



S.C. III MEDICA  
SERVIZIO di DIABETOLOGIA &  
MALATTIE METABOLICHE

AZIENDA OSPEDALIERO-UNIVERSITARIA  
OSPEDALE di CATTINARA - TRIESTE



XII SESSIONE  
AFERESI & ENDOTELIO

*Inquadramento delle Malattie Dismetaboliche*  
*Luigi Cattin*



**XV CONGRESSO NAZIONALE DELLA  
SOCIETA' ITALIANA DI EMAFERESI E  
MANIPOLAZIONE CELLULARE**

**XVI CORSO DI AGGIORNAMENTO IN  
EMAFERESI PER PERSONALE  
INFERMIERISTICO E TECNICO**

**3° SIMPOSIO “CELLULE STAMINALI:  
DALLA BIOLOGIA ALLA CLINICA”**

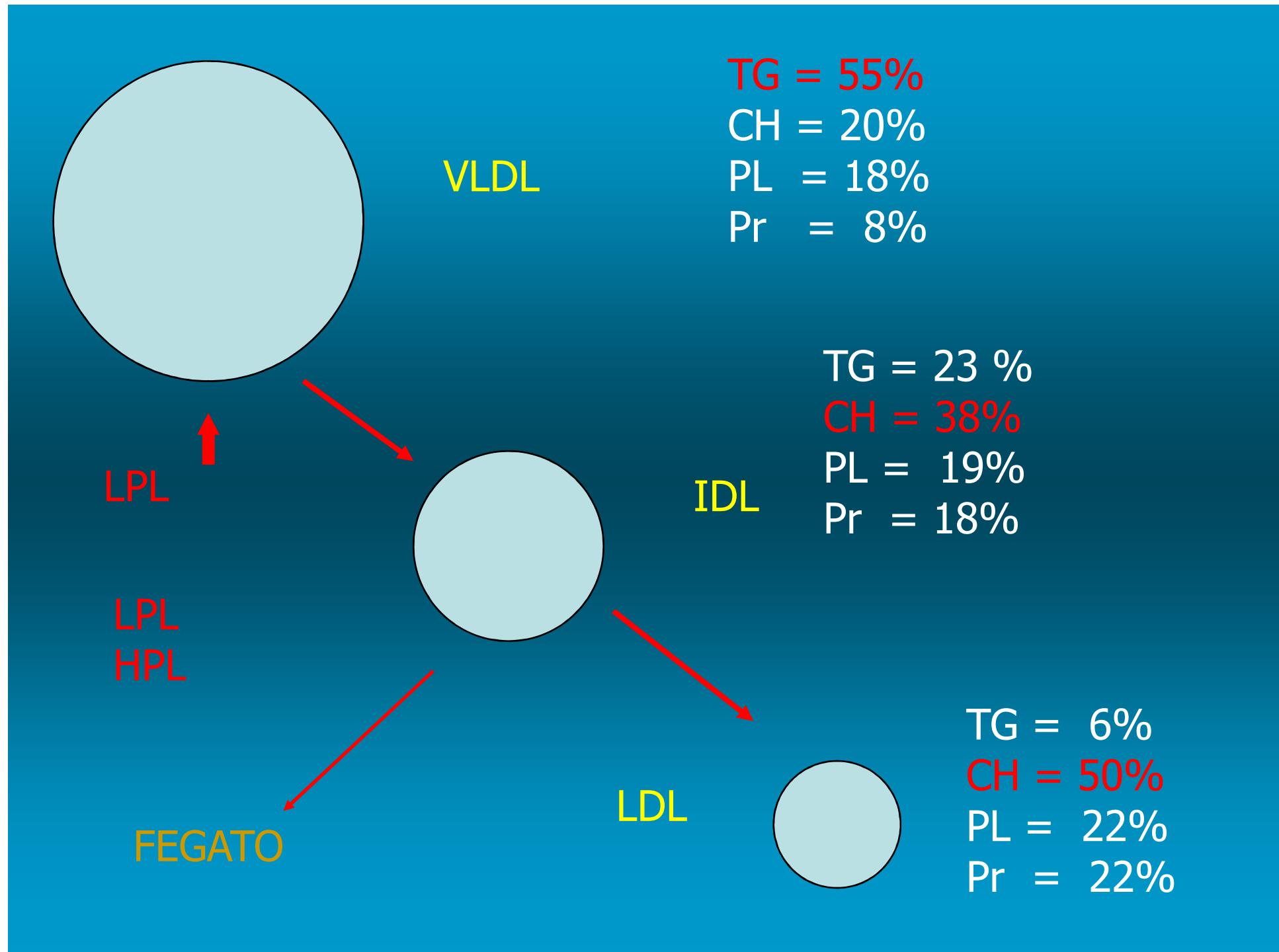
**Torino, 9-12 novembre 2011**  
Centro Congressi Lingotto

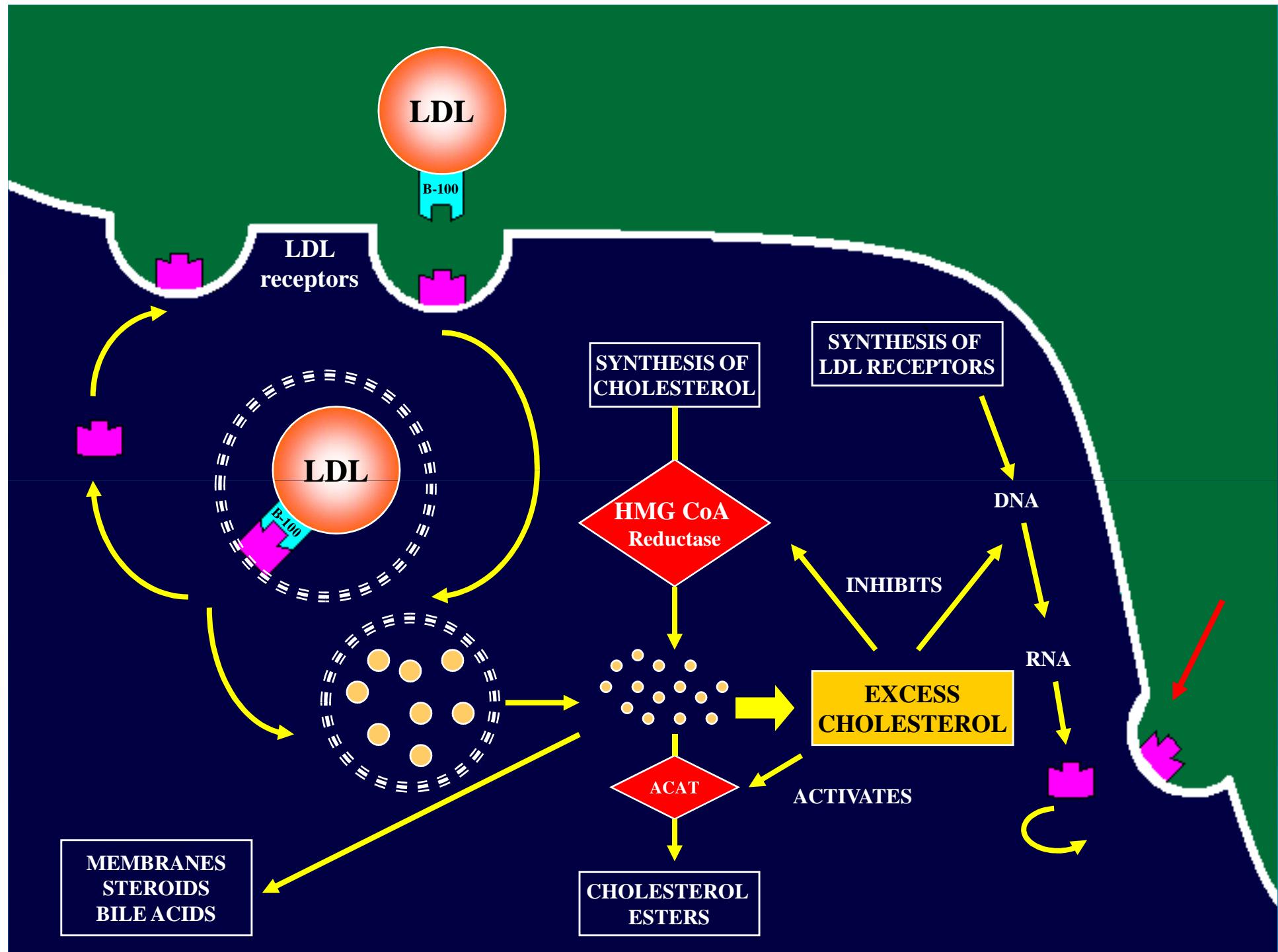


# The Number of Familial Dyslipidaemias in Italy

• Familial hypercholesterolemias	230.000
• <i>Familial combined hyperlipidemia</i>	> 600.000
• Type III hyperlipidemia	10.000 ?
• Severe hypertriglyceridemias	200 ?
• <i>Familial hypertriglyceridemias</i>	?
• Familial hypobetalipoproteinemias	20.000
• Combined hypolipidemia	?
• Abeta & Chylomicron Retention Disease	50-100
• Familial hypoalphalipoproteinemias	?
• Familial hyperalphalipoproteinemias	?







# INHERITED MONOGENIC HYPERCHOLESTEROLEMIAS

- Genetic disorders due to mutations of a single gene (monogenic / Mendelian disease)
- Biochemical phenotype:    **LDL-C >95th percentile**
- Clinical phenotype:
  - Tendon and cutaneous xanthomatosis
  - Premature coronary artery disease (pCAD)

# INHERITED MONOGENIC HYPERCHOLESTEROLEMIAS

## Dominant transmission

- Heterozygote      LDL-C ↑↑  
(One mutant allele)
- Homozygote      LDL-C ↑↑↑↑  
(Two mutant alleles)
- Gene dosage effect

## Recessive transmission

- Heterozygote      LDL-C ↔  
(One mutant allele)
- Homozygote      LDL-C ↑↑↑  
(Two mutant alleles)

# DOMINANT HYPERCHOLESTEROLEMIAS

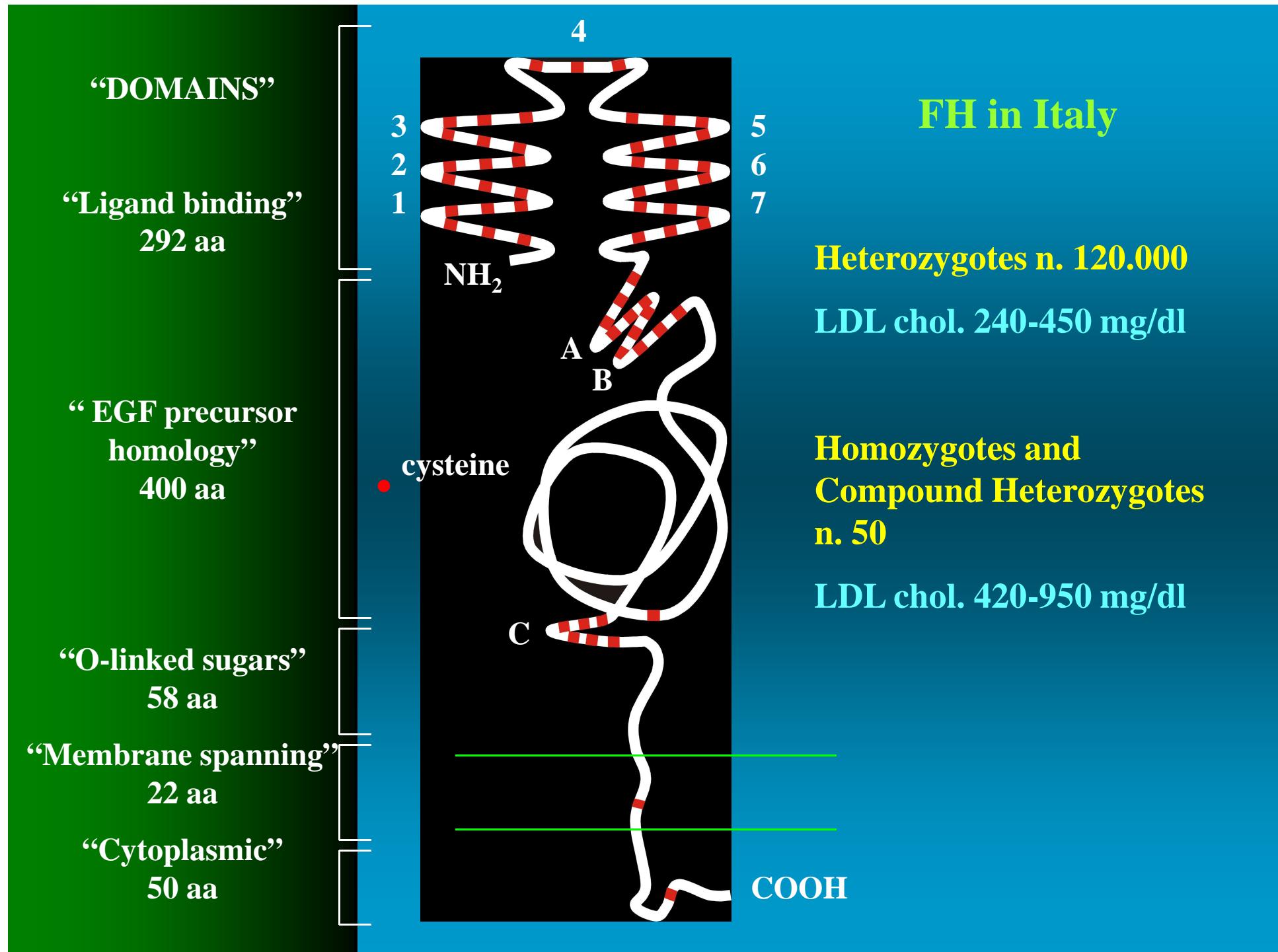
## (Familial hypercholesterolemia phenotype)

Disease	Gene	Prevalence in population
FH-1 (Classic FH)	LDL-R	
Heterozygous		1 per 500
Homozygous		1 per million
FH-2 (FDB)	Apo B-100	
Heterozygous		1 per 1000
Homozygous		>1 per million
FH-3 (ADH)	PCSK9	
Heterozygous		?
Homozygous		?

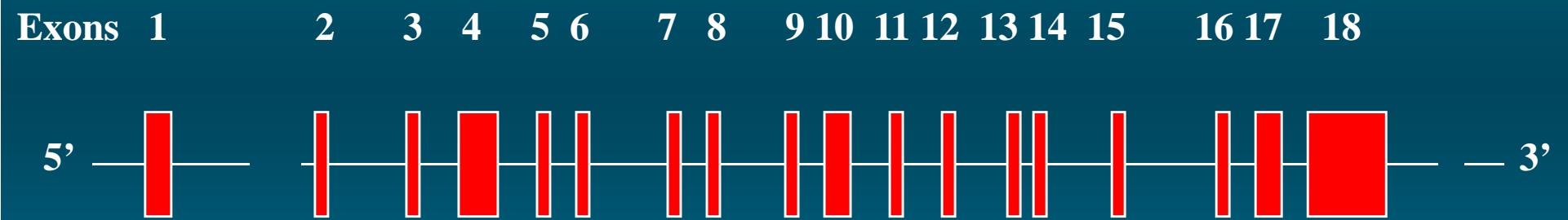
# DOMINANT HYPERCHOLESTEROLEMIAS

## (Familial hypercholesterolemia phenotype)

Disease	Gene	Prevalence in population
FH-1 (Classic FH)	LDL-R	<u>1 per 500</u>
Heterozygous		1 per million
FH-2 (FDB)	Apo B-100	1 per 1000
Heterozygous		>1 per million
FH-3 (ADH)	PCSK9	?
Heterozygous		?



## LDL receptor gene (chr. 19p13)



## MUTATIONS OF LDL-R GENE IN ITALIAN FH-1 PATIENTS

Major Rearrangements 17

Minute Deletions/Insertion 17

Point Mutations

Coding sequence

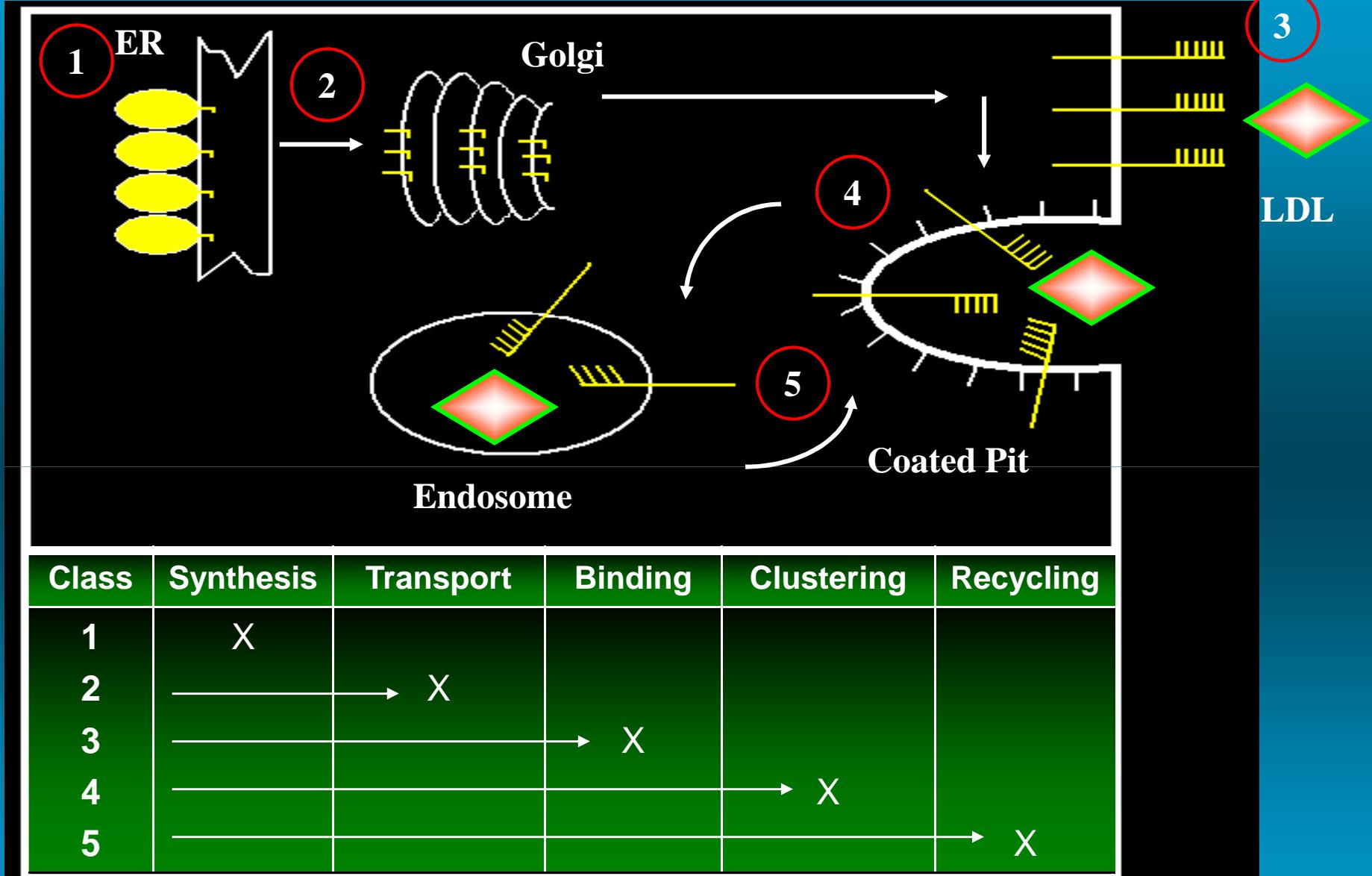
Deletions/insertions 7

Missense 63

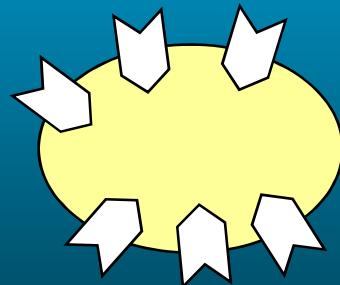
Nonsense 14

Splice junctions 13

TOTAL 131

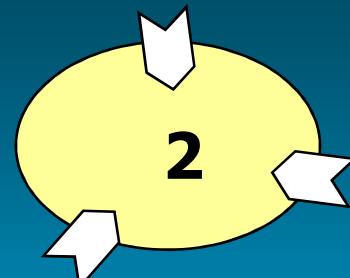
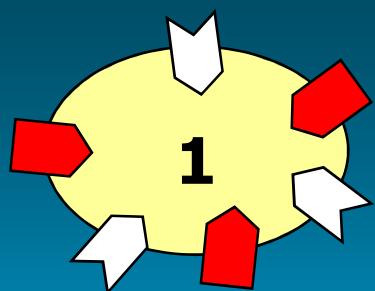


Normale

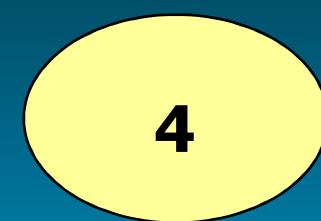
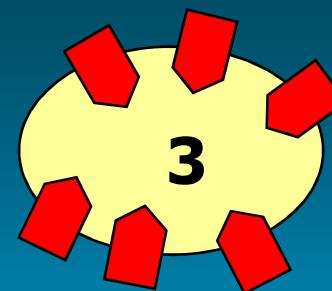


LDL-R

FH- eterozigote



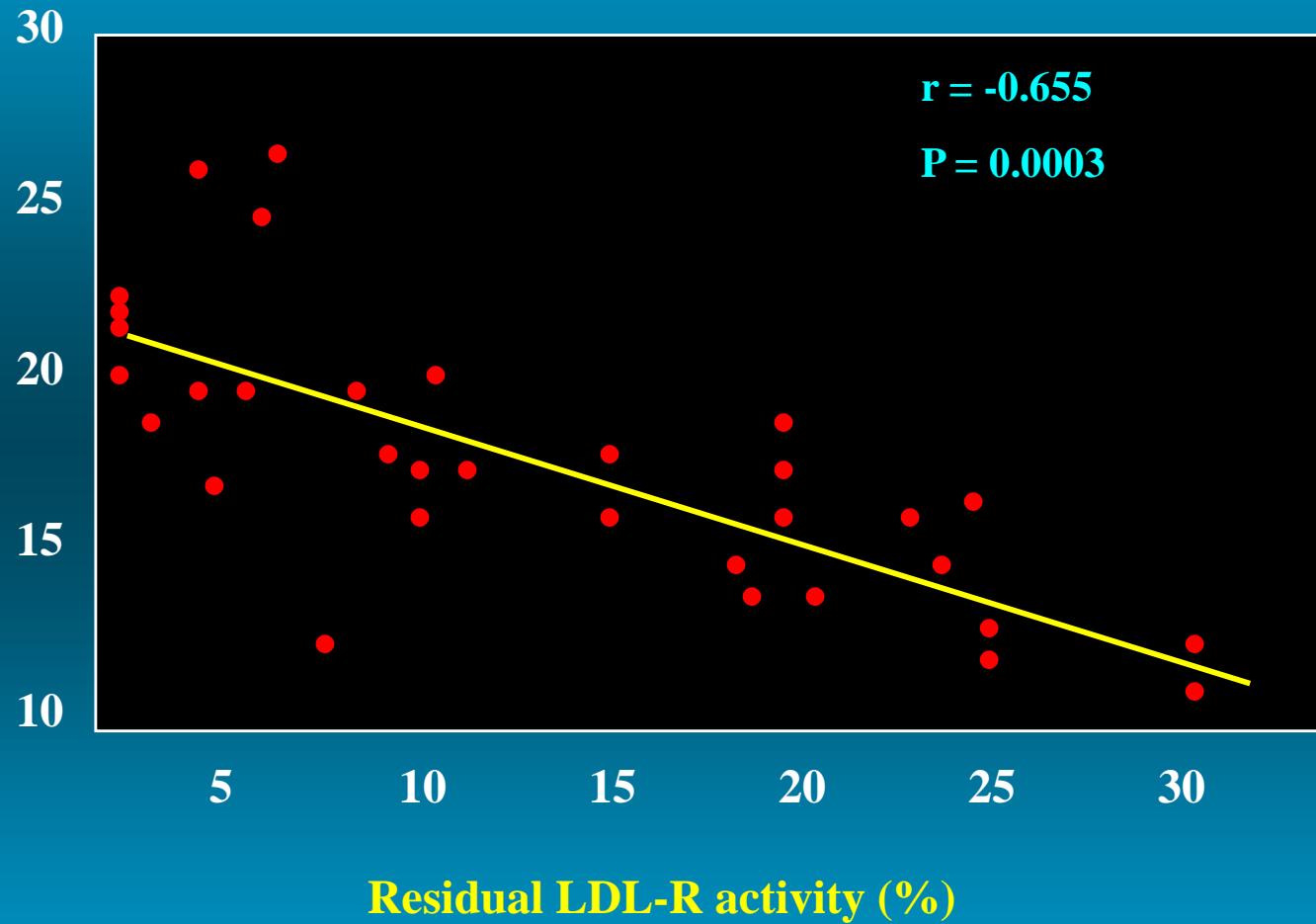
FH- omozigote



1 e 3 Fenotipo RECETTORE-DIFETTIVO

2 e 4 Fenotipo RECETTORE-NEGATIVO

**LDL-CH (mmol/L)**

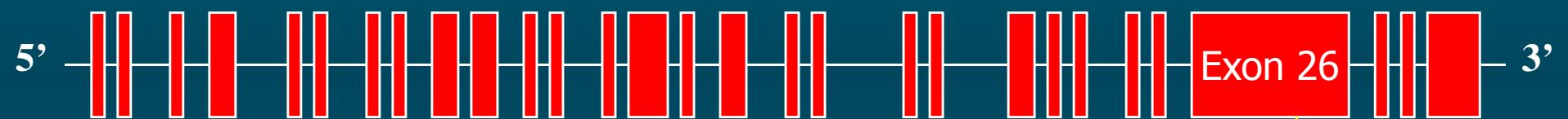


# DOMINANT HYPERCHOLESTEROLEMIAS

## (Familial hypercholesterolemia phenotype)

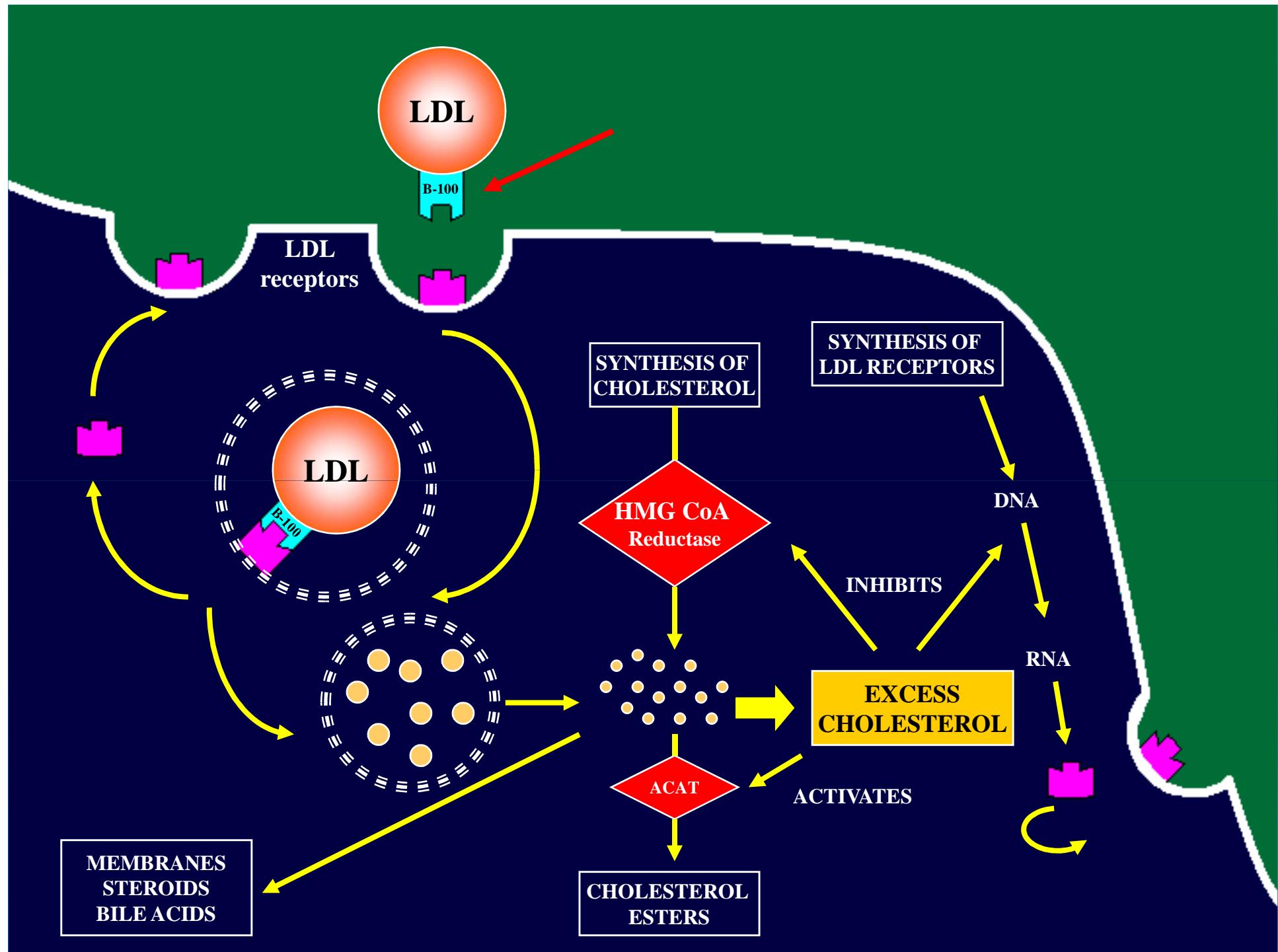
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FH-2 (FDB)	Apo B-100	
Heterozygous		1 per 1000
Homozygous		<1 per million
FH-3 (ADH)	PCSK9	?
Heterozygous		?
Homozygous		?

## Apolipoprotein B gene (chr. 2p24)



ApoB-100: 4536 aa.

CGG → TGG Arg3480 > Trp  
CGG → CAG Arg3500 > Gln  
CGG → TGG Arg3500 > Trp  
CGC → TGC Arg3531 > Cys  
CAC → TAC Arg3543 > Tyr

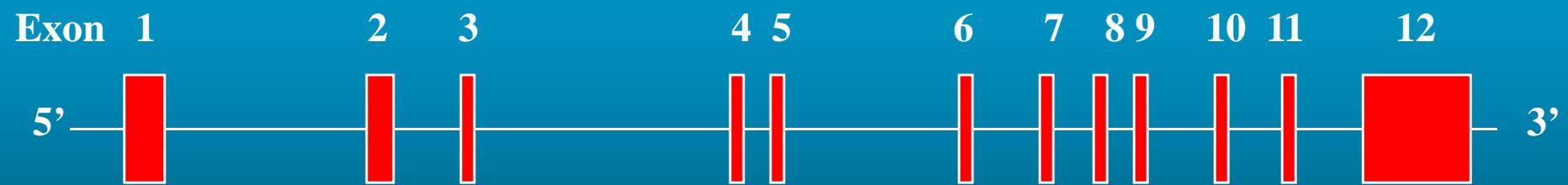


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Heterozygous		?
Homozygous		?

# PCSK9 gene (1p32)



mRNA (3636 nt)

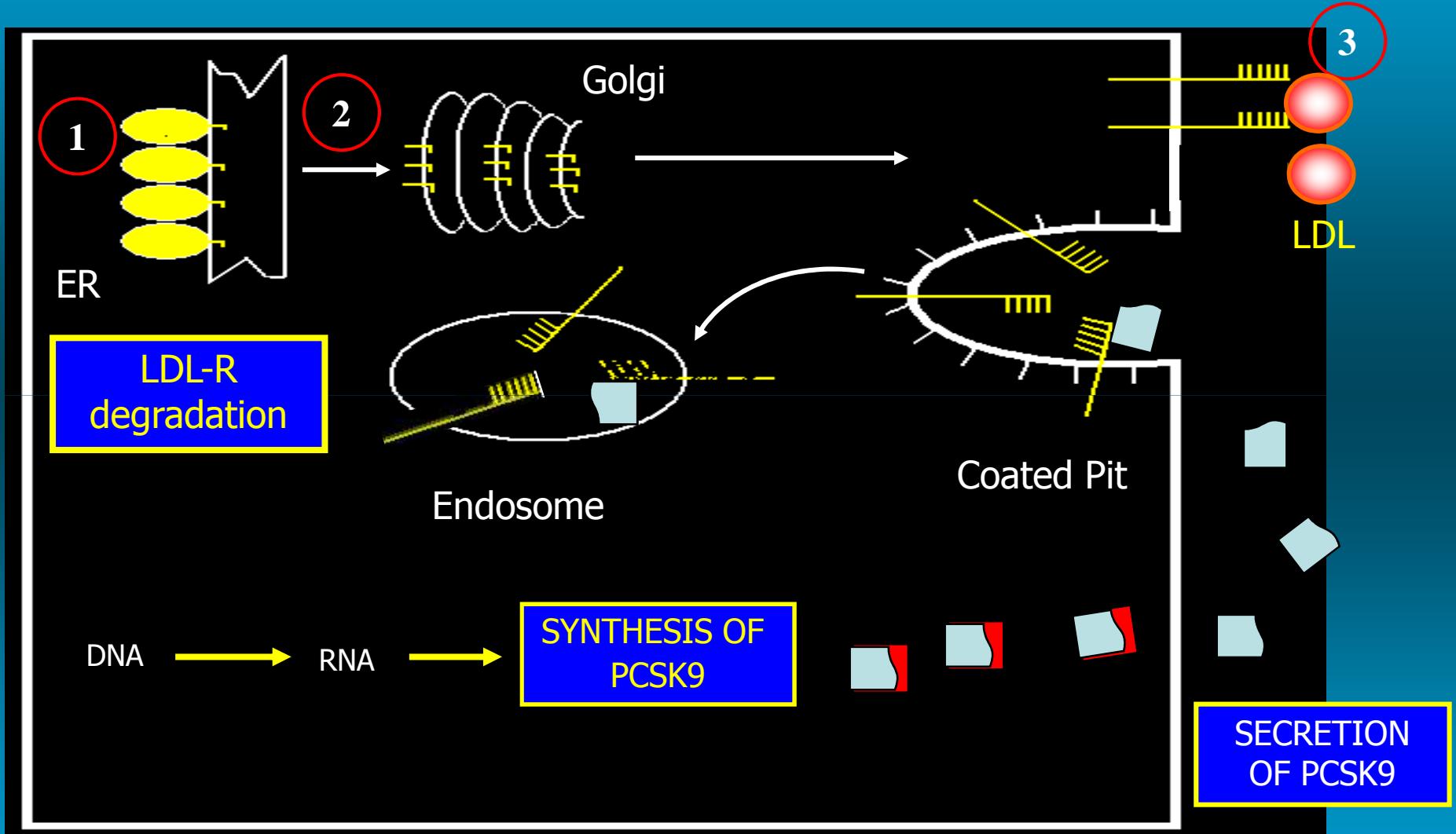
PCSK9 protein (692 aa)

Proprotein Convertase Subtilisin/hexin type 9 Serine Protease

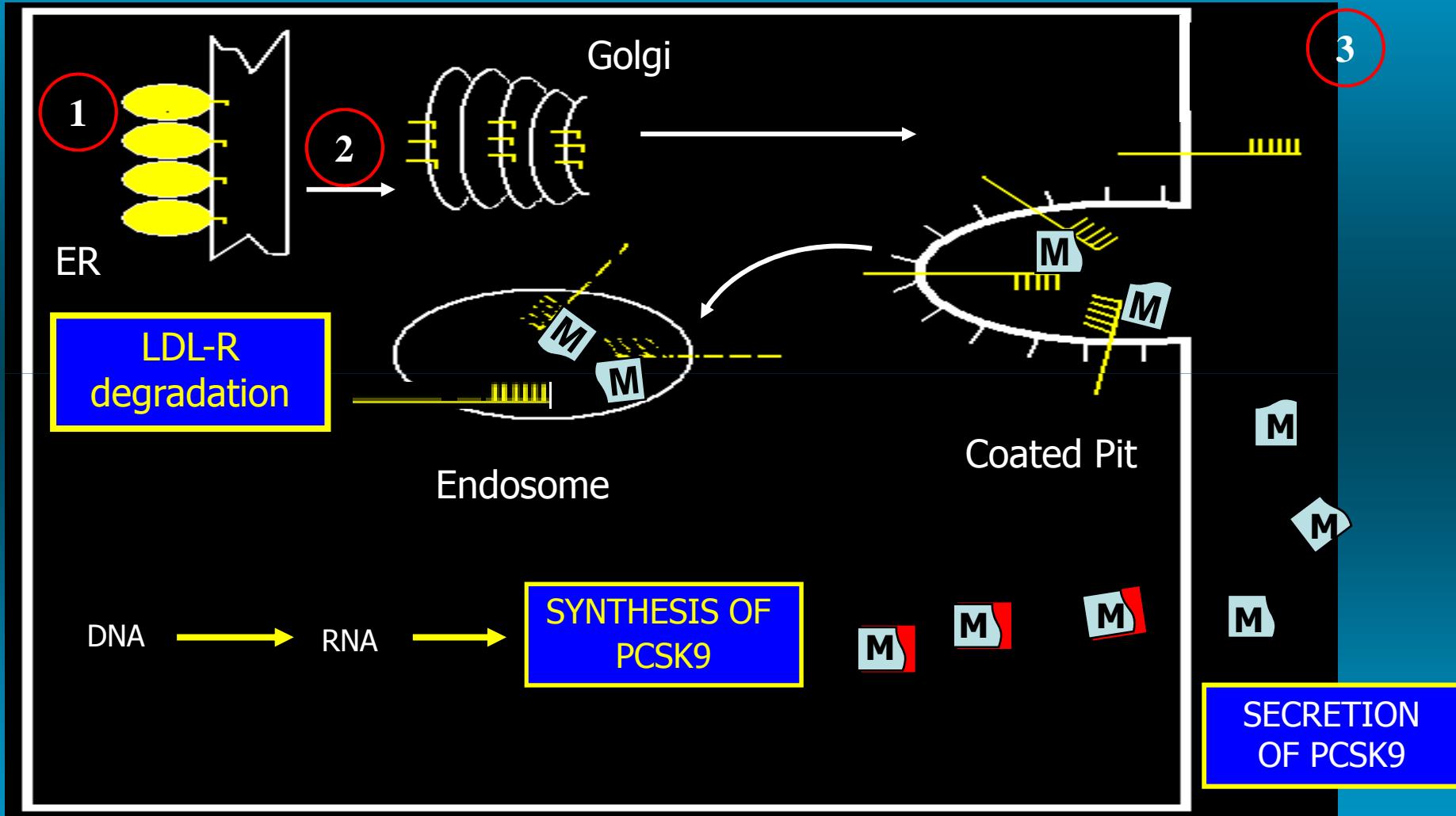


Synthesized by the liver and secreted into the circulation

# Regulation of LDL-R number by PCSK9 activity



# Reduction of LDL-R number induced by increased PCSK9 activity



# INHERITED MONOGENIC HYPERCHOLESTEROLEMIAS

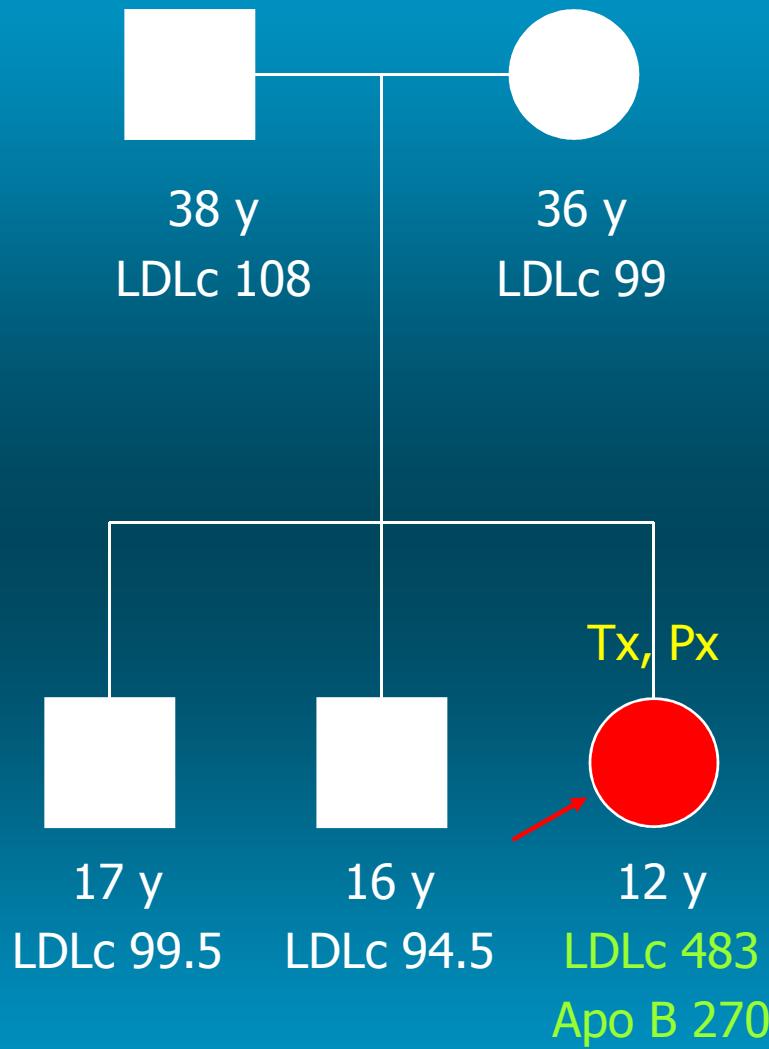
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## Recessive transmission

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(One mutant allele)
- Homozygote      LDL-C ↑↑↑  
(Two mutant alleles)

## M. Family (from Sardinia)



*LDL-R activity in fibroblasts*

*Normal*

*Affinity of LDL for LDL-R*

*Normal*

*FCR of LDL in vivo*

*Reduced as in homozygous FH*

Clinical diagnosis: Pseudo-homozygous FH

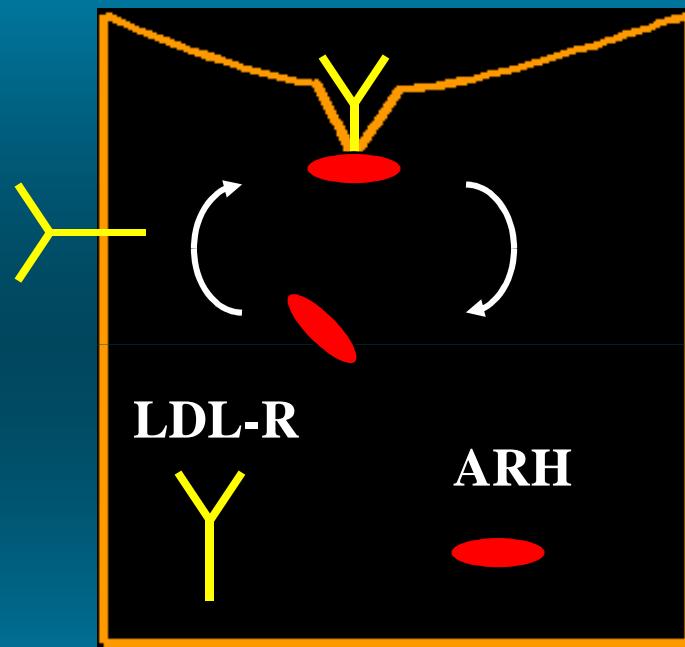
LDLc = mg/dl

# Chromosome 1p

ARH 1p36-35



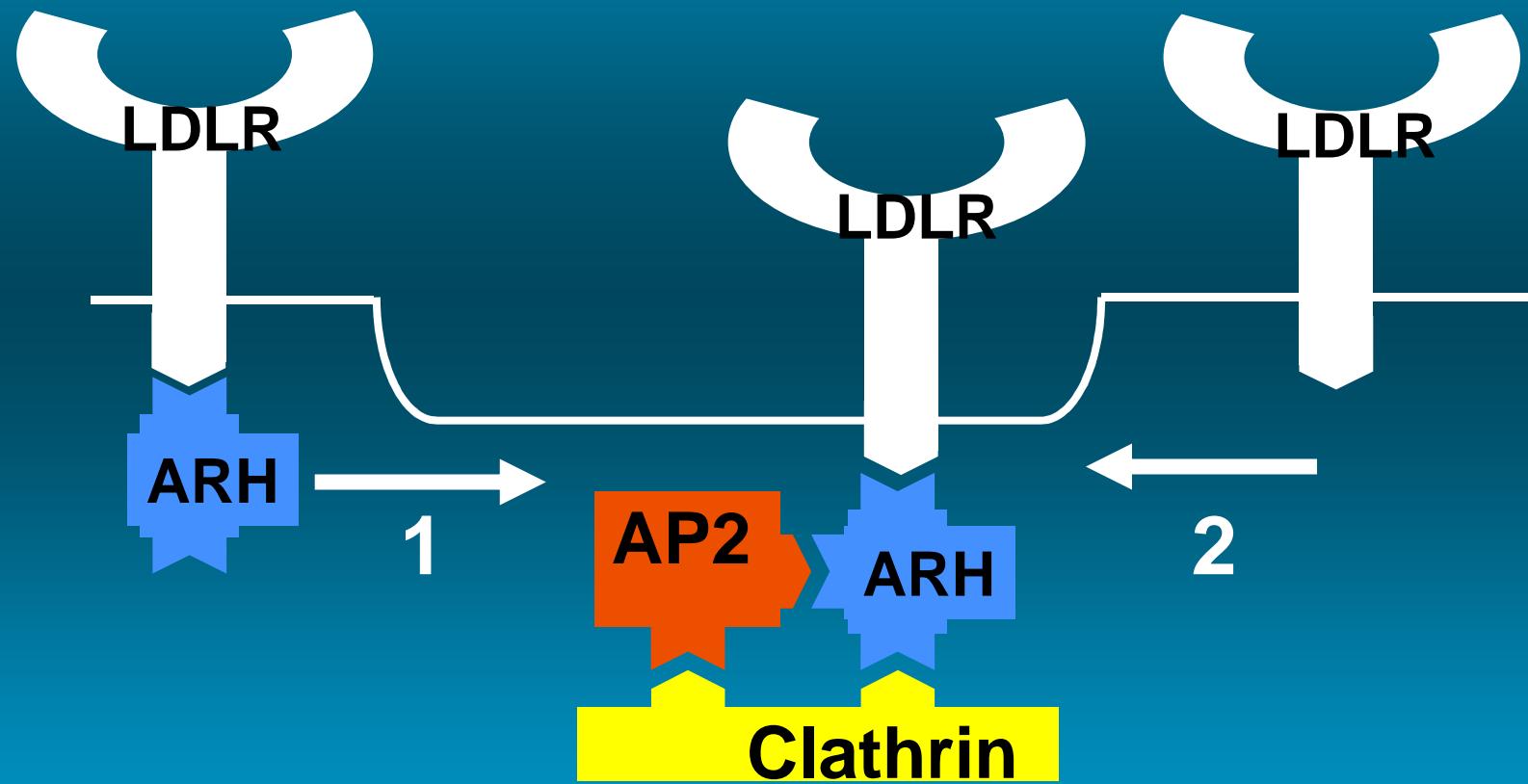
*blood*



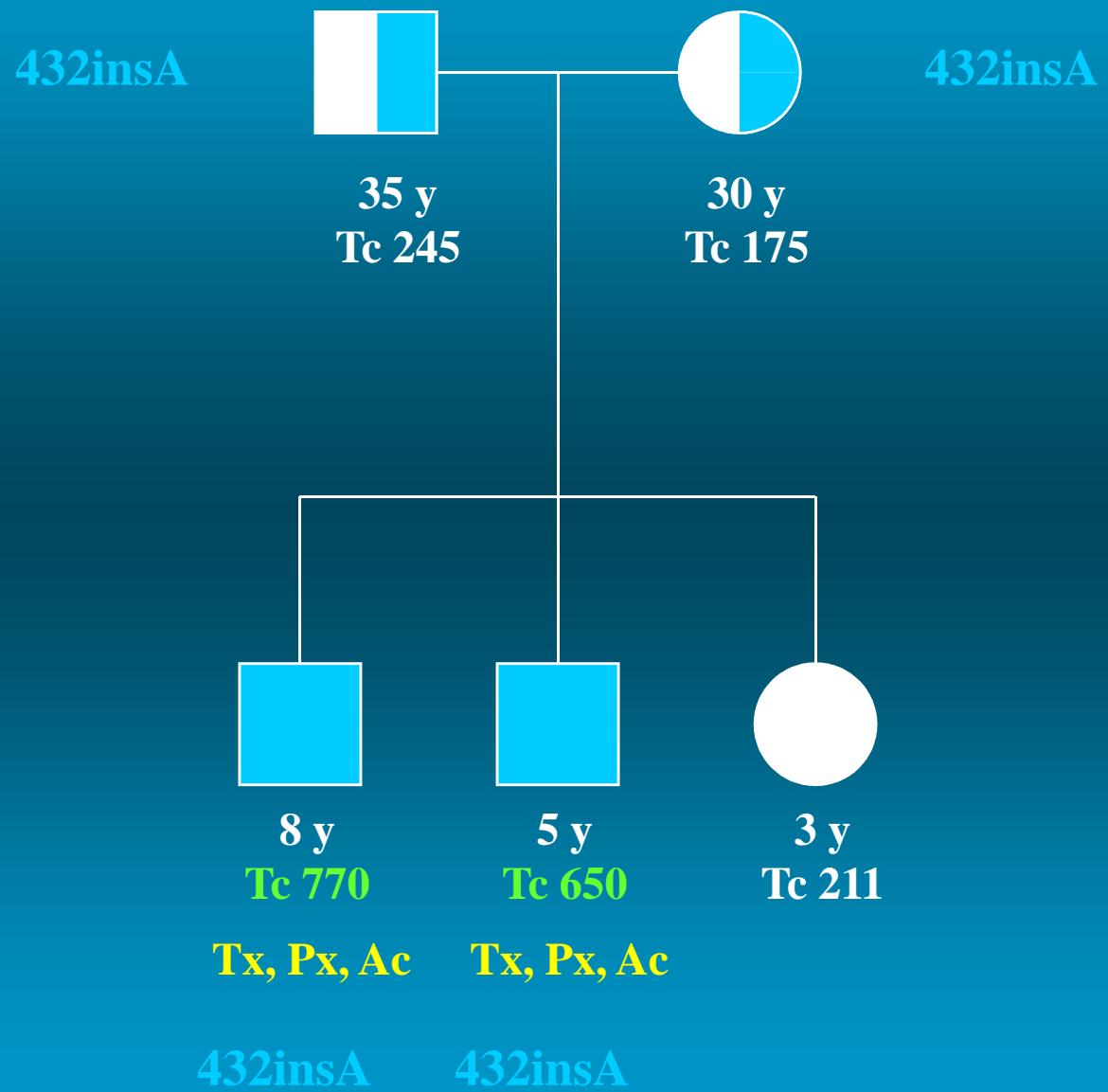
*hepatocite*

*bile*

# Hypothesis: ARH Clusters LDLR in Coated Pits



## S. Family





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# DISLIPIDEMIE

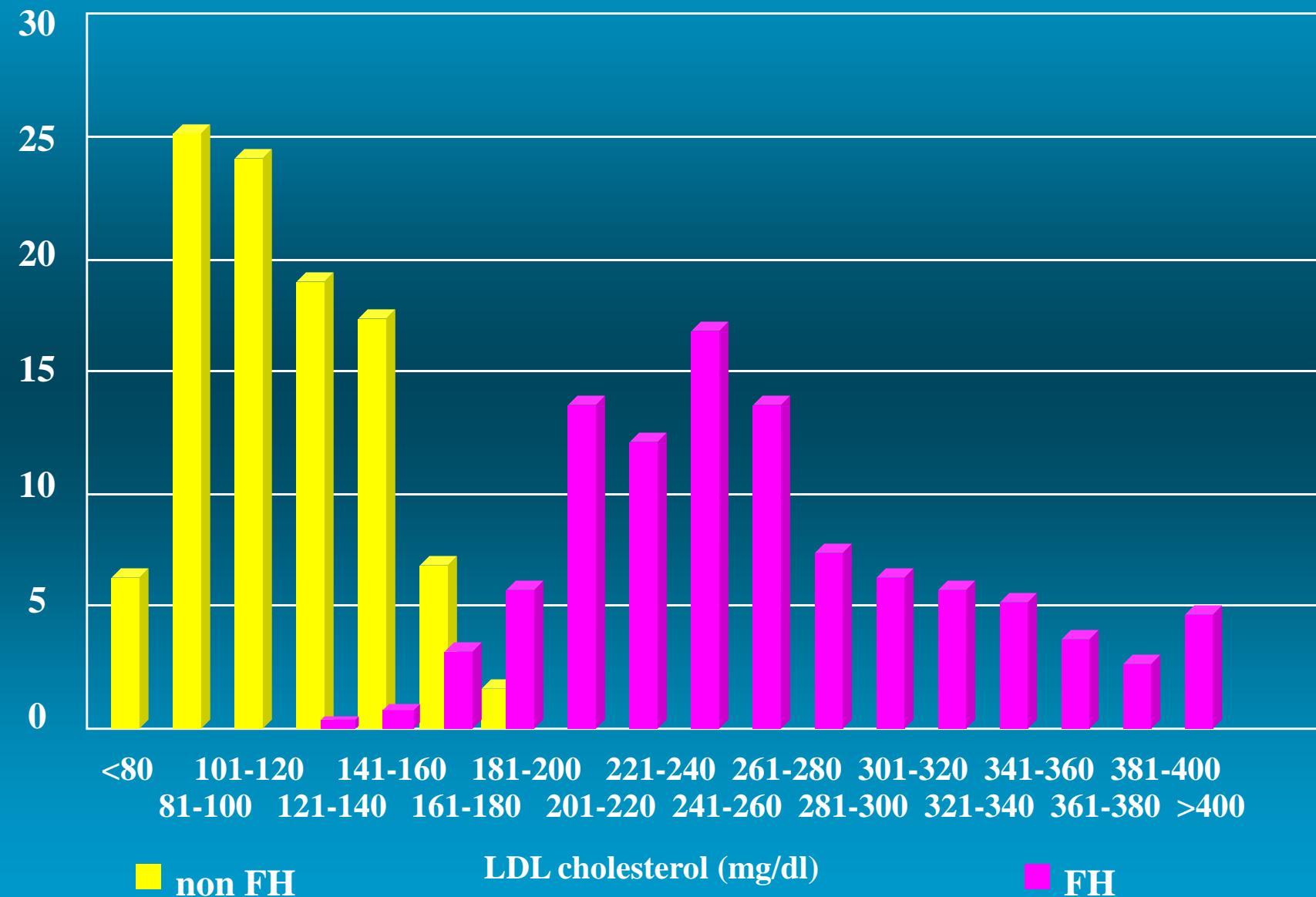


# CLINICAL DIAGNOSIS OF FAMILIAL HYPERCHOLESTERolemia

- Plasma LDL-C level
- Clinical history
- Family history
- Clinical score

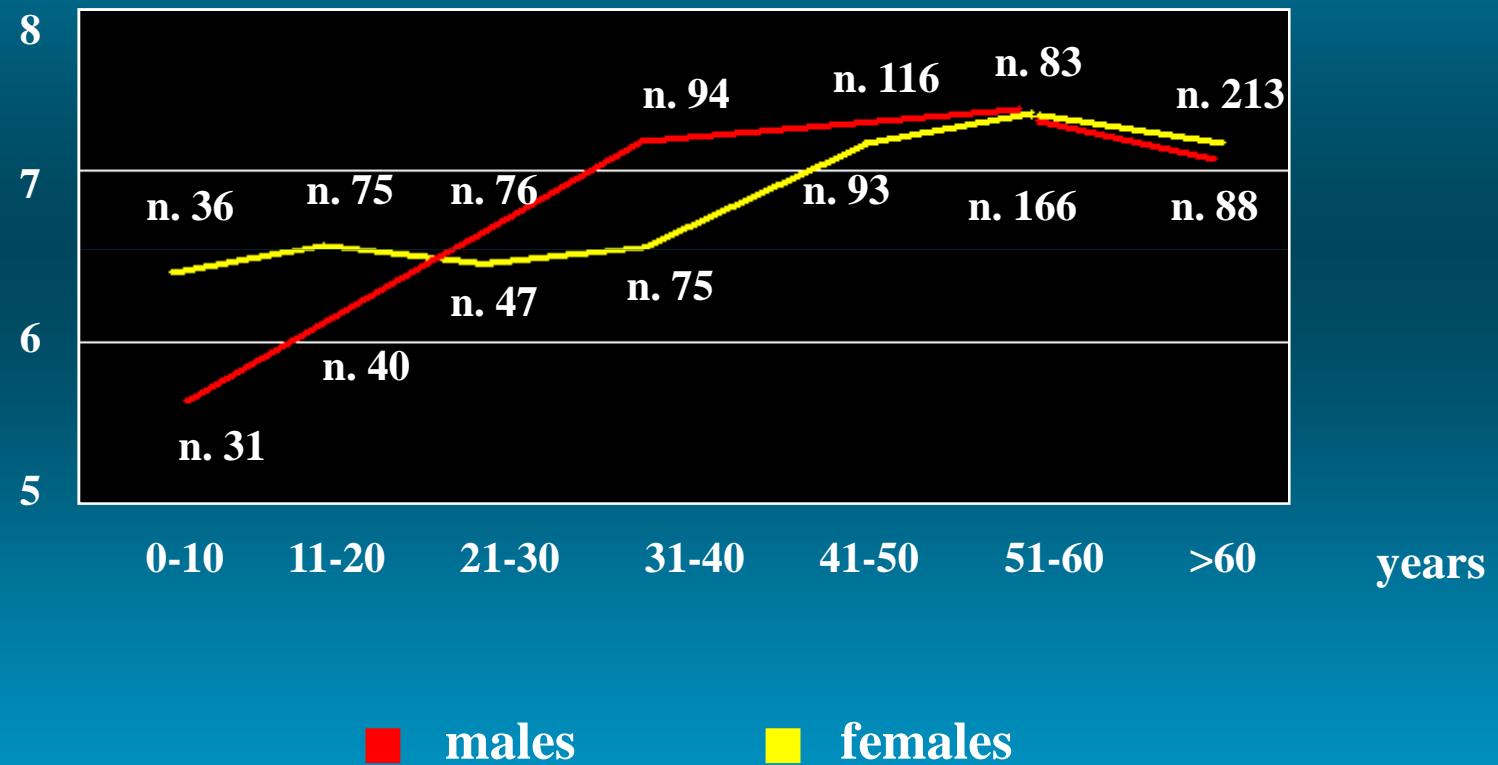
## LDL cholesterol distribution in FH families

Frequency (%)



## LDL cholesterol levels in FH patients according to age

mmol/l

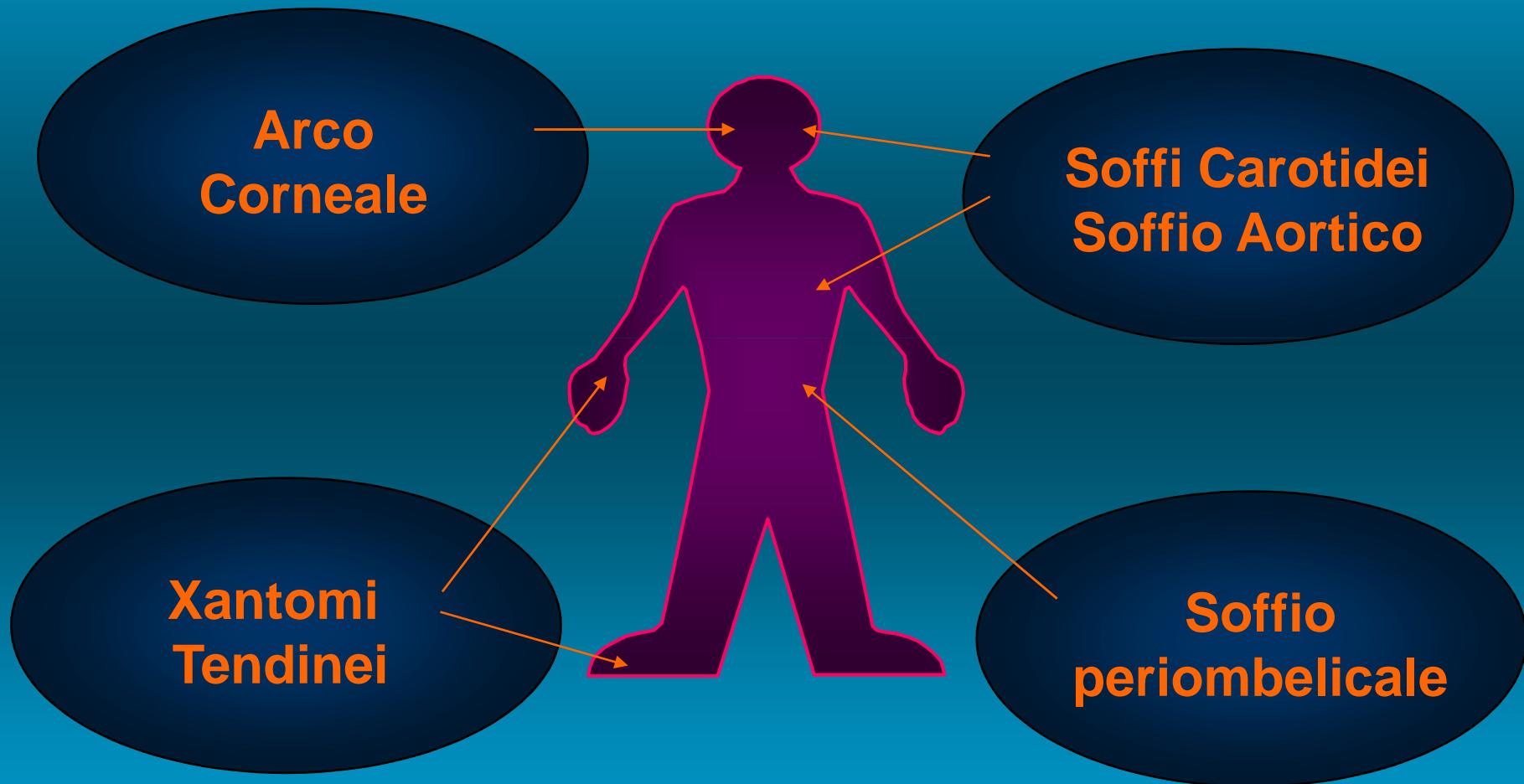


# CLINICAL DIAGNOSIS OF FH

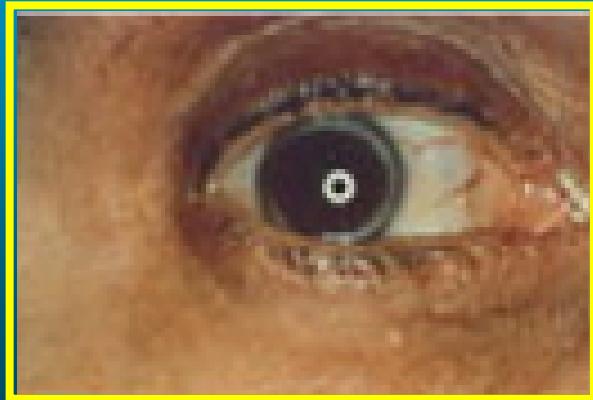
## Plasma LDL-C level

- LDL-C > 8.5 \* > 330 \*\* = 8 p.
- LDL-C 6.5-8.4 250 -329 = 5 p.
- LDL-C 5.0-6.4 190-249 = 3 p.
- LDL-C 4.0-4.9 155-198 = 1 p.
- \* mmol/ \*\* mg/dl

## Approccio clinico al paziente FH 1



**Gerontoxon**



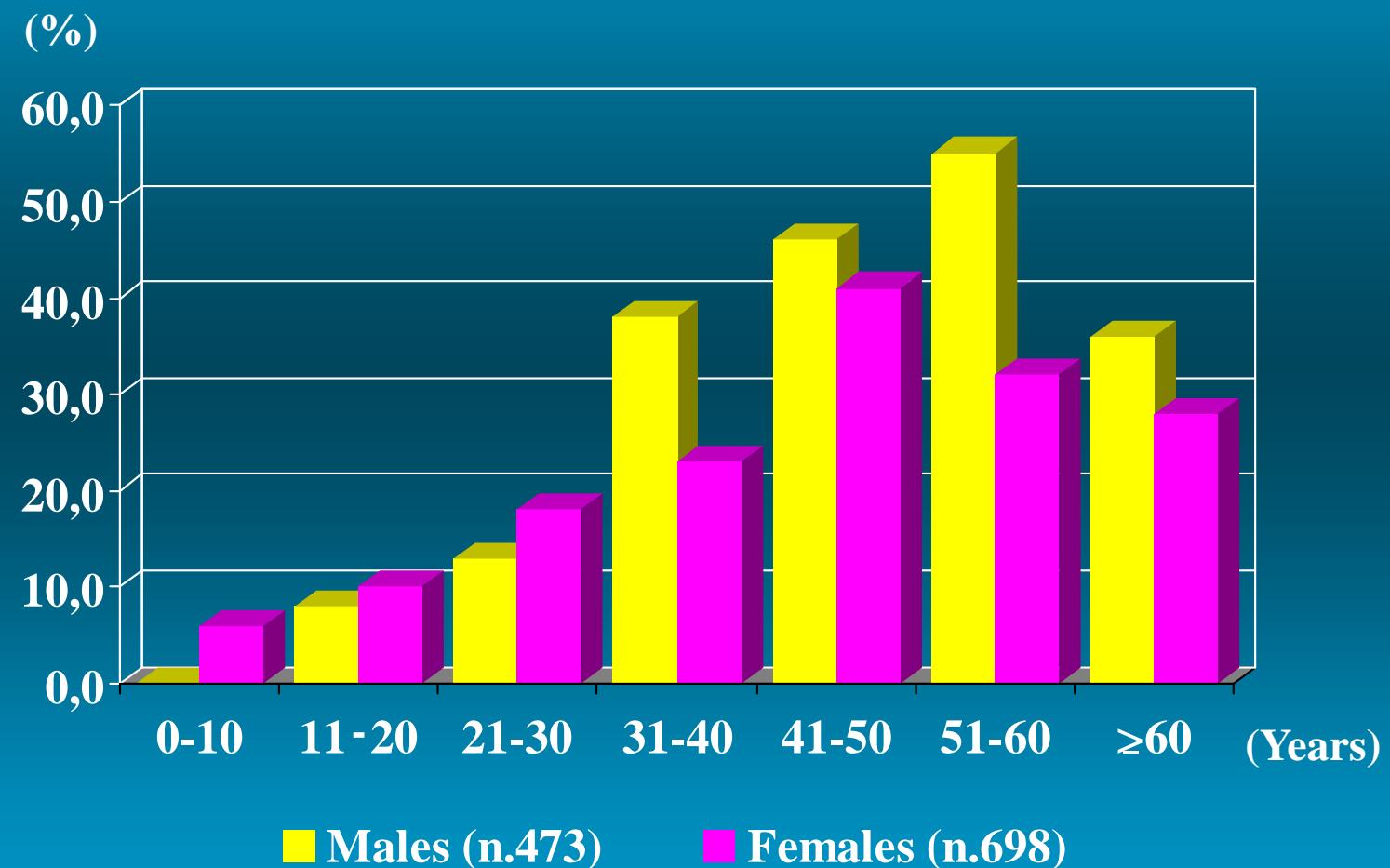
**Xanthelasmas**



**Tendon xanthomas**



## Prevelence of Tendon Xanthomas in FH heterozygous patients

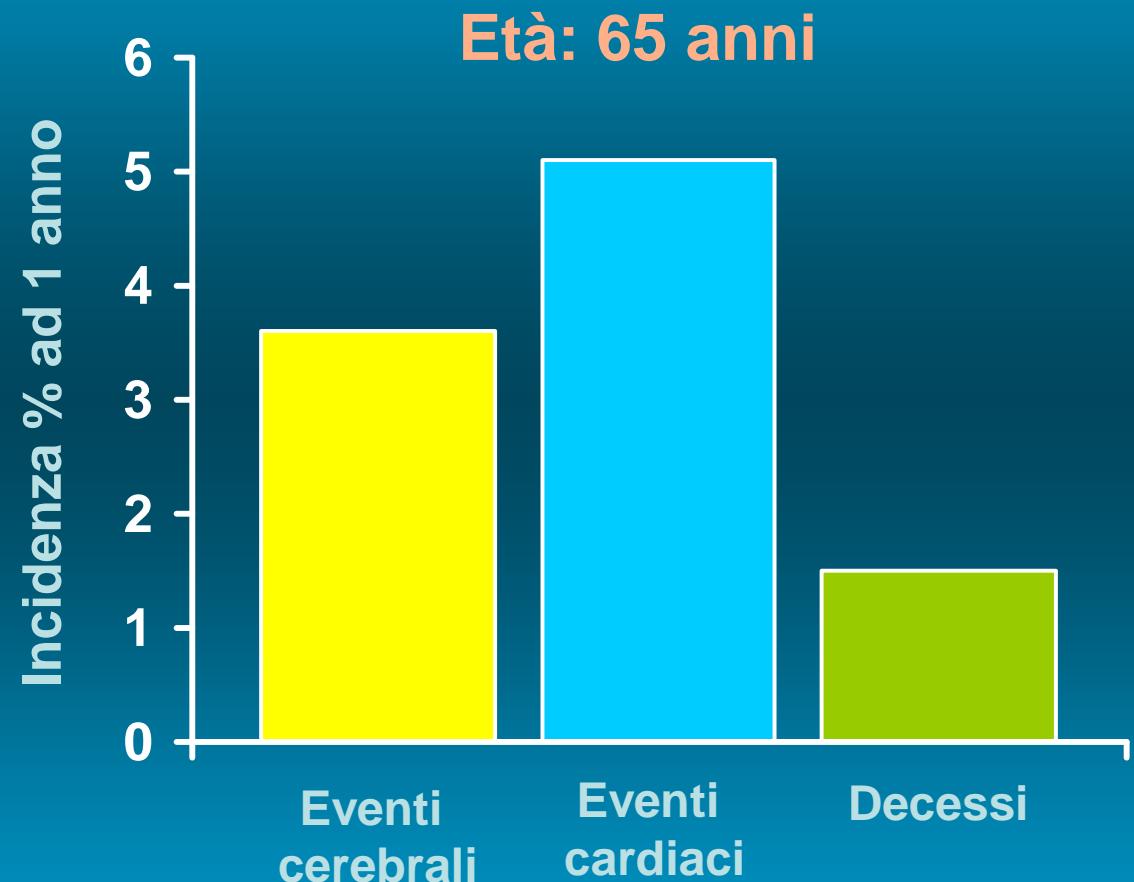
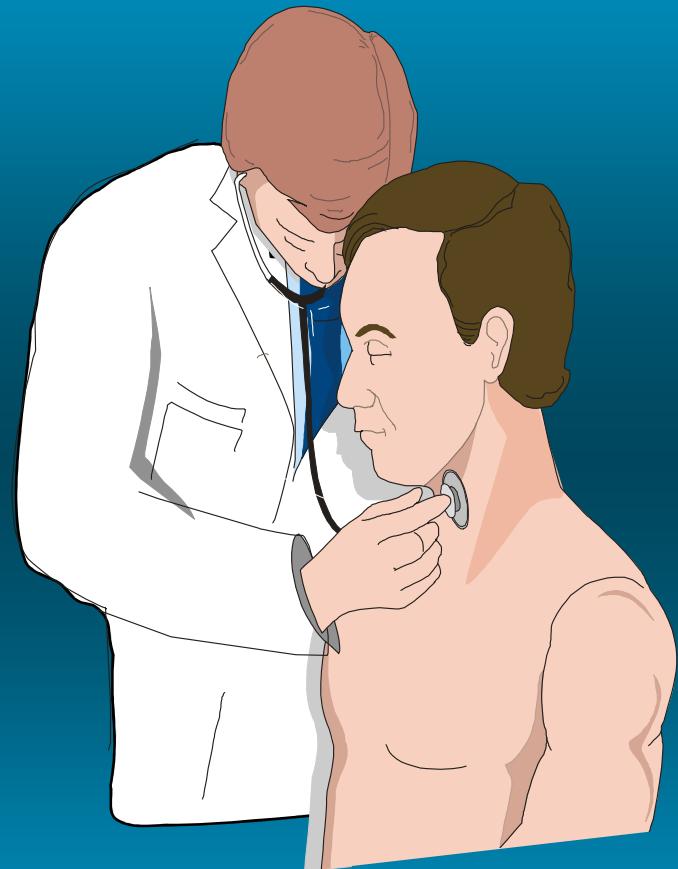


## Coefficienti di correlazione r tra aree TC dei tendini e parametri esaminati

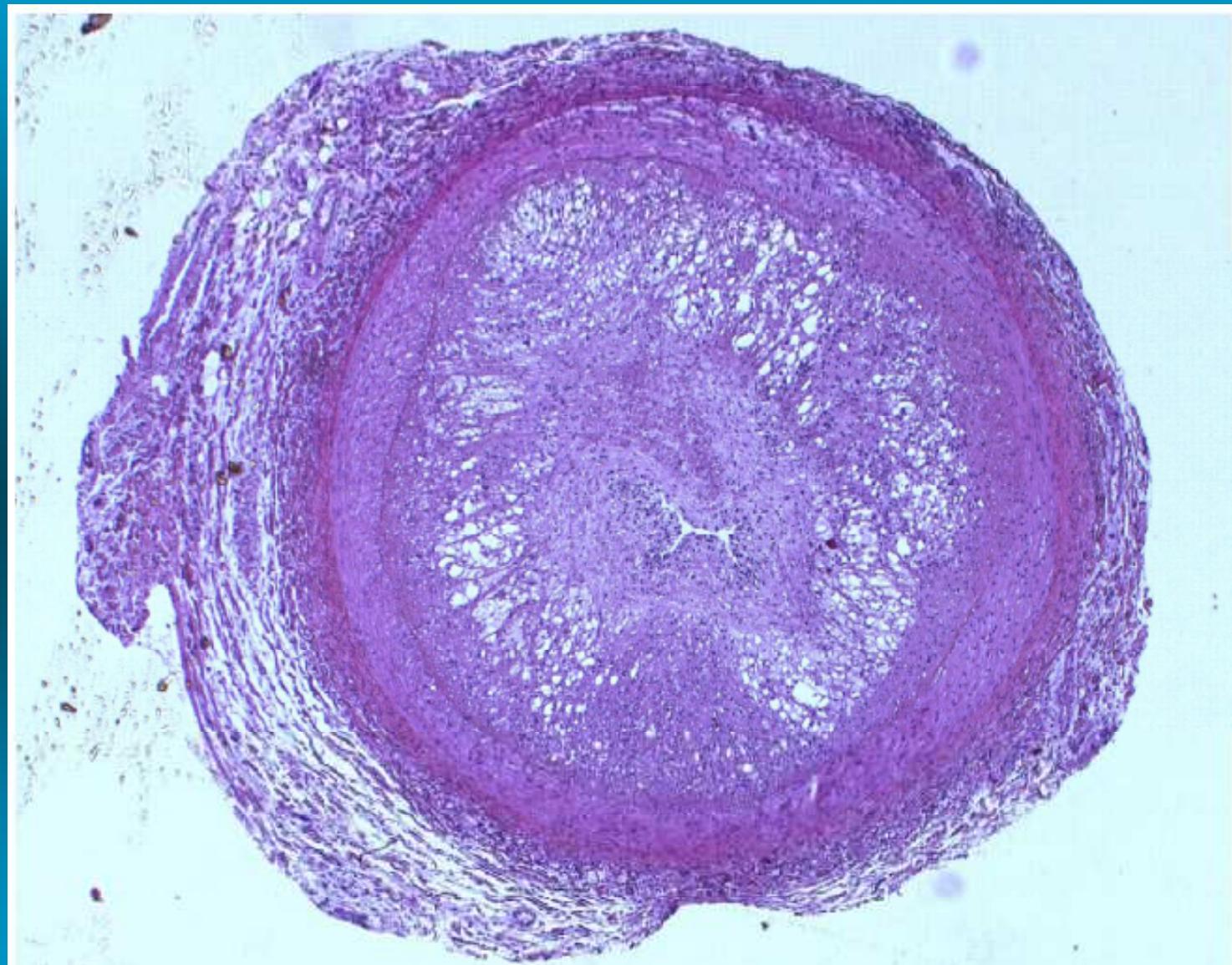
• ETA'	0.22	p< 0.05
• BMI	0.20	0.067
• CT	0.47	p< 0.001
• LDL-C	0.53	p< 0.001
• HDL-C	-0.35	p< 0.005
• CT/HDL-C	0.63	p< 0.001
• TG	0.01	0.45



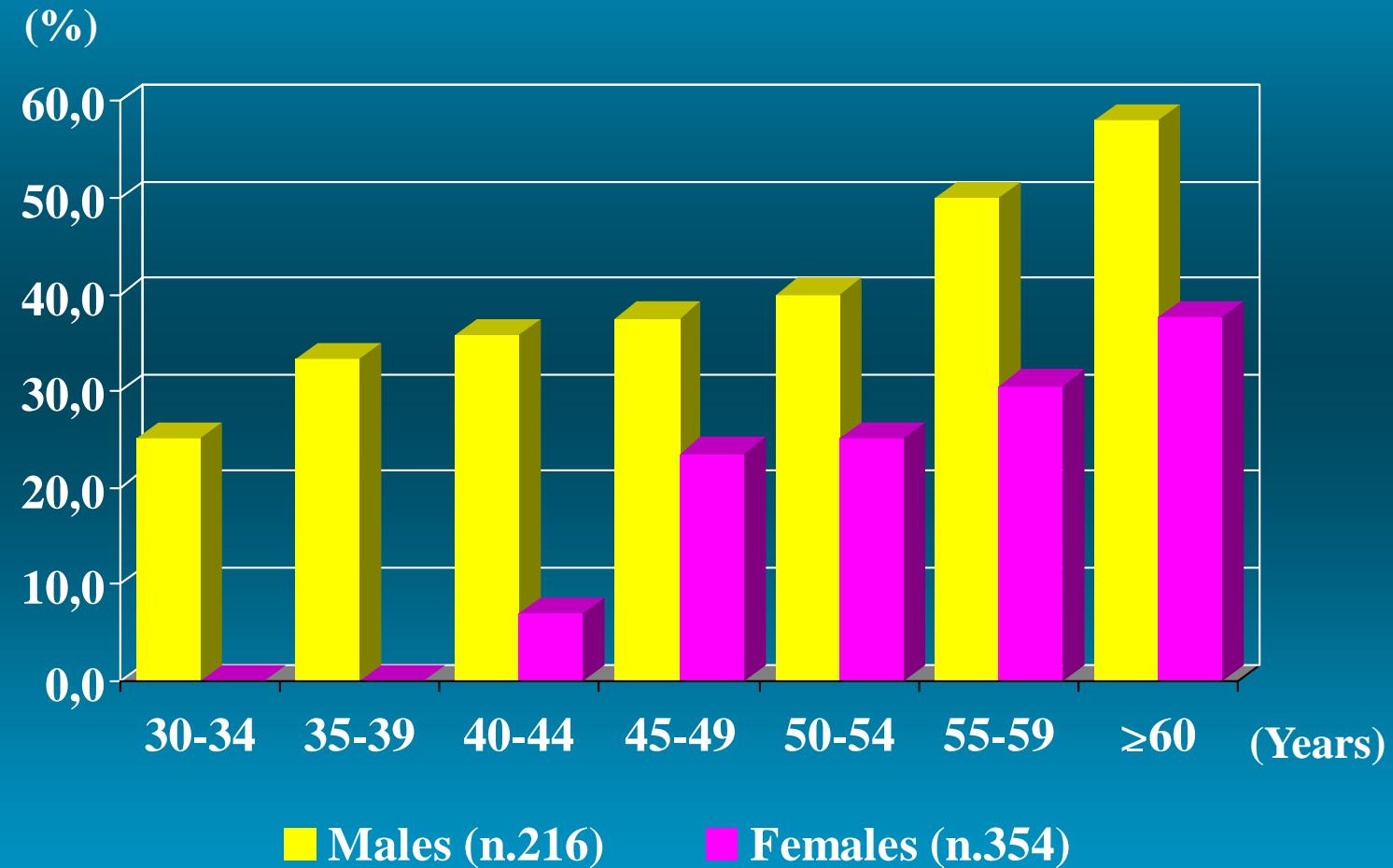
# Soffi carotidei



Chambers BR et al. N Engl J Med 1986;315:860-865



## CHD in Italian FH heterozygotes

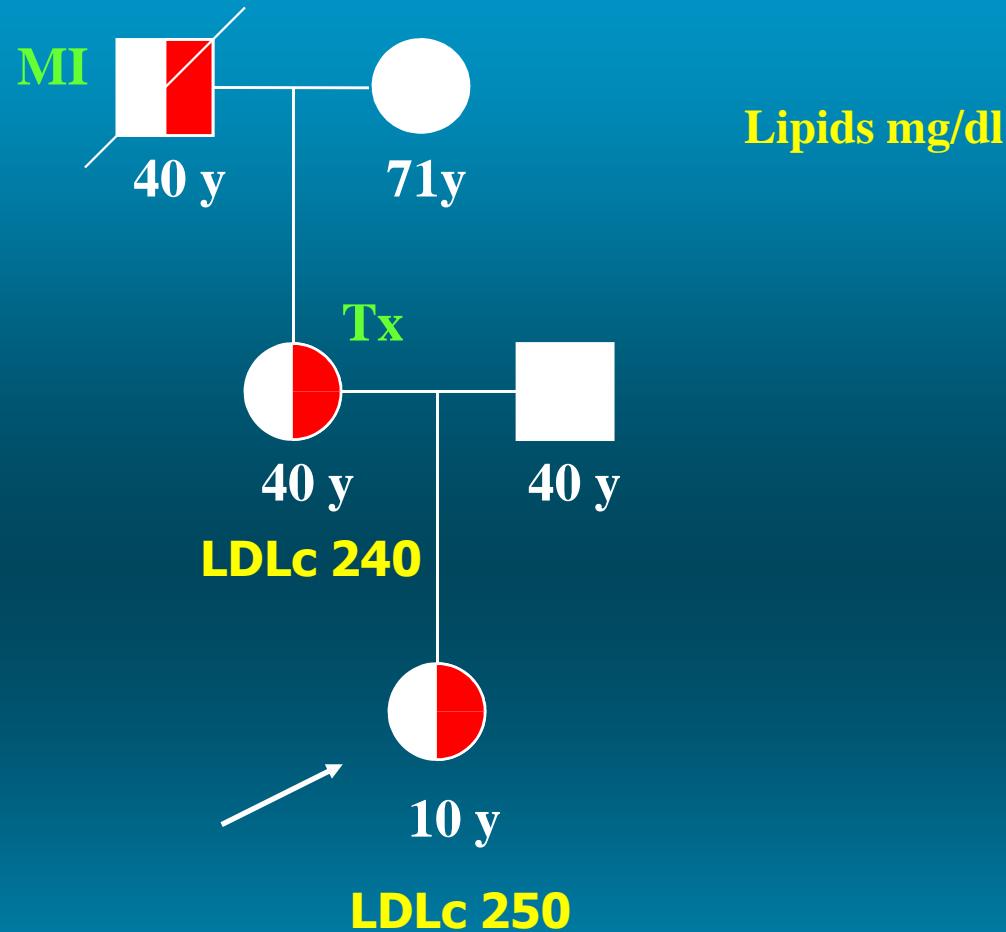


# CLINICAL DIAGNOSIS OF FH

## Clinical History

- Patient with pCAD = 2 p.
- Patient with premature cerebral/peripheral artery disease = 1 p.
- Tendon Xanthomas = 6 p.
- Arcus Cornealis = 4 p.

## CLINICAL DIAGNOSIS OF DEFINITE FH



**Tx = tendon xanthomas**

**MI = myocardial infarction**

# CLINICAL DIAGNOSIS OF FH

## Family History

- First degree relative with pCAD = 1 p.
- First degree relative with LDL-C > 95° = 1 p.
- First degree relative with Tx = 2 p.
- Children <18 with LDL-C >95° = 2 p.

# CLINICAL DIAGNOSIS OF FH

## Final Score

- Definite FH score > 8 p.
  - Probable FH score 6 – 8 p.
  - Possible FH score 3 – 5 p.
  - No diagnosis score < 3 p.



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DISLIPIDEMIA FAMILIARE  
COMBINATA  
FCHL



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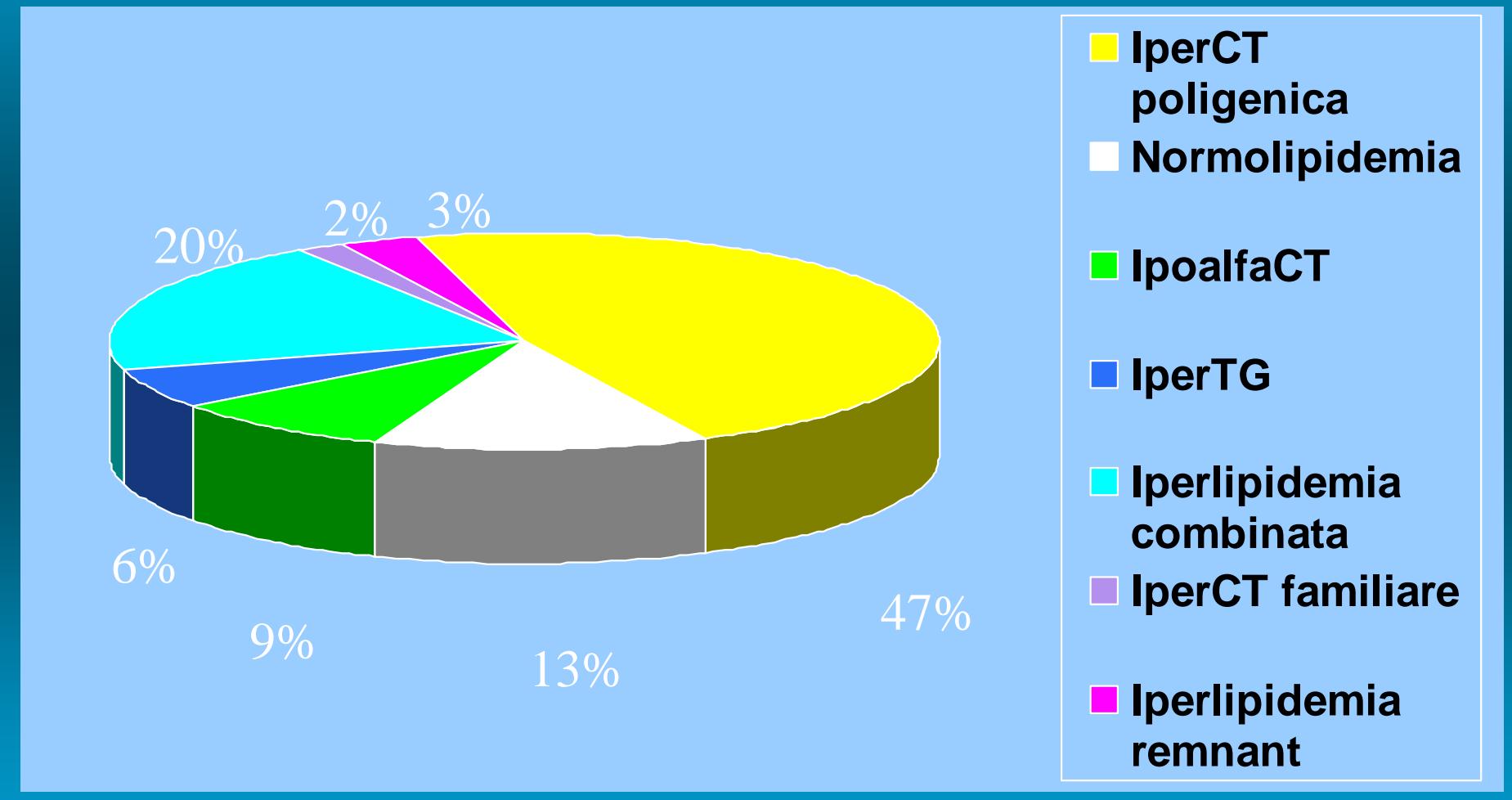


# **Hyperlipidemia in Coronary Heart Disease**

## **II. GENETIC ANALYSIS OF LIPID LEVELS IN 176 FAMILIES AND DELINEATION OF A NEW INHERITED DISORDER, COMBINED HYPERLIPIDEMIA**

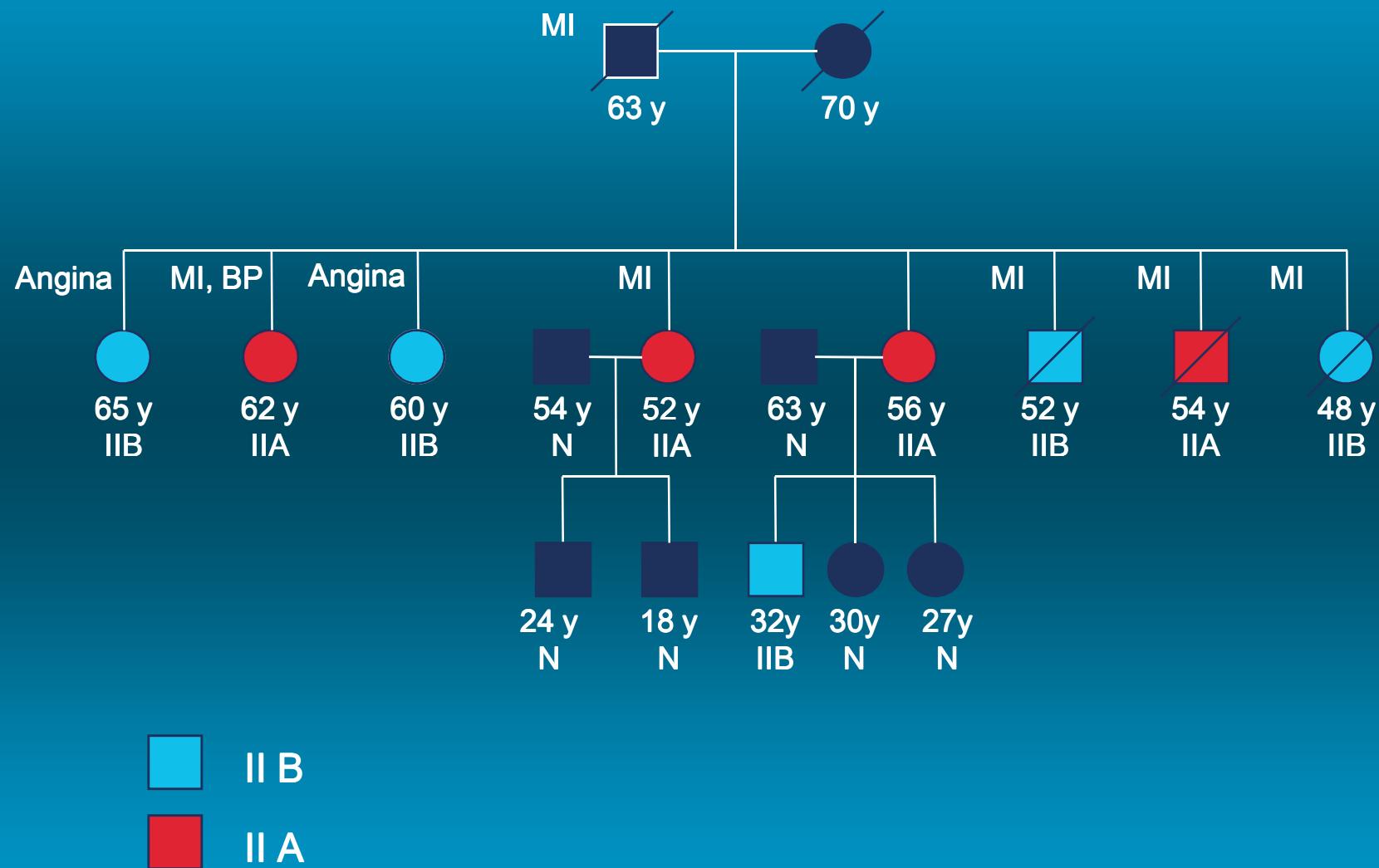
**JOSEPH L. GOLDSTEIN, HELMUT G. SCHROTT, WILLIAM R. HAZZARD,  
EDWIN L. BIERNAN, and ARNO G. MOTULSKY with the technical  
assistance of ELLEN D. CAMPBELL and MARY JO LEVINSKI**

# DISLIPIDEMIA nei sopravvissuti a IMA (n=807)

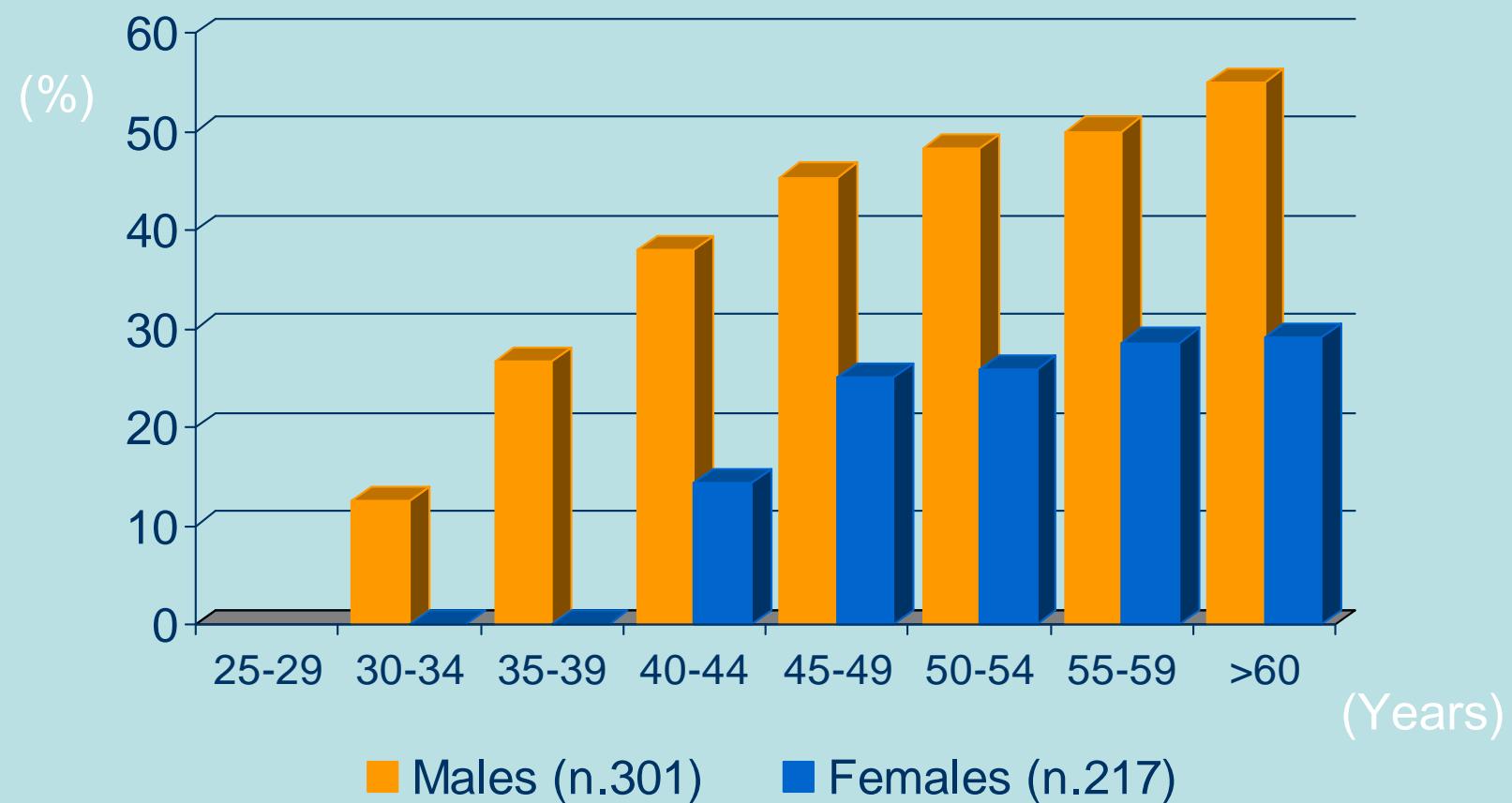


# FAMILIAL COMBINED HYPERLIPIDAEMIA (FCHL)

- ↑ Common (1-3:100)
- ↑ Cholesterol and/or triglyceride levels are elevated, usually to only a moderate extent, reflecting increased LDL and/or VLDL
- ↑ **FAMILIAL**: 1st degree relatives with different phenotypes
- ↑ HDL may be low
- ↑ Characterized by overproduction of apoB-100
- ↑ Coronary heart disease risk increased



# CHD in Italian FCHL



# FCHL nei bambini

---

- La FCHL rappresenta la più frequente (47%) dislipidemia genetica in bambini che afferiscono ad una “lipid clinic” pediatrica.
- La manifestazione fenotipica più frequente è rappresentata da un aumento combinato di CT e TG
- I livelli dei lipidi plasmatici nei bambini con FCHL sono molto “sensibili” al peso corporeo, all’insulinemia (i.e. TG) e all’età (i.e. CT).
- Queste osservazioni suggeriscono che il controllo del peso corporeo può essere un utile approccio terapeutico alla FCHL in età pediatrica.

# Caratteristiche dei bambini

	Famiglie FCHL	Famiglie Normali	
	Affetti (n = 98)	Non affetti (n = 9)	(n = 28)
Età	9.0 ± 4.3	9.9 ± 4.6	8.2 ± 4.3
BMI (kg/m <sup>2</sup> )	19.0 ± 3.7	19.7 ± 2.7	18.3 ± 3.2
Lipidi (mg/dl)			
Colesterolo	216.4 ± 39.7***	158.7 ± 6.6†	148.1 ± 14.7
HDL-C	51.3 ± 14.3	44.0 ± 7.5†	52.0 ± 9.7
Trigliceridi	131.7 ± 74.2***	63.3 ± 16.0	59.9 ± 17.1
LDL-C	138.6 ± 42.6**	102.0 ± 11.9††	84.1 ± 16.5
Apo B	97.6 ± 30.0**	71.8 ± 8.4 †	59.2 ± 11.6
Glicemia	82.5 ± 12.0	83.5 ± 21.7 ††	64.9 ± 9.2
Insulina (μU/ml)	8.7 ± 6.8	8.9 ± 6.4	5.7 ± 4.0

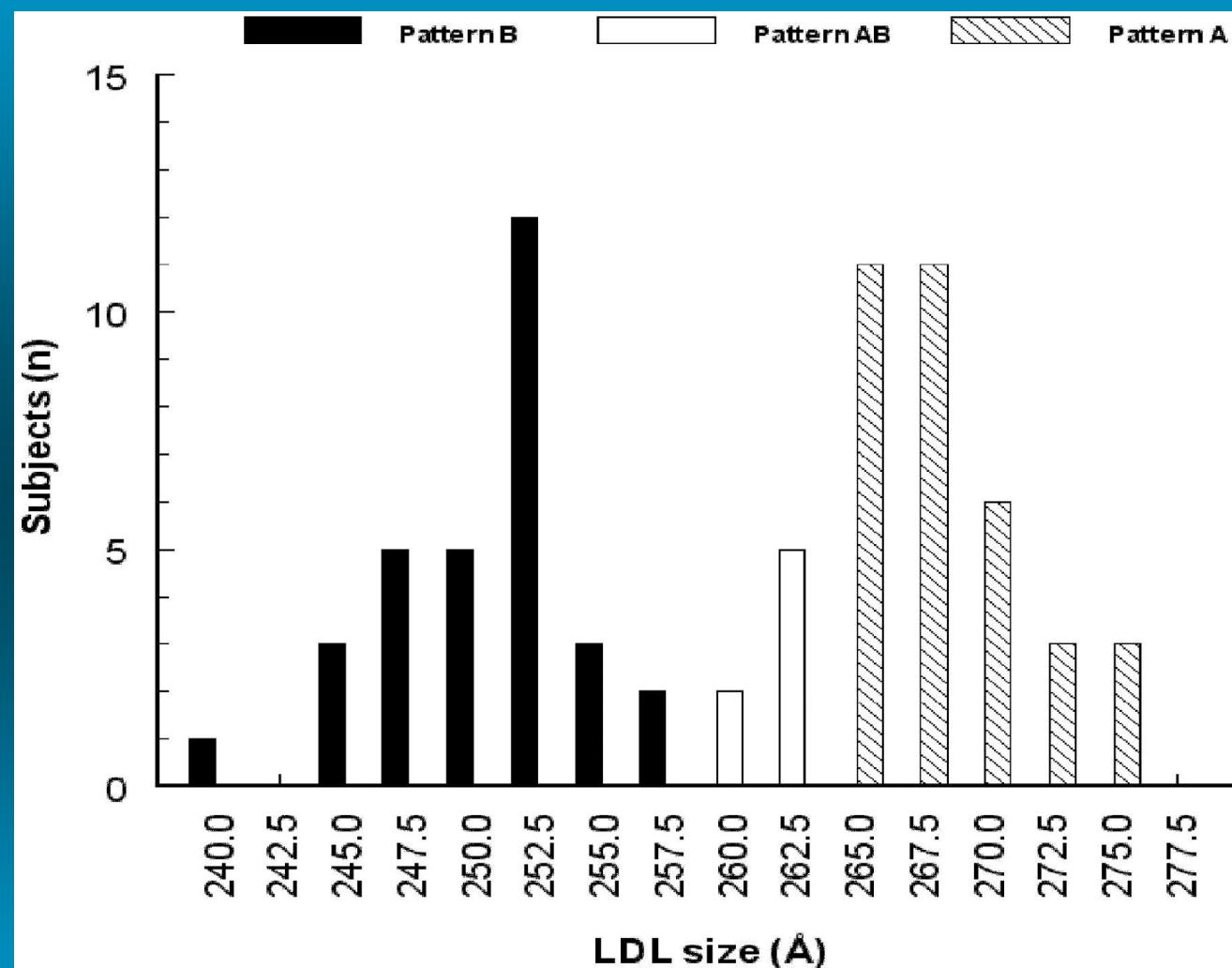
\*\*p<0.001; \*\*\*p<0.001 affetti FCHL vs non affetti    †p<0.05; ††p<0.01 non affetti FCHL vs Normali

# Caratteristiche dei genitori

	Famiglie FCHL	Famiglie Normali	
	Affetti (n = 61)	Non affetti (n = 51)	(n = 38)
Età	<b>40.1 ± 5.9</b>	<b>39.3 ± 4.6</b>	<b>37.5 ± 6.0</b>
BMI (kg/m <sup>2</sup> )	<b>26.8 ± 3.7***</b>	<b>24.1 ± 3.4</b>	<b>24.7 ± 3.9</b>
Lipidi (mg/dl)			
Colesterolemia	<b>246.5 ± 58.7***</b>	<b>186.3 ± 28.2</b>	<b>181.8 ± 21.4</b>
HDL-C	<b>47.1 ± 17.0</b>	<b>48.7 ± 9.4††</b>	<b>55.7 ± 15.6</b>
Trigliceridemia	<b>265.9 ± 183.6***</b>	<b>91.1 ± 28.7</b>	<b>85.8 ± 32.5</b>
LDL-C	<b>150.1 ± 65.1**</b>	<b>119.3 ± 24.8†</b>	<b>109.0 ± 24.6</b>
Apo B	<b>140.5 ± 50.0**</b>	<b>91.8 ± 19.1</b>	<b>83.8 ± 18.9</b>
Glicemia	<b>93.0 ± 31.1*</b>	<b>81.4 ± 15.4†††</b>	<b>69.0 ± 13.1</b>
Insulinemia (uU/ml)	<b>10.0 ± 6.1</b>	<b>8.6 ± 13.8</b>	<b>6.6 ± 4.7</b>

\*p<0.05; \*\*p<0.01; \*\*\*p<0.001 affetti vs non affetti    †p<0.05; ††p<0.01; †††p<0.001 non affetti vs normali

## Frequency distribution of LDL size in hyperlipidemic FCHL relatives

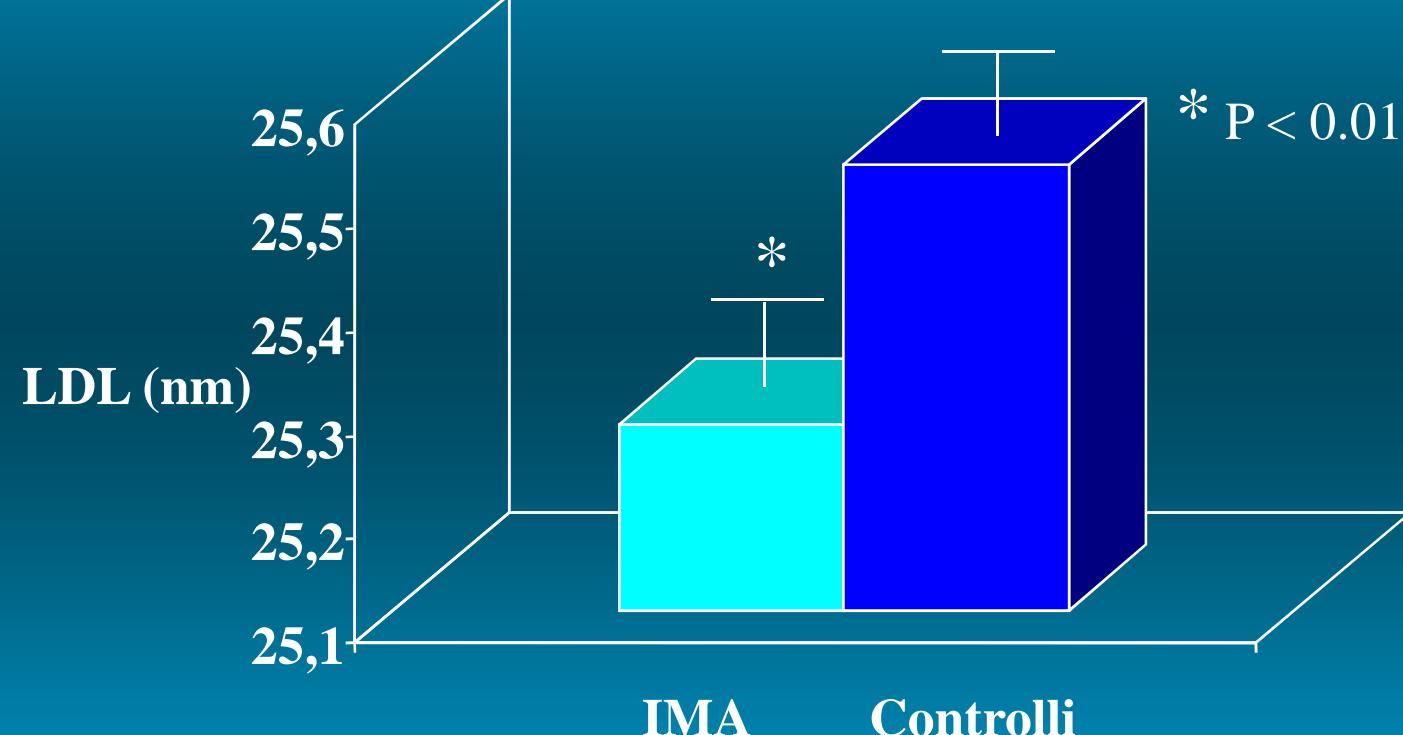


Georgieva, A.M. et al. Arterioscler Thromb Vasc Biol 2004;24:744-749

Arteriosclerosis, Thrombosis,  
and Vascular Biology

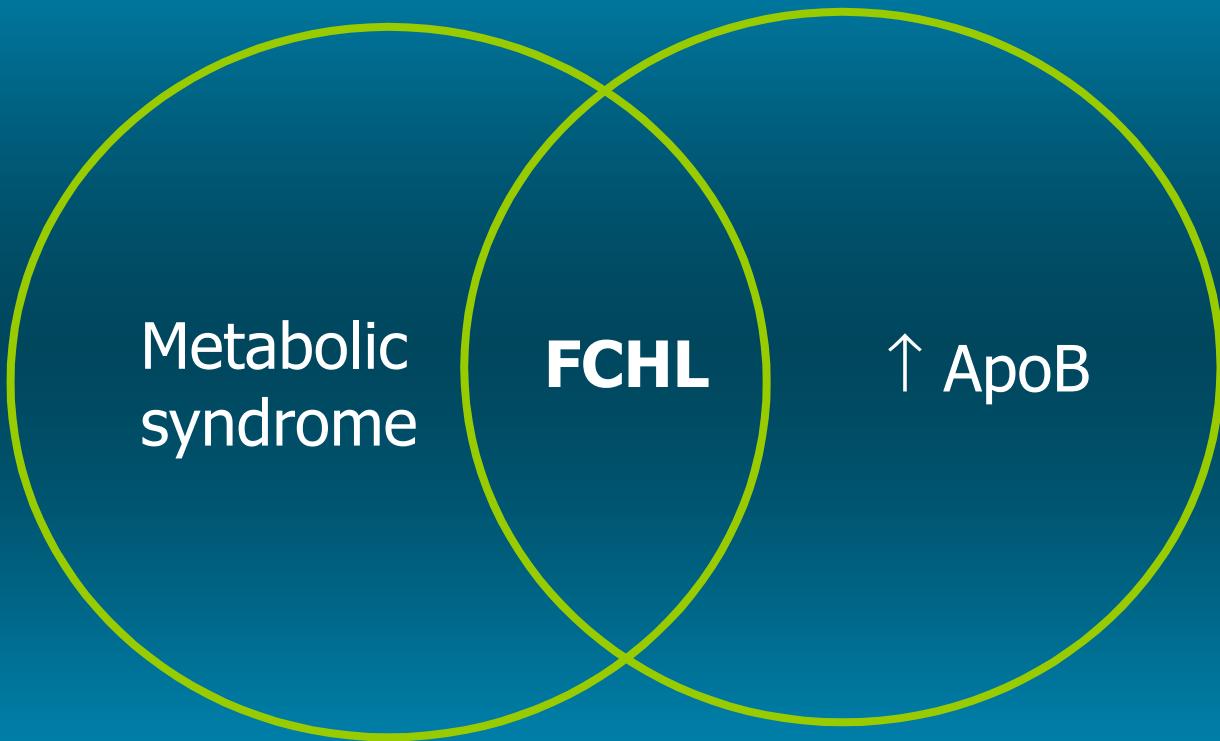


# Dimensioni delle LDL negli infartuati e nei controlli



Valori corretti per colesterolemia HDL, trigliceridemia, colesterolemia totale, sesso, diabete, ipertensione arteriosa, familiarità cardiovascolare





*Arteriosclerosis, Thrombosis, and Vascular Biology.* 2003;23:1289



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# LIPOPROTEINA(a) Lp(a)



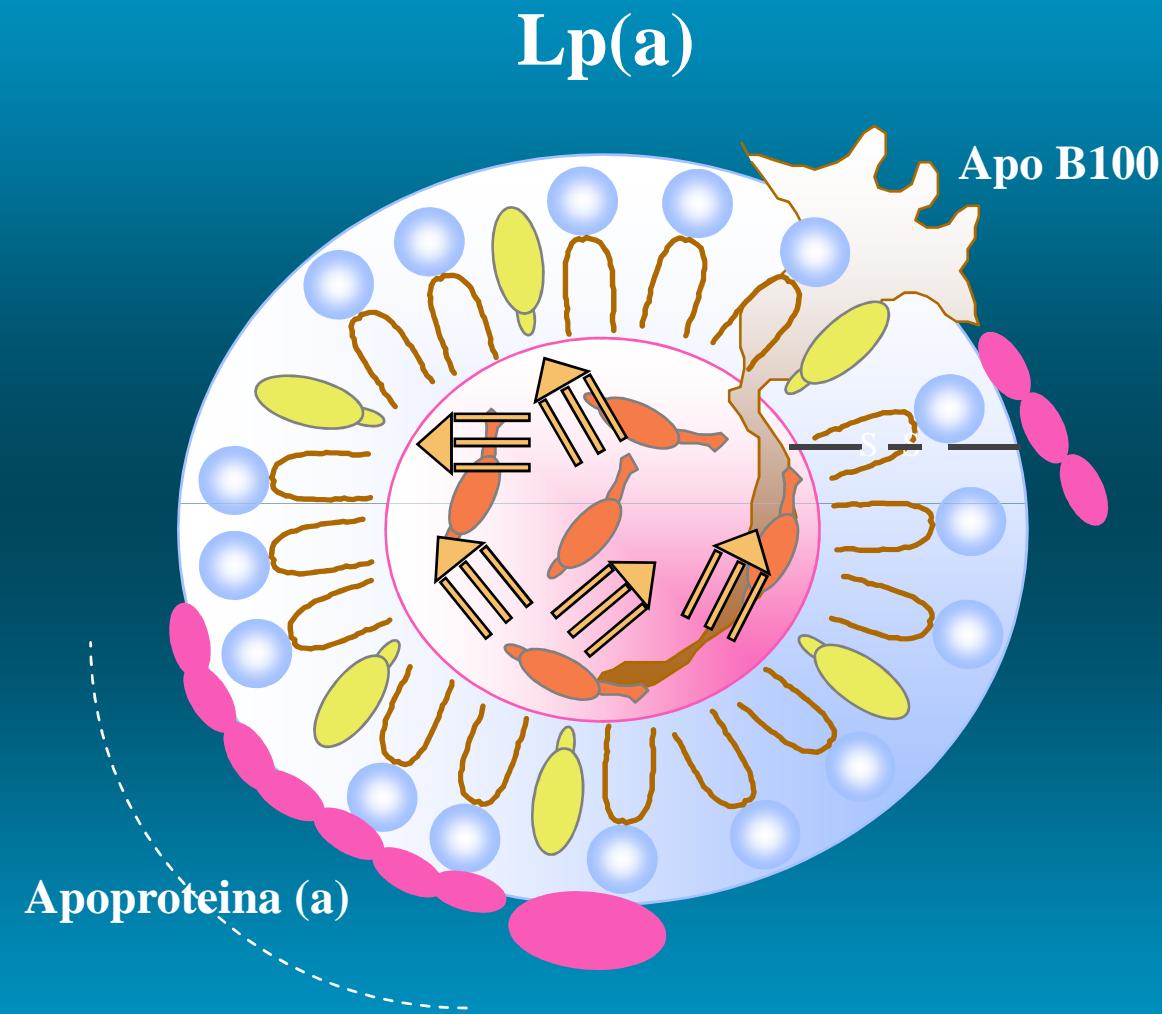
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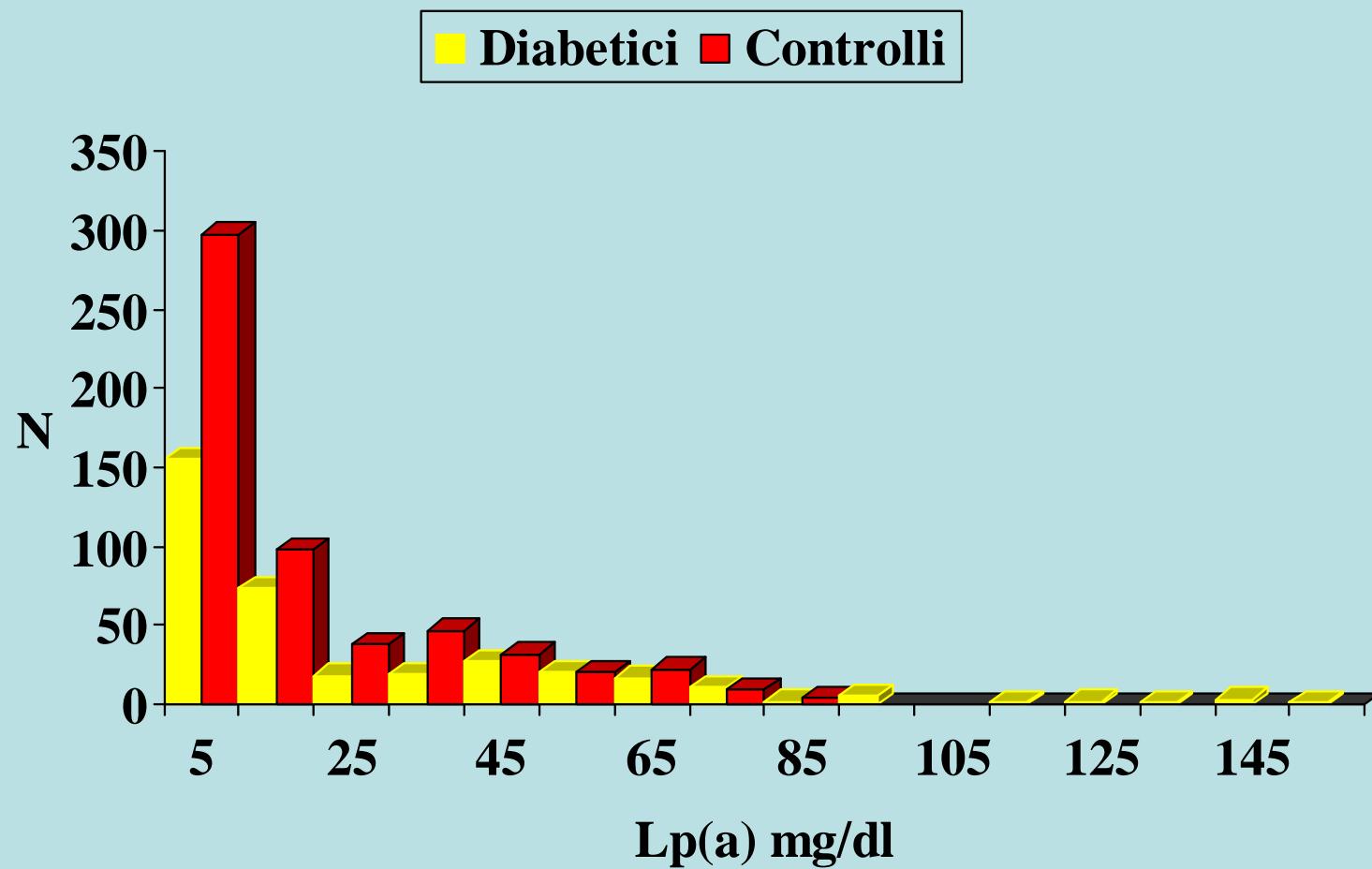




**Plasminogeno**

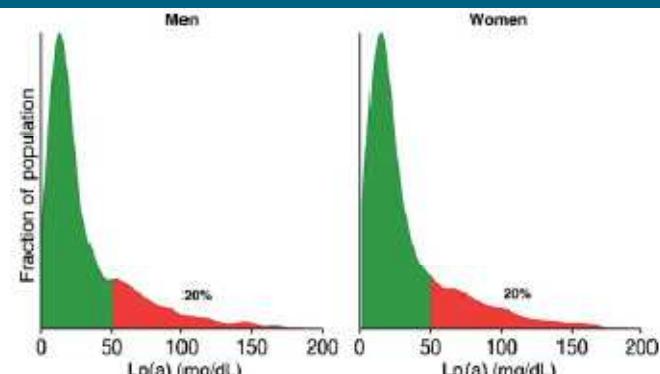
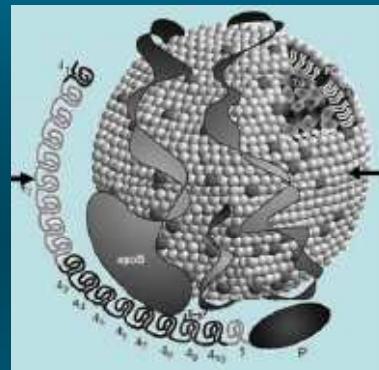


# DISTRIBUZIONE DI Lp(a) IN DIABETICI (N=355) E CONTROLLI SANI (N=568)

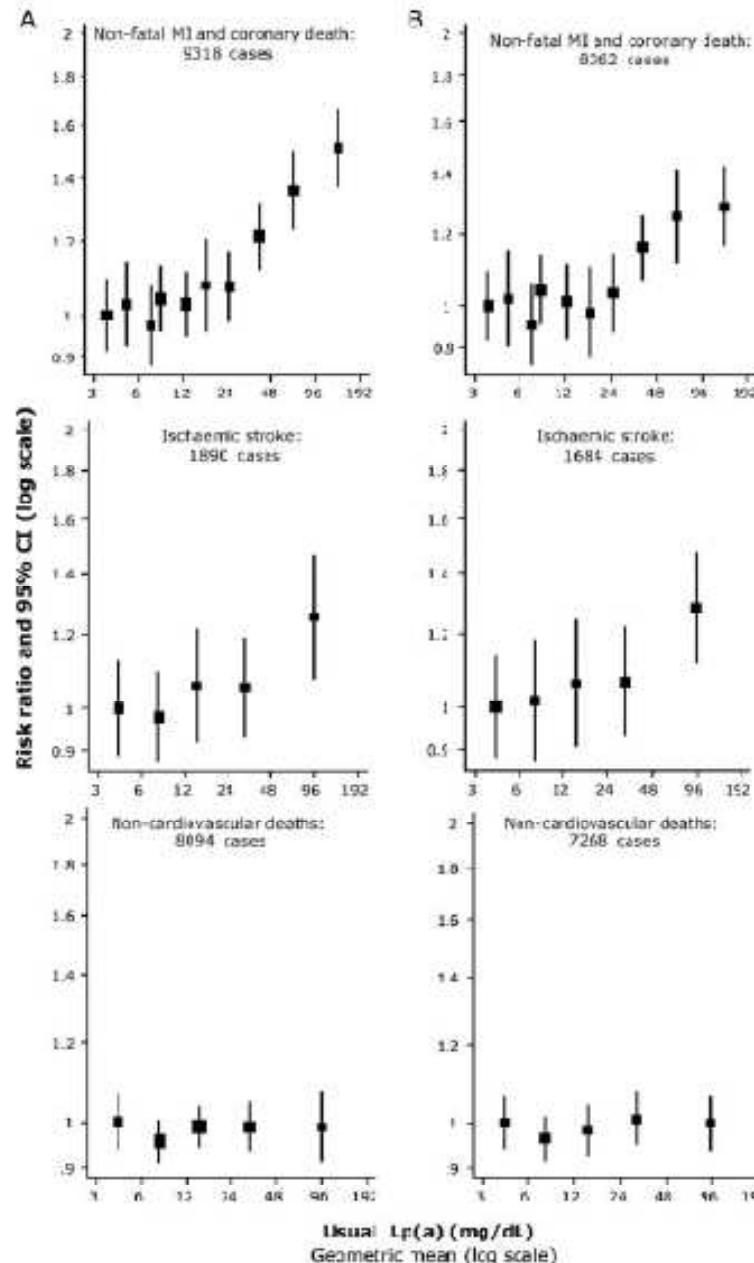


## Lipoprotein(a) as a cardiovascular risk factor: current status

Børge G. Nordestgaard<sup>1\*</sup>, M. John Chapman<sup>2</sup>, Kausik Ray<sup>3</sup>, Jan Borén<sup>4</sup>, Felicita Andreotti<sup>5</sup>, Gerald F. Watts<sup>6</sup>, Henry Ginsberg<sup>7</sup>, Pierre Amarenco<sup>8</sup>, Alberico Catapano<sup>9</sup>, Olivier S. Descamps<sup>10</sup>, Edward Fisher<sup>11</sup>, Petri T. Kovánen<sup>12</sup>, Jan Albert Kuivenhoven<sup>13</sup>, Philippe Lesnik<sup>2</sup>, Luis Masana<sup>14</sup>, Zeljko Reiner<sup>15</sup>, Marja-Riitta Taskinen<sup>16</sup>, Lale Tokgozoglu<sup>17</sup>, and Anne Tybjærg-Hansen<sup>18</sup>, for the European Atherosclerosis Society Consensus Panel<sup>†</sup>

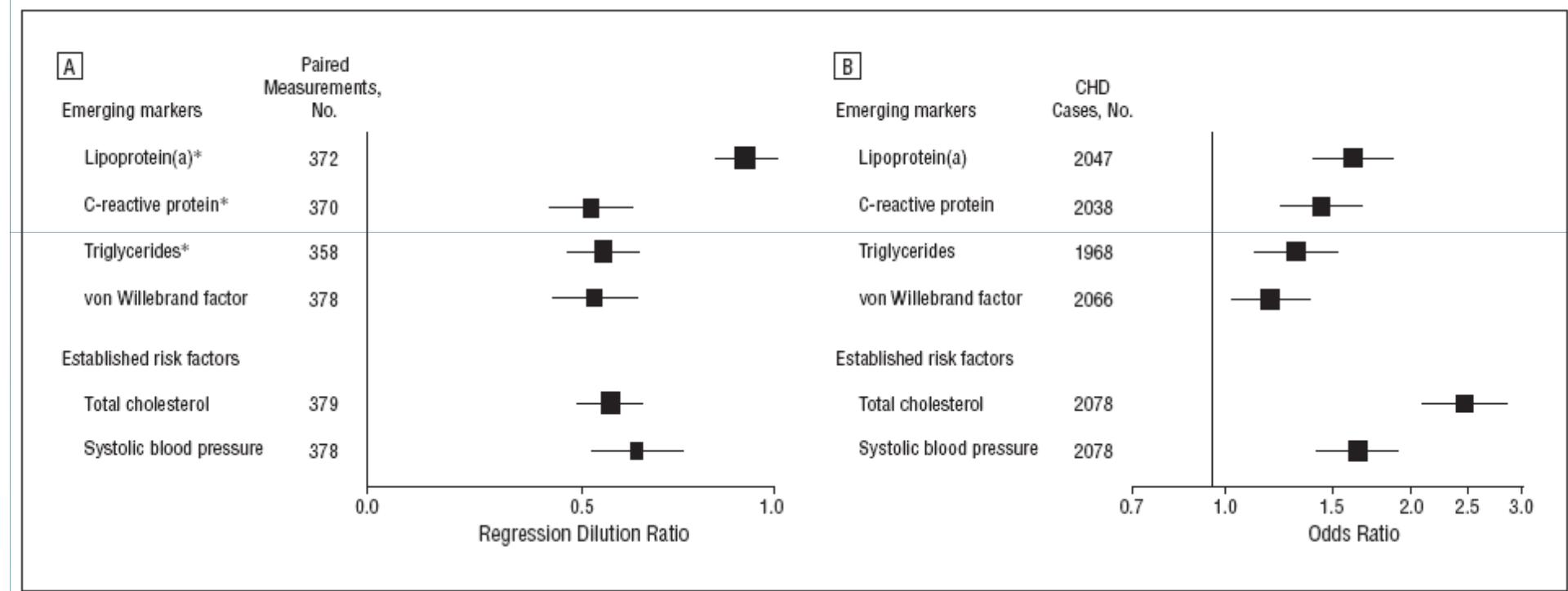


**Figure 3** Risk ratios of coronary heart disease, ischaemic stroke and non-cardiovascular death by quantiles of usual lipoprotein(a) levels. CI, confidence interval. Sizes of data markers are proportional to the inverse of the variance of the risk ratios. (A) Adjustment for age and sex only. (B) Further adjustment for systolic blood pressure, smoking status, history of diabetes, body mass index, and total cholesterol. MI, myocardial infarction. Modified from The Emerging Risk Factors Collaboration.<sup>3</sup>



# Lipoprotein(a) Levels and Risk of Future Coronary Heart Disease

*Large-Scale Prospective Data*



Comparison of Lp(a) values, established cardiovascular risk factors, and emerging markers across 12 years

ORIGINAL ARTICLE

## Genetic Variants Associated with Lp(a) Lipoprotein Level and Coronary Disease

Robert Clarke, F.R.C.P., John F. Peden, Ph.D., Jemma C. Hopewell, Ph.D., Theodosios Kyriakou, Ph.D., Anuj Goel, M.Sc., Simon C. Heath, Ph.D., Sarah Parish, D.Phil., Simona Barlera, M.S., Maria Grazia Franzosi, Ph.D., Stephan Rust, Ph.D., Derrick Bennett, Ph.D., Angela Silveira, Ph.D., Anders Malarstig, Ph.D., Fiona R. Green, Ph.D., Mark Lathrop, Ph.D., Bruna Gigante, M.D., Karin Leander, Ph.D., Ulf de Faire, M.D., Udo Seedorf, Ph.D., Anders Hamsten, F.R.C.P., Rory Collins, F.R.C.P., Hugh Watkins, F.R.C.P., and Martin Farrall, F.R.C.Path., for the PROCARDIS Consortium\*

ABSTRACT

### METHODS

We used a novel gene chip containing 48,742 single-nucleotide polymorphisms (SNPs) in 2100 candidate genes to test for associations in 3145 case subjects with coronary disease and 3352 control subjects. Replication was tested in three independent populations involving 4846 additional case subjects with coronary disease and 4594 control subjects.

### CONCLUSIONS

We identified two *LPA* variants that were strongly associated with both an increased level of Lp(a) lipoprotein and an increased risk of coronary disease. Our findings provide support for a causal role of Lp(a) lipoprotein in coronary disease.





N Engl J Med 2009;361:2518-28.

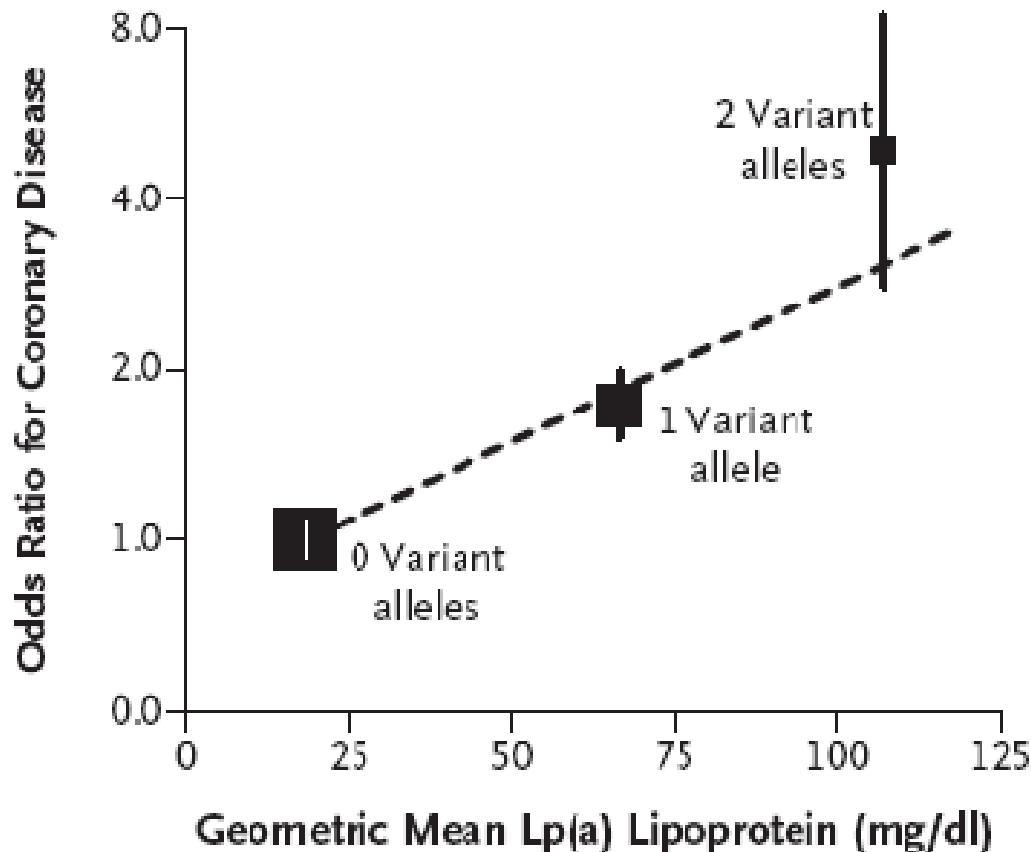
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ABSTRACT



**Figure 3. Association of the LPA Genotype Score with the Lp(a) Lipoprotein Level and the Risk of Coronary Disease in the PROCARDIS Cohort.**

# CONCLUSIONI

- identificazione precoce dell'Ipercolesterolemia Familiare attraverso anamnesi, esame obiettivo ed assetto lipidico esteso ai familiari di 1° grado;
- identificazione della Dislipidemia Familiare Combinata attraverso anamnesi, esame obiettivo ed assetto lipidico esteso ai familiari di 1° grado;
- valutazione vascolare periodica al fine di identificare l' arteriosclerosi in fase precoce;



ottimizzazione della terapia farmacologica

↓  
**LDL aferesi**

