



MOLECULAR PATHOLOGY OF ATHEROSCLEROSIS

Dubrovnik 2013

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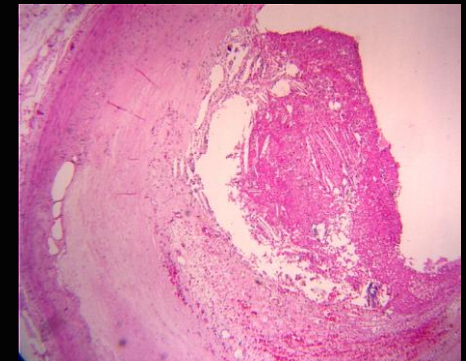
IIB-Sant Pau, Hospital de la Santa Creu i Sant Pau

ATHEROSCLEROSIS – ATHEROTHROMBOSIS – CLINICAL EVENTS

STROKE
TIAs

ACS

PAD



CARDIOVASCULAR DISEASE - ATHEROTHROMBOSIS



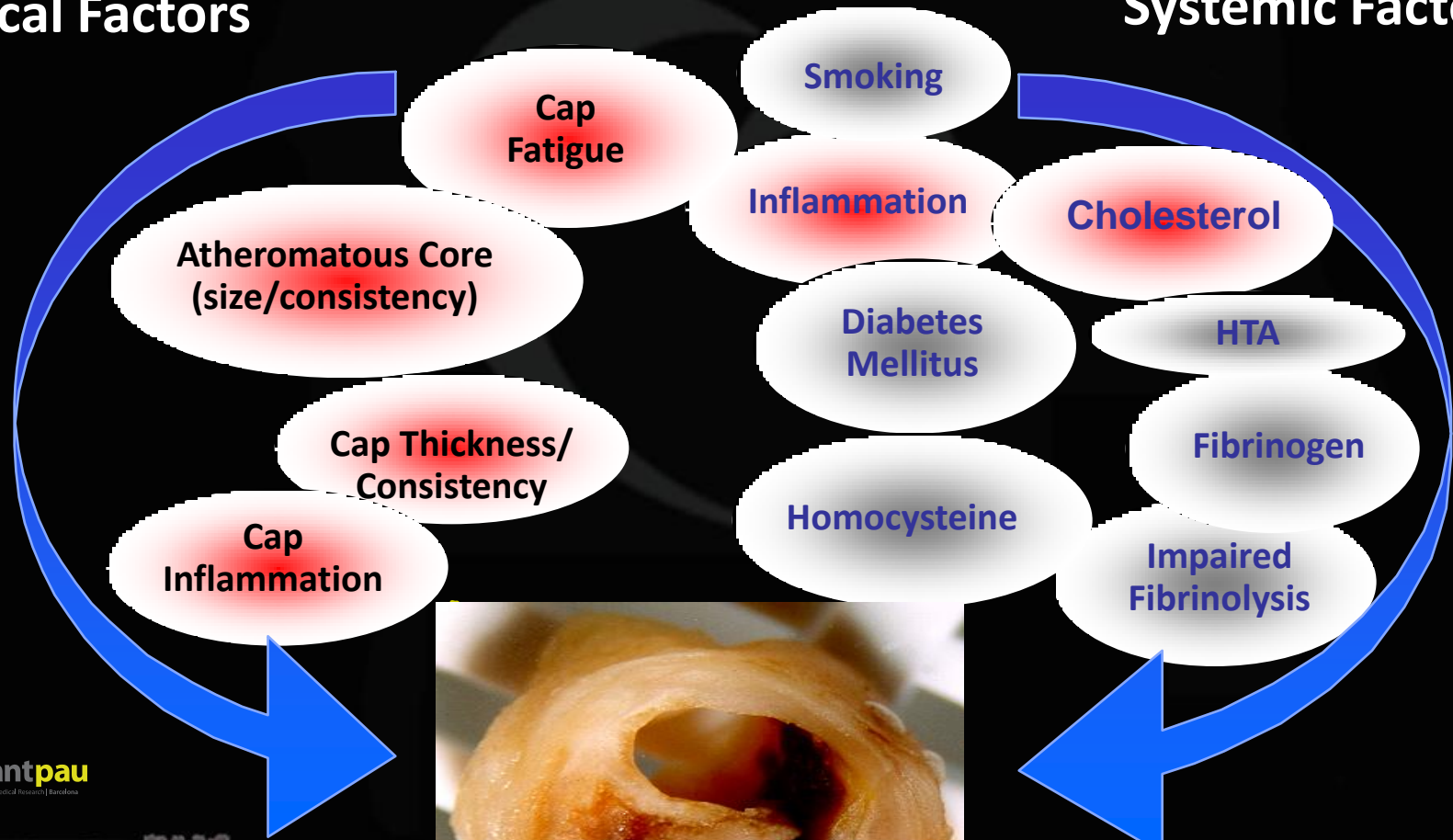
Acute Ischemic Syndromes

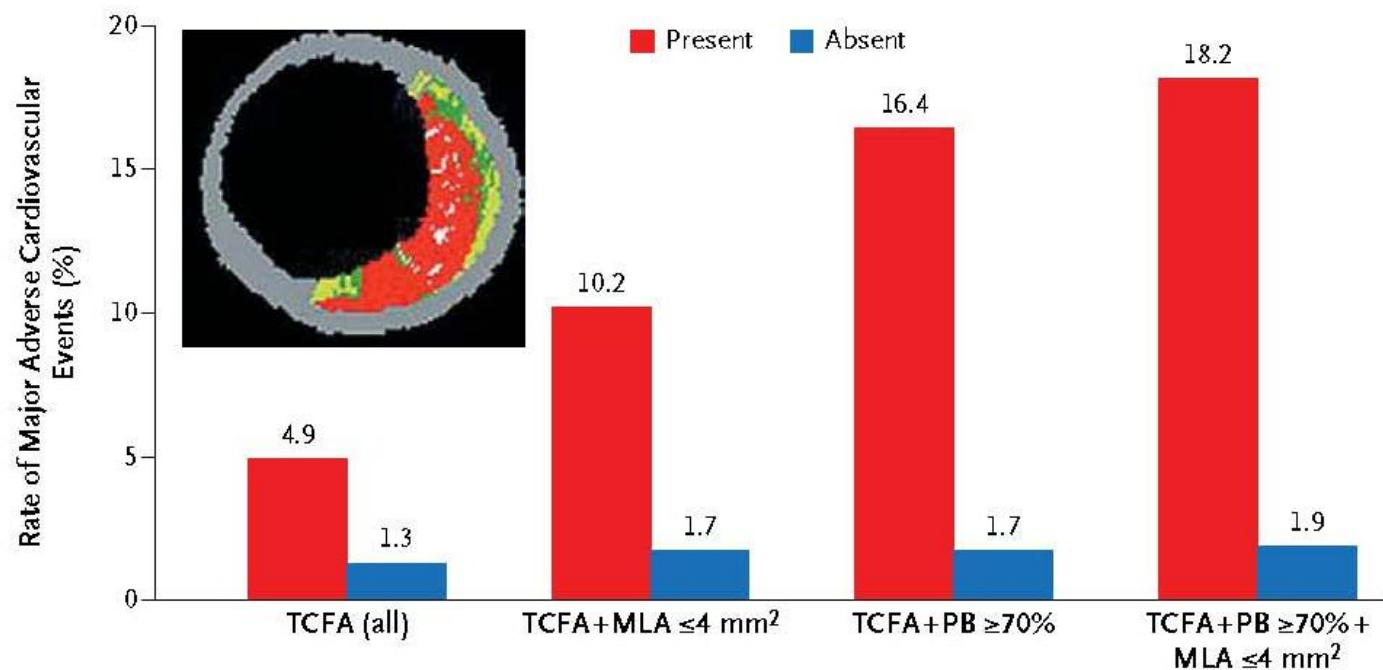
- **Contenders:**
 - Platelets
 - White cells & RBCs
 - EPCs
 - Inflammation
 - Microparticles

Risk Factors for Plaque Progression and Clinical Complication

Local Factors

Systemic Factors



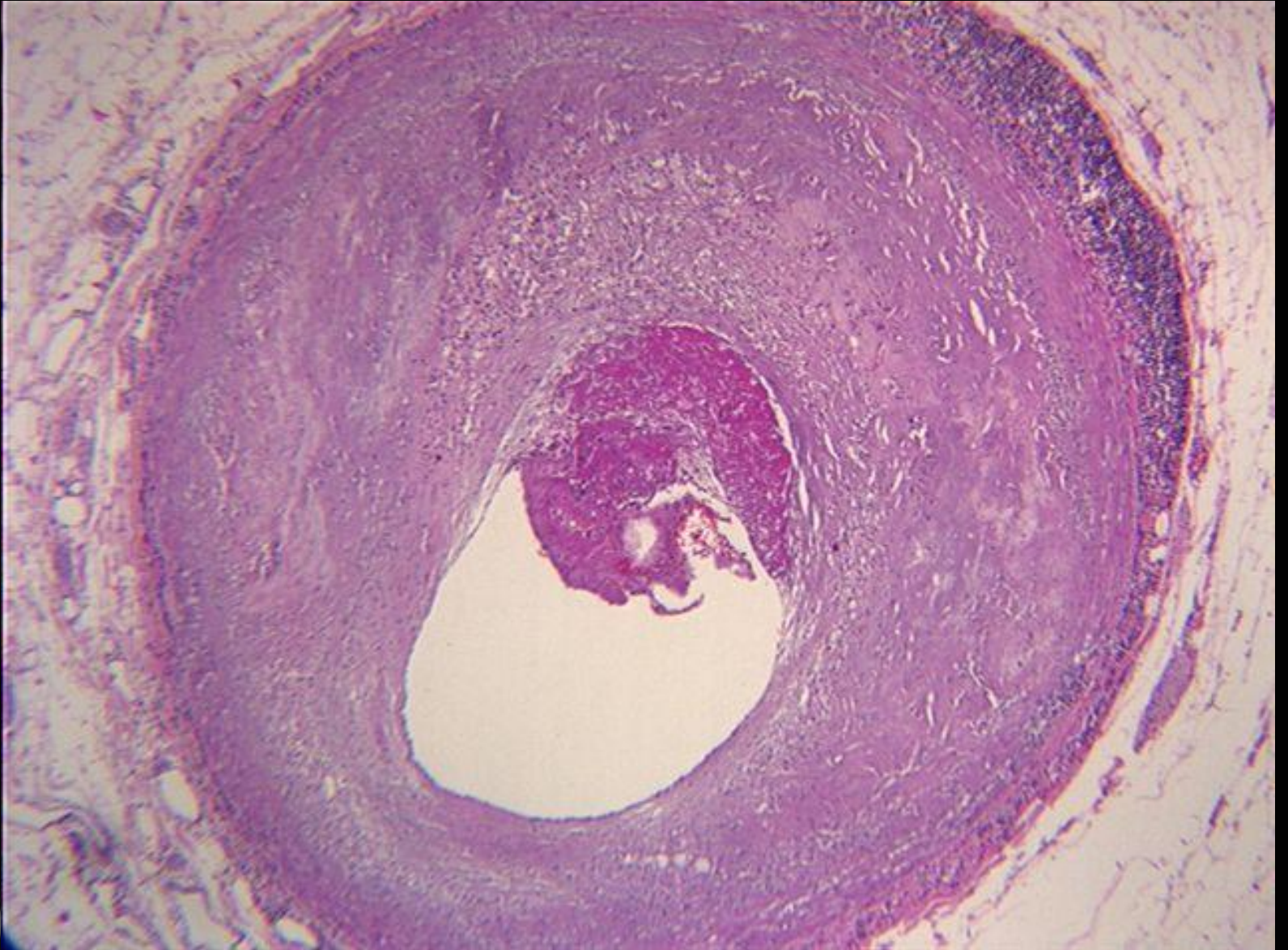


Lesion hazard ratio (95% CI)	3.90 (2.25–6.76)	6.55 (3.43–12.51)	10.83 (5.55–21.10)	11.05 (4.39–27.82)
P value	<0.001	<0.001	<0.001	<0.001
Prevalence (%)	46.7	15.9	10.1	4.2

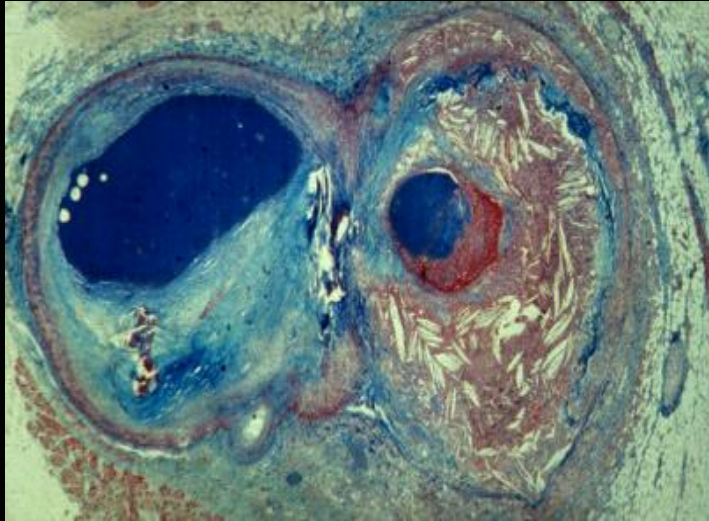
Figure 2. Event Rates for Lesions That Were and Those That Were Not Thin-Cap Fibroatheromas, at a Median Follow-up of 3.4 Years.

Event rates associated with 595 nonculprit lesions that were characterized as thin-cap fibroatheromas (TCFA) and 2114 that were not by means of radiofrequency intravascular ultrasonographic imaging are shown according to minimal luminal area (MLA) and plaque burden (PB) as detected on gray-scale intravascular ultrasonography. The inset shows an example of a thin-cap fibroatheroma imaged by radiofrequency ultrasonography. Data on prevalence are for one or more such lesions per patient. Lesions in patients with indeterminate events were excluded. (For additional details, see Table 6 in the Supplementary Appendix.) CI denotes confidence interval.

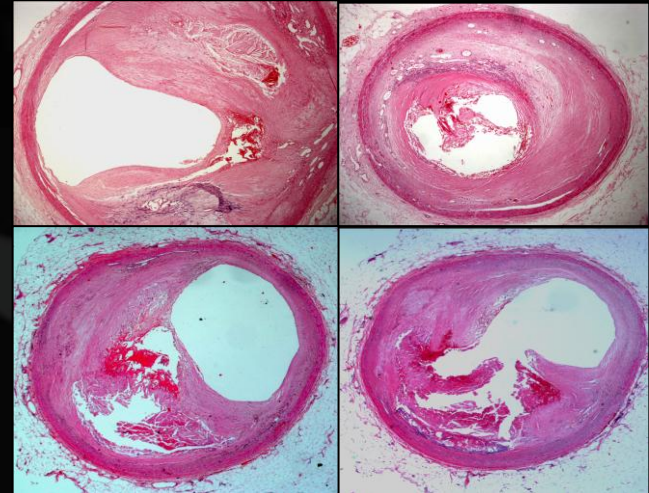




HIGH RISK PLAQUES AND STRUCTURAL CHALLENGES

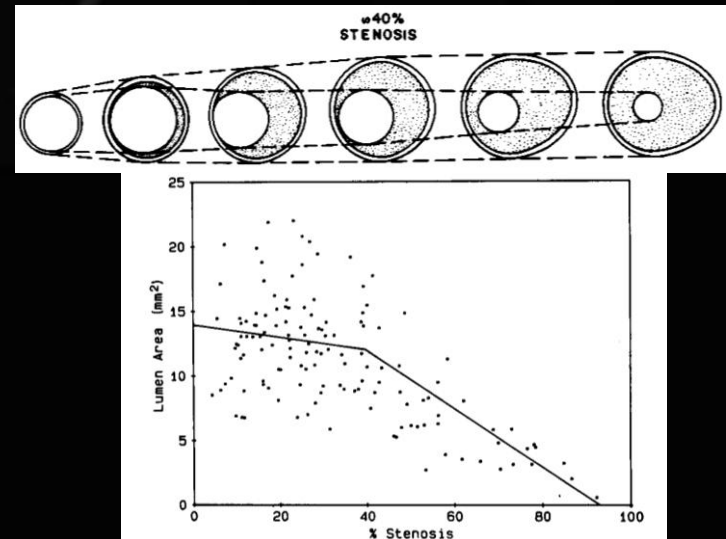


Falk E et al



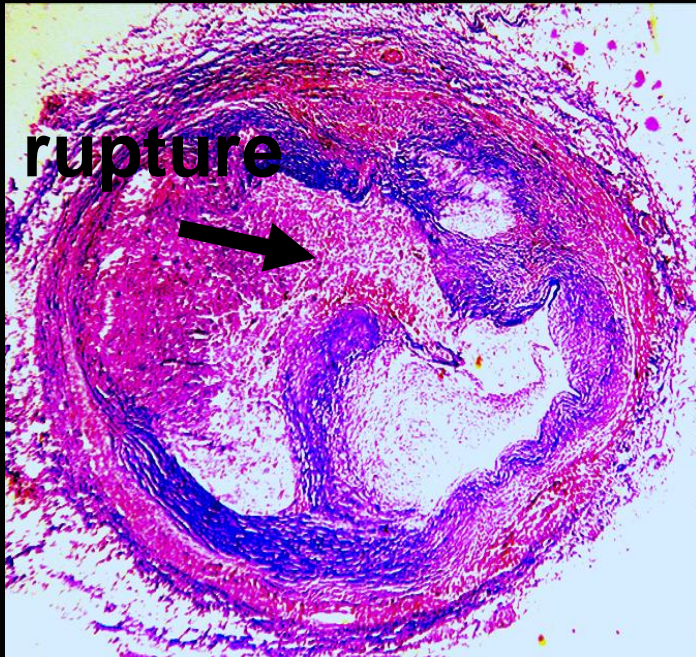
Badimon L, Juan O.

- Unstable plaque
 - Low in collagen
 - Necrotic Core
(rich in **lipids**, inflammatory cells, lymphocytes, microcalcifications)
 - Thin fibrous cap (<65um)



Glagov S, N Engl J Med 1987 May 28, 316(22)1371-5

HIGH RISK PLAQUES



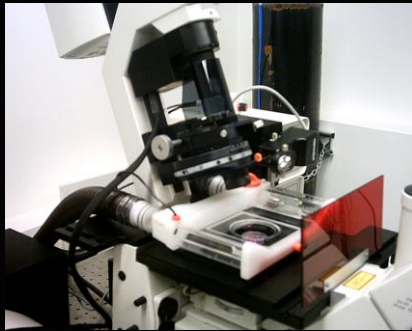
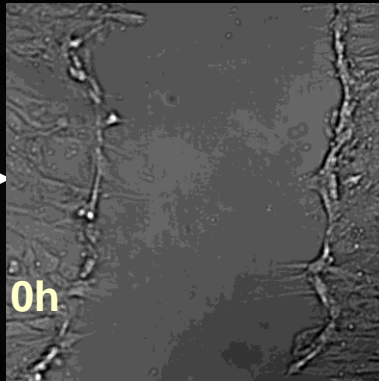
- ↑ lipid core
- ↓ smooth muscle cells
- ↓ collagen fibers
- ↑ inflammatory cells
- ↑ necrotic core/thin fibrous plaque

CV RISK FACTORS



Infiltrated lipids impair human coronary VSMC repair mechanisms

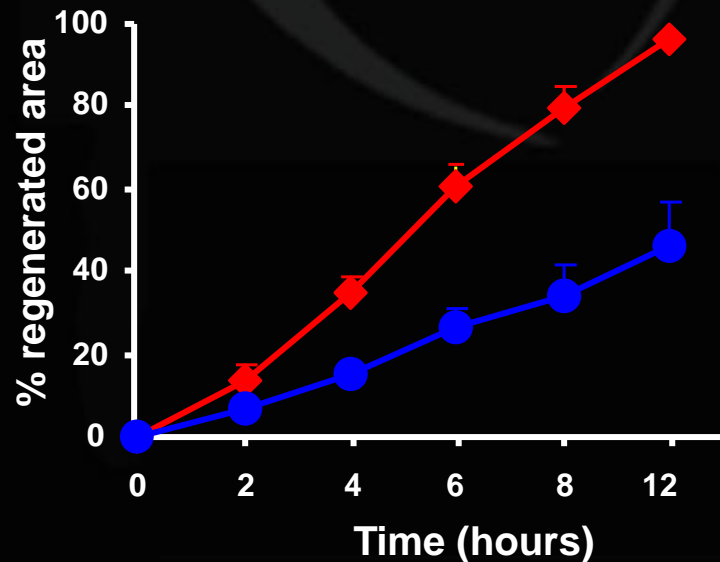
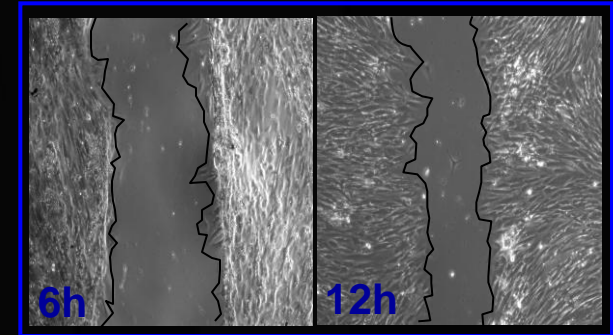
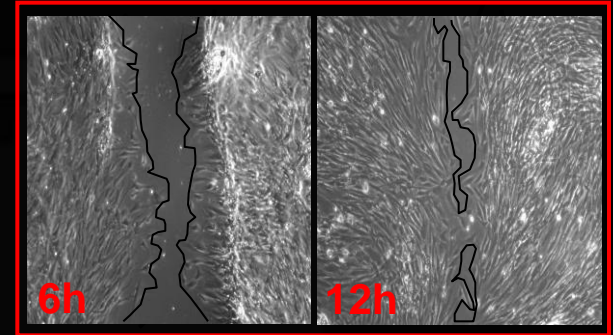
VSMC
+/- agLDL
(16 hours)



double sided
scrape-wound

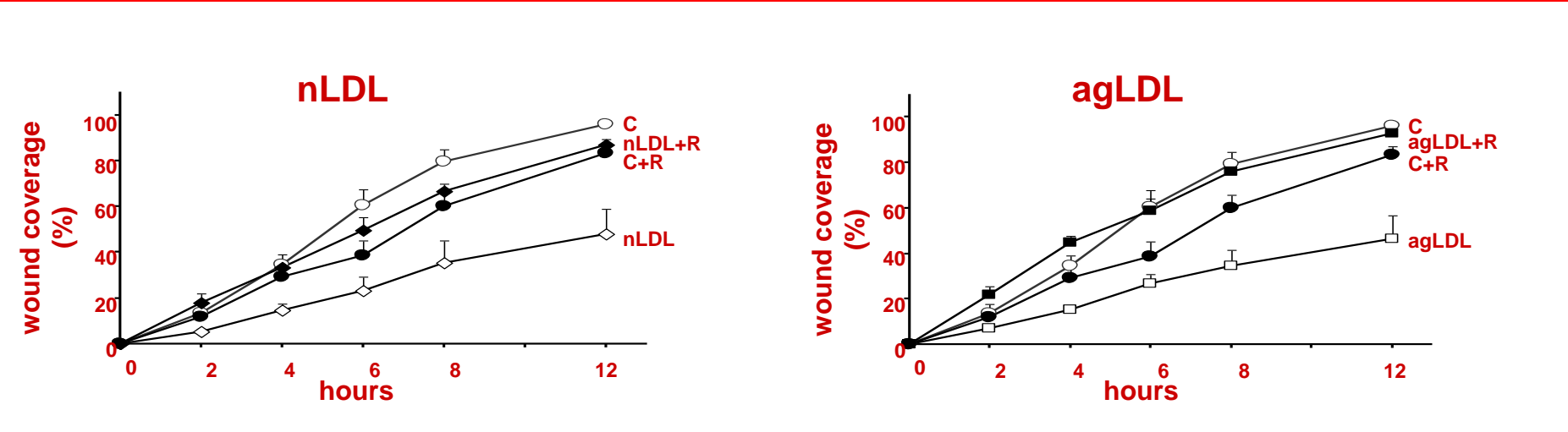
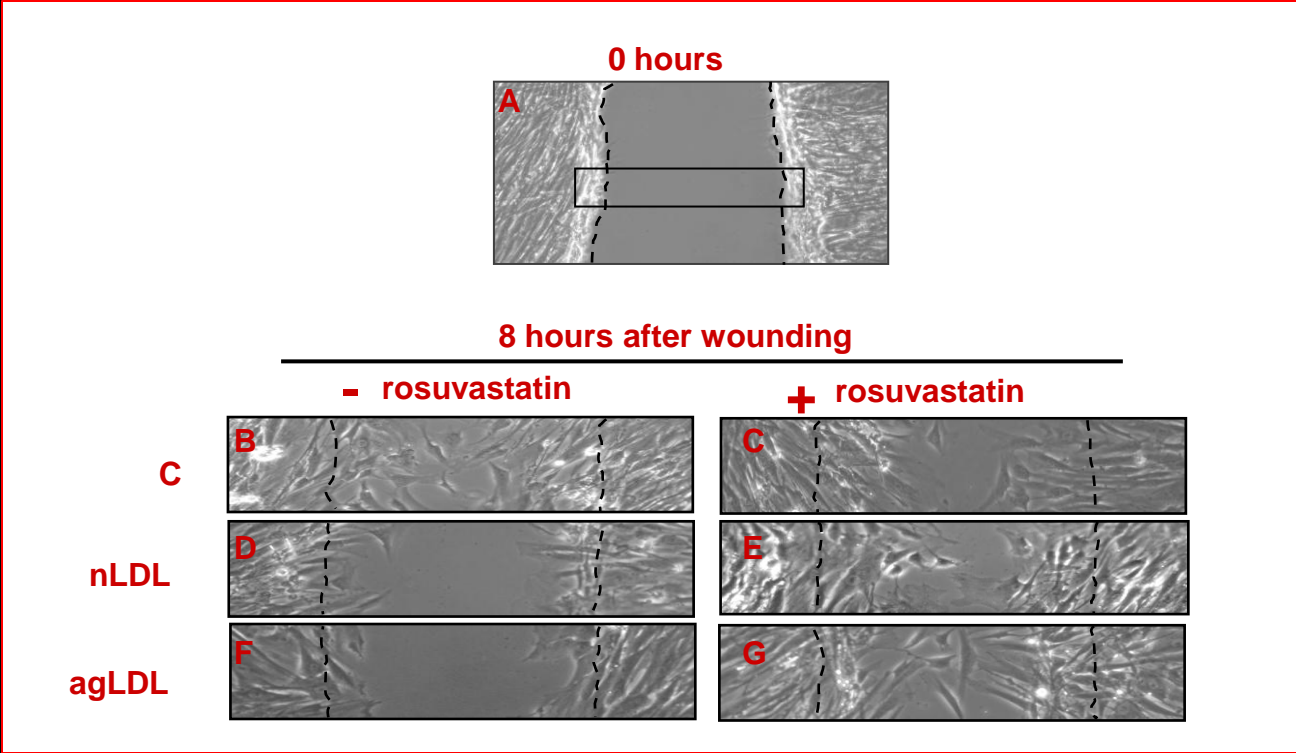
control

agLDL

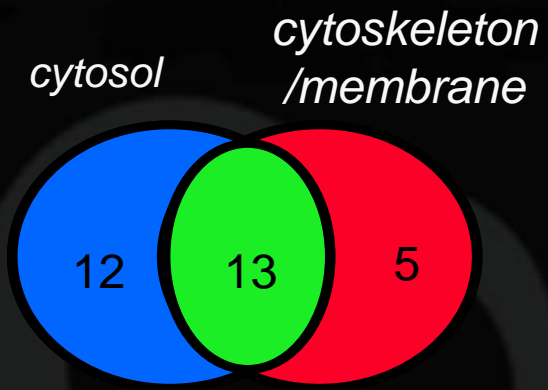


control

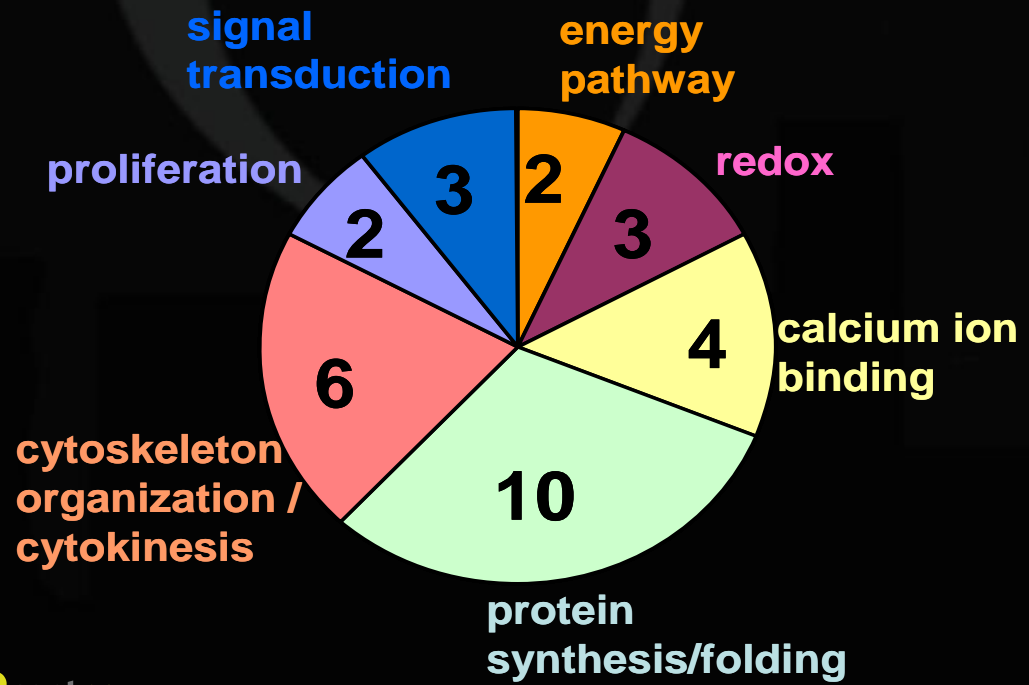
agLDL



Infiltrated lipids impair human coronary VSMC repair mechanisms



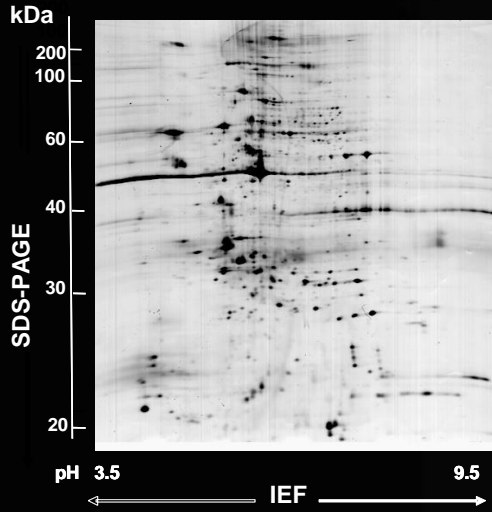
- MRLC
- MLC
- caldesmon
- actinin
- tropomyosin
- transgelin



Proteome of human coronary SMC

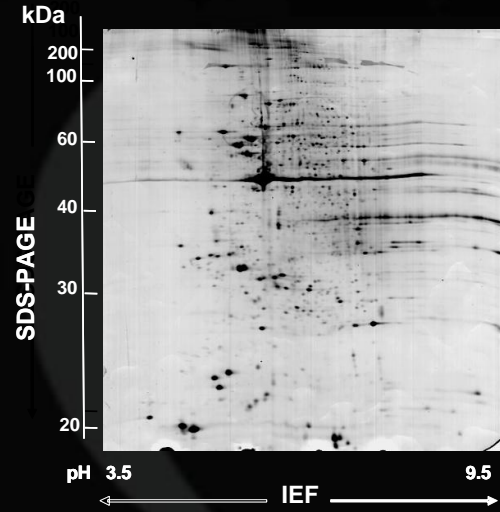
Cytosol

Tris Fraction

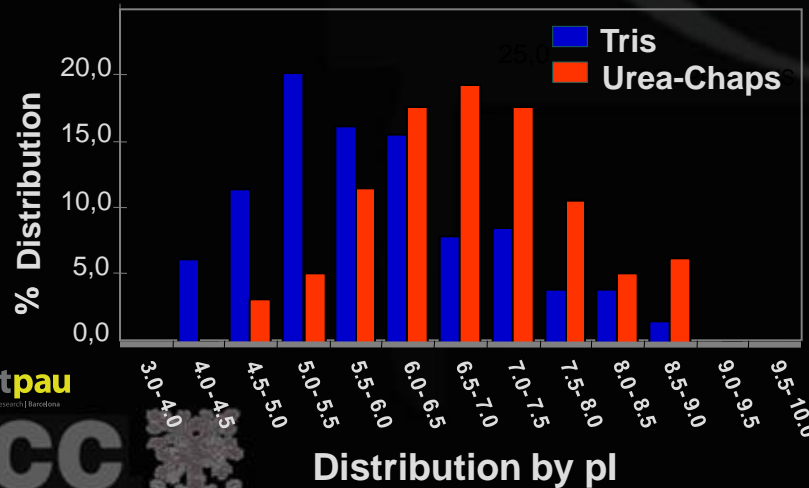
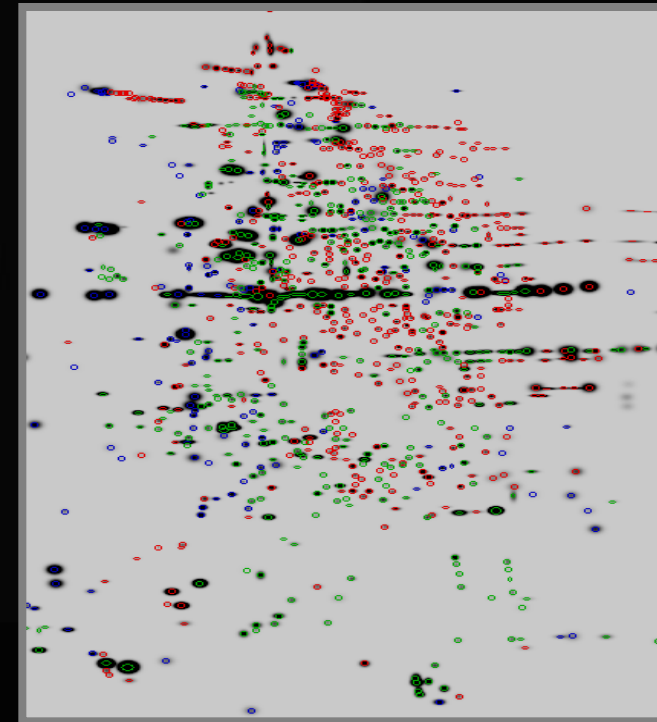


Cytosk./membrane

Urea-Chaps Fraction

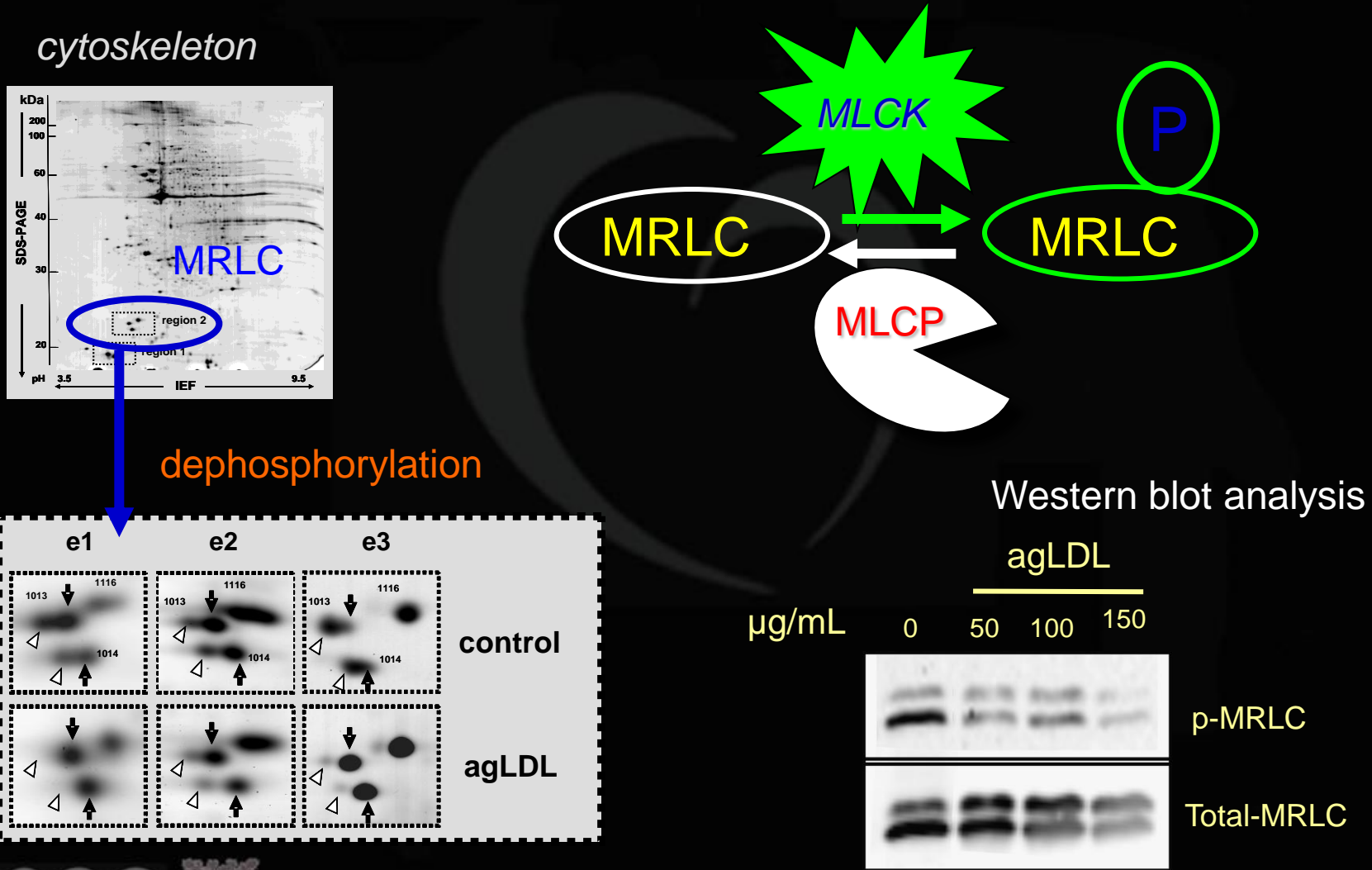


Total spots 880 ± 176

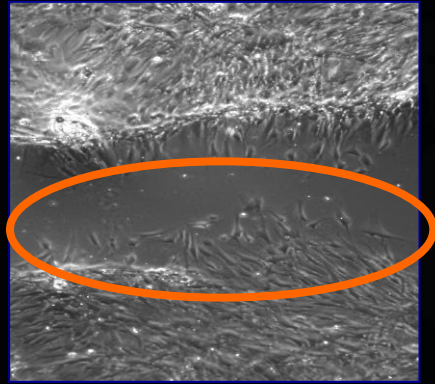


- Fracción Tris (16%)
- Fracción Urea-Chaps (48%)
- Both Fractions (36%)

agLDL induce changes in the proteomic profile of myosin regulatory light-chain (MRLC) in SMC



Localization of MRLC in VSMC at the migrating front



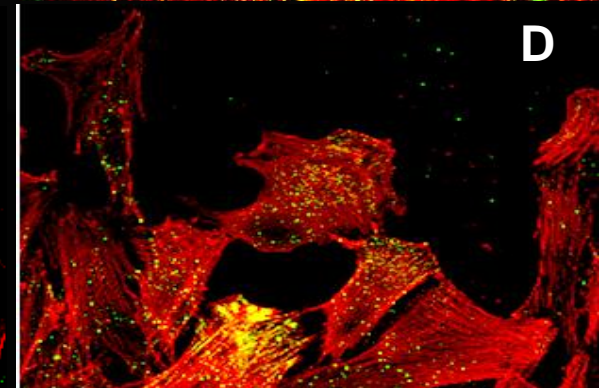
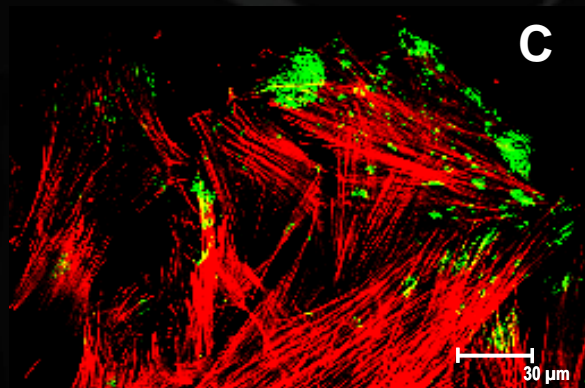
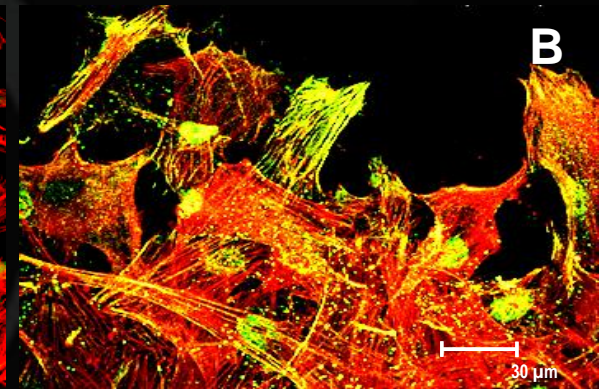
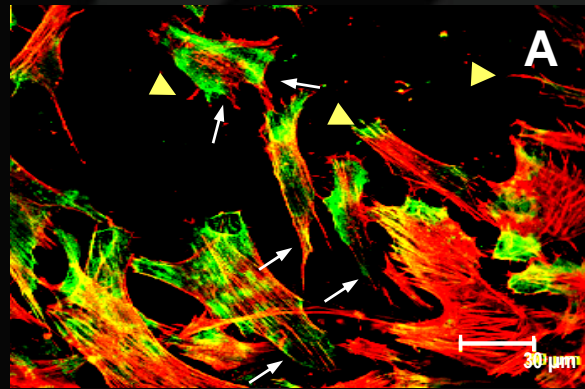
control

agLDL

Confocal microscopy

Total MRLC

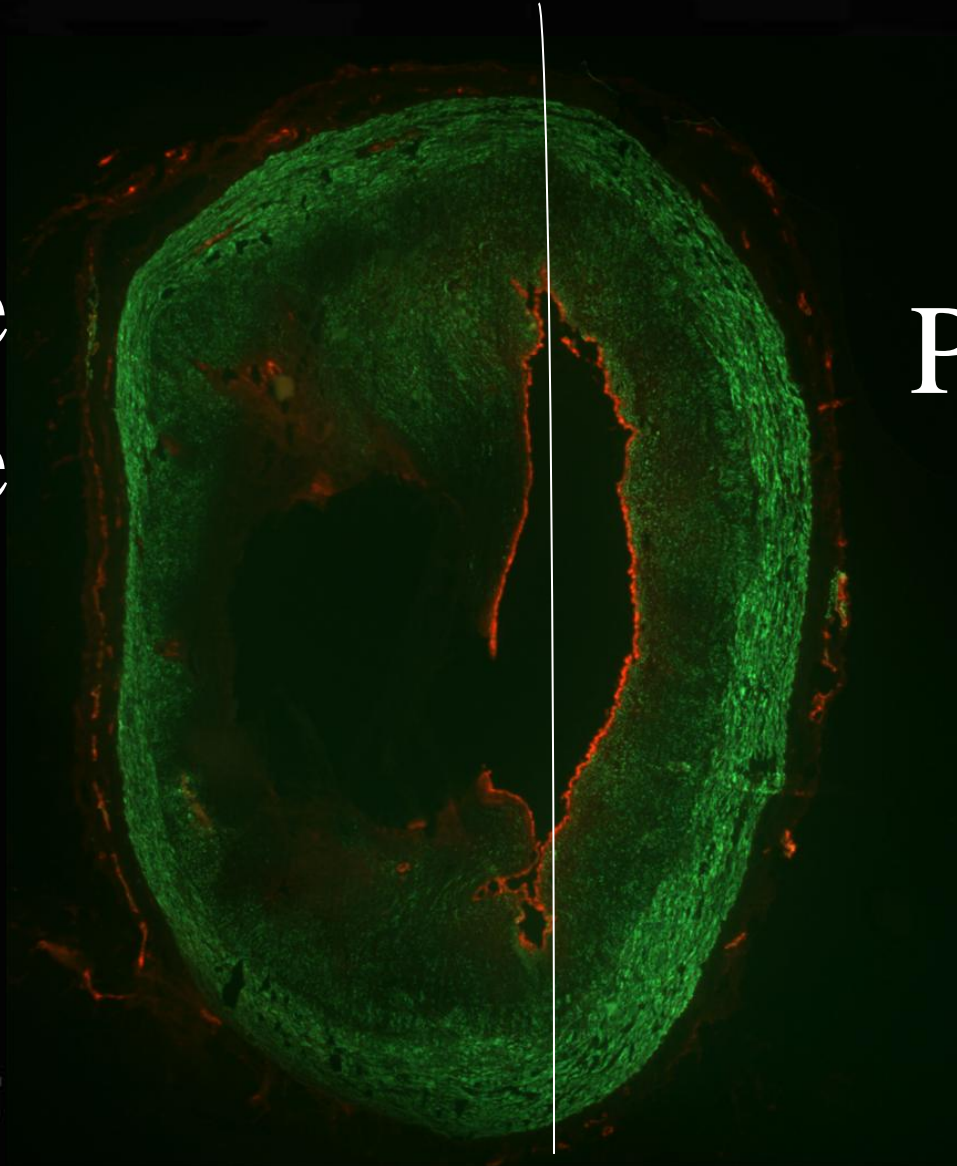
P-MRLC



F-Actin (red): Alexa -594 Phalloidin
MRLC ; P-MRLC (green): FITC

TRANSCRIPTOMICS – HUMAN CORONARY ARTERIES

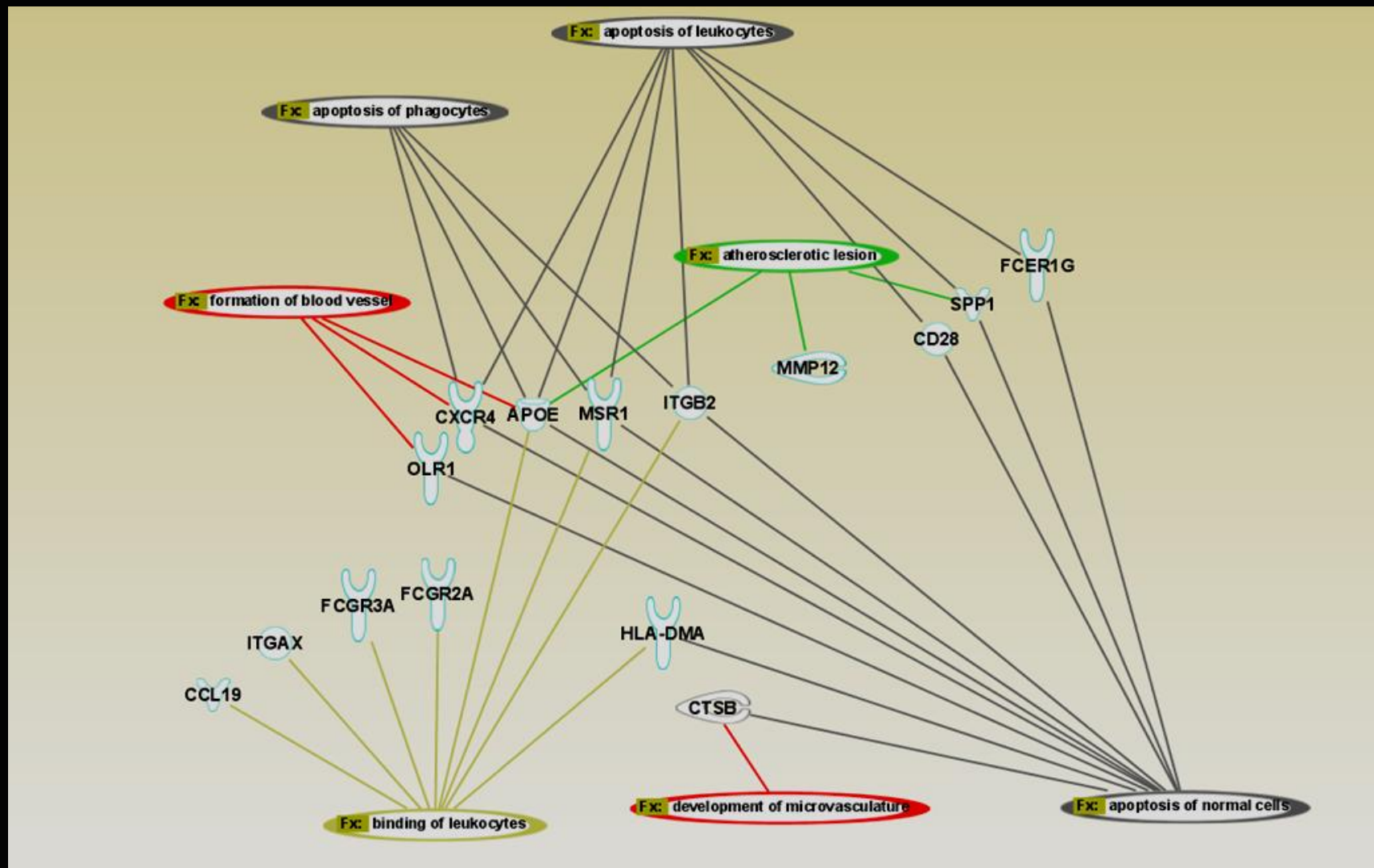
Severe
Plaque

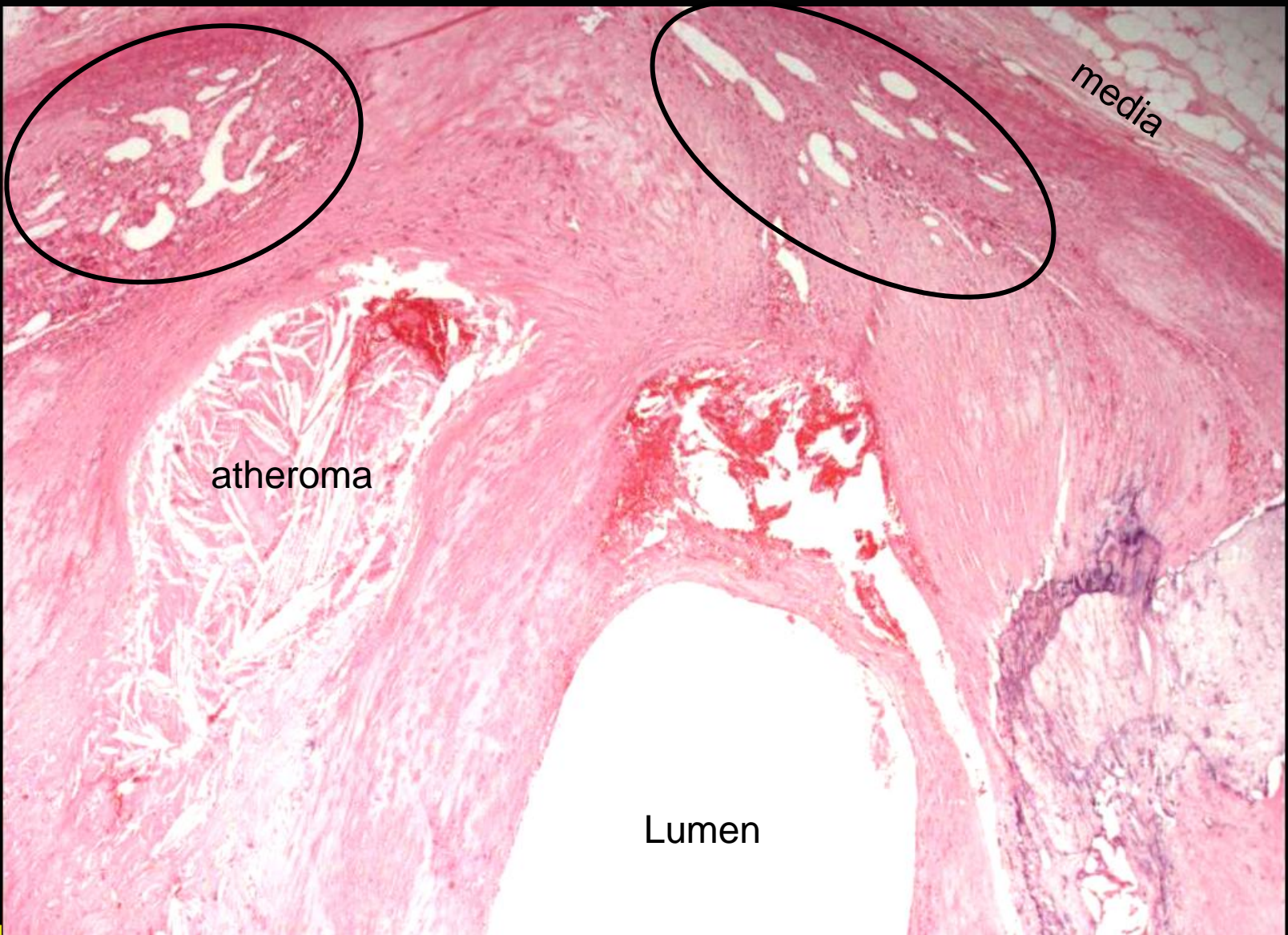


Plaque-IT

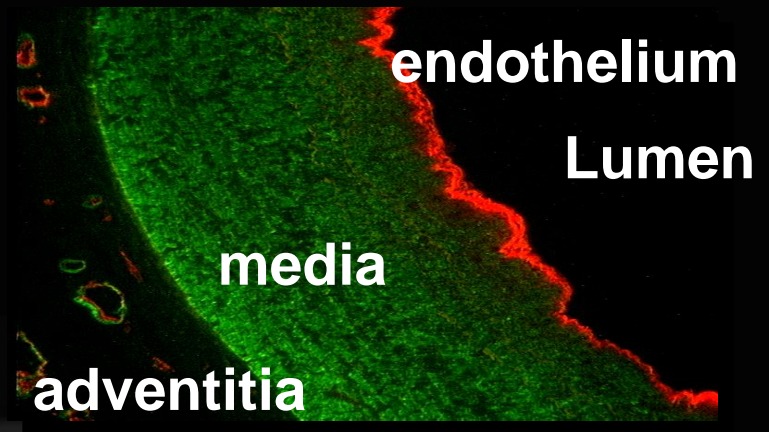
genesymbol	noDM>DM	noHTA>HTA	noIAM>IAM	P<O	P<O HTA	P<O IAM	genesdescription
SCD	■	■					stearoyl-CoA desaturase (delta-9-desaturase)
ITLN1			■	■		■	intelectin 1 (galactofuranose binding)
MYOC					■	■	myocilin, trabecular meshwork inducible glucocorticoid response
TSPAN8					■	■	tetraspanin 8
TTN							titin
DYNLL1							dynein, light chain, LC8-type 1
KLF15		■					Kruppel-like factor 15
GSTT1	■	■					glutathione S-transferase theta 1
genesdescription	DM	HTA	IAM	P>O	P>O HTA	P>O IAM	genbank
IRF1	■	■	■	■			interferon regulatory factor 1
TNC	■	■	■	■			tenascin C (hexabrachion)
CXCL9	■	■	■	■			chemokine (C-X-C motif) ligand 9
CXCL10	■	■	■	■			chemokine (C-X-C motif) ligand 10
CCL8	■	■	■	■			chemokine (C-C motif) ligand 8
PTX3	■	■	■	■			pentraxin-related gene, rapidly induced by IL-1 beta
SPP1	■	■	■	■			secreted phosphoprotein 1 (osteopontin, bone sialoprotein I, early T-lymphocyte activation 1)
SELE	■	■	■	■			selectin E (endothelial adhesion molecule 1)
IGJ	■	■	■	■	■	■	immunoglobulin J polypeptide, linker protein for immunoglobulin alpha and mu polypeptides
IBSP	■	■	■	■	■	■	integrin-binding sialoprotein (bone sialoprotein, bone sialoprotein II)
CCL19	■	■	■	■	■	■	chemokine (C-C motif) ligand 19
CCL2	■	■	■	■	■	■	chemokine (C-C motif) ligand 2
CAPG	■	■	■	■	■	■	capping protein (actin filament), gelsolin-like

PO functions

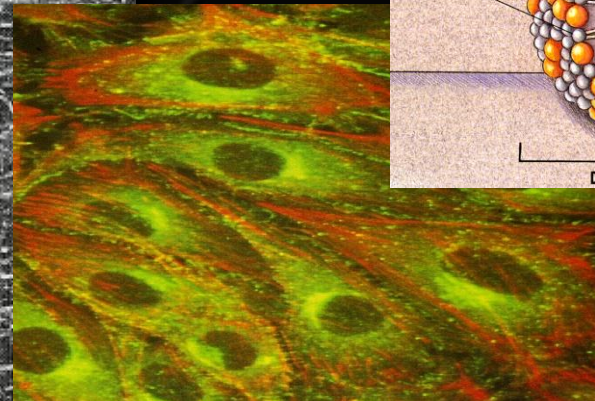
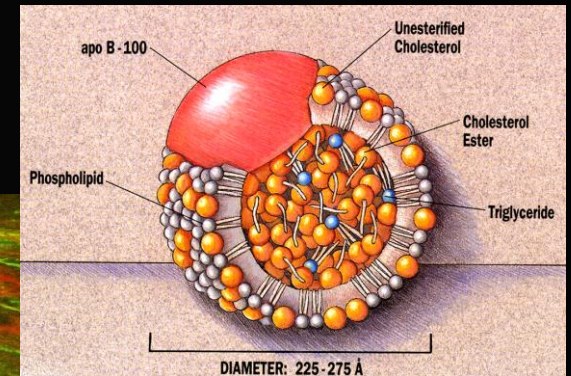
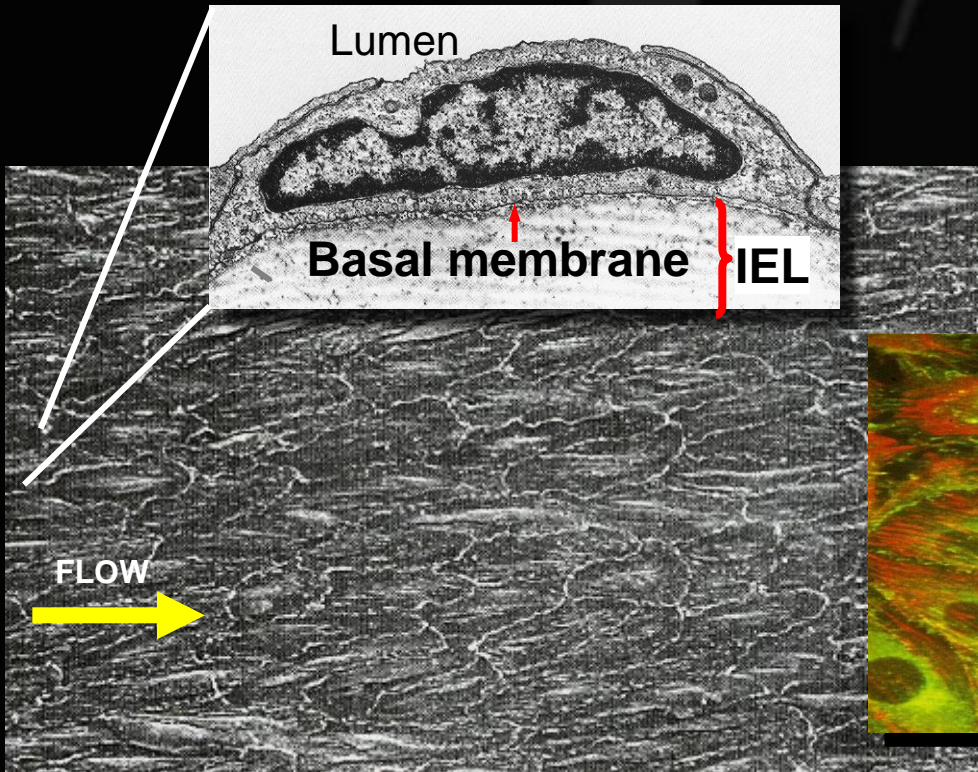




THE VASCULAR WALL AND THE ENDOTHELIUM



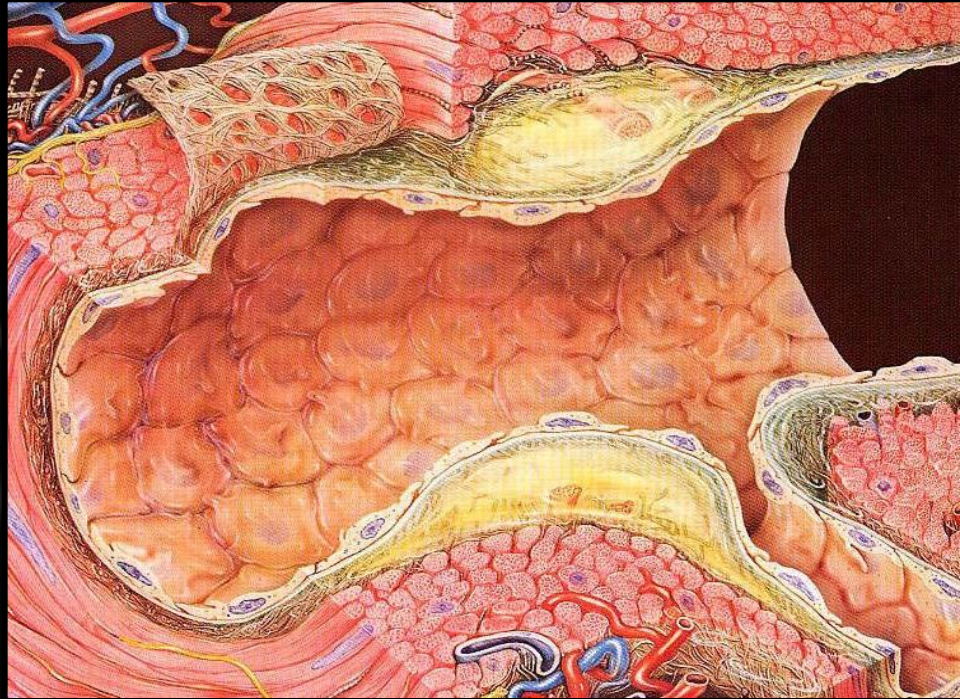
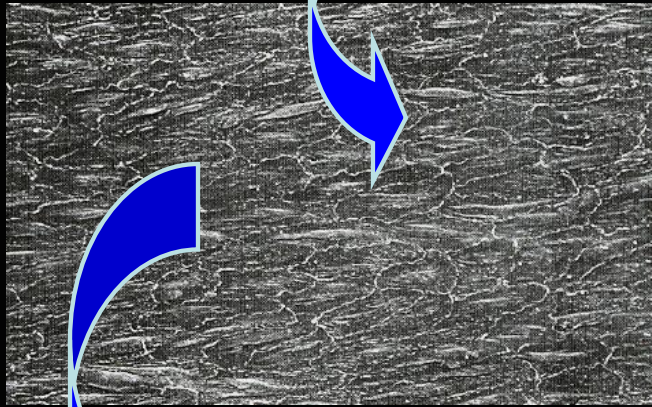
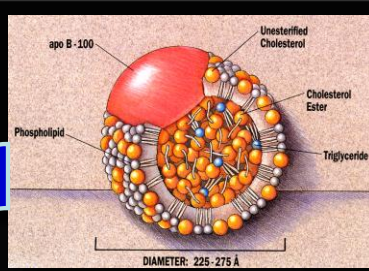
- Endothelium (**vWF**)
- VSMCs (**alfa actin**)



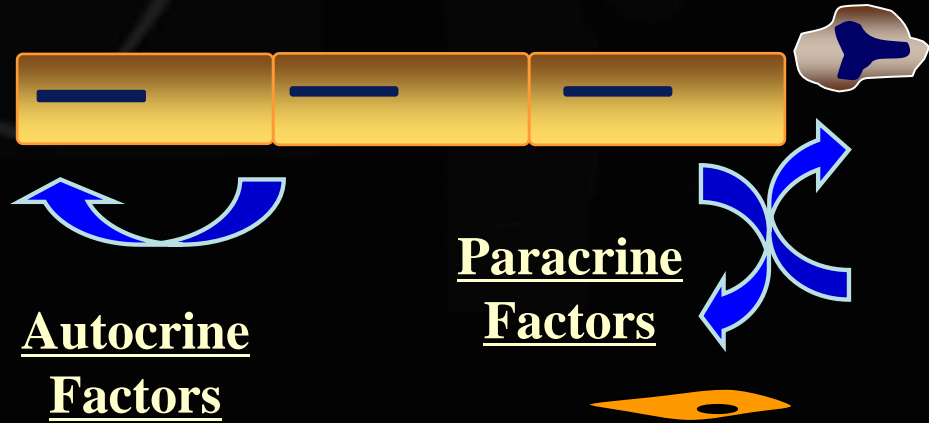
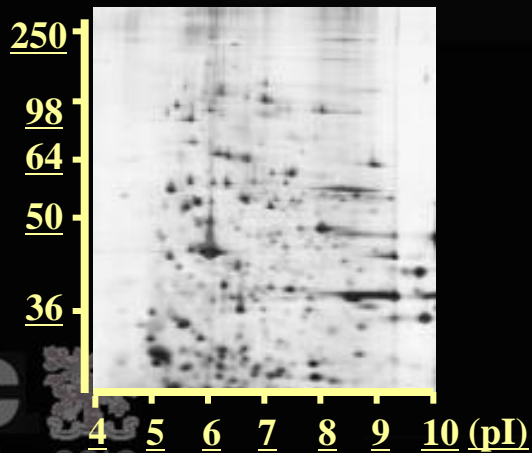
■ Faloidin

Barcelona Endothelium: SEM

Modified from Badimon L, 2005

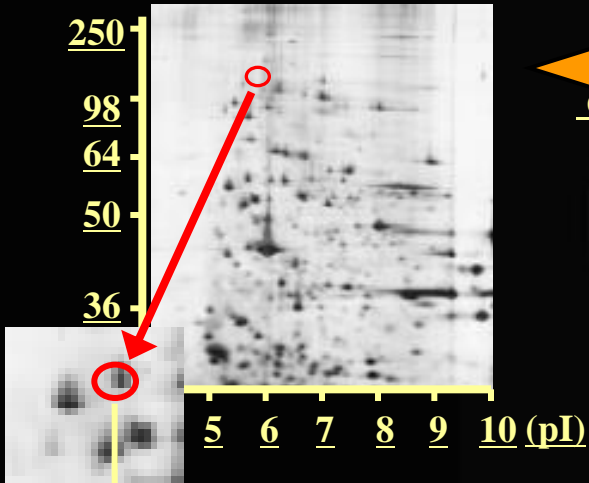


2D- gel electroforesis

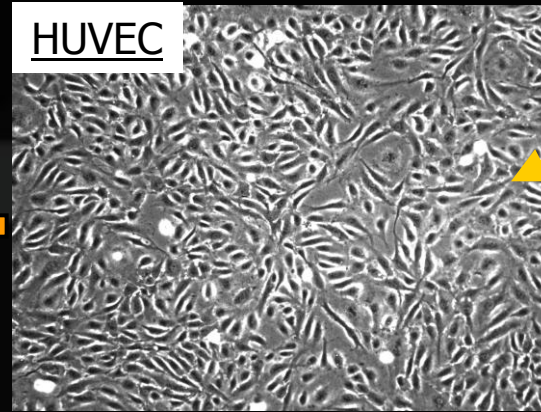


PROTEOMICS

2D- gel electroforesis

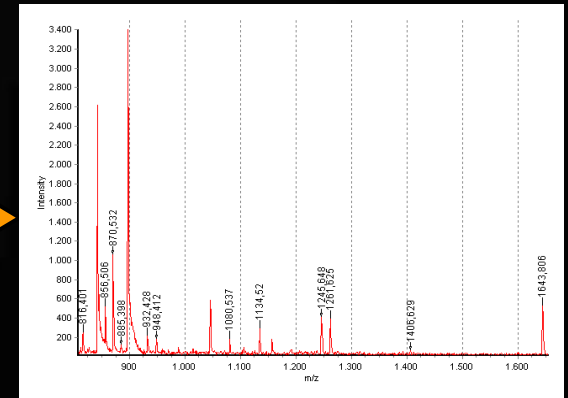


Sequential
protein
extraction



-/+LDL
180 mg/dl

MALDI - ToF

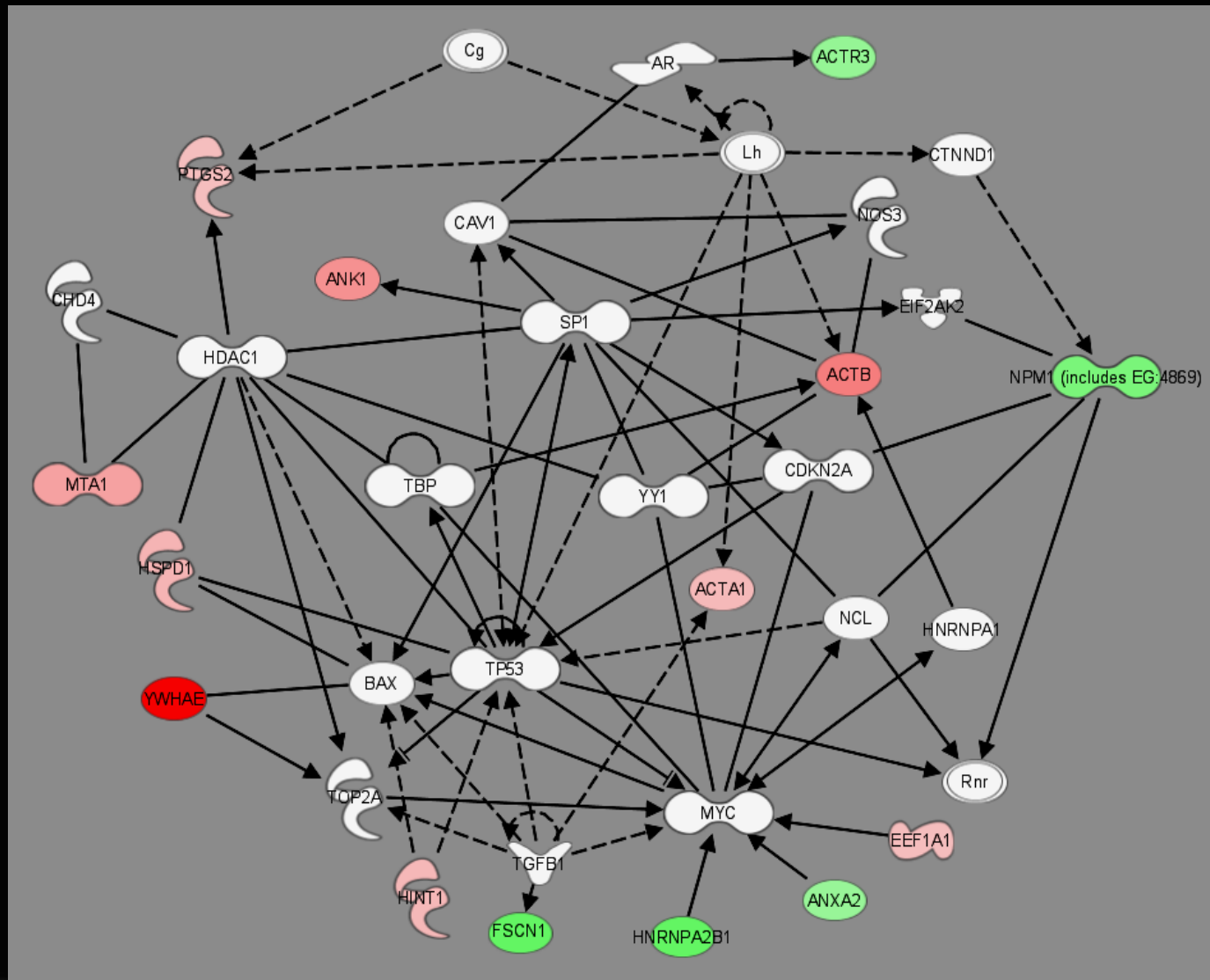


Acquisition
and spectral
analysis

1. Trypsin
digestion

2. Peptide
extraction

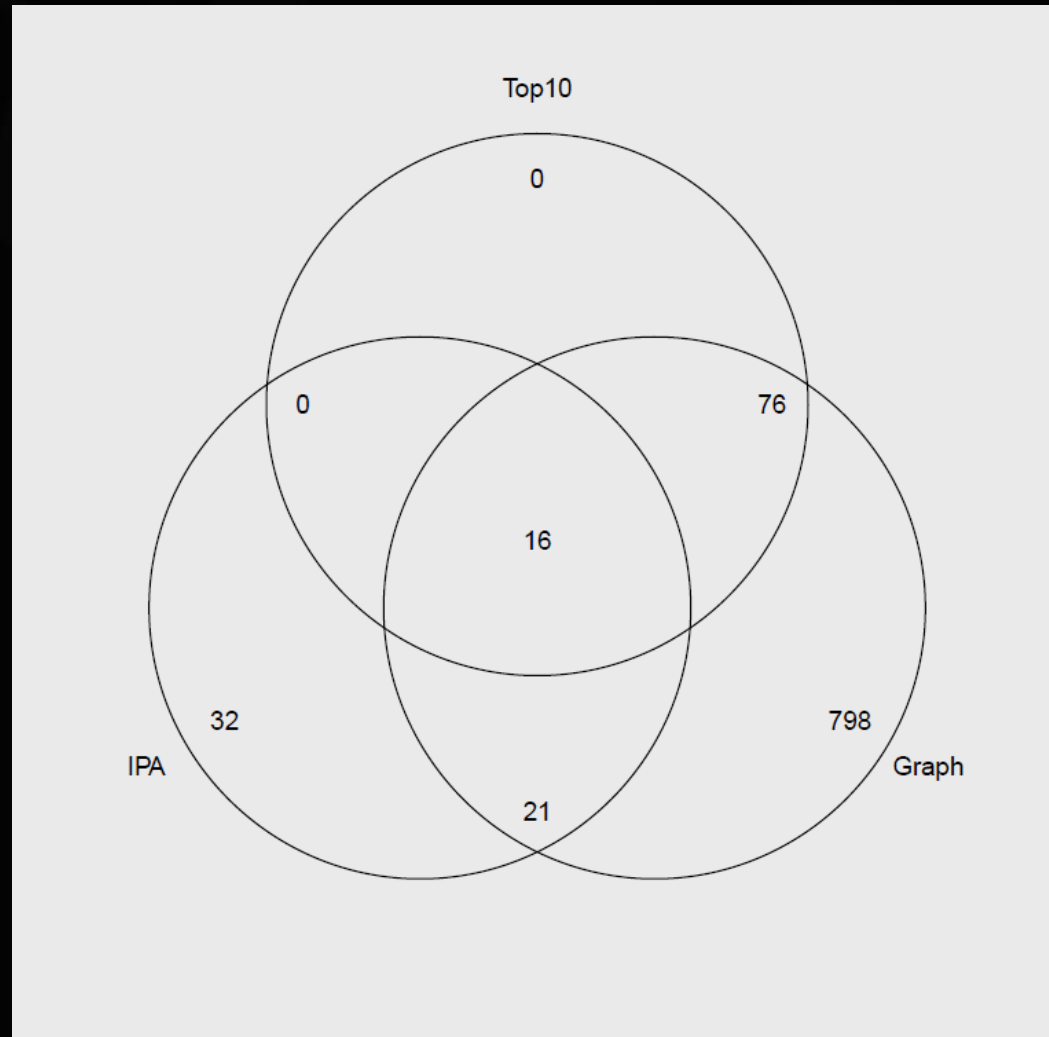
INDUCED ENDOTHELIAL CELL CYTOSOLIC PROTEOME



RED, upregulated proteins; GREEN, downregulated proteins; WHITE, IPA-generated protein
 Color intensity, level of regulation

BADIMON L, ALARCON JL, CARDUS A, PADRO T. UNPUBLISHED OBSERVATIONS

INDUCED ENDOTHELIAL CELL CYTOSOLIC PROTEOME



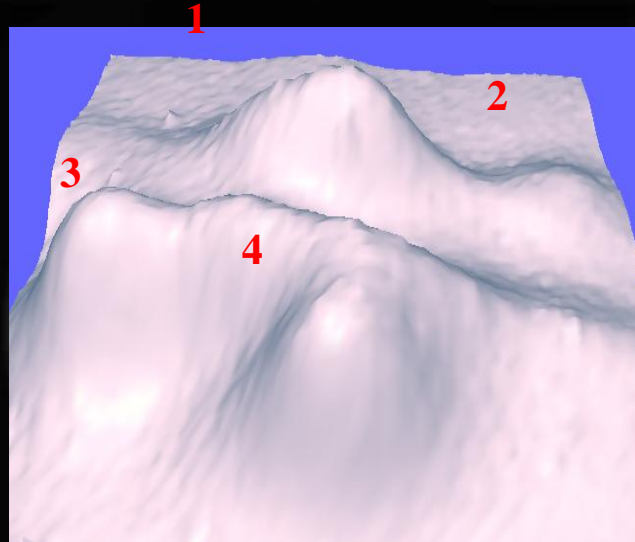
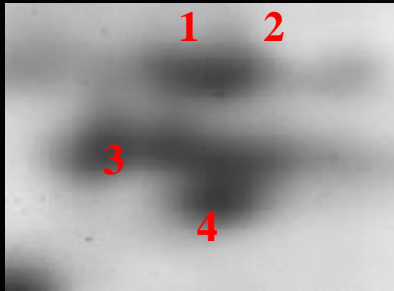
Badimon L, Alarcon JL, Cardus A, Padro T. Unpublished observations

TOP 16 PROTEINS: IPA-GRAPH-CONNECTIVITY

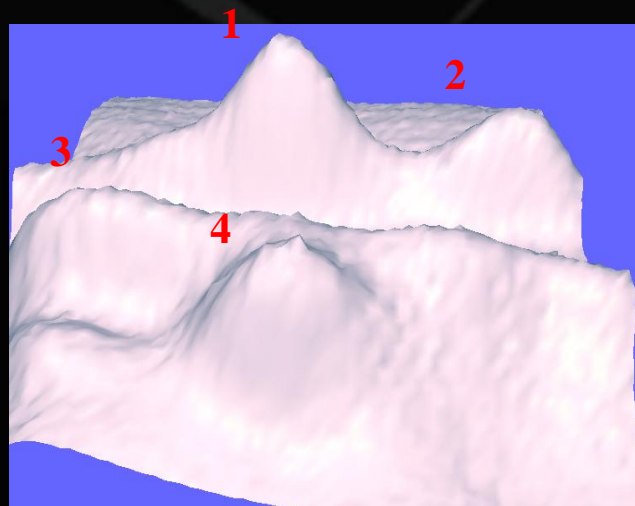
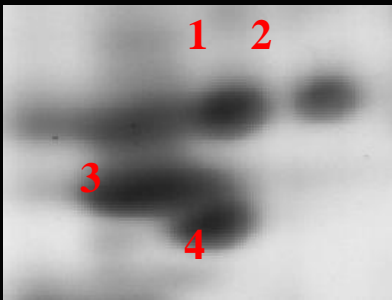
Symbol	Protein name
YWHAE	14-3-3 protein epsilon
EEF1A1	Elongation factor 1-alpha 1
ACTB	Actin, cytoplasmic 1
TP53	Cellular tumor antigen p53
GRB2	Growth factor receptor-bound protein 2
HDAC1	Histone deacetylase 1
MTA1	Metastasis-associated protein MTA1
ACTA1	Actin, alpha skeletal muscle
HSPD1	60 kDa heat shock protein, mitochondrial
CAV1	Caveolin-1
ANXA2	Annexin A2
APP	Amyloid beta A4 protein
AR	Androgen receptor
CDKN2A	Cyclin-dependent kinase inhibitor 2A, isoforms 1/2/3
SHC1	SHC-transforming protein 1
EIF2AK2	Interferon-induced, double-stranded RNA-activated protein kinase

14-3-3 Proteins

CTRL



+ LDL 180 mg/dl



Mean Values in 2D

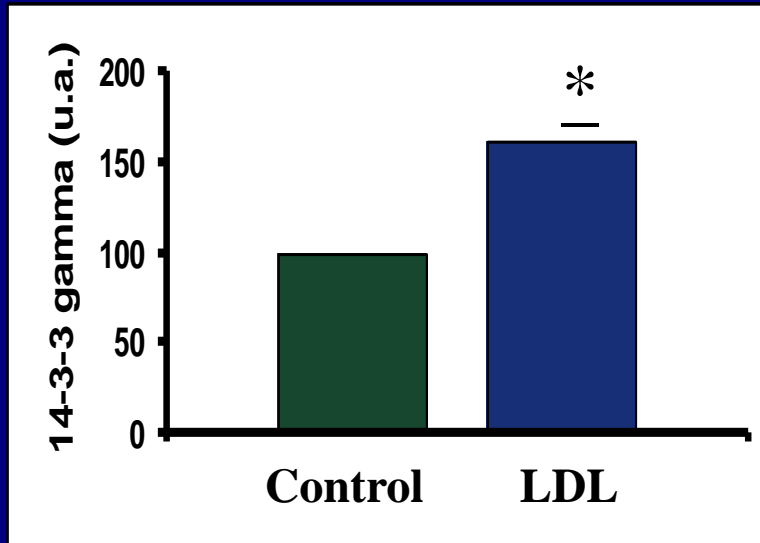
		Ratio +LDL/ctrl
1	gamma	2,94
2	epsilon	4,30
3	tsheta	0,83
4	teta/ delta	0,93

14-3-3 γ Proteins

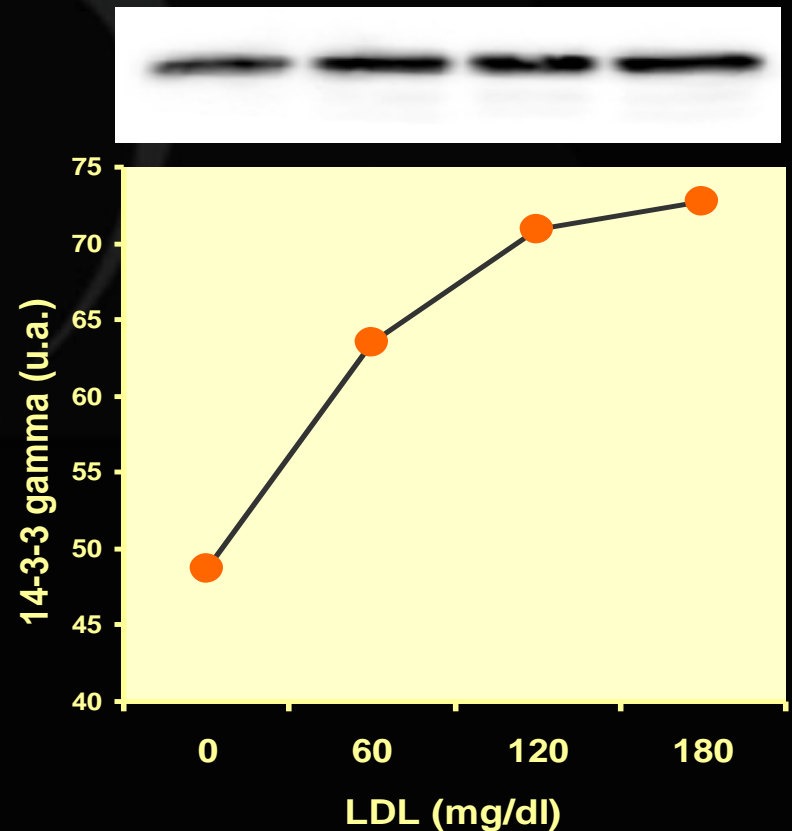
Western blot



33 KDa →



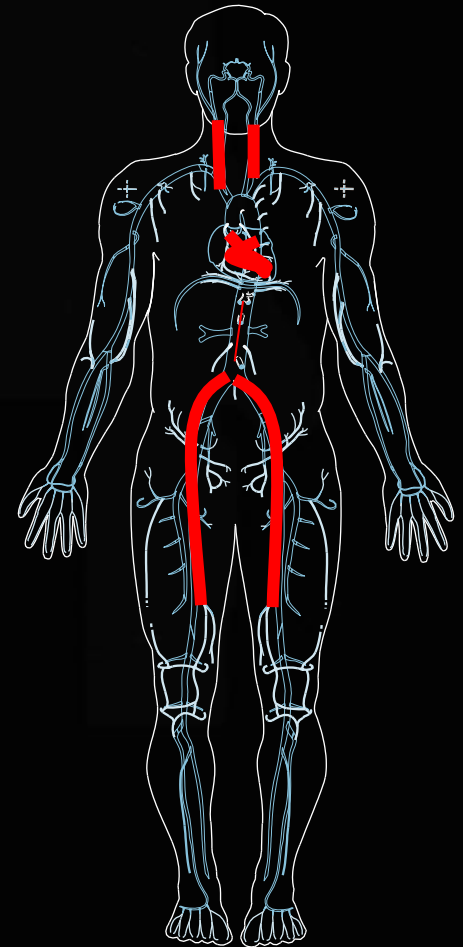
Dose-response effects of LDL on 14-3-3 gamma levels



LDL: 180 mg/dl

ATHEROSCLEROSIS

- VESSEL REMODELING
- NEOVESSEL FORMATION
- INFLAMMATION
- REGULATORY EFFECTS
ENDOTHELIAL CELLS



CARDIOVASCULAR RESEARCH CENTER

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