing of blood extraction in epidemiologic and proteomic studies: results and proposals from the PANKRAS II Study**

Miquel Porta · José Pumarega · Olga Ferrer  
European Journal of Epidemiology 2007

There are no consensus guidelines or standards for epidemiologic and '-omics' studies using blood biomarkers on how to report the timing of extraction of blood samples. However, disease-induced changes in blood concentrations of exogenous and endogenous compounds may bias studies.

Results suggest ways to report intervals involving blood biomarkers and may contribute to develop consensus guidelines and standards on the collection of blood samples in epidemiologic and '-omics' research.

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2. Based on your study hypotheses and design, select indicators and collect information.
3. Conduct quantitative analyses:
  - Do the indicators vary among the groups of subjects under comparison?
  - Do associations exist between the indicators and the concentrations of the organochlorine compounds?
  - Do the indicators induce effect modification or confounding of the causal estimators of the relation between organochlorine concentrations and your clinical or biological end-point?
4. Assess to what extent results are coherent with and contribute to the existing knowledge on issues such as:
  - Half-life and physico-chemical characteristics of the compound.
  - Influence of the specific cancer on the metabolism of the compound.
  - Biodegradation, bioaccumulation, and interactions.
  - Timing and dose required to influence the biological or clinical end-point.
5. Develop and test quantitative standards for the relevant indicators.

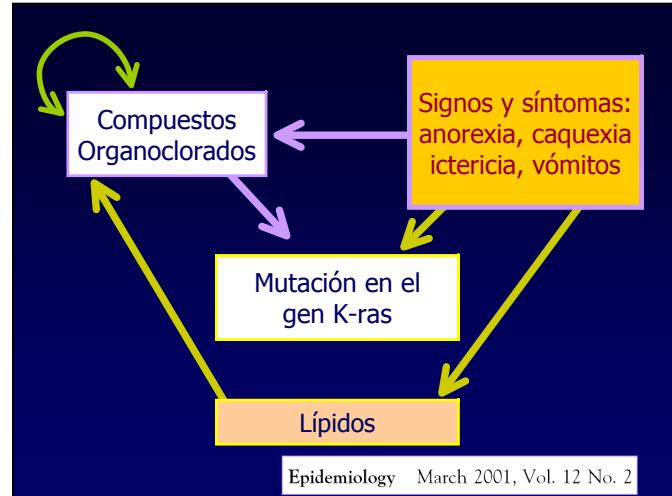
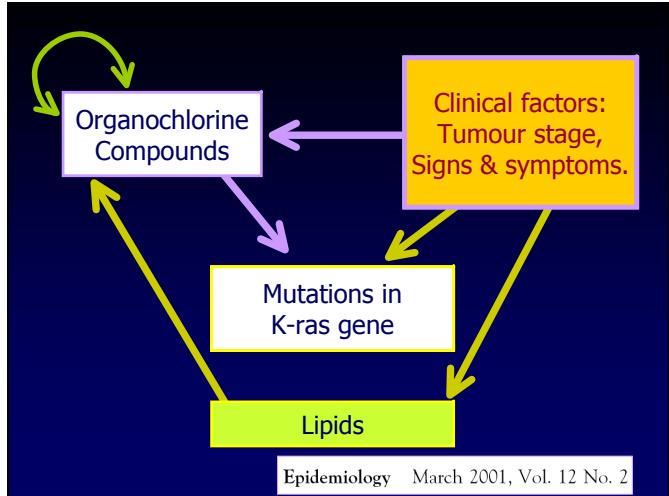
Epidemiology March 2001, Vol. 12 No. 2

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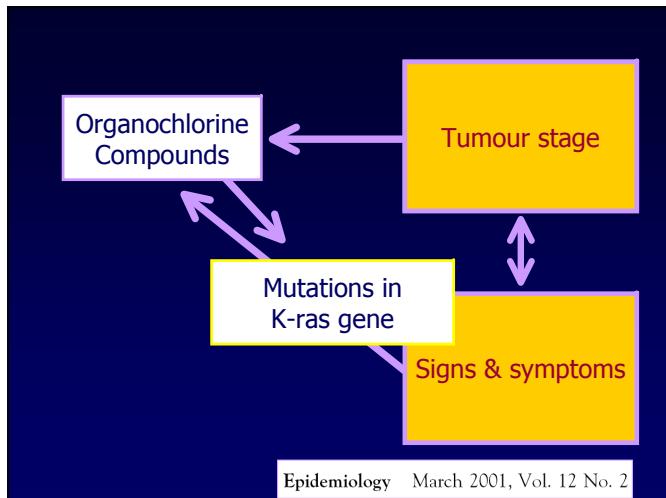
**Journal of Clinical Epidemiology 61 (2008)**

In exocrine pancreatic cancer clinical factors  
 were related to timing of blood extraction  
 and influenced serum concentrations of lipids

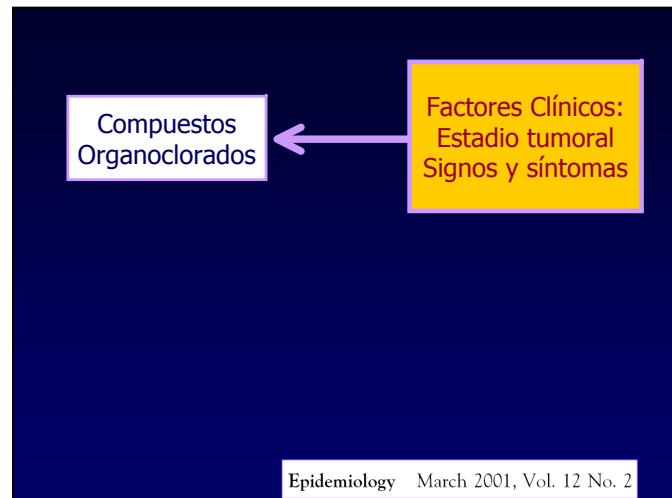
Miquel Porta<sup>a-c\*</sup>, Olga Ferrer-Armengou<sup>a</sup>, José Pumarega<sup>a</sup>, Tomàs López<sup>a</sup>

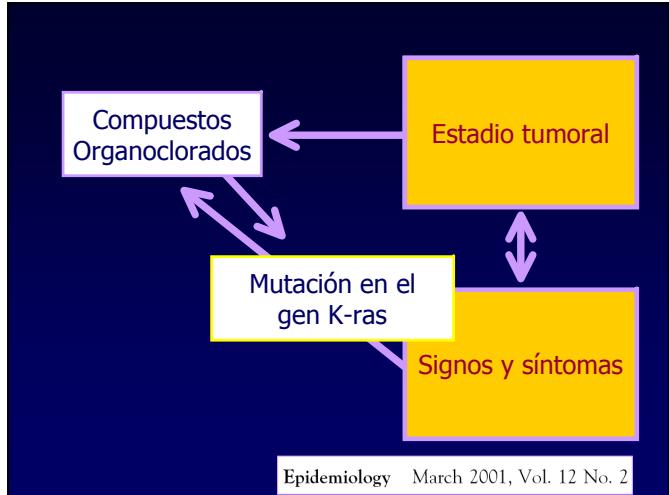


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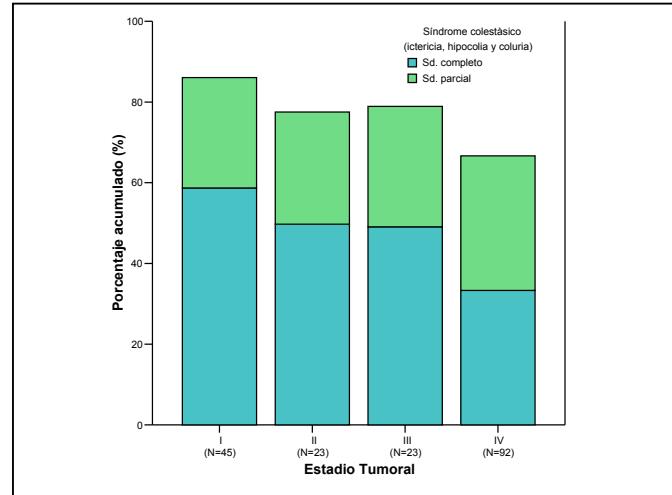


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**clinical epidemiology ↔ proteomics**

*Clin Transl Oncol. 2005;7(5):195-204*

**Exocrine pancreatic cancer: symptoms at presentation and their relation to tumour site and stage**

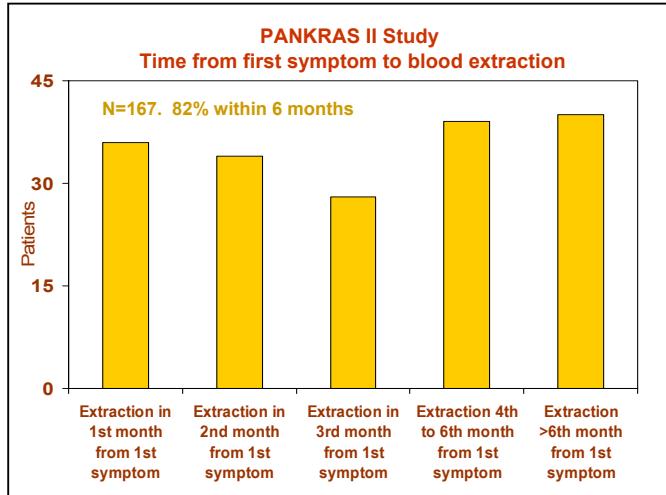
Miquel Porta<sup>a,b</sup>, Xavier Fabregat<sup>b,c</sup>, Núria Malats<sup>a</sup>, Luisa Guarner<sup>d</sup>, Alfredo Carrato<sup>c</sup>, Ana de Miguel<sup>c</sup>,

There was a clear trend towards more localized tumours with increasing numbers of cholestatic signs ( $p<0.001$ ).

**Indicators**

- Time from symptom onset to sample extraction (TSOSE)
  - The lower, the better...

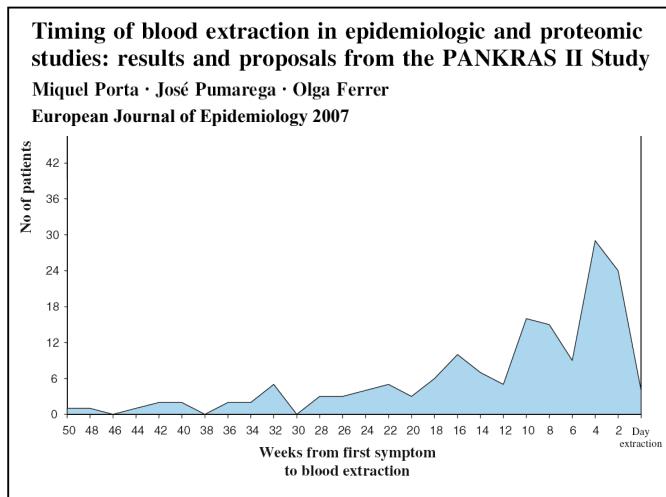
but TSOSE depends on patient, referral system, hospital admission, diagnosis, study inclusion...



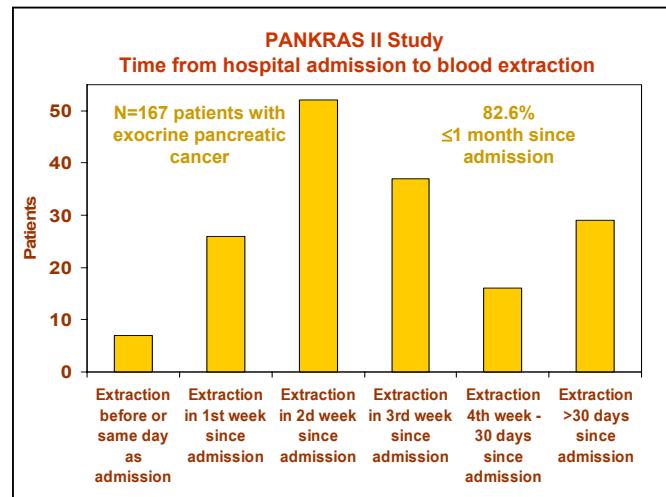
## Indicators

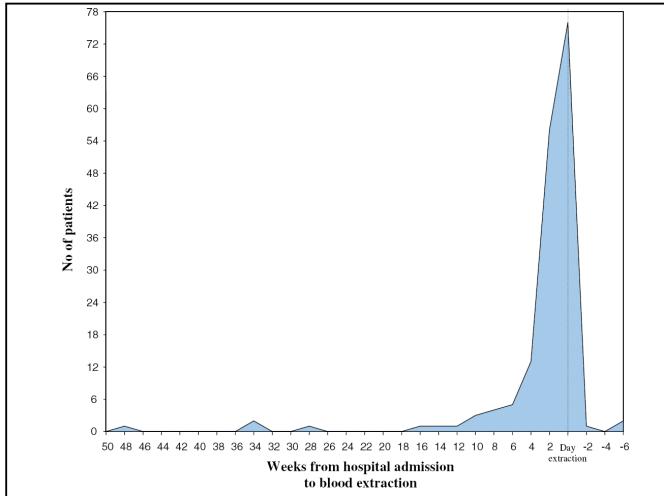
- Time from
- symptom onset
- hospital admission
- diagnosis
- study inclusion
- treatment onset...
- to sample extraction.

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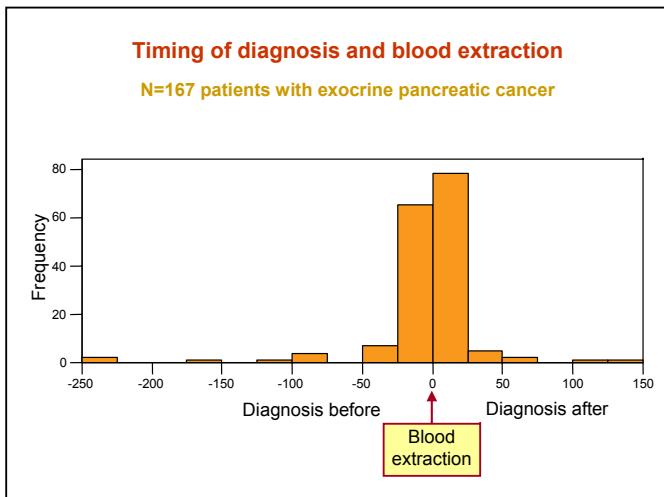


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## Indicators

- Time from symptom onset to sample extraction (TSOSE)
  - Is TSOSE + related to OCs levels? if so...
  - Is TSOSE related to the outcome? if so...

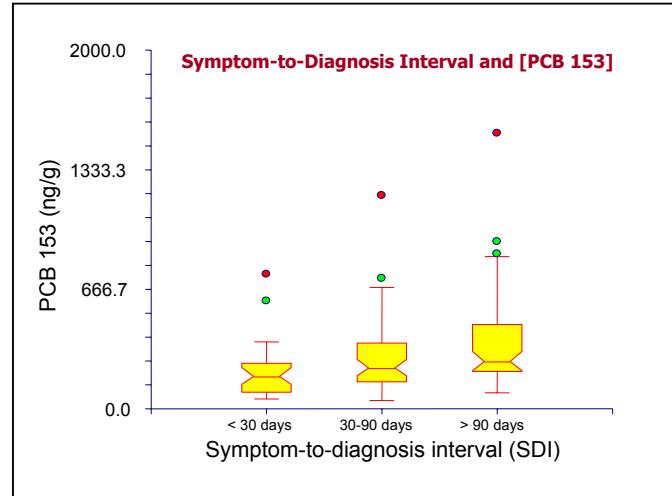
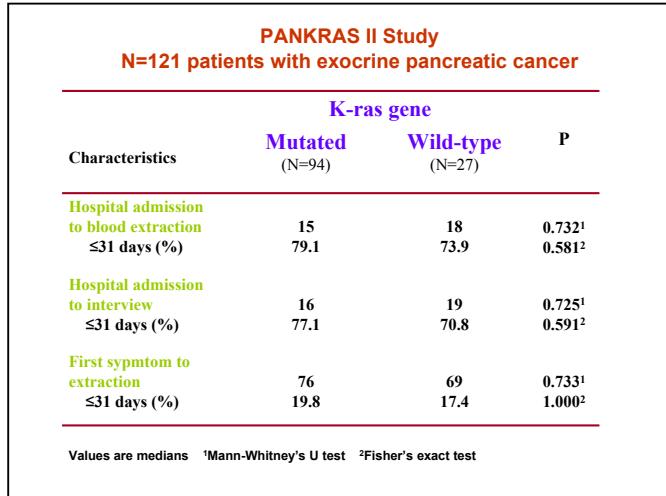
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### PANKRAS II Study

N=121 patients with exocrine pancreatic cancer

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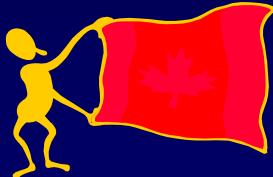
Characteristics	K-ras gene		P
	Mutated (N=94)	Wild-type (N=27)	
<hr/>			
First symptom to extraction			
≤31 days (%)	76 19.8	69 17.4	0.733 <sup>1</sup> 1.000 <sup>2</sup>
<hr/>			
Values are medians	1Mann-Whitney's U test	2Fisher's exact test	



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## Indicators

- Time from symptom onset to diagnosis (SDI) (duration of symptoms, 'delay')
  - Is SDI + related to OCs levels? if so...



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PART 2 – 32

## Indicators

- Symptoms and clinical status at time of sample extraction.
  - Malnourishment, weight loss / changes in BMI, anorexia-cachexia, jaundice...
  - Are symptoms associated with biological outcome...? if so...



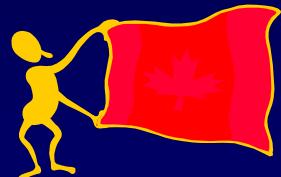
K-ras gene			
Signs & symptoms	Mutated (N=94)	Wild-type (N=27)	P <sup>1</sup>
<b>Constitutional syndrome</b>	<b>70.2</b>	<b>59.3</b>	<b>0.469</b>
Asthenia	86.0	85.2	1.000
Anorexia	81.7	77.8	0.781
Weight loss	86.0	81.5	0.550
Cachexia	12.9	7.4	0.734
<b>Cholestatic syndrome</b>	<b>38.3</b>	<b>40.7</b>	<b>0.751</b>
Jaundice	51.6	44.4	0.662
Choluria	52.7	55.6	0.830
Hypocholia	50.5	44.4	0.664
<b>Vomiting</b>	<b>31.2</b>	<b>37.0</b>	<b>0.642</b>

<sup>1</sup>Fisher's exact test

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## Indicators

- Symptoms and clinical status at time of sample extraction.
  - Malnourishment, weight loss / changes in BMI, anorexia-cachexia, jaundice...
  - Are symptoms associated with OCs levels...? if so...



Relationship between cholestatic syndrome and serum concentrations of p,p'-DDE				
	Cholestatic syndrome			
	Absent (N=56)	Partial (N=27)	Complete (N=61)	P
p,p'-DDE (median)	4.067,70	3.479,24	2.405,69	0,037 <sup>a</sup> 0,018 <sup>b</sup>

OCs levels lipid adjusted (ng/g).

N: Number of patients.

P: Statistical significance.

<sup>a</sup>: Kruskal-Wallis' test.

<sup>b</sup>: Spearman's correlation between OC levels and number of cholestatic symptoms.

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## Relationship between complete cholestatic syndrome and serum concentration of 4 organochlorine compounds

	B	95% CI	P
p,p'-DDE	-0,533	-0,948 -0,118	0,012
PCB 153	-0,372	-0,634 -0,110	0,006
PCB 180	-0,377	-0,653 -0,102	0,009 <sup>a</sup>
$\beta$ -HCH	-0,340	-0,631 -0,049	0,024 <sup>a</sup>

Linear regression adjusted by age, sex and tumour stage.

OCs lipid-corrected and log transformed.

Reference category: total absence of syndrome.

B: Regression coefficient (Beta).

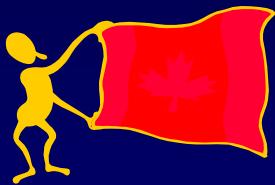
95% CI : Beta coefficient 95% CI.

P: Statistical significance Beta coefficient.

<sup>a</sup> : Statistical significance conservative F.

## Indicators

- Stage at diagnosis
  - Is stage + related to the biological outcome / end-point? if so...



## Indicators

- Stage at diagnosis
  - Is stage + related to OCs levels? if so...



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K-ras gene			
Characteristics	Mutated (N=94)	Wild-type (N=27)	P
SDI mean	94.6	131.8	0.268 <sup>1</sup>
median	70.0	72.0	0.932 <sup>2</sup>
≤31 days (%)	20.2	22.2	0.793 <sup>3</sup>
Stage			0.256 <sup>3</sup>
I	18.5	22.7	
II	21.0	4.5	
III	13.6	9.1	
IV	46.9	63.6	

<sup>1</sup>Student's t-test <sup>2</sup>Mann-Whitney's U test <sup>3</sup>Fisher's exact test

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### Relationship between age, tumour stage and [organochlorine compounds]

	B	95%CI of B	P	Model parametres
p,p'-DDT				
→ Age	0.025	0.006 0.043	0.009	R <sup>2</sup> : 0.074
→ Tumour stage II vs. I	0.065	-0.729 0.859	0.872	F: 2.723
→ Tumour stage III vs. I	-0.661	-1.448 0.126	→ 0.099	P <sup>b</sup> : 0.032
→ Tumour stage IV vs. I	-0.016	-0.595 0.563	0.956	Levene: 0.209
HCB				
→ Age	0.025	0.015 0.034	<0.001	R <sup>2</sup> : 0.193
→ Tumour stage II vs. I	-0.048	-0.472 0.376	0.823	F: 8.168
→ Tumour stage III vs. I	-0.537	-0.958 -0.117	0.013	P <sup>b</sup> : <0.001
→ Tumour stage IV vs. I	-0.201	-0.510 0.109	0.202	Levene: 0.110
β-HCH				
→ Age	0.025	0.015 0.034	<0.001	R <sup>2</sup> : 0.194
→ Tumour stage II vs. I	-0.024	-0.450 0.403	0.913	F: 8.265
→ Tumour stage III vs. I	-0.566	-0.988 -0.143	0.009	P <sup>b</sup> : <0.001
→ Tumour stage IV vs. I	-0.257	-0.568 0.054	0.104	Levene: 0.111

General linear model.

OCs lipid-corrected and log-transformed.

B: Regression coefficient (Beta)

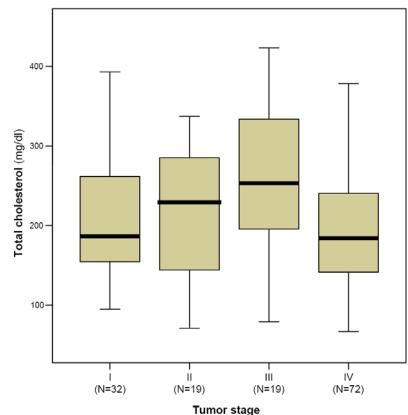
P: Statistical significance Beta coefficient.

R<sup>2</sup>: Determination coefficient of the global model.

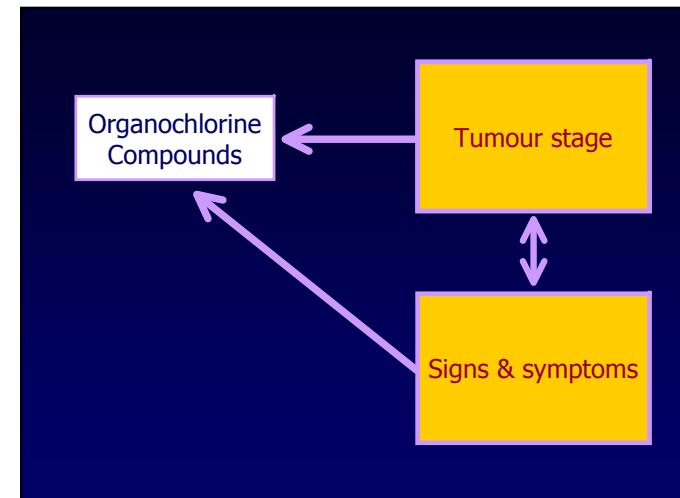
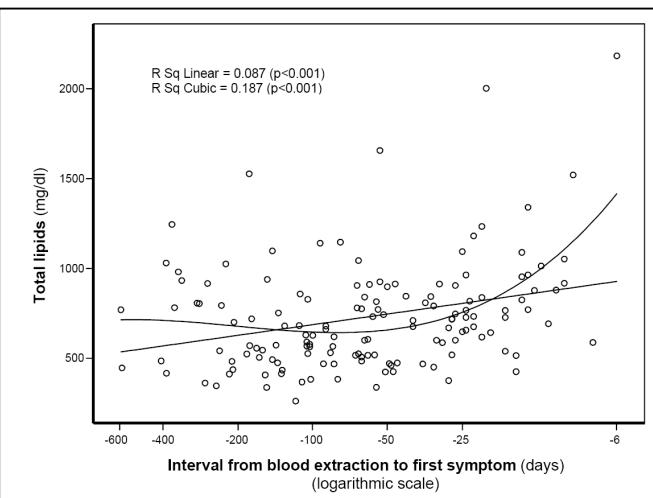
F: Fisher-Snedecor's F (degrees of freedom v<sub>1</sub>=4, v<sub>2</sub>=137).

P<sup>b</sup>: Statistical significance degree of the model.

Levene: Levene's test statistical significance variance homogeneity.



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### Symptoms do influence tumour stage

Sign or symptom, or SDI	Tumour stage			P
	I N=45 (%)	II-III N=46 (%)	IV N=92 (%)	
Any pain	71,1	80,4	89,0	0,034
No cholestatic symptoms	13,3	28,3	48,9	<0,001
One or two symptoms	15,6	19,6	23,9	<0,001 <sup>a</sup>
All three symptoms	71,1	52,2	27,2	
Constitutional syndrome	91,1	93,5	95,6	0,558
SDI (median, in months)	1,5	2,3	2,6	0,048 <sup>b</sup>

N: Numbers of patients.

P: Fisher's exact test.

<sup>a</sup> Spearman's correlations between stage and numbers of cholestatic symptoms.

<sup>b</sup> Kruskal-Wallis' test.

**Constitutional syndrome, age, sex, and tumour stage  
and concentrations of p,p'-DDT**

	B	95% CI	P	Model parameters	
p,p'-DDT					
Partial syndrome	0.901	-0.033	1.835	0.059	R <sup>2</sup> : 0.109
Complete syndrome	0.873	0.010	1.735	0.047	F: 2.348
Age (each year)	0.021	0.002	0.040	0.034	P <sup>a</sup> : 0.027
Sex (male vs. female)	0.231	-0.266	0.728	0.360	Levene: 0.595
Tumour stage II vs. I	0.075	-0.719	0.869	0.852	
Tumour stage III vs. I	-0.598	-1.382	0.186	0.134	
Tumour stage IV vs. I	0.025	-0.555	0.604	0.933	

General lineal model.

OCs lipid-adjusted and log transformed.

B: Regression significance Beta.

P: Statistical significance for each parameter.

R<sup>2</sup>: Determination coefficient of the model.

F: Fisher-Snedecor's F with degree of freedom v<sub>1</sub>=7, v<sub>2</sub>=134.

P<sup>a</sup>: Statistical significance of the model.

Levene: Levene's test statistical significance variances homogeneity.

**Indicators**

- Samples drawn before or after some diagnostic tests, surgery, other treatments?
  - Are OCs levels before ≠ after? if so...



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**Indicators**

- Stage at diagnosis
  - Is stage + related to OCs levels? if so...
  - Are ORs homogeneous across stage strata?

**Indicators**

- Survival (more clinically aggressive tumors may both ↑ OCs levels and ↓ survival).
  - Is survival – related to OCs levels? if so...



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### Effects of cholestatic symptomatology on [OC]

Sign, symptom or syndrome	p,p'-DDT	p,p'-DDE	PCB 138	PCB 153	PCB 180	HCB	$\beta$ -HCH
Jaundice (symptom)				↓	↓		
Jaundice (sign)				↓	↓		
Hypocholia		↓		↓	↓		↓
Choluria	↓			↓	↓		↓
Pruritus	↓			↓	↓		↓
Partial cholestatic syndrome	↓		↓	↓	↓	↓	
Complete cholestatic syndrome	↓		↓	↓	↓		↓

General linear model, adjusted by age, sex and tumour stage.

Journal of Clinical Epidemiology 61 (2008)

### Factors influencing serum concentrations of total cholesterol.

Model*	$\beta$ †	95% CI	Partial p value‡	R²	Model p value
6. Interval from blood extraction to first symptom (IES); jaundice, weight loss, and number of invasive diagnostic tests performed before blood extraction (IDT)				0.285	<0.001
IES 40-120 days vs. IES<40 days	-47.6	-91.5, -3.7	0.034		
IES>120 days vs. IES<40 days	-57.9	-102.9, -12.8	0.012		
Jaundice (present vs. absent)	67.2	29.4, 105.0	0.001		
Weight loss (present vs. absent)	53.5	6.5, 100.4	0.026		
1 test vs. 0 tests	-61.6	-101.9, -21.2	0.003		
≥2 tests vs. 0 tests	-57.8	-107.5, -8.1	0.023		
Tumor stage II vs. stage I	75.0	16.9, 133.1	0.012		
Tumor stage III vs. stage I	83.9	28.4, 139.4	0.003		
Tumor stage IV vs. stage I	46.3	0.9, 91.8	0.046		

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### Effects of constitutional syndrome on [OC]

Sign, symptom or syndrome	p,p'-DDT	p,p'-DDE	PCB 138	PCB 153	PCB 180	HCB	$\beta$ -HCH
Asthenia							
Anorexia							
Loss of weight				↑	↑		
Cachexia						↑	
Partial toxic syndrome	↑						
Complete toxic syndrome	↑						

General linear model, adjusted by age, sex and tumour stage.

### The effect of PCB 153 on the probability of a mutated (vs. wild-type) tumour.

PCB 153	OR1	P-value (OR 95% CI)	OR2	P-value (OR 95% CI)
≤187 (ng/g)	<b>1.0</b>	<b>0.017*</b>	<b>1.0</b>	<b>0.012*</b>
188 – 313	<b>1.7</b>	<b>(0.6-5.0)</b>	<b>1.6</b>	<b>(0.5-4.8)</b>
>313	<b>5.2</b>	<b>(1.3-20.5)</b>	<b>6.0</b>	<b>(1.5-24.8)</b>
≤305 (ng/g)	<b>1.0</b>	<b>0.026</b>	<b>1.0</b>	<b>0.017</b>
>305	<b>4.4</b>	<b>(1.2-16.0)</b>	<b>5.1</b>	<b>(1.3-19.1)</b>

OR1: adjusted by age and sex

OR2: age + sex + cholestatic syndrome

\*Mantel's extension test for linear trend.

## clinical epidemiology ↔ proteomics

*Clin Transl Oncol.* 2005;7(5):195-204

### Exocrine pancreatic cancer: symptoms at presentation and their relation to tumour site and stage

Miquel Porta<sup>a,b</sup>, Xavier Fabregat<sup>b,c</sup>, Núria Malats<sup>a</sup>, Luisa Guarner<sup>d</sup>, Alfredo Carrato<sup>a</sup>, Ana de Miguel<sup>c</sup>,

There was a clear trend towards more localized tumours with increasing numbers of cholestatic signs ( $p<0.001$ ).

An increased symptom-to-diagnosis interval was associated with more advanced stage ( $p=0.048$ ).

## clinical epidemiology ↔ proteomics

*Clin Transl Oncol.* 2005;7(5):195-204

First, the analysis of signs and symptoms is one practical way to assess whether early manifestations of the disease were related to a specific pattern of proteins<sup>43,44</sup>.

And second, since test results may vary with the spectrum of disease/patients<sup>43-47</sup>, such information would aid to judge the properties of the proteomic test (e.g., its predictive values) in each stage of disease and patient spectrum.

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## clinical epidemiology ↔ proteomics

*Clin Transl Oncol.* 2005;7(5):195-204

**Conclusions.** Proper attention to signs and symptoms, especially cholestasis, may help identify patients with pancreatic cancer at an earlier stage. Results also provide a current picture of the semiology of pancreatic cancer which could be of use in studies on the potential of proteomic tests in the early detection of this neoplasm.

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*Clin Transl Oncol.* 2005;7(5):195-204

Although the study of cancer signs and symptoms might appear as just a traditional concern for clinicians, at the moment such analysis is also most relevant to appraise the biological and clinical significance of the emerging proteomic technologies.

In particular, proteomic pattern diagnostics are currently under investigation to improve early detection of different cancers<sup>40,42</sup>. This naturally includes the analysis of such patterns in the different stages of cancer<sup>40,43</sup>.

Yet, there is only scant information on the correlation between semiology, cancer stage and the ‘protein ion signatures’.

Such technologies may provide new insights into the biologic changes that occur in the different phases of tumourigenesis<sup>40-44</sup>. However, some aspects remain unclear; for instance, the discriminatory peaks identified by mass spectrometry and bioinformatic systems may represent molecules released into the circulation as a result of the disease, or reflect cancer epiphenomena (malnutrition, inflammation, immunodeficiency)<sup>44</sup>.

Proteomic patterns may also be particular to the spectrum of cases under study.

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Because thousands of proteins and protein fragments are analysed at once, bioinformatic analyses often must overlook the identity of the proteins, as well as the mechanisms by which proteins are released into the circulation, their concentration in biologic fluids, their metabolism, and their relationships within the human host<sup>44</sup>. In this context signs and symptoms constitute an important referent; for instance, a combined analysis of semiology and tumour stage may enable to assess in what spectrum of patients the proteomic test may detect disease early (i.e., to select patients for the test), and to judge what pathophysiological processes (weight loss and malnutrition, inflammation, metabolic changes associated with obstructive jaundice), as reflected in signs and symptoms, may be associated with the putatively discriminatory proteomic pattern<sup>45,44</sup>.

## clinical epidemiology ↔ proteomics



Journal of Clinical Epidemiology 56 (2003) 815-819

Journal of  
Clinical  
Epidemiology

Semiology, proteomics, and the early detection of symptomatic cancer

Miquel Porta<sup>a,b,c,\*</sup>, Esteve Fernandez<sup>d,e</sup>, Joan Alguacil<sup>f</sup>

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<sup>c</sup>School of Public Health, University of North Carolina at Chapel Hill, Chapel Hill, NC, USA

<sup>d</sup>Institut Català d’Oncologia, L’Hospitalet de Llobregat, Barcelona E-08907, Spain

<sup>e</sup>Universitat de Barcelona, Barcelona, Spain

<sup>f</sup>Division of Cancer Epidemiology & Genetics, National Cancer Institute, Bethesda, MD 20892, USA

the new wave

of genomic and proteomic analyses of early-stage cancers might provide new insights into changes that occur in early phases of tumorigenesis; it is already offering new candidate biomarkers for early-stage disease [37–39].

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Studies that profile proteomic patterns in body fluids should adhere to the methodologic standards that are usual in clinical epidemiology but less widely applied in basic science [40–42]. Rigorous pathophysiologic and clinical thinking should guide studies that pretend to correlate molecular abnormalities, symptoms, diagnostic performance, and clinical course—as exemplarily illustrated by *auxometric* measures of tumor growth and other works of Alvan Feinstein [8–17,43]. Whether the degree of correlation between proteomic patterns and classic semiology will be strong or weak remains to be seen (“weak” is our informed guess). Such analysis is of interest in itself. The respective relations of proteomic and semiologic patterns with cancer prognosis should be assessed in properly designed clinicoepidemiologic studies.

#### **BOX 5 Ethical, legal and social implications (ELSI).**



Today's genomics research and applications rest on more than a decade of valuable investigation into their ethical, legal and social implications. As the application of genomics to health increases along with its social impact, it becomes ever more important to expand on this work. There is an increasing need for focused ELSI research that directly informs policies and practices. One can envisage a flowering of 'translational ELSI' research that builds on the knowledge gained from prior and forthcoming 'basic ELSI research', which would provide knowledge for direct use by researchers, clinicians, policy-makers and the public. Examples include:

◆ The development of models of genomics research that use attention to these ELSI issues

for enhancing the research, rather than viewing such issues as impediments

- ◆ The continued development of appropriate and effective genomics research methods and policies that promote the highest levels of science and of protecting human subjects
- ◆ The establishment of crosscutting tools, analogous to the publicly accessible genomic maps and sequence databases that have accelerated other genomics research (examples of such tools might include searchable databases of genomic legislation and policies from around the world, or studies of ELSI aspects of introducing clinical genetic tests)
- ◆ The evaluation of new genetic and genomic tests and technologies, and effective oversight of their implementation, to ensure that only those with confirmed clinical validity are used for patient care

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