

Scientific session of the
Institut d'Investigació Sanitària de Palma (IdISPa)

The search for Biomarkers in coronary artery disease

Jaume Marrugat

Research Group on Cardiovascular Epidemiology & Genetics,
IMIM, Barcelona, Spain.

Spanish Cardiovascular Research Program HERACLES (RIC-RETICS-ISCIII)



PROGRAMA HERACLES
Red de Investigación
Cardiovascular

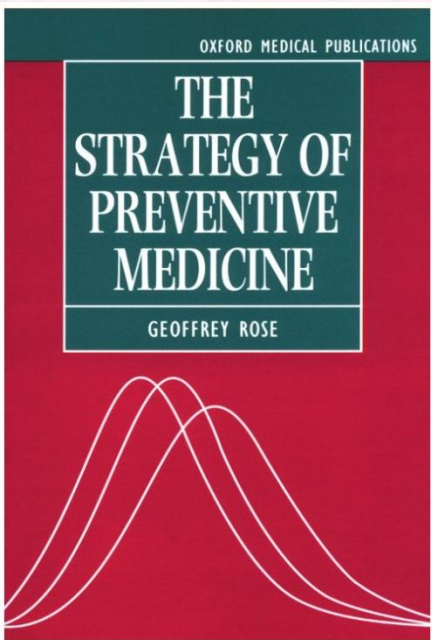
The problem

- Each year cardiovascular disease (CVD) causes over **4.3 million deaths** in Europe.
- CVD causes nearly **half of all deaths in Europe (48%)**.
- CVD is the main cause of the **disease burden** in Europe (**23%** of all the disease burden).
- Overall CVD is estimated to cost the EU economy **€192 billion a year**.

Justification of primary prevention of coronary heart disease

- Greatest cause of death in developed countries. In ~35% of cases its onset symptom is sudden death.
- Most cases are related to lifestyle & other modifiable factors, whose improvement results in reduced CHD incidence.

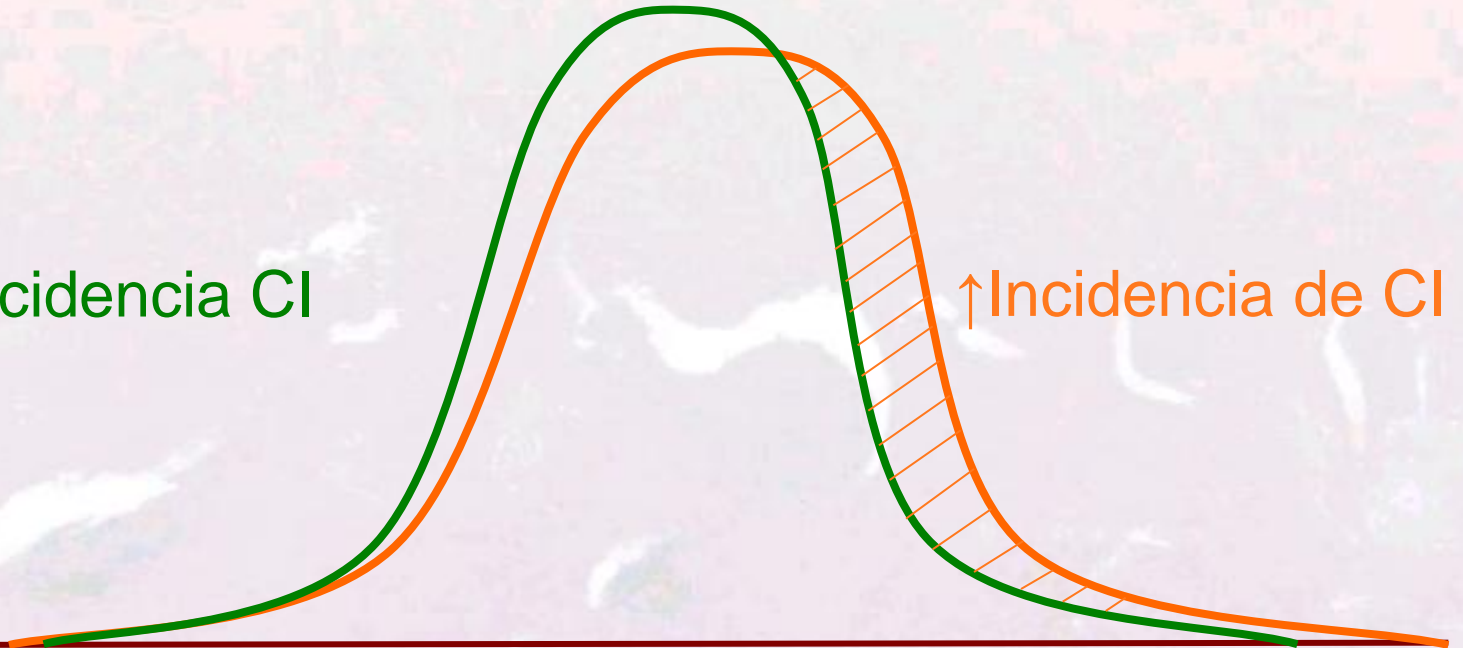
Modificación de la distribución poblacional de un factor de riesgo e incidencia de cardiopatía isquémica: el ejemplo del colesterol



↓ Incidencia CI

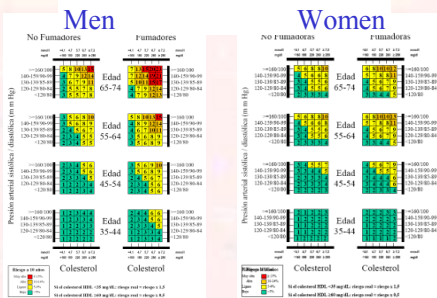
↑ Incidencia de CI

200 220
mg/dL



Disease prevention

Screening

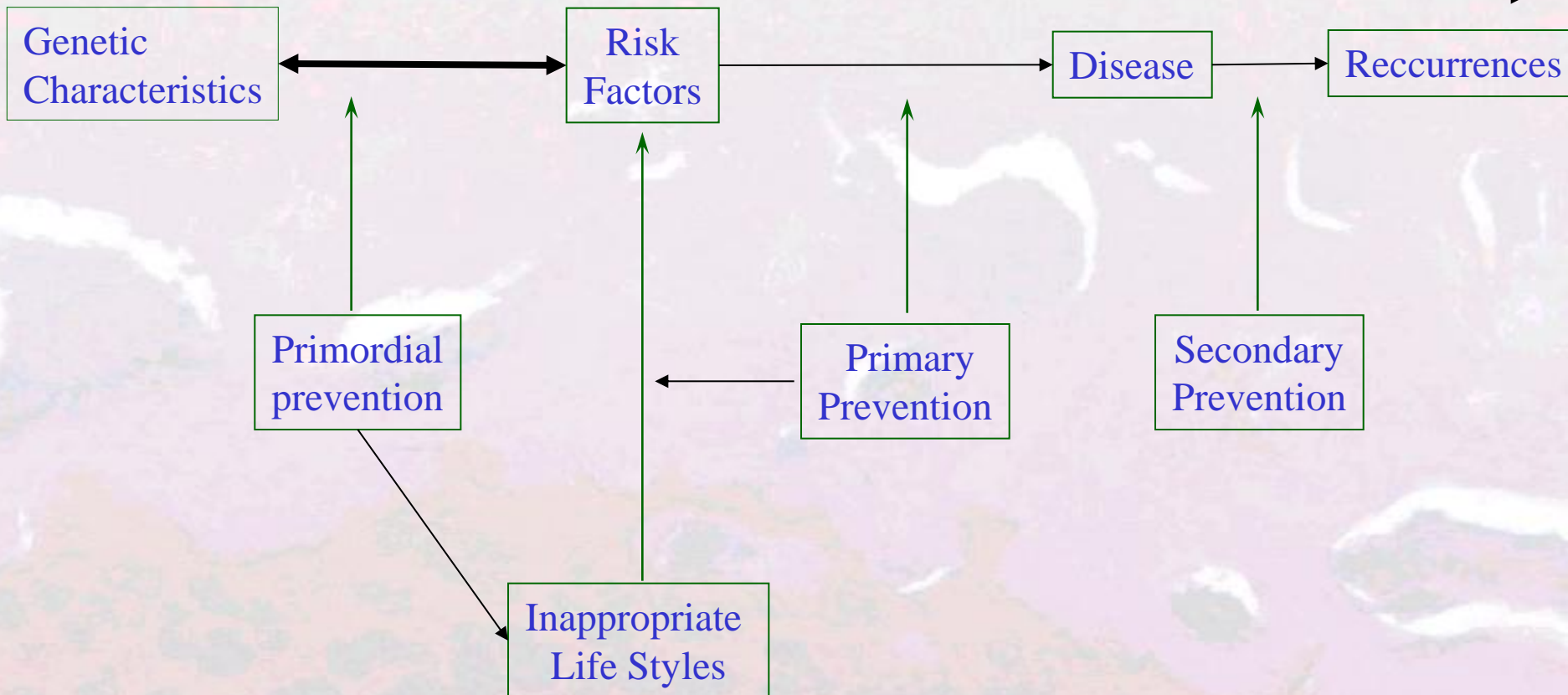


POPULATION

Population individuals

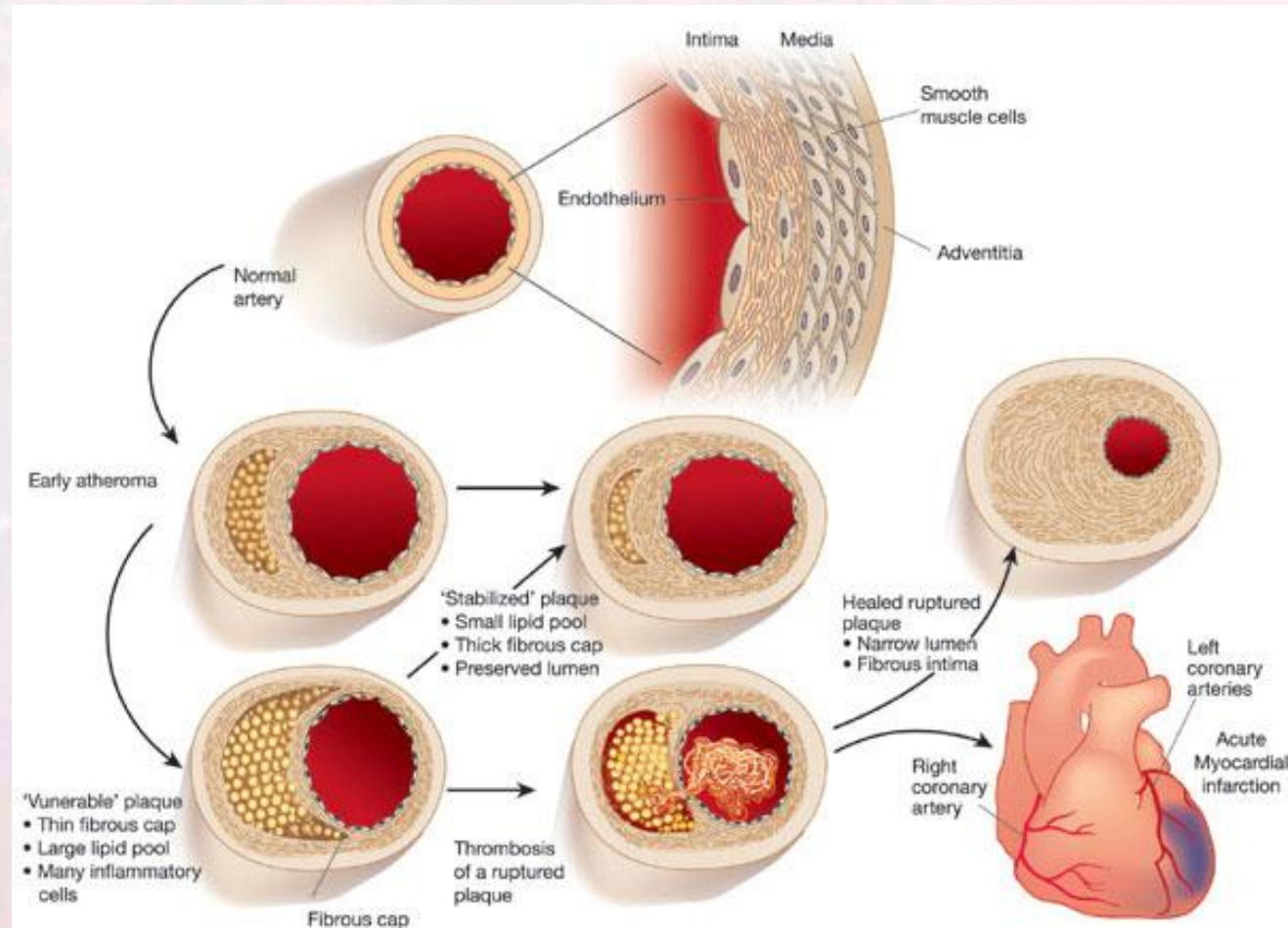
Patients

Patients



Reto para el futuro

- Identificación pacientes con placas vulnerables (estenóticas o no) susceptibles de rotura



Potential use of biomarkers

- Diagnostic tests
 - To clarify etiology of symptoms
 - During acute phase of disease
- To assess risk
 - Risk of developing symptomatic disease: Primary prevention (10 years)
 - Prognosis of established diseases (days, months)

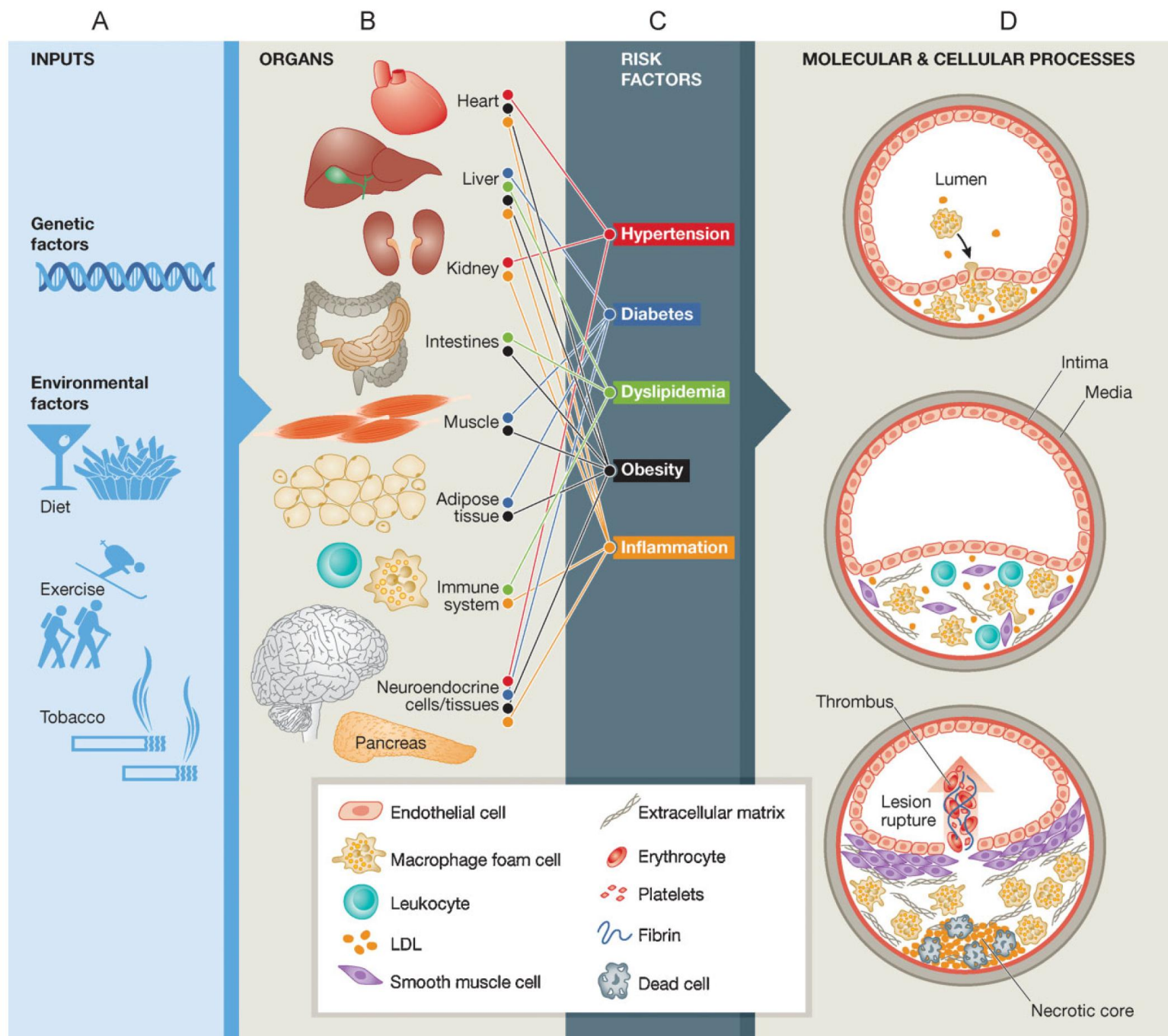
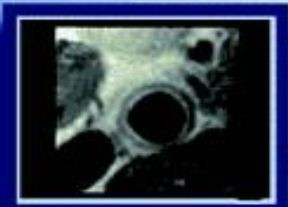


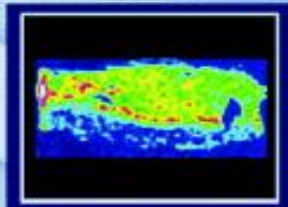
Figure 1. The pathophysiology of atherosclerosis involves interacting systems at multiple levels.

Types of biomarkers

- Life styles & cardiovascular risk factros
- Imaging
- Molecular
 - Biochemical
 - Genetic



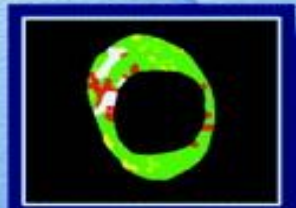
MRI



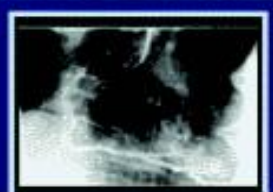
NIR



CT



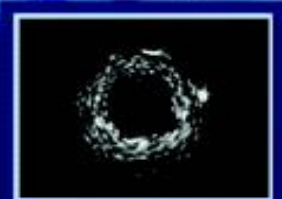
VIRTUAL HISTOLOGY



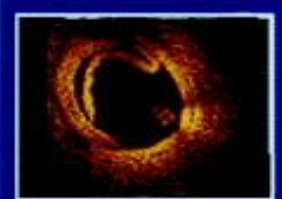
ANGIOGRAPHY



THERMOGRAPHY



IVUS



OCT

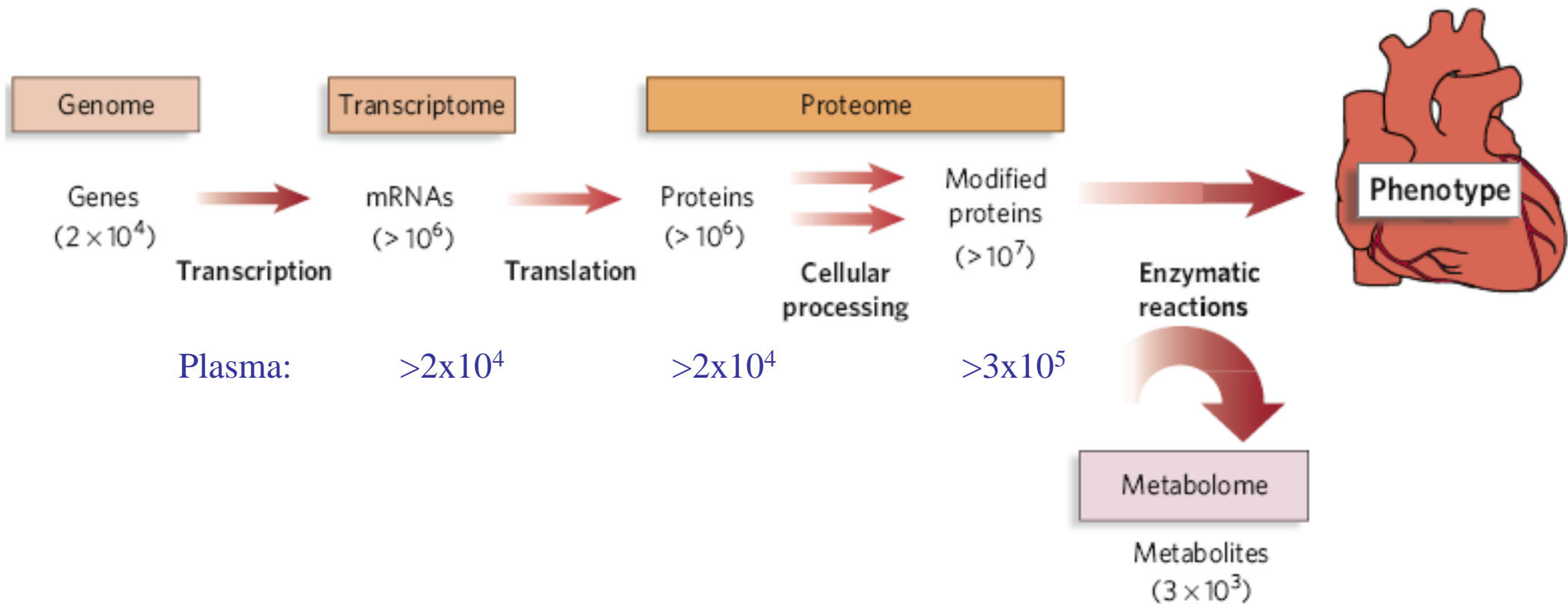


ANGIOSCOPY



Gopu

Molecular Biomarkers



Potential biomarkers in CV diseases

The 22 more abundant proteins, including albumin & immunoglobulins, constitute 99% of the plasmatic proteome mass.

Many biologically interesting molecules circulate in very small concentrations:

- Troponin: **nanomolar** (10^{-6}) concentration
- Insulin: **picomolar** (10^{-9}) concentration
- TNF: **femtomolar** (10^{-12}) concentration

Anderson, N. L. et al. The human plasma proteome: a nonredundant list developed by combination of four separate sources. Mol. Cell. Proteomics 3, 311–326 (2004).

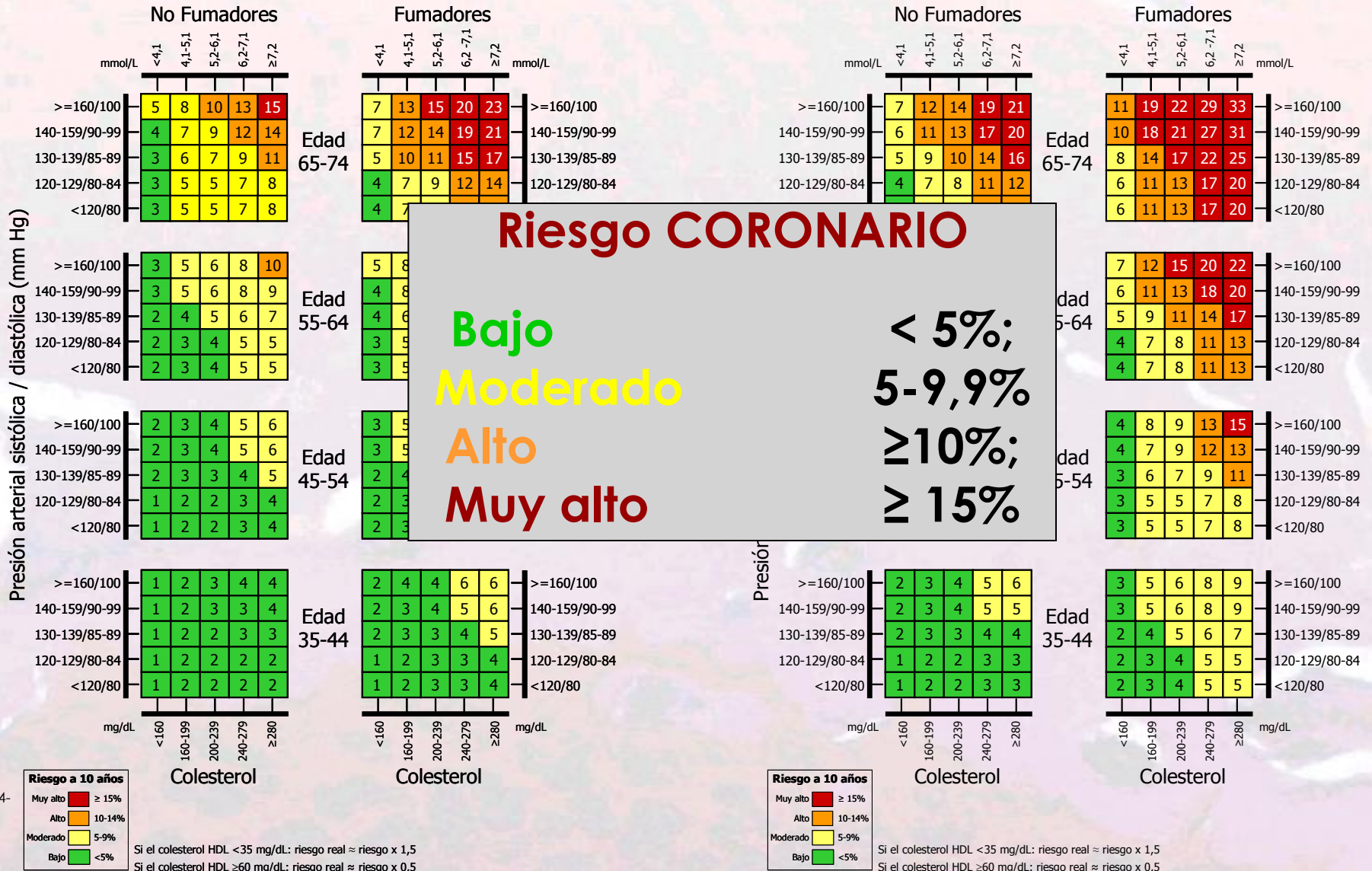
Cardiovascular risk functions

- Mathematical equations to estimate the probability of developing the disease in the future.
- This estimation is based on:
 - exposure to the different risk factors included in the function;
 - the effect size of the association between each risk factor and the disease;
 - the incidence rate of the disease in the population.

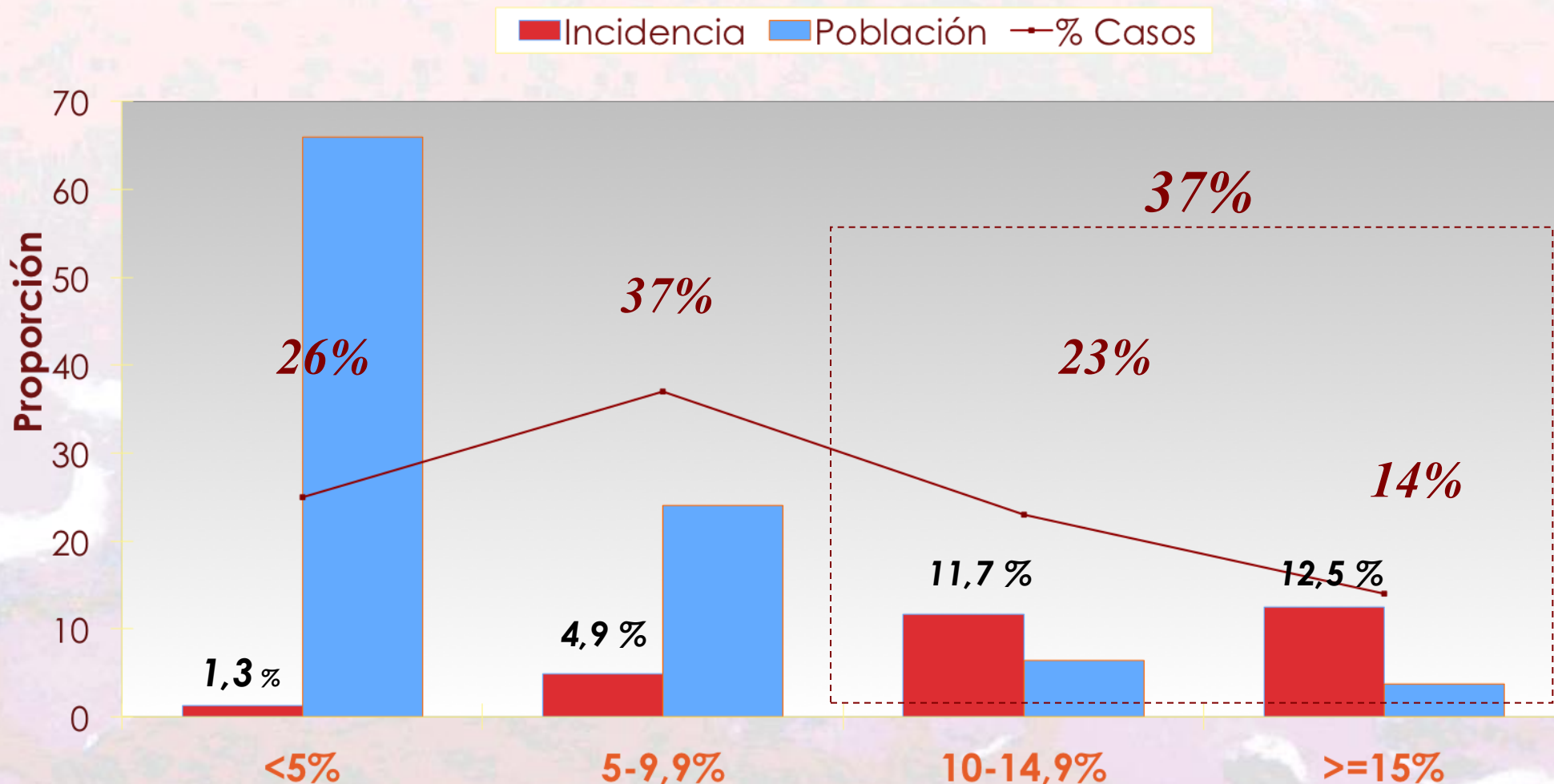
Riesgo REGICOR: Instrumento básico para el cribado poblacional

Hombres sin diabetes

Hombres con diabetes



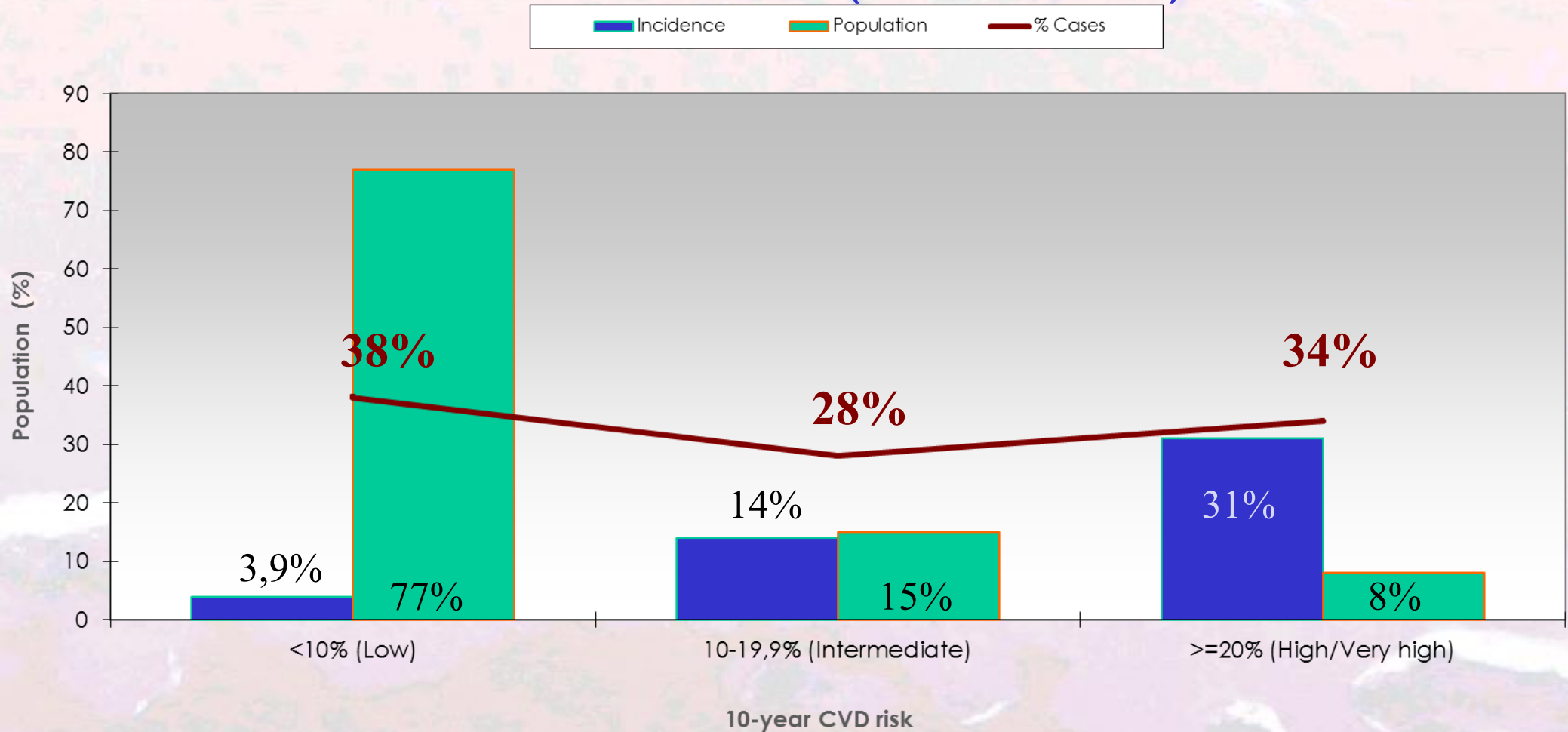
Riesgo coronario e incidencia de acontecimientos coronarios a 10 años en el estudio REGICOR (n=3724)

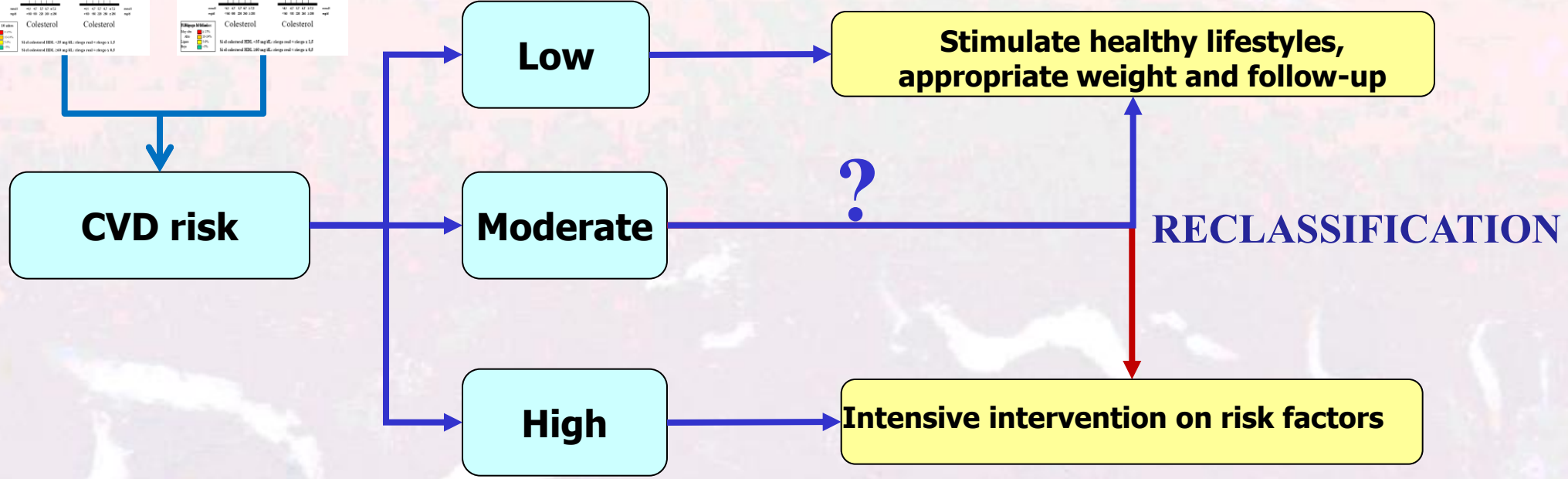
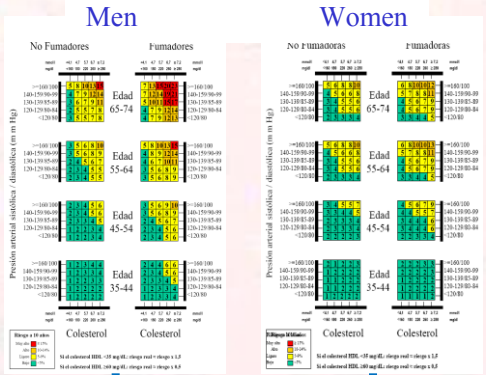


Riesgo a 10 años

Marrugat J et al. Rev Esp Cardiol 2011; 64: 385-94

Coronary artery disease risk and 10-year CAD event incidence in the USA The Emerging Risk Factors Collaboration* (n=246,669)





Posibles factores para la reclasificación de candidatos a prevención CV primaria

- Proteína C reactiva -as > 1g/l, Lp(a) > 30 UI
- Historia familiar de ECV precoz,
- Obesidad (IMC > 30), o cintura > recomendaciones,
- Microalbuminuria o insuficiencia renal,
- Dieta inadecuada (cuestionario corta auto-administrado)
- Ejercicio insuficiente (cuestionario corta auto-administrado)
- Perfil genético adverso (predisposición/carga genética)
- Grosor de la íntima media Carotídea
- Índice tobillo brazo
- Calcio intracoronario

**Enfermedad
subclínica**

Evaluación de un biomarcador

Prueba de concepto—Se asocia a la enfermedad

Estudios de asociación, caso-control: OR

¿pueden presentar la enfermedad?

Validación prospectiva—Predice la enfermedad en el seguimiento

Estudios de cohorte poblacional: RR, estadígrafo c

¿pueden presentar la enfermedad?

Valor añadido—Mejora la capacidad predictiva de las funciones clásicas?

Estudios de cohorte poblacional: mejora estadígrafo c, reclasificación NRI

Evaluación de un biomarcador

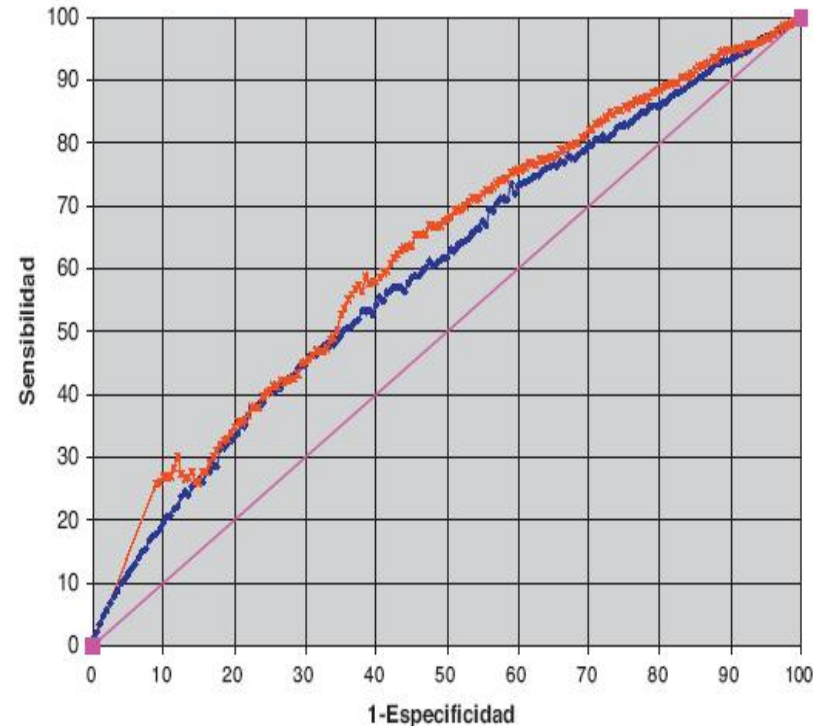
- Utilidad clínica—Los cambios implican cambios en las curvas de decisión
- Acontecimientos clínicos—El biomarcador implica cambios en la incidencia de los acontecimientos clínicos
- Coste-efectividad—Está justificada la determinación de un biomarcador?

Calcio intracoronario

Added value of CA cardiovascular ev

Sanne A. E. Peters, Marina Bakker, He
Julius Center for Health Sciences and Prim

- Mejora estadígrafo c
 - Mejora del estadígrafo c
- Mejora reclasificación (4 estudios)
 - NRI: 14 – 25%



Some advantages of determining the cardiovascular risk genotype over other biomarkers

Genetic characteristics :

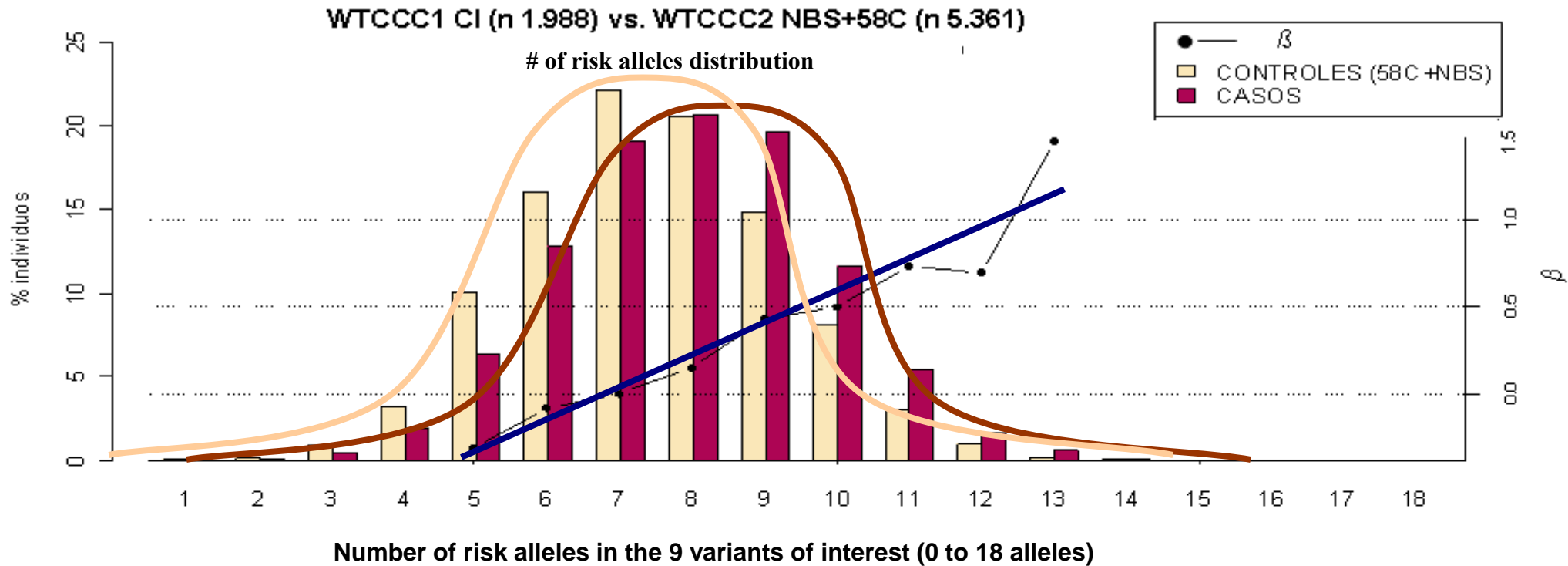
- Does not change with age or sex
- Does not change with food
- Does not change with drugs
- Does not have intra-individual variability
- Need to be determined only once in life (or until new markers are found)
- Can be incorporated to CV risk functions

$$\text{prob}(\text{event}_i | \text{CRF}_{p,i}, \text{SNP}_{j,i}) = 1 - \hat{S} \exp \left[\sum_{p=1}^P \beta_{\text{CRF}_p} * \text{CRF}_{p,i} + \sum_{j=1}^J \beta_{\text{SNP}_j} * \text{SNP}_{j,i} - \sum_{p=1}^P \beta_{\text{CRF}_p} * \text{CRF}_p - \sum_{j=1}^J \beta_{\text{SNP}_j} * \text{SNP}_j \right]$$

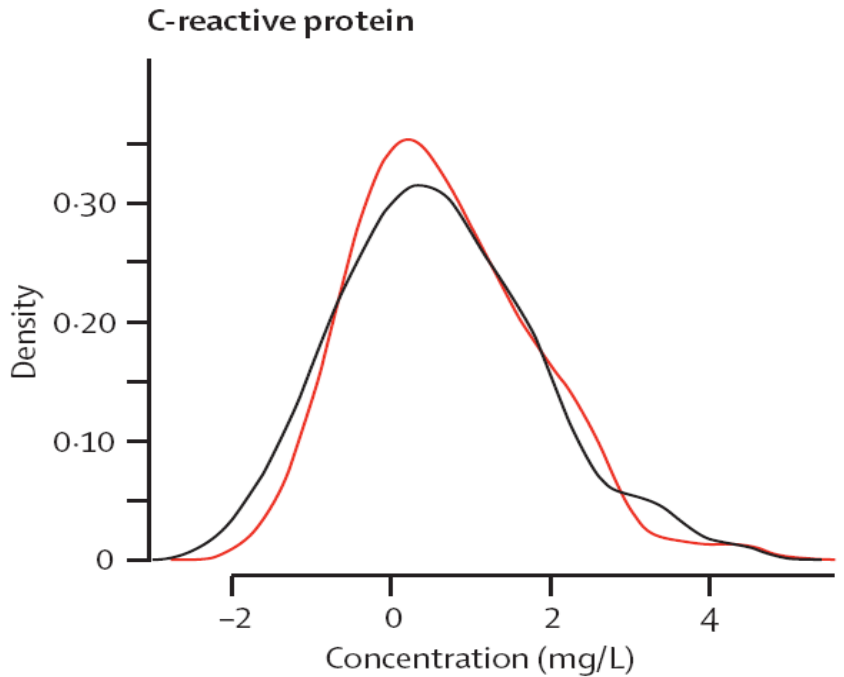
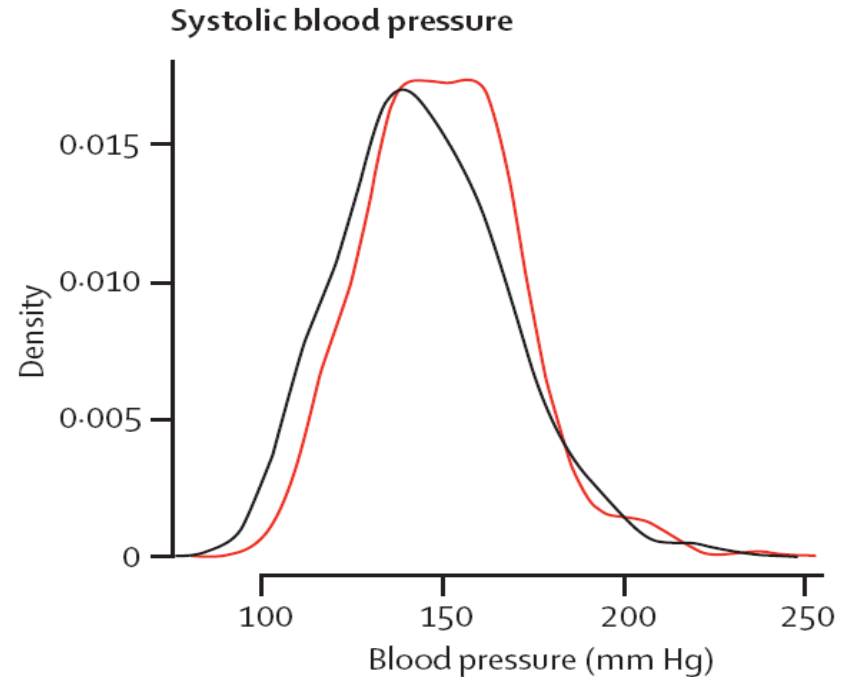
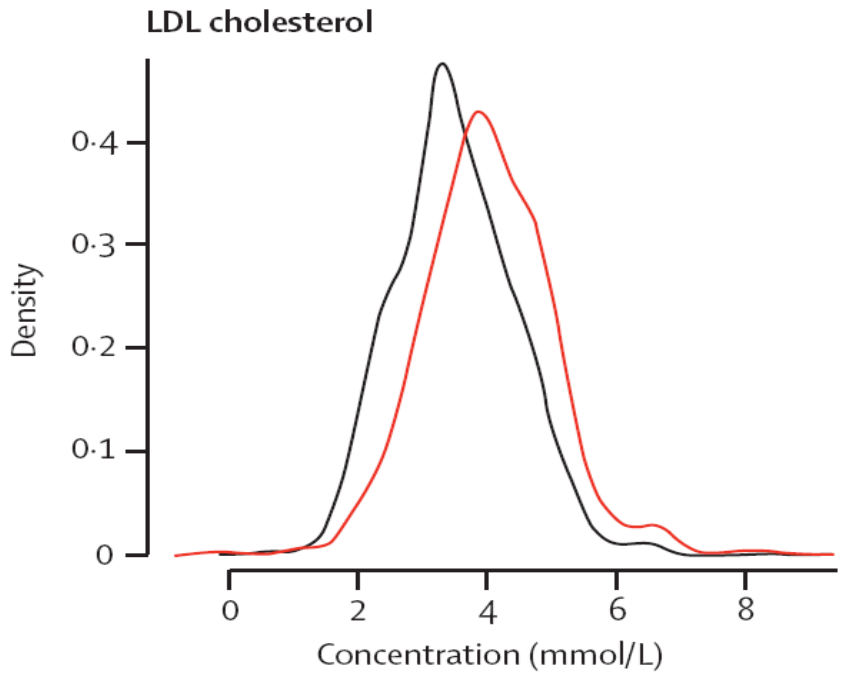
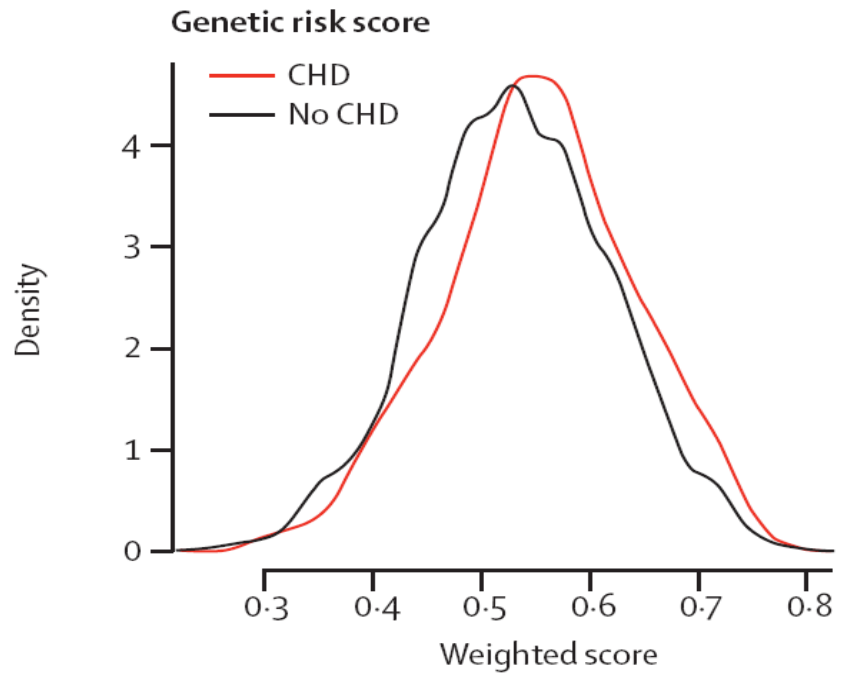
- Proof of concept—Do novel genetic markers, non associated with classical risk factors, differ between subjects with and without CHD?
 - In silico case-control study: on the Wellcome Trust Case Control Consortium public data
- Prospective validation—Incremental value:
 - Cohort studies:
 - REGICOR + Framingham
 - ...

Hlatky MA et al. *Circulation* 2009;119:2408-16.

Distribution of the number of alleles in the Welcome Trust CHD cases and controls consortium



OR per allele = 1.18

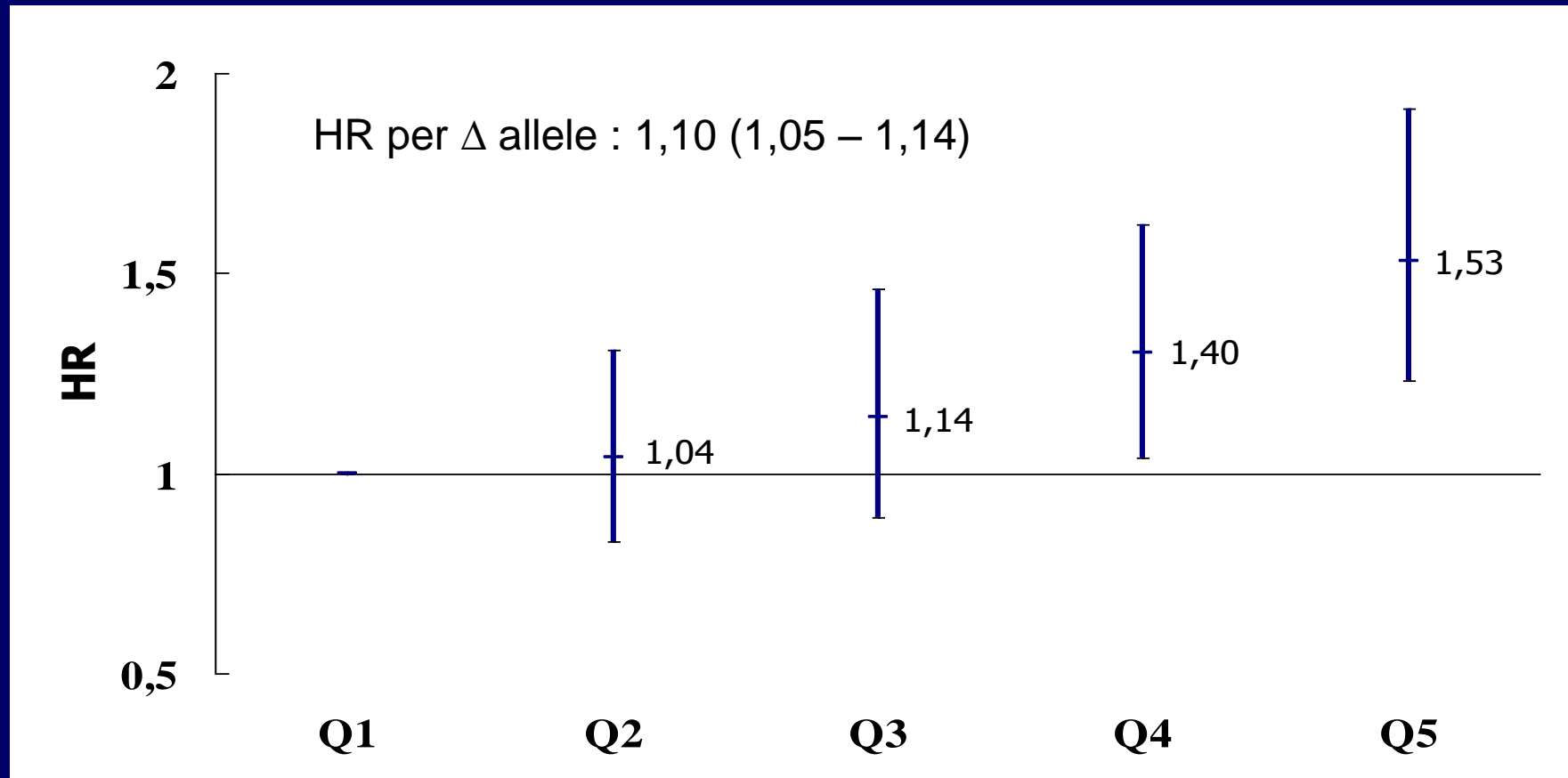


- Proof of concept—Do novel genetic markers, non associated with classical risk factors, differ between subjects with and without CHD?
 - In silico case-control study: on the Wellcome Trust Case Control Consortium public data
- **Prospective validation—Incremental value:**
 - Cohort studies:
 - REGICOR + Framingham

Hlatky MA et al. *Circulation* 2009;119:2408-16.

Quintiles of genetic risk score and 10-year CHD incidence

Meta-analysis of the REGICOR, Spain & Framingham, US, separate cohorts (Preliminary results)



Coronary events

REGICOR

Framingham

Classical risk factors + Genetic Score

Classical risk factors	Low risk	Intermediate-low risk	Intermediate-high risk	High risk
<u>Cases</u>				
Low risk	24	6	0	0
Intermediate-low risk	1	22	8	1
Intermediate-high risk	0	4	10	5
High risk	0	0	1	21
<u>Non-cases</u>				
Low risk	1415	105	1	0
Intermediate-low risk	115	339	64	9
Intermediate-high risk	0	36	69	30
High risk	0	5	20	40

Classical risk factors + Genetic Score

Classical risk factors	Low risk	Intermediate-low risk	Intermediate-high risk	High risk
<u>Cases</u>				
Low risk	60	11	0	0
Intermediate-low risk	7	36	5	0
Intermediate-high risk	0	4	30	9
High risk	0	0	8	84
<u>Non-cases</u>				
Low risk	2014	50	0	0
Intermediate-low risk	57	444	49	1
Intermediate-high risk	0	47	207	30
High risk	0	0	34	350

	REGICOR-Spain		FRAMINGHAM		Meta-analysis	
	All	Interm Risk	All	Interm Risk	All	Interm Risk
NRI for CHD (%)	12	25	3	14	6	17

Life-time CV risk representation (35 to 74 y) by REGICOR risk function with CRF and with CAD genetics alone

$$\text{prob}(\text{event}_i | \text{CRF}_{p,i}, \text{SNP}_{j,i}) = 1 - S \exp \left[\sum_{p=1}^P \beta_{\text{CRF}_p} * \text{CRF}_{p,i} + \sum_{j=1}^J \beta_{\text{SNP}_j} * \text{SNP}_{j,i} - \sum_{p=1}^P \beta_{\text{CRF}_p} * \overline{\text{CRF}_p} - \sum_{j=1}^J \beta_{\text{SNP}_j} * \overline{\text{SNP}_j} \right]$$

REGICOR Framingham, PROCAM, QRISK, SCORE

**Roberto Elosua
Maria Grau
Ana Redondo
Griselda Gonzalez
Joan Vila**

**Gavin Lucas
Carla Lluís
Isaac Subirana
Marta Tomás
Mariano Sentí**

WWW.REGICOR.ORG

**Funció de risc
per ios i Android**



**MI Covas, M Fitó, D Muñoz, S Heredia, G Blanchart, S Gaixas, J Peñafiel
S Tello, M Cabañero, L Franco, H Sanz, Y Ferrer, A Blasco, E Gomez, S Farré**

**Girona: J Sala, R Masiá, R Ramos, R Martí,
P Solanas, I Ramió, M Piqué, M Sirera, B Camps**

CHD/CVD screening

- Population
 - ✓ **Risk charts: screening basic system**
- Individuals
 - ✓ **First level of Reclassification**
 - ✓ Biomarkers
 - ✓ Genetic predisposition
 - ✓ **Second level of reclassification**
 - ✓ Exercise test, ABI, C-IMT
 - ✓ Non-invasive angiography
- Patients
 - ✓ **Diagnostic Confirmation**
 - ✓ Coronary angiography
- Patients
 - ✓ **PCI / other revascularization**

Simplified figure for potential use of emerging risk factors and imaging techniques in CV primary prevention

