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# Inflammation in the formation and rupture of intracranial aneurysms

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ABC Win meeting 2016



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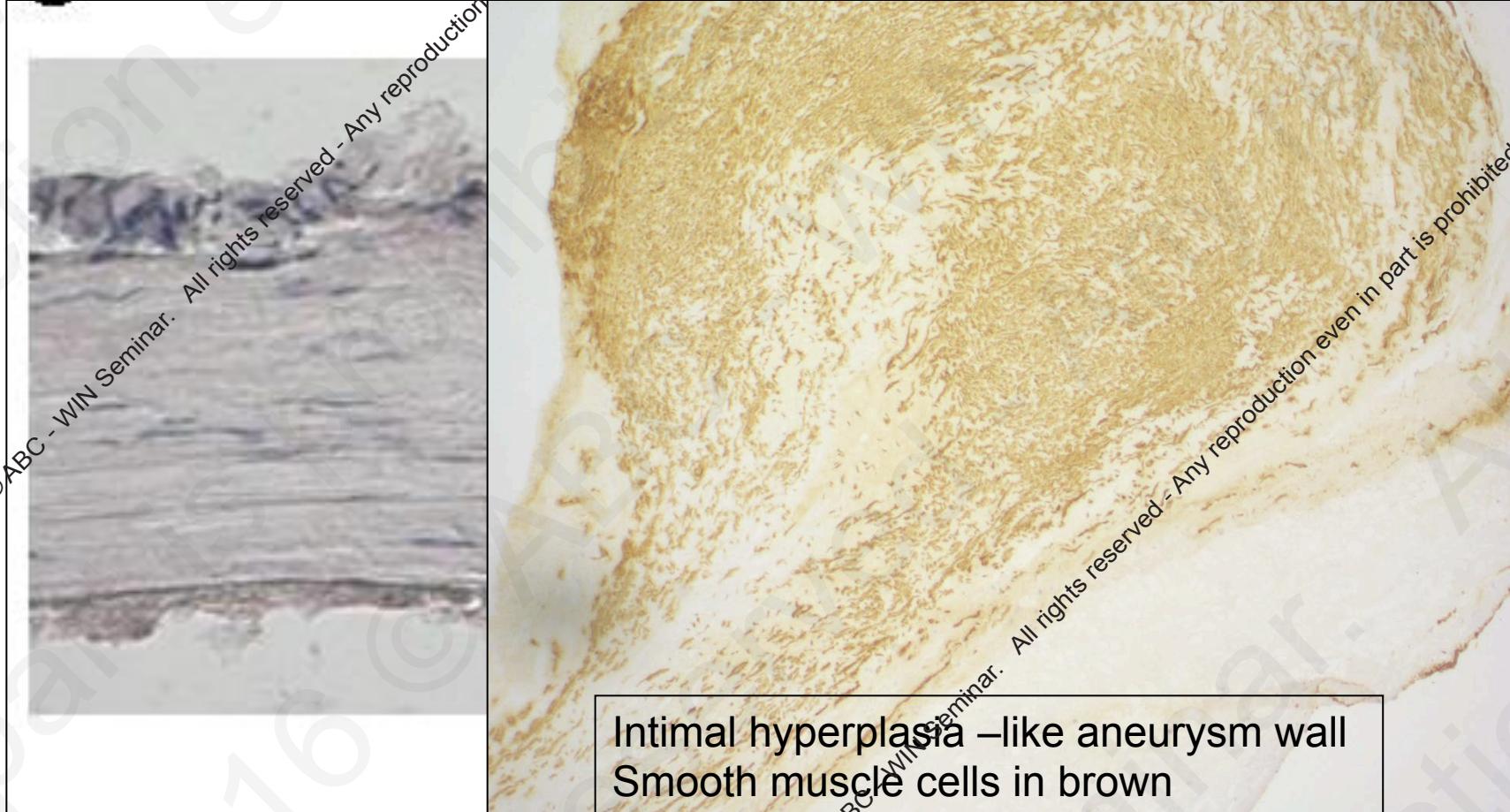


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# Aneurysm formation ≠ Aneurysm rupture

- Not all aneurysms rupture -  
(autopsy studies, prospective follow-up studies, clinical experience.)
- In a life-long follow-up study of unruptured aneurysms, only 1/3 ruptured (Korja et al. Stroke 2014)
- Rupture is a rare event in animal models of induced aneurysms (*in the classical Hashimoto model 3% in 3months*)

# Aneurysm wall is not "inert rubber" – it can grow, remodel, and thicken



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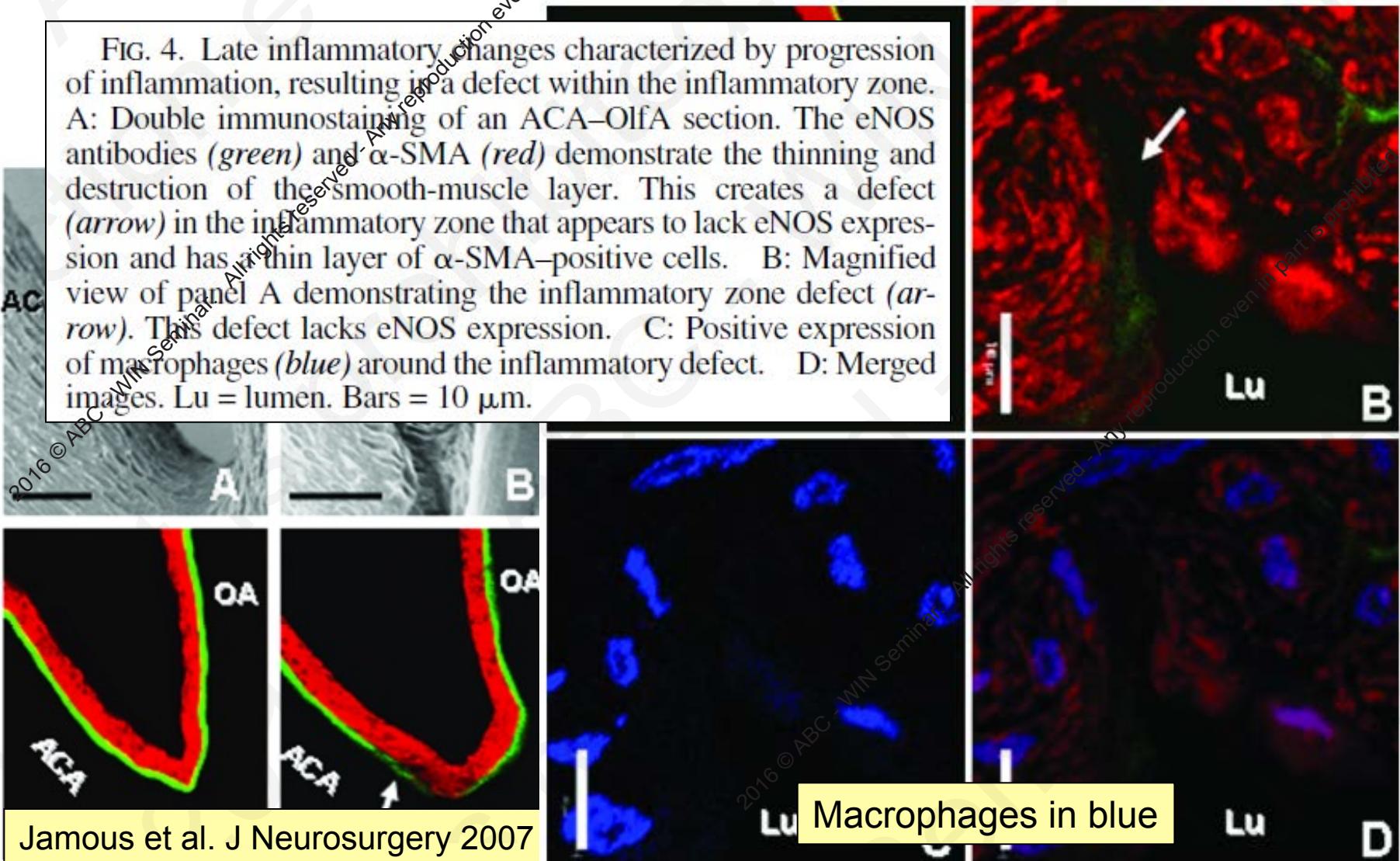
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# Data for the role of inflammation during aneurysm formation – animal models

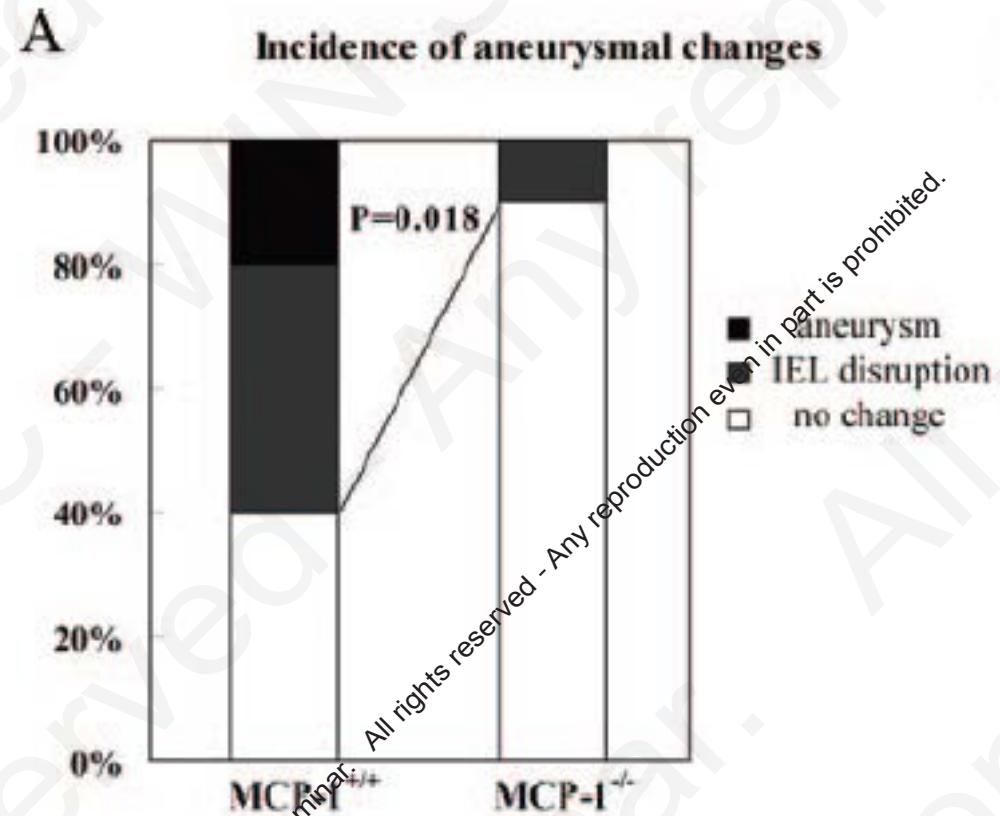
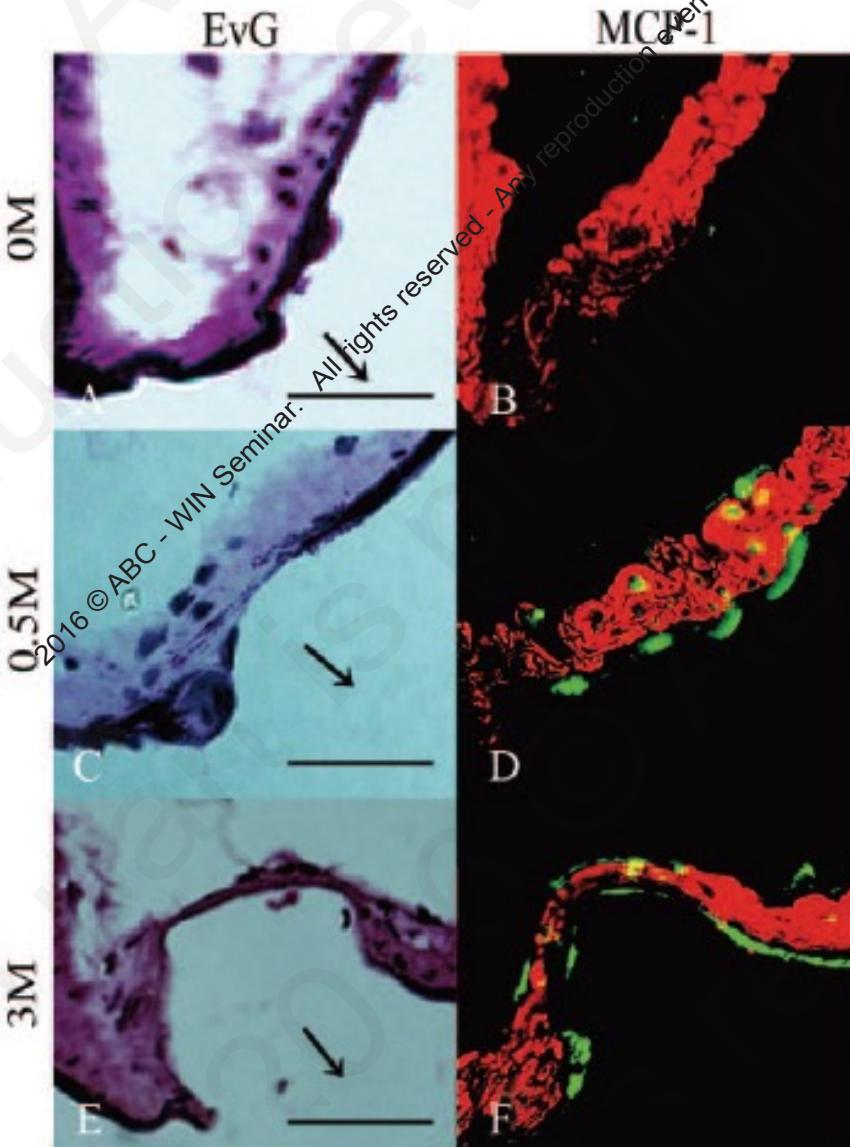
- Hashimoto N. group's model (Kyoto): Rupture rare
  - **Induced hypertension** (salt + renal artery ligation)
  - Inhibition of collagen crosslinking
  - *Unilateral ligation of carotid artery*
- Hashimoto T. group's model (UCSF): Rupture common
  - **Induced hypertension** (angiotensin II)
  - *Elastase injection into cisterna magna*

# Cerebral artery wall is infiltrated by macrophages during formation of intracranial saccular aneurysms (unilateral carotid artery ligation model)

FIG. 4. Late inflammatory changes characterized by progression of inflammation, resulting in a defect within the inflammatory zone. A: Double immunostaining of an ACA–OlfA section. The eNOS antibodies (green) and  $\alpha$ -SMA (red) demonstrate the thinning and destruction of the smooth-muscle layer. This creates a defect (arrow) in the inflammatory zone that appears to lack eNOS expression and has a thin layer of  $\alpha$ -SMA–positive cells. B: Magnified view of panel A demonstrating the inflammatory zone defect (arrow). This defect lacks eNOS expression. C: Positive expression of macrophages (blue) around the inflammatory defect. D: Merged images. Lu = lumen. Bars = 10  $\mu$ m.

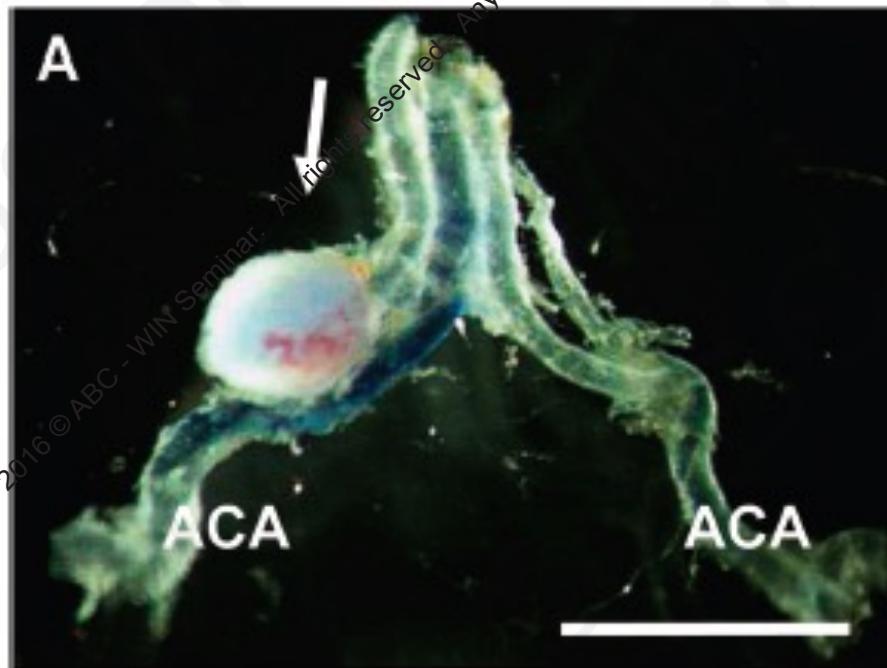


# Inhibition of macrophage recruitment via MCP-1 inhibits aneurysm formation (carotid ligation model)

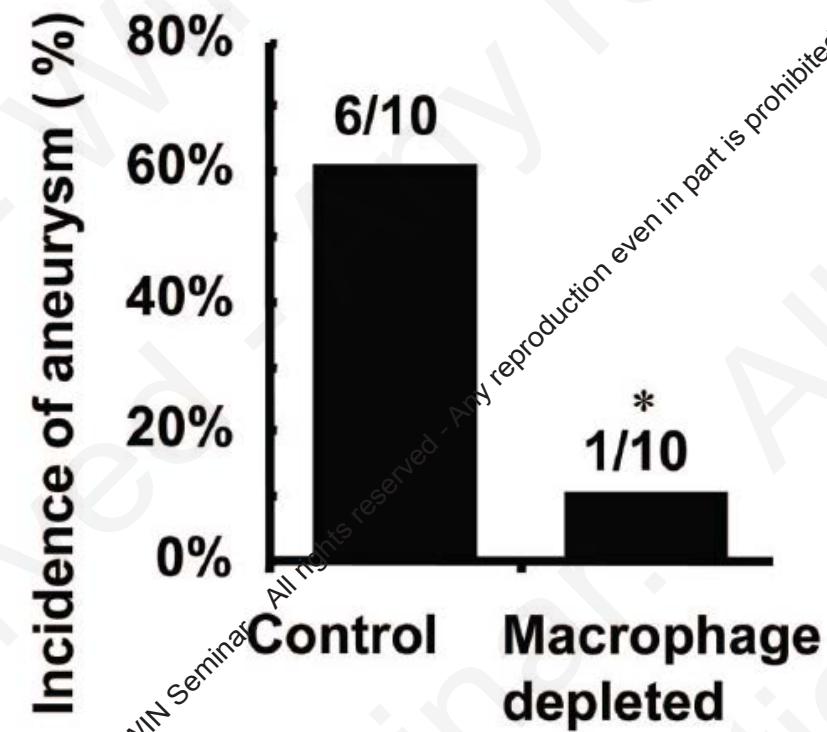


Aoki T et al. Stroke 2009

# Depletion of macrophages inhibits aneurysm formation (elastase model)



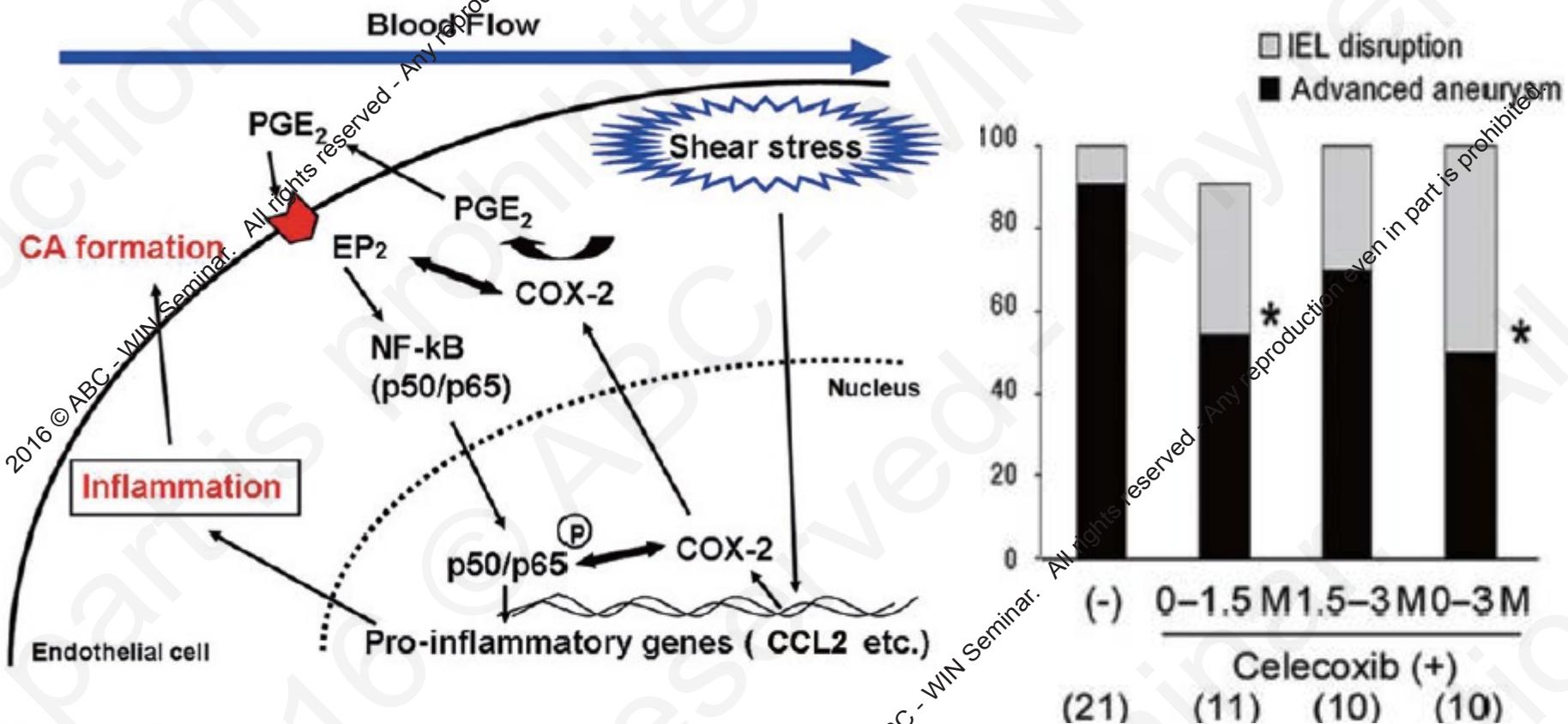
Kanematsu et al. Stroke 2011



# How macrophages cause aneurysm formation?

- Secretion of matrix degrading proteases  
(e.g. MMP-2 and MMP-9) (*Aoki et al. Stroke 2007*)
- Induction of oxidative stress – iNOS and reactive oxygen species  
(*Eukuda et al. Circulation 2000* and *Aoki et al. Lab Invest 2009*)
- *Secretion of inflammatory cytokines (! COX2 and TNFa signaling)*  
– phenotype change in smooth muscle and endothelial cells
- *Potentially induction of cell death in the aneurysm wall (Fas-signaling and TNFa signaling)*

# Induction of macrophage activation via COX-2 and EP2 positive feed back loop during formation of intracranial aneurysms (carotid ligation model)



**Figure 5**

A proposed model for the formation of cerebral aneurysm induced by shear stress.

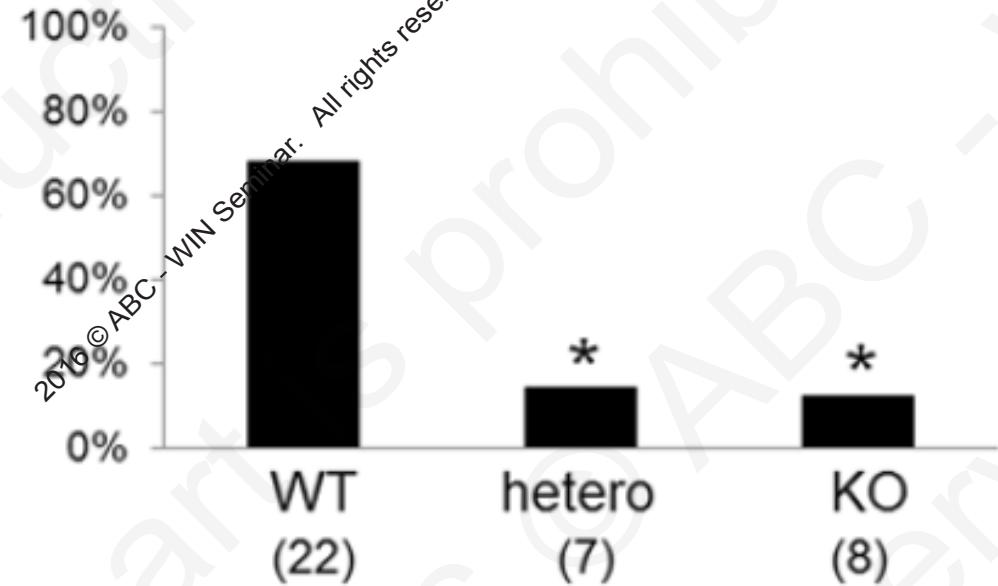
Aoki T et al. Br J Pharmacology 2011

Inhibition of TNFa signaling (TNFaR1 knock out)

1) reduces aneurysm formation and

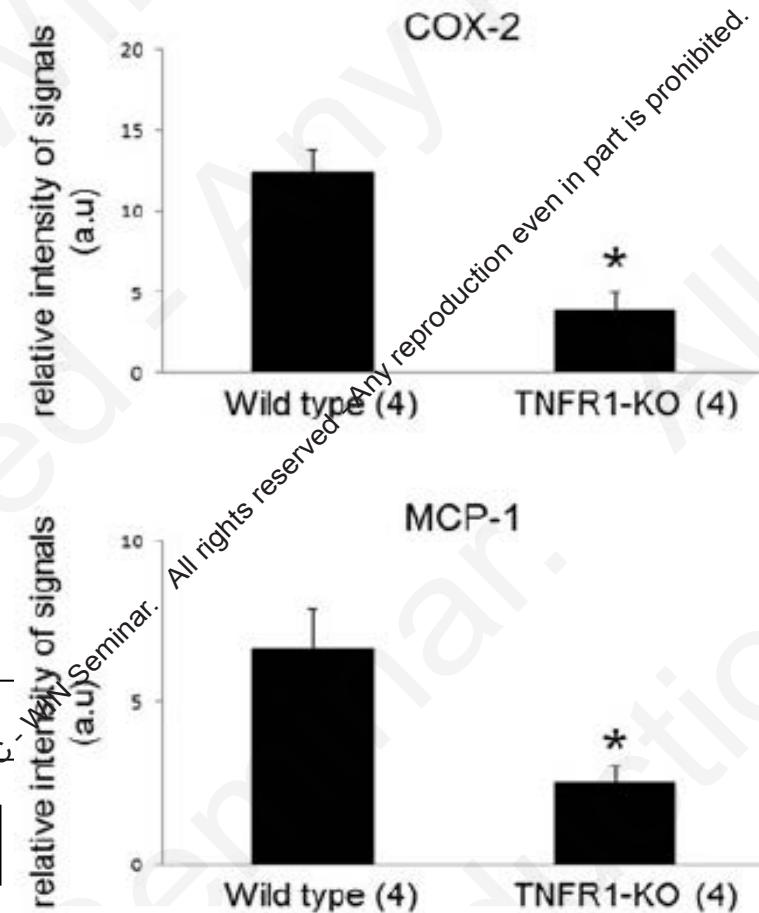
2) MCP-1 and COX-2 expression in the aneurysm wall

### Incidence of intracranial aneurysm



TNFaR1 knock out mice (KO) vs. wild type (WT)

Aoki et al. Acta Neuropathologica Comm 2014



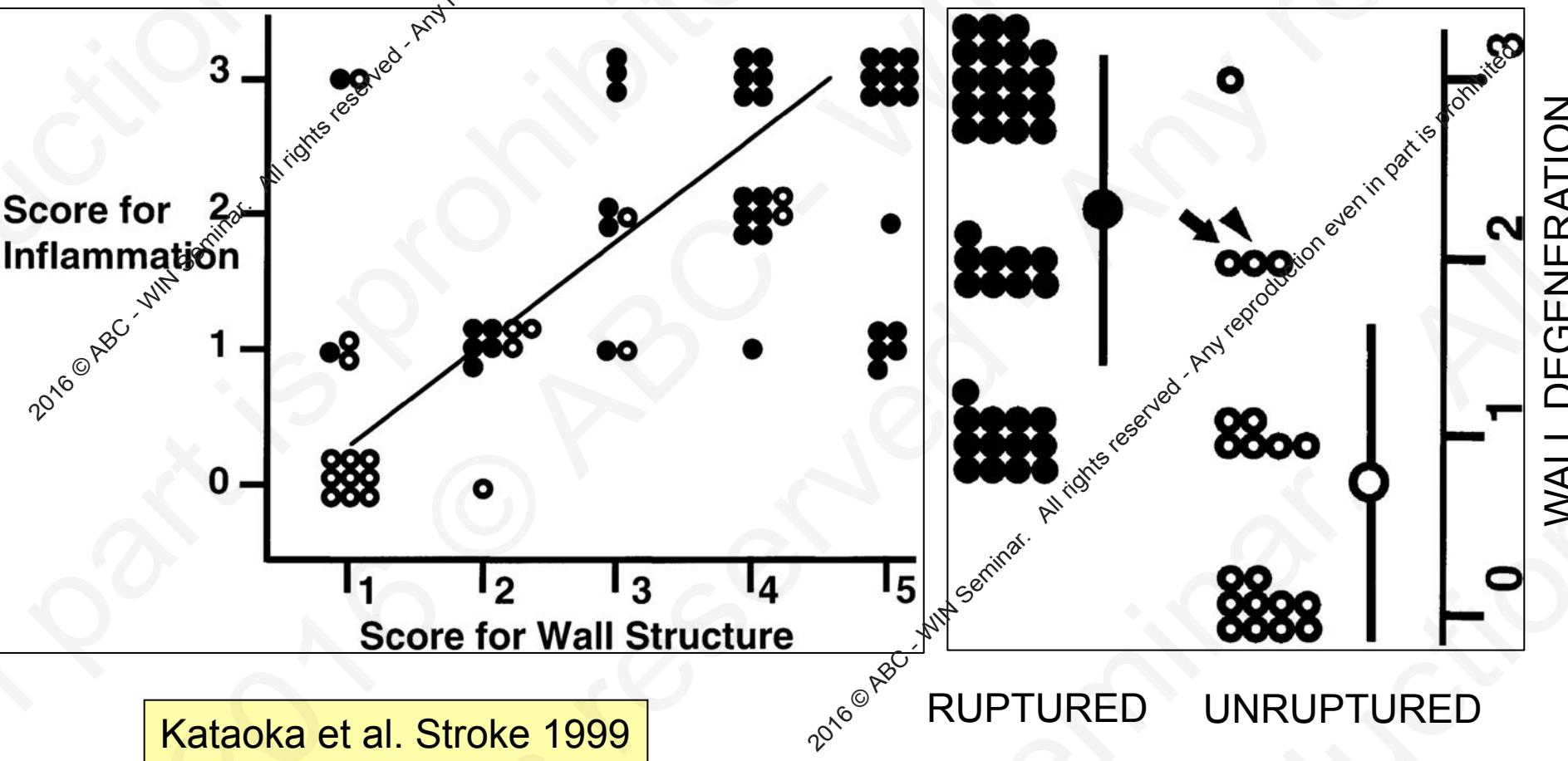
# Conclusion: Formation of intracranial aneurysms is mediated by macrophages

1. Macrophage infiltration associates with aneurysm formation
2. Blocking macrophage infiltration prevents aneurysm formation
3. Inhibition of macrophage signaling (TNFa and COX-2) prevents aneurysm formation

# What about aneurysm rupture?

- Data from human tissue samples available (ruptured vs. unruptured aneurysms)
- Various animal models – none that for certain replicates the human pathobiology of aneurysm rupture
  - Induced hypertension + elastase injection model
  - Decellularized aortic graft model

# Macrophage infiltration associates with wall degeneration and rupture in human intracranial aneurysms

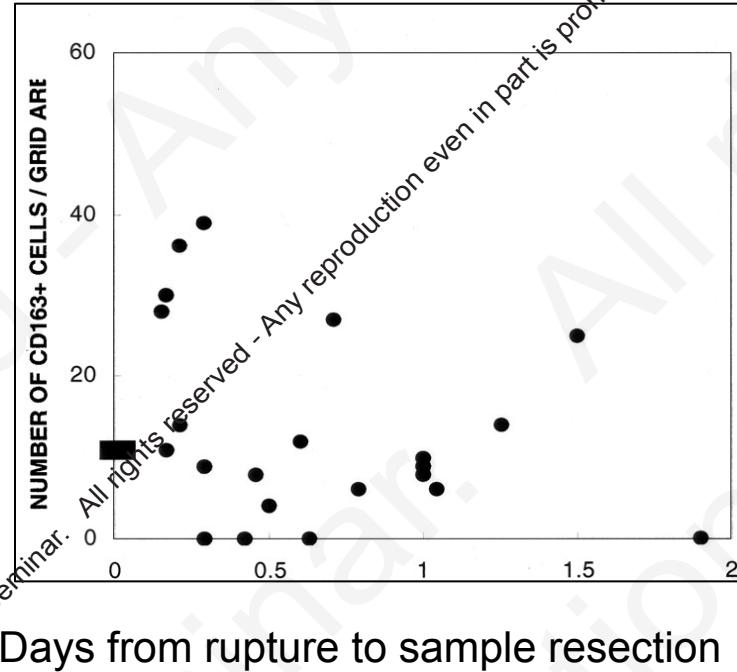
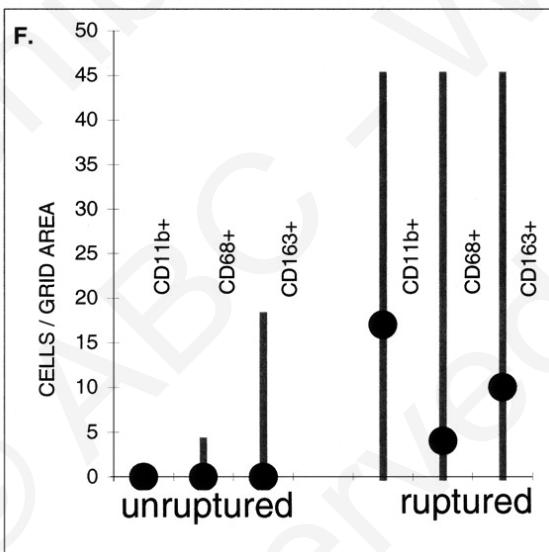
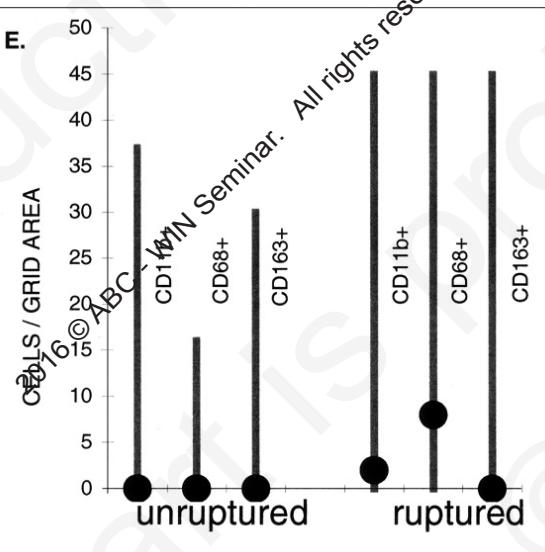


# Macrophage infiltration increases with rupture and is not just a reaction to rupture

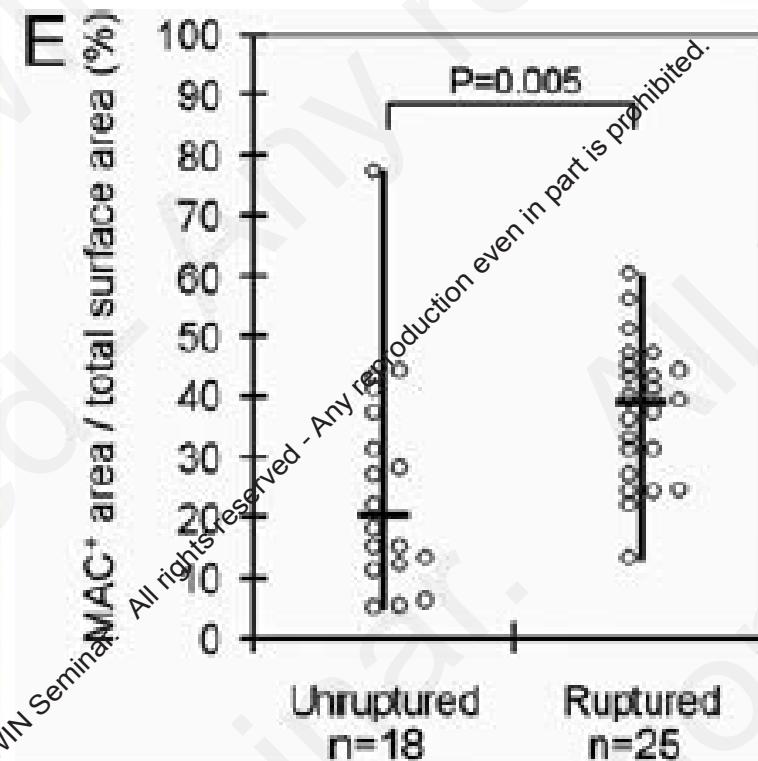
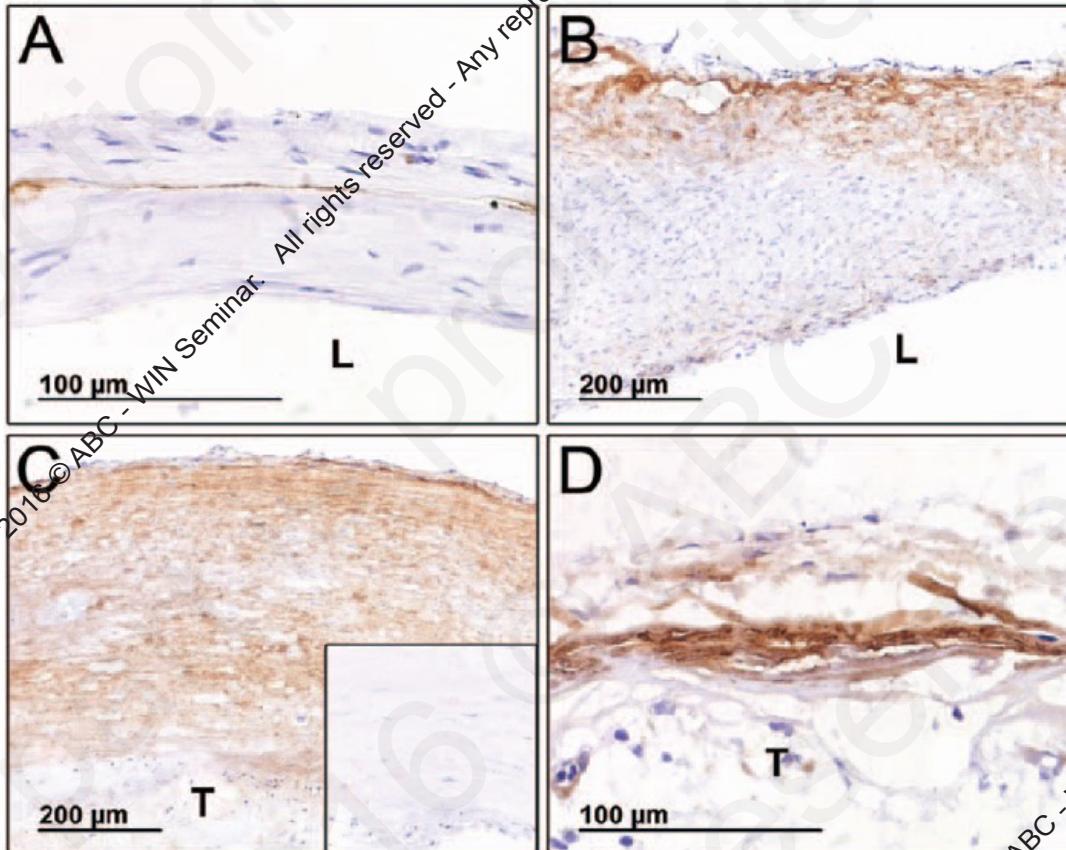
WALL AREAS OTHER THAN MYINTIMAL HYPERPLASIA / ORGANIZING THROMBOSIS

AREAS OF MYINTIMAL HYPERPLASIA / ORGANIZING THROMBOSIS IN THE ANEURYSM WALL

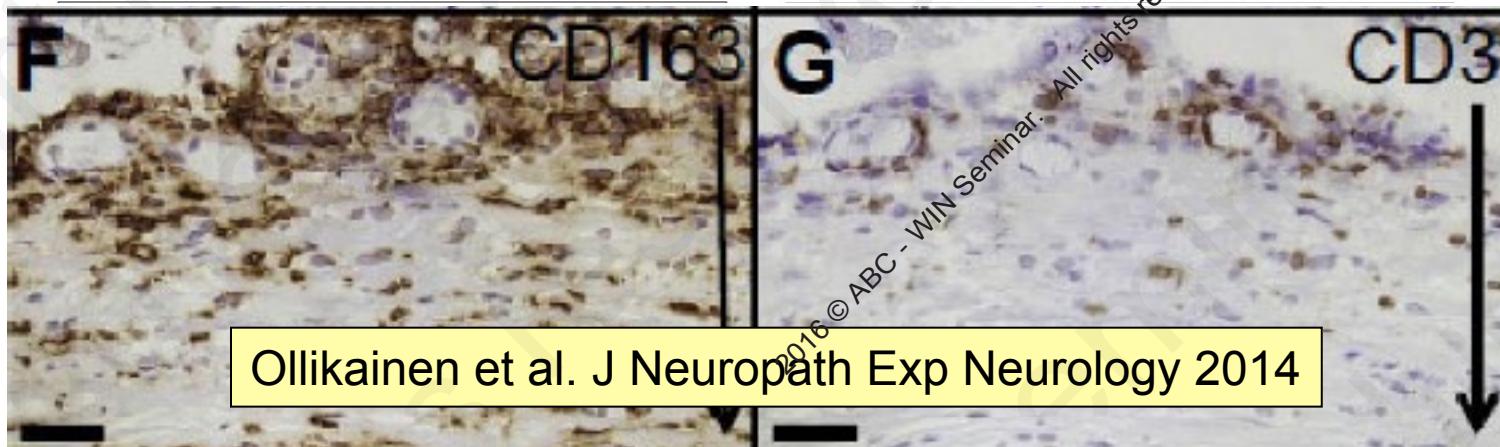
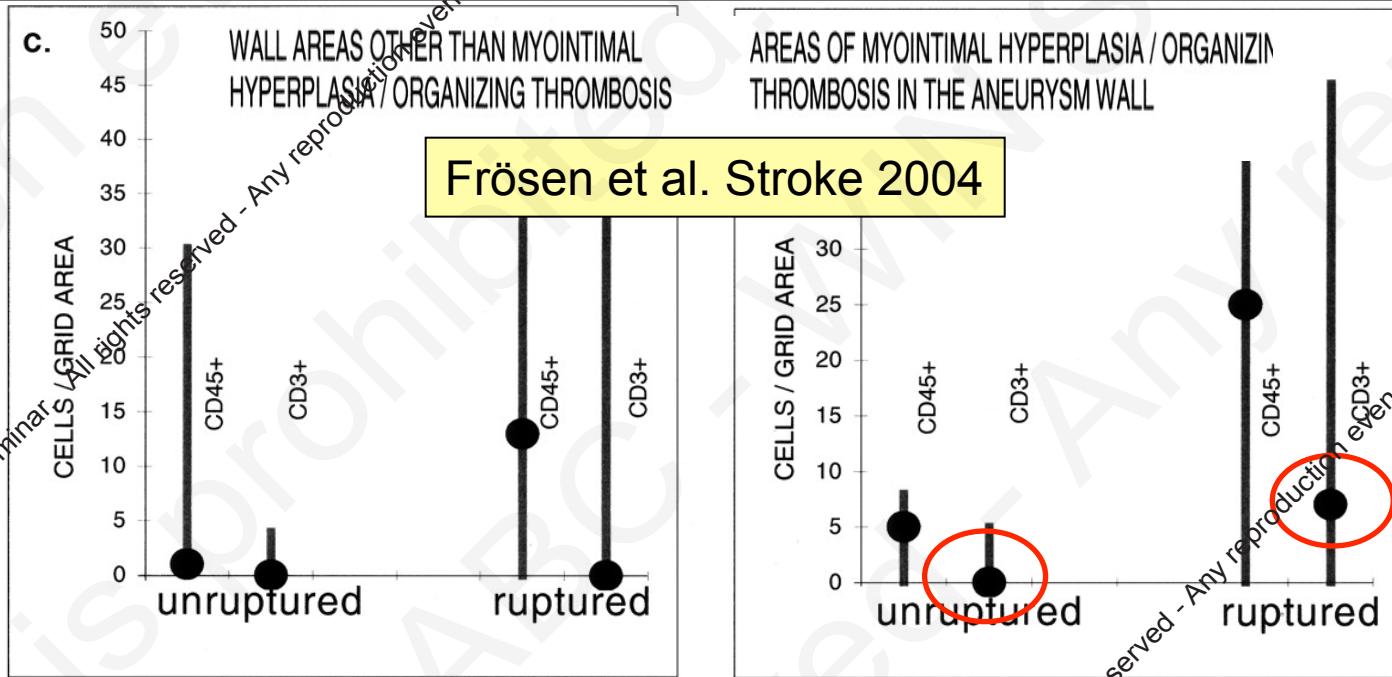
No correlation with time from rupture and macrophage infiltration



# In addition to macrophage infiltration – Aneurysm wall degeneration and rupture associates with activation of the complement (humoral immune system)



# Also T-cells (CD3) are present in the aneurysm wall – and associate with rupture



...and also, *mast cells* are present in the aneurysm wall and associate with accumulation of iron and lipids as well as with macrophage and T-cell infiltration

TABLE 4. Association of Histologic Characteristics of sIAs With MCs

| Variable   | MCs             |                 | p       |
|--|-----------------|-----------------|---------|
|  | Absent (n = 27) | Present (n = 9) |         |
| Histology of sIA wall  |                 |                 |         |
| Endothelium not intact, n (%)*   | 17/27 (63)      | 9/9 (100)       | 0.09†   |
| Intraluminal fresh thrombosis present, n (%)*                                    | 10/26 (38)      | 0/7 (0)         | 0.073   |
| Intraluminal organizing thrombosis present, n (%)*                               | 9/26 (35)       | 2/7 (29)        | 1.000   |
| Lipid present in the ECM, n (%)*   | 21/27 (78)      | 7/9 (78)        | 1.000   |
| Ratio of lipid-rich area, median (range), %‡                                     | 9 (0–37)        | 37 (0–69)       | 0.022†  |
| Neovessels present, n (%)*   | 6/27 (22)       | 9/9 (100)       | <0.001† |
| Distribution density of neovessels, median (range), neovessels/mm <sup>2</sup> ‡ | 0 (0–15)        | 2 (0–50)        | <0.001† |
| Iron deposition present, n (%)*  | 3/27 (11)       | 6/9 (67)        | 0.003†  |
| Strong iron deposition present, n (%)*   | 1/27 (4)        | 5/9 (56)        | 0.002†  |
| Total number of inflammatory cells   |                 |                 |         |
| CD68-positive macrophages, median (range)‡§                                      | 15 (0–527)      | 24 (42–297)     | 0.012†  |
| CD163-positive macrophages, median (range)‡§                                     | 72 (5–417)      | 158 (35–443)    | 0.110   |
| CD3-positive T lymphocytes, median (range)‡§                                     | 11 (0–357)      | 52 (25–137)     | 0.010†  |
| Further analysis of MC-positive sIAs   |                 |                 |         |
| Total number of MCs, median (range)¶   |                 | 23 (3–170)      |         |
| Total number of degranulated MCs, median (range)¶                                |                 | 15 (2–110)      |         |
| Ratio of degranulated MCs to all MCs, median (range), %¶                         |                 | 67 (43–96)      |         |
| Distribution density of MCs, median (range), MCs/mm <sup>2</sup>                 |                 | 4 (1–69)        |         |
| Total number of MCs in standard-sized areas, median (range)§                     |                 | 7 (2–32)        |         |

# Genome wide comparison of gene expression in ruptured and unruptured human aneurysms

– inflammation strongly associates with rupture

**TABLE 2. Biological Processes in Ruptured Saccular Intracranial Aneurysm Wall Samples<sup>a</sup>**

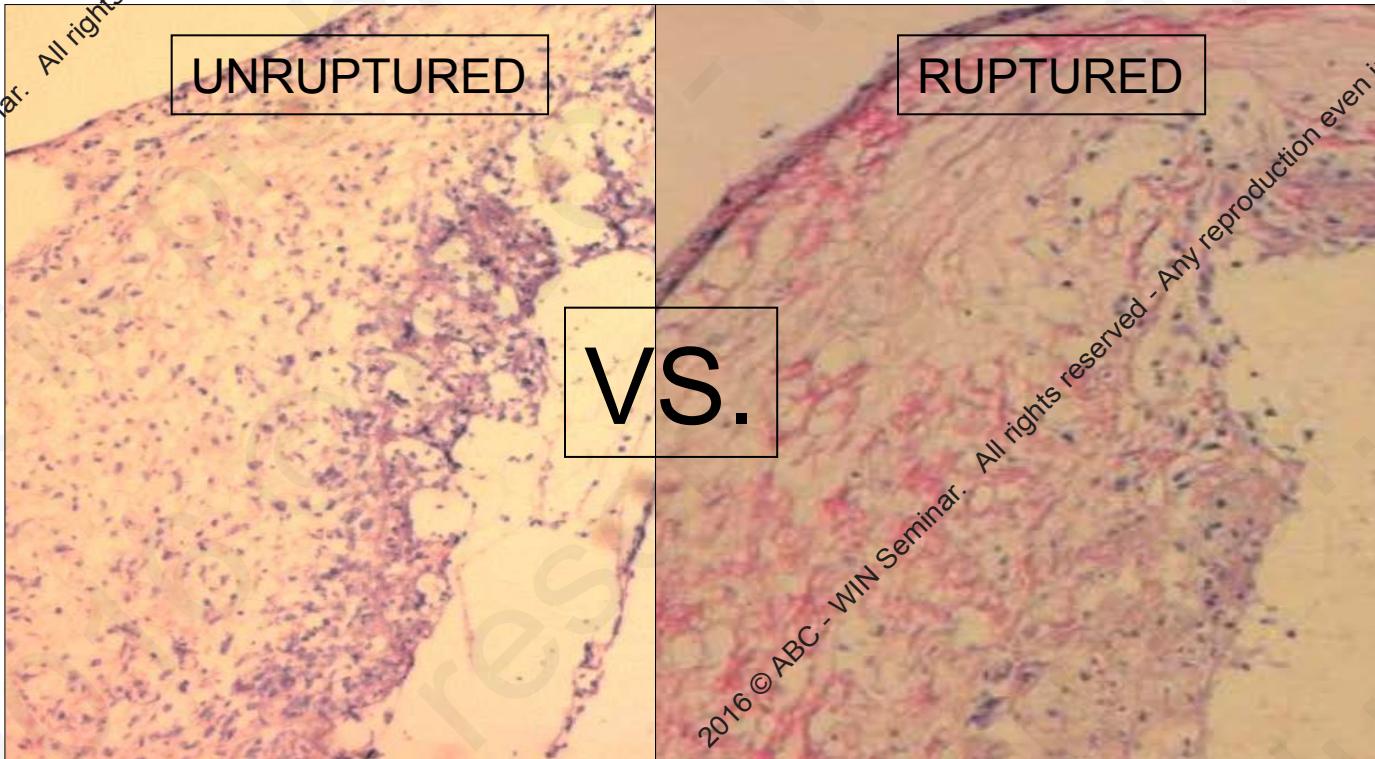
| Gene Ontology (GO) Biological Processes                  | Upregulated Genes  |                |                  |     |                    |                   |
|--|--------------------|----------------|------------------|-----|--------------------|-------------------|
|  | GO ID <sup>b</sup> | P <sup>c</sup> | FDR <sup>d</sup> | OR  | Count <sup>e</sup> | Size <sup>f</sup> |
| Chemotaxis   | GO:0006935         | 2.60E – 14     | 5.66E – 11       | 7   | 30                 | 125               |
| Immune response  | GO:0006955         | 5.70E – 14     | 6.26E – 11       | 4   | 49                 | 333               |
| Response to external stimulus                            | GO:0009605         | 1.00E – 13     | 7.41E – 11       | 3   | 7                  | 651               |
| Inflammatory response                                    | GO:0006954         | 1.40E – 13     | 7.52E – 11       | 4.2 | 46                 | 296               |
| Locomotory behavior                                      | GO:0007626         | 1.50E – 11     | 6.46E – 09       | 4.5 | 35                 | 208               |
| Response to stress                                       | GO:0006950         | 3.60E – 09     | 1.31E – 06       | 2.1 | 99                 | 1224              |
| Response to other organism                               | GO:0051707         | 4.70E – 08     | 1.49E – 05       | 5.3 | 18                 | 87                |
| Positive regulation of tumor necrosis factor production  | GO:0032760         | 5.80E – 08     | 1.60E – 05       | 7.7 | 6                  | 7                 |
| Locomotion   | GO:0040011         | 1.30E – 07     | 3.18E – 05       | 4.7 | 20                 | 114               |
| Cytokine production                                      | GO:0001816         | 2.50E – 05     | 5.49E – 03       | 3.8 | 16                 | 108               |
| Phosphate metabolic process                              | GO:0006796         | 5.30E – 05     | 1.06E – 02       | 1.8 | 66                 | 893               |
| Positive regulation of interleukin-6 production          | GO:0032755         | 5.90E – 05     | 1.08E – 02       | 42  | 4                  | 6                 |
| Regulation of cell proliferation                         | GO:0042127         | 1.80E – 04     | 3.10E – 02       | 1.9 | 44                 | 550               |
| Intracellular lipid transport                            | GO:0032365         | 2.60E – 04     | 3.55E – 02       | 21  | 4                  | 8                 |
| Neutrophil chemotaxis                                    | GO:0030593         | 2.70E – 04     | 3.55E – 02       | 12  | 5                  | 14                |
| Regulated secretory pathway                              | GO:0045055         | 2.70E – 04     | 3.55E – 02       | 12  | 5                  | 14                |
| Protein amino acid phosphorylation                       | GO:0006468         | 2.70E – 04     | 3.55E – 02       | 1.8 | 47                 | 613               |
| Regulation of cytokine biosynthetic process              | GO:0042035         | 3.90E – 04     | 4.55E – 02       | 4.2 | 10                 | 61                |
| Purine ribonucleoside monophosphate biosynthetic process | GO:0009168         | 3.90E – 04     | 4.55E – 02       | 11  | 5                  | 15                |

So  
**inflammation associates with rupture in  
human aneurysms**

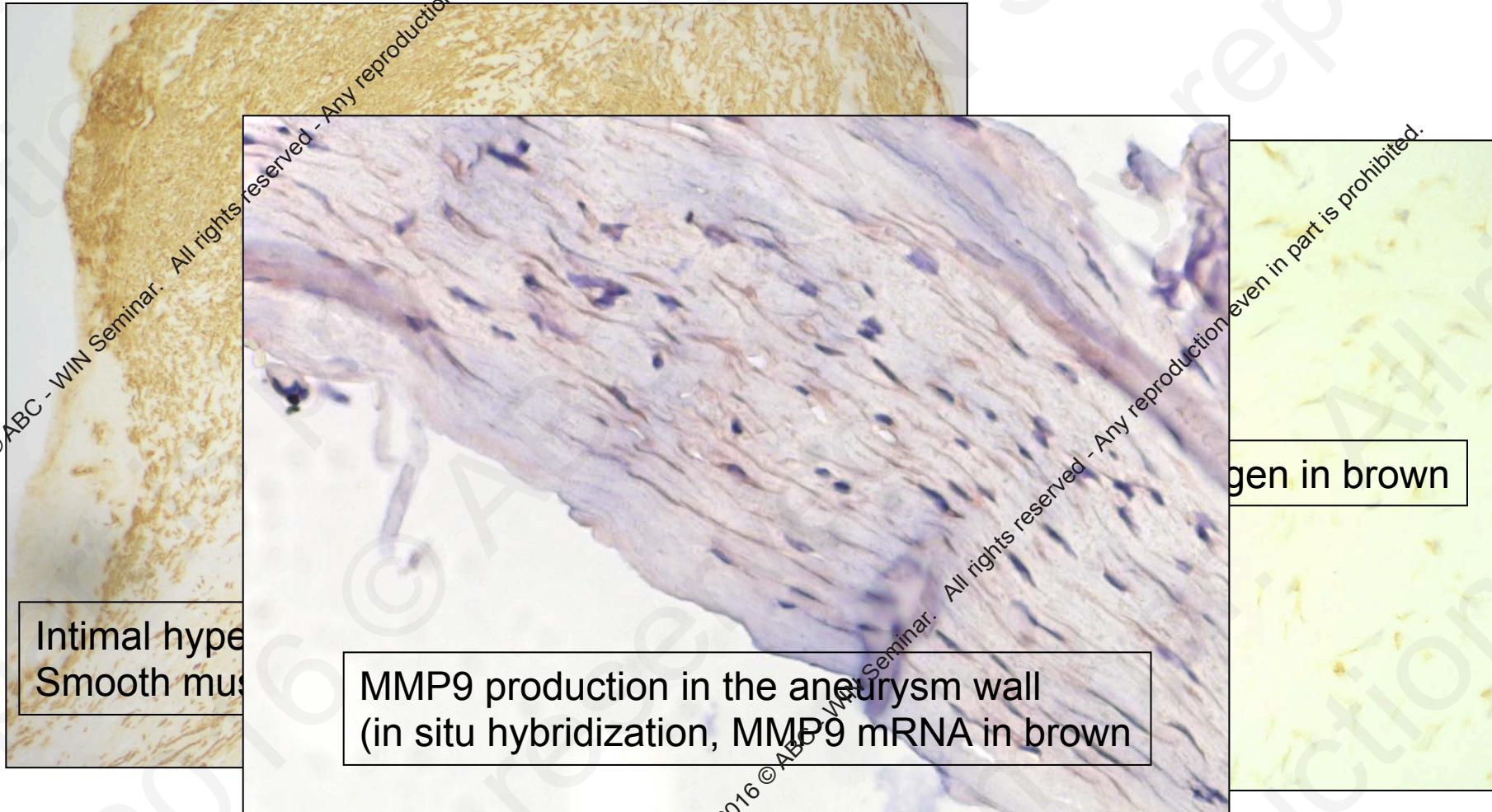
**– but is it the cause of rupture?**

# Degenerative remodeling of aneurysm wall – What characterizes a ruptured aneurysm wall

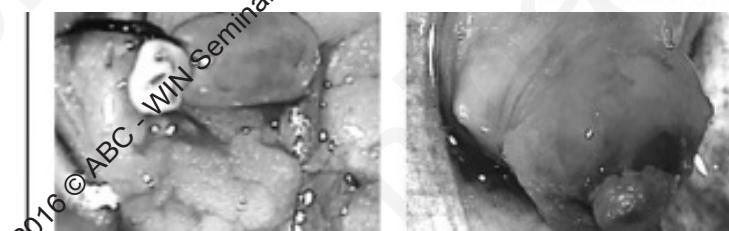
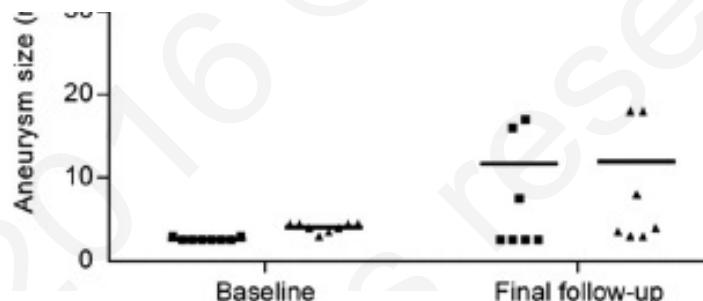
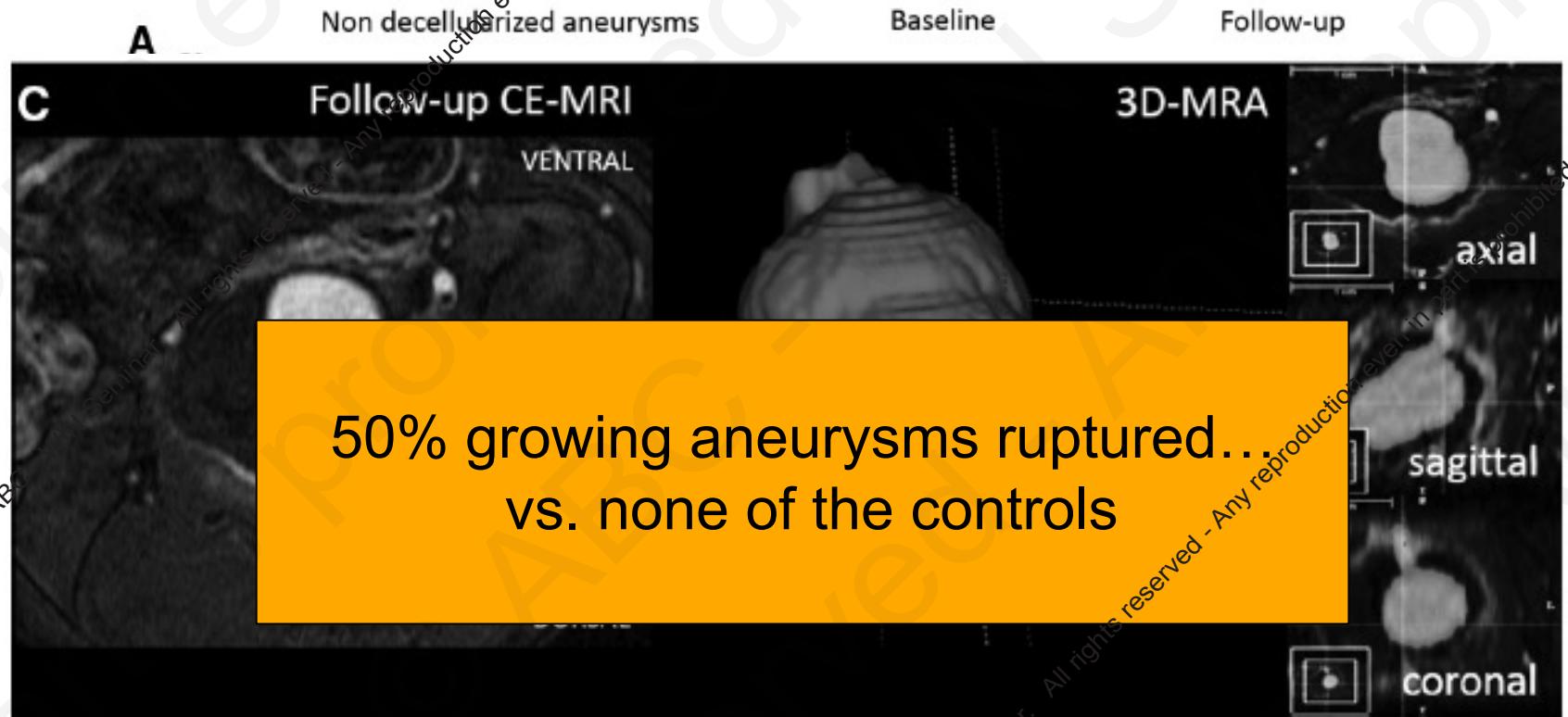
→ Loss of smooth muscle cells



# Why smooth muscle cells are important? Capacity for matrix synthesis & adaptive remodeling

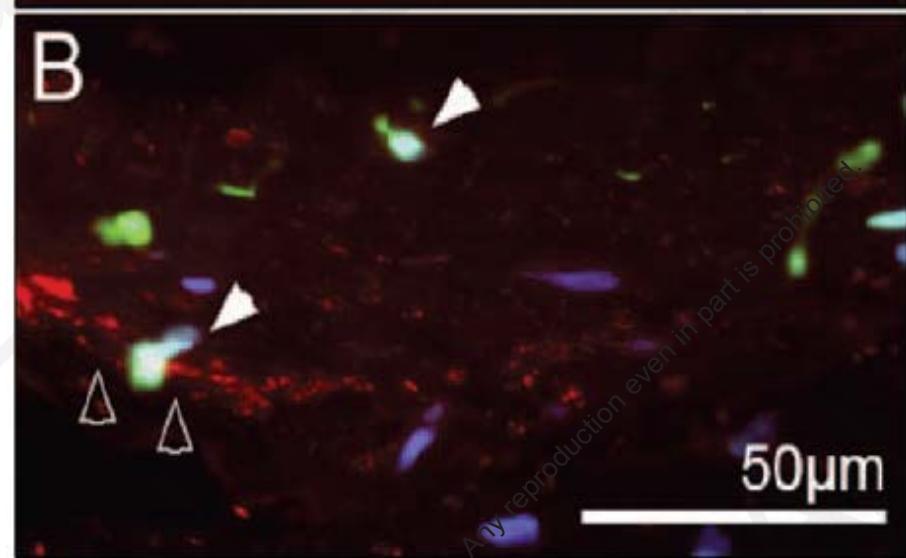
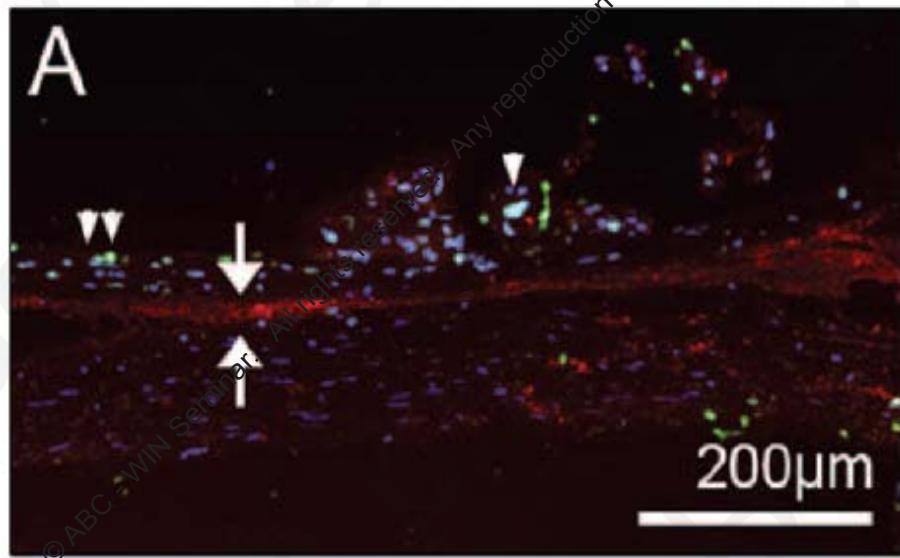


# Loss of mural cells is what causes aneurysm growth and rupture...



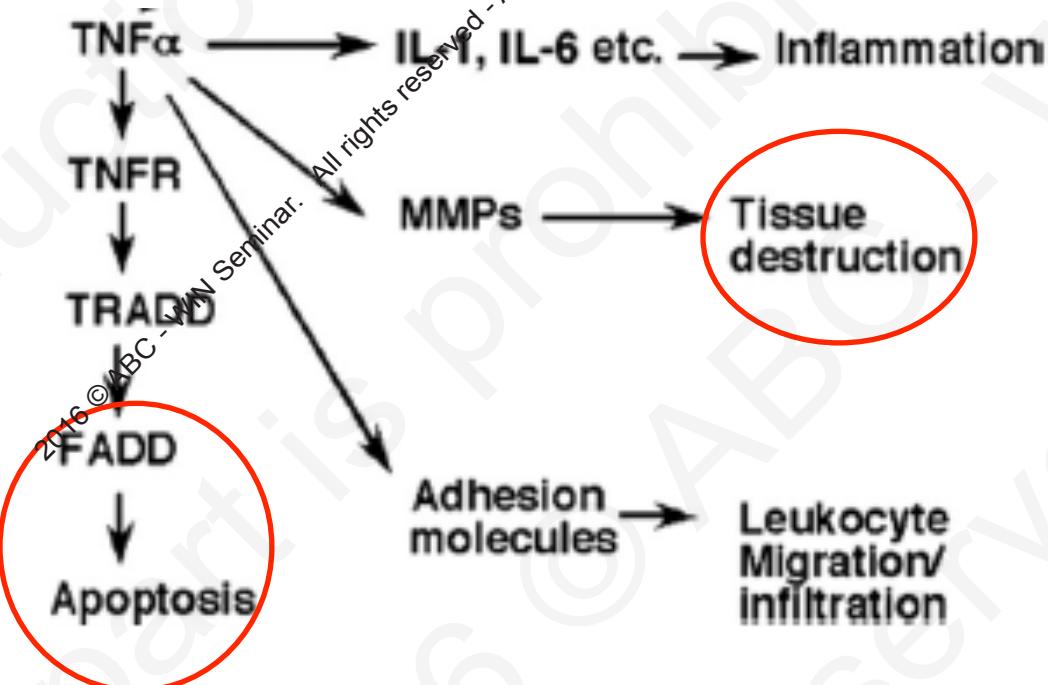
Marbacher et al. Stroke 2014

# No evidence that complement acts directly on smooth muscle cells

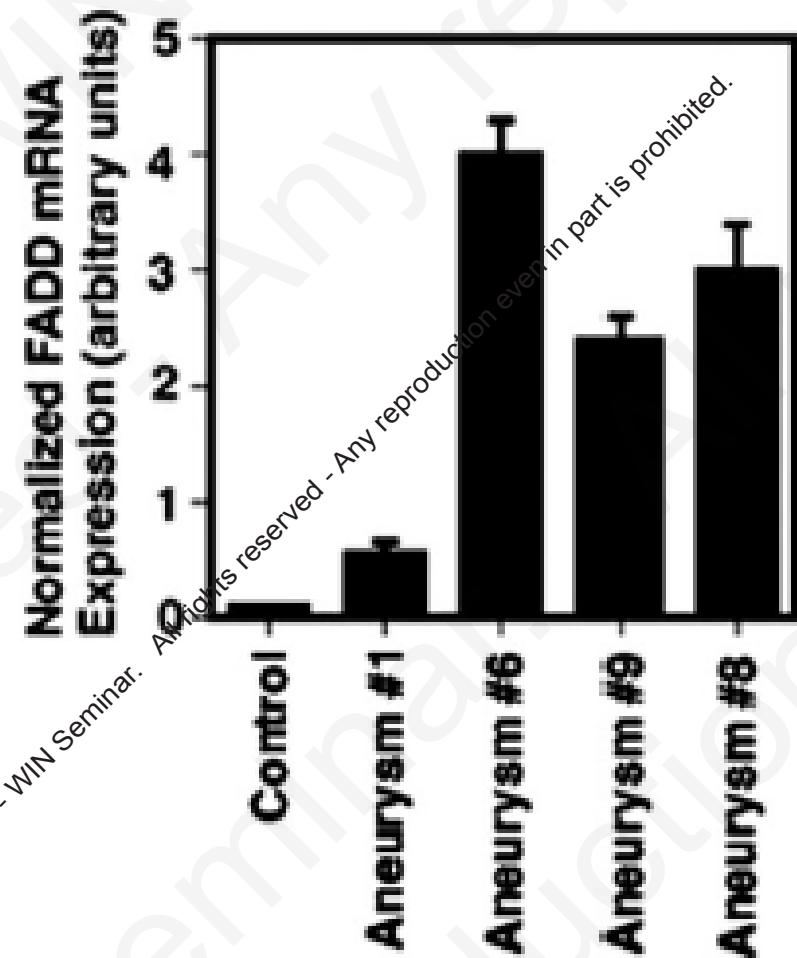


Complement activation (**in red**) and cell death (**green/turquoise**)  
DO NOT colocalize in the aneurysm wall

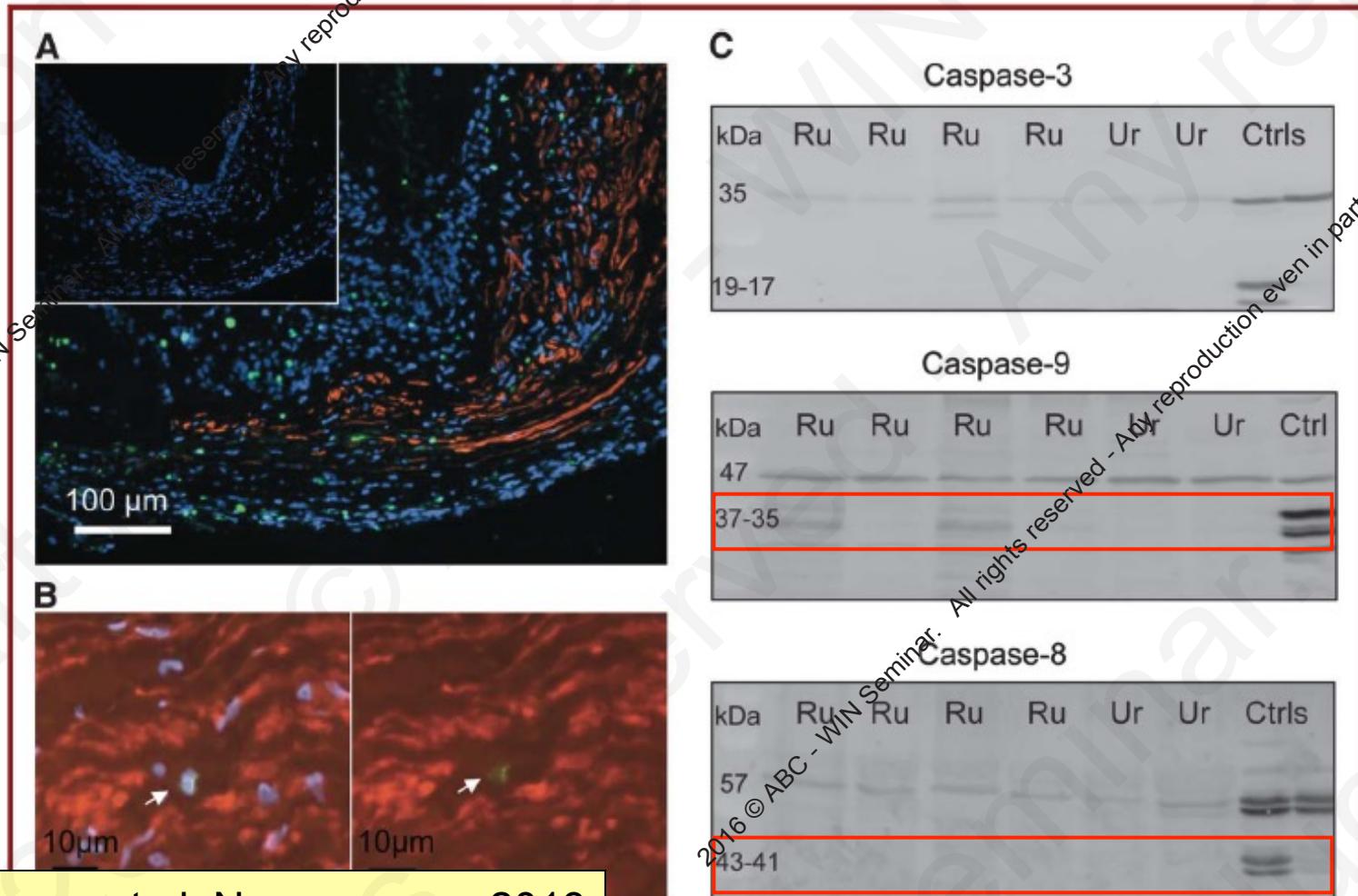
# Macrophage derived cytokines – TNF $\alpha$ implicated in aneurysm formation in mice ... what about rupture in human aneurysms?



Jayaraman et al. Neurosurgery 2006



# No evidence for apoptosis induced by inflammatory cells (Caspase-8 activation) in human aneurysm wall – other factors implicated (Caspase-9 activation)

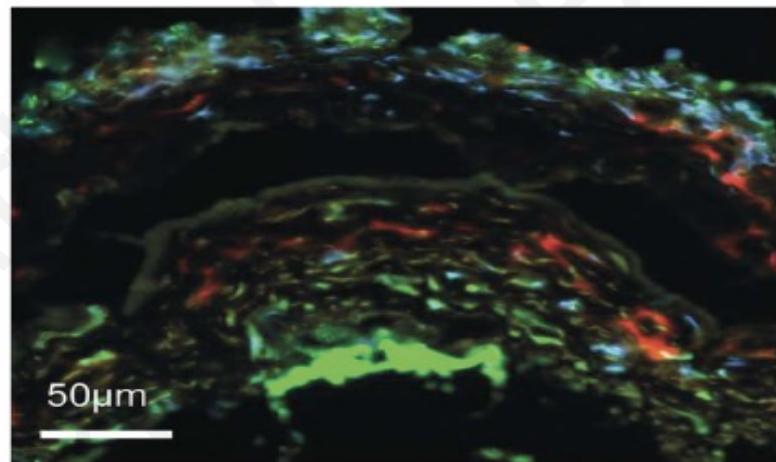


... instead apoptosis & inflammatory cell infiltration in the aneurysm wall associates with response to oxidative stress  
- and some inflammatory cells express hemeoxygenase-1

A

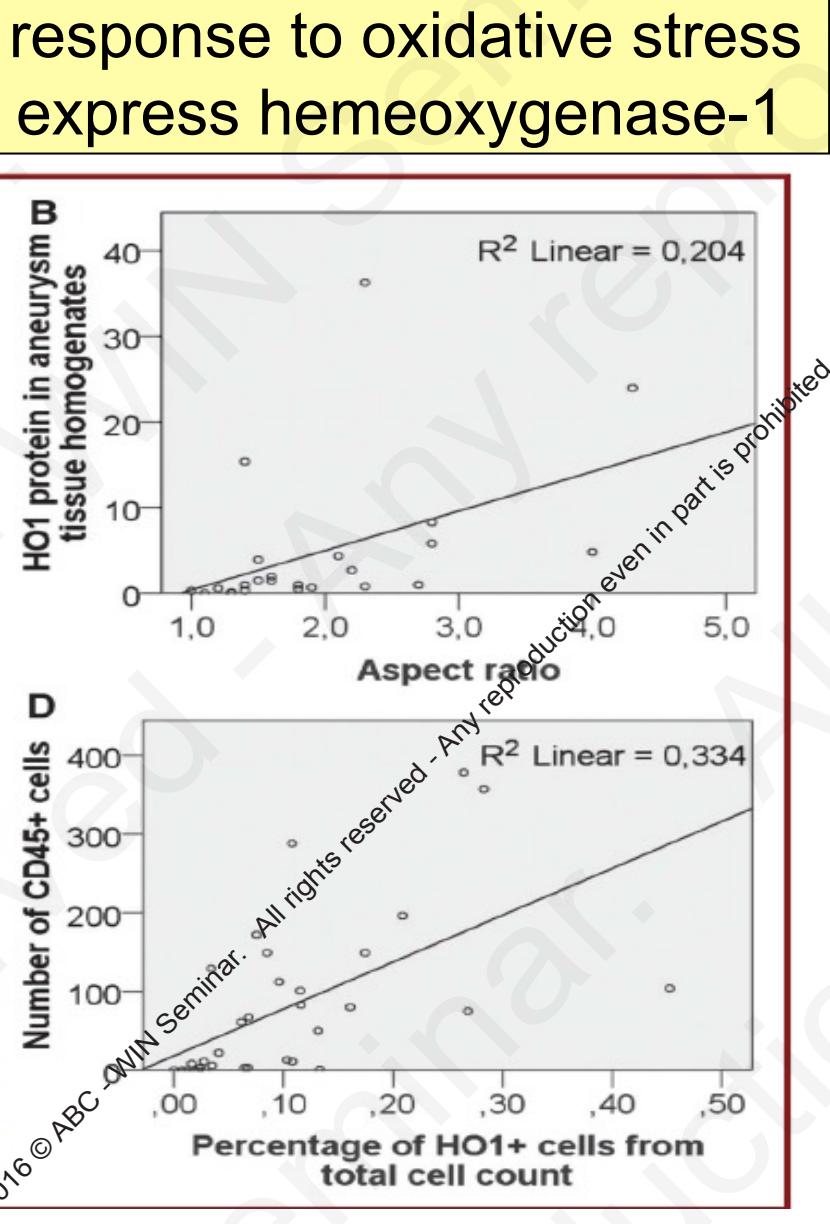


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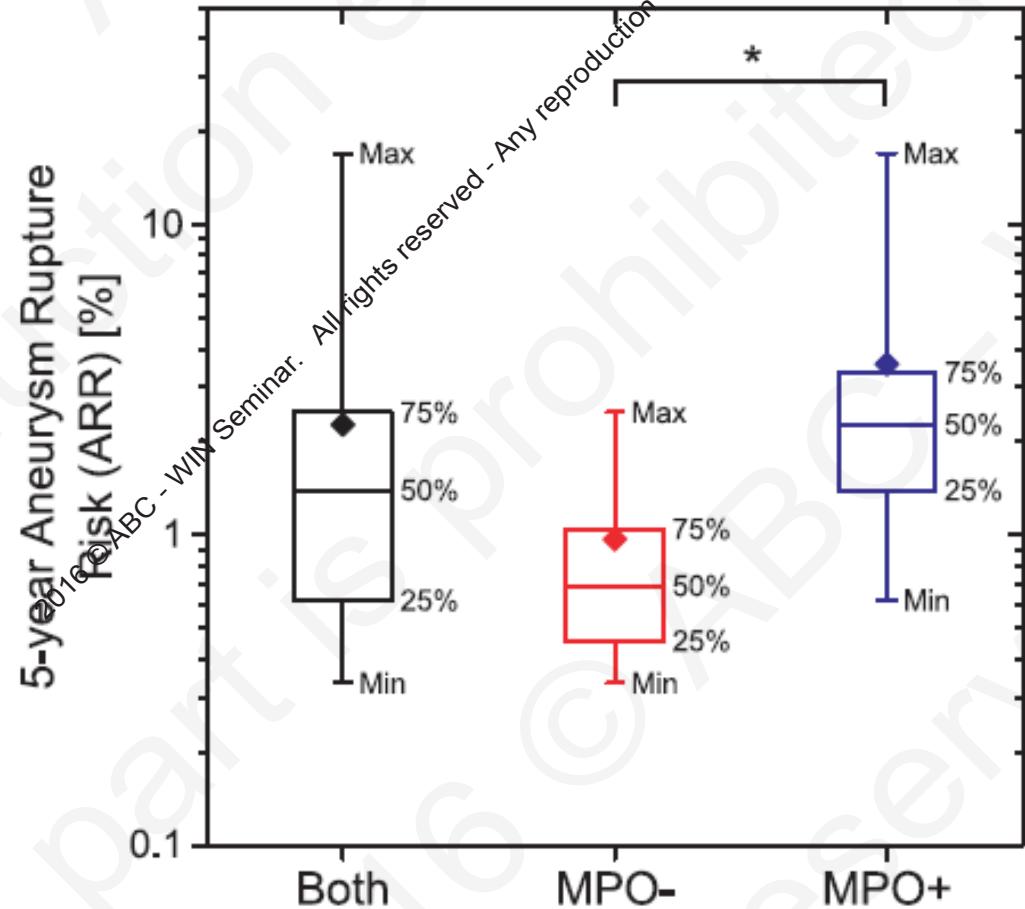


HO1 protein in aneurysm tissue homogenates

Number of CD45+ cells

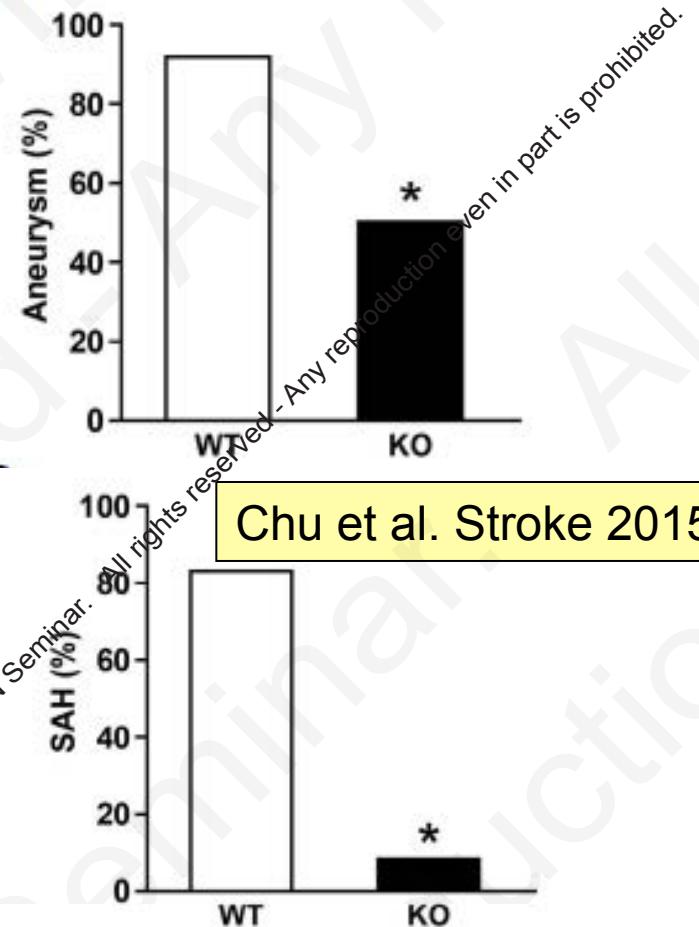


# Myeloperoxidase (derived from neutrophils & macrophages) causes oxidative stress – and associates with rupture risk in aneurysms



Gounis et al. Stroke 2014

Aneurysm formation & SAH in myeloperoxidase knock out mice



Chu et al. Stroke 2015

1) So other causes – perhaps oxidative stress- rather than macrophage induced cell death are the primary cause of smooth muscle cell loss

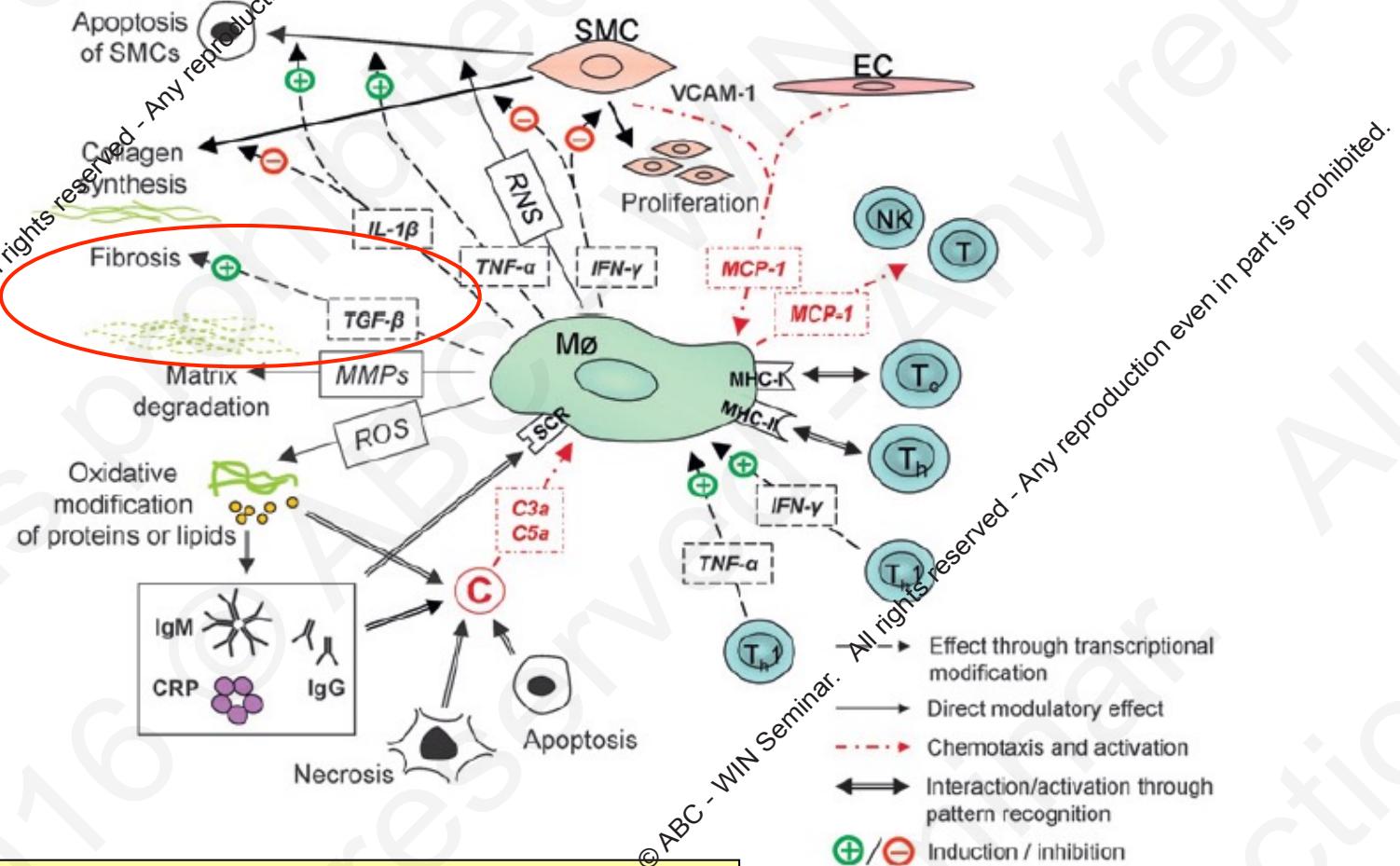
and

2) Some of the inflammatory cells perhaps cause oxidative stress

– whereas others are a response to it?

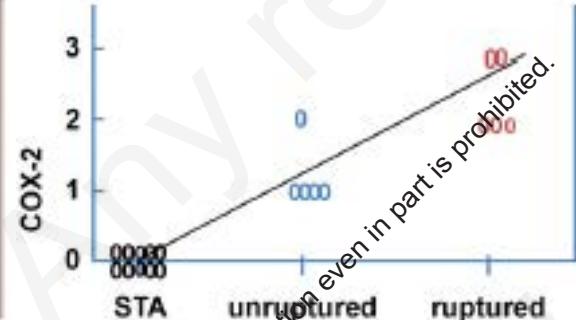
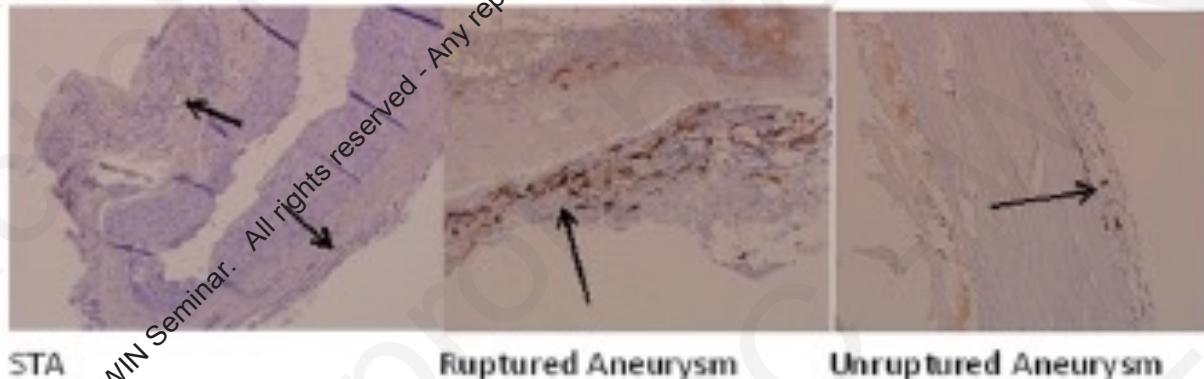
... not all inflammatory cells or macrophages in the aneurysm wall necessarily have the same function?

# Macrophages affect smooth muscle cell function in multiple ways – and can even promote fibrosis

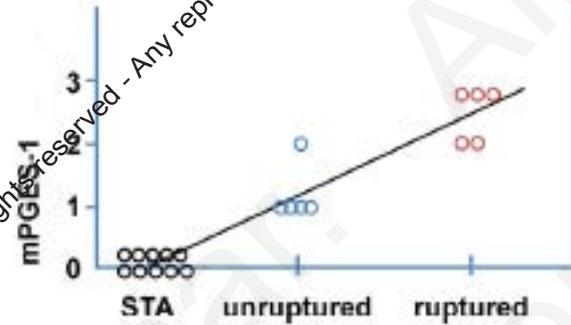
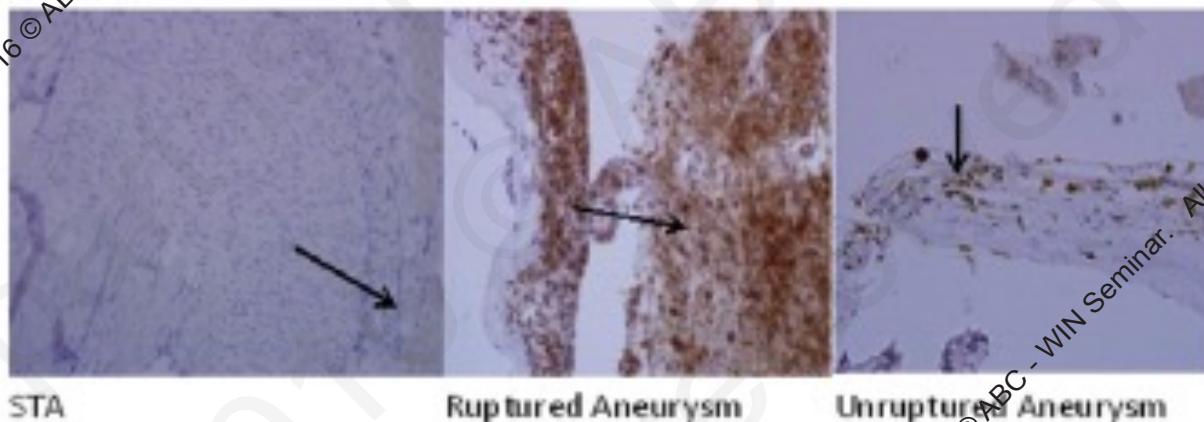


# Macrophage derived cytokines – COX2 signaling implicated in aneurysm formation ... what about rupture in human aneurysms?

A COX-2:



B mPGES-1:



There is evidence that COX inhibition reduces  
 1) inflammation and  
 2) risk of aneurysm rupture in humans

**Table 5. Nested Case-Control: Hemorrhages and Control Subjects (Matched by Site and Size)—Frequency of Aspirin Use**

|                                     | Control Subjects |         | Cases |         | OR   | <i>P</i> for Linear Association<br>Conditional Odds* |
|-------------------------------------|------------------|---------|-------|---------|------|--|
|                                     | No.              | Percent | No.   | Percent |      |  |
| Use of aspirin (grouped)            |                  |         |       |         |      | 0.025  |
| Never                               | 109              | 73.6    | 39    | 26.4    | 1    |  |
| Less than or equal to once a mo     | 23               | 79.3    | 6     | 20.7    | 0.80 |  |
| More than once a mo to 2 times a wk | 14               | 77.8    | 4     | 22.2    | 0.86 |  |
| Three times a wk to daily           | 67               | 88.2    | 9     | 11.8    | 0.40 |  |

Hasan et al. Stroke 2011

Hasan et al. JAH 2013

## **Conclusion from studies on patients samples:**

**Inflammation does not seem to directly kill aneurysm wall cells but affects rupture risk**

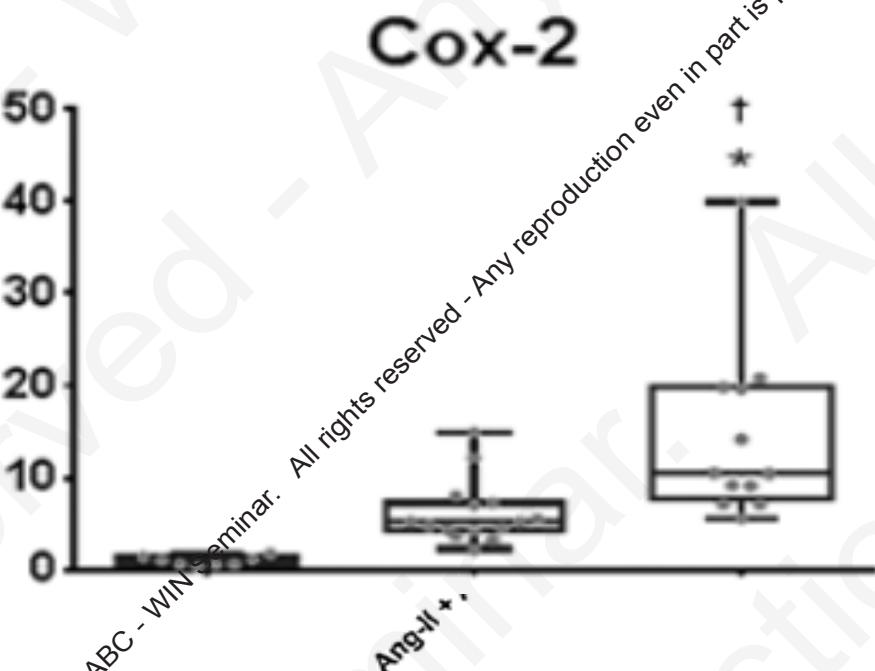
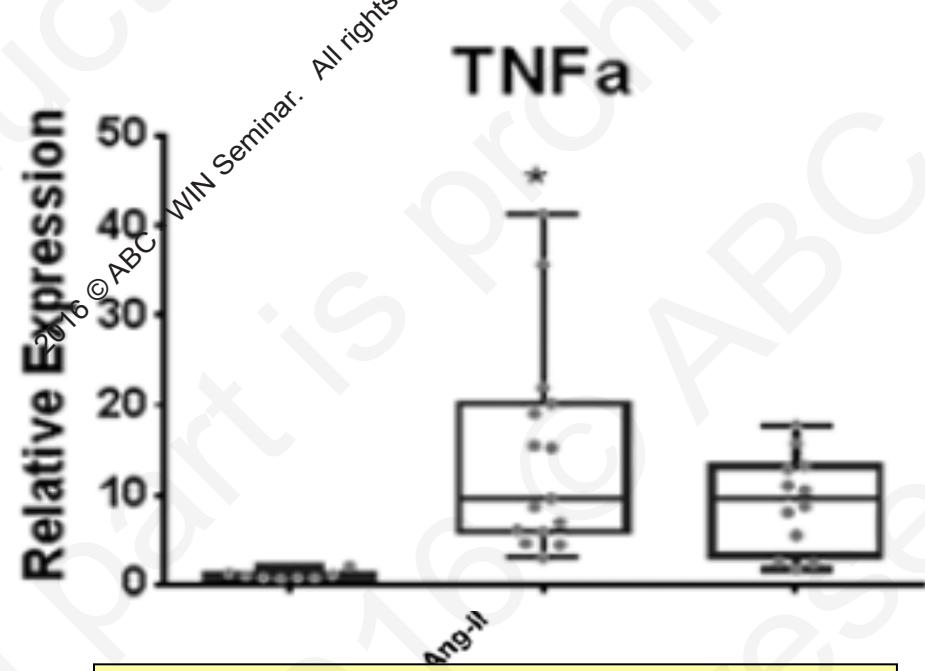
# **What about animal models – does inflammation predispose to rupture?**

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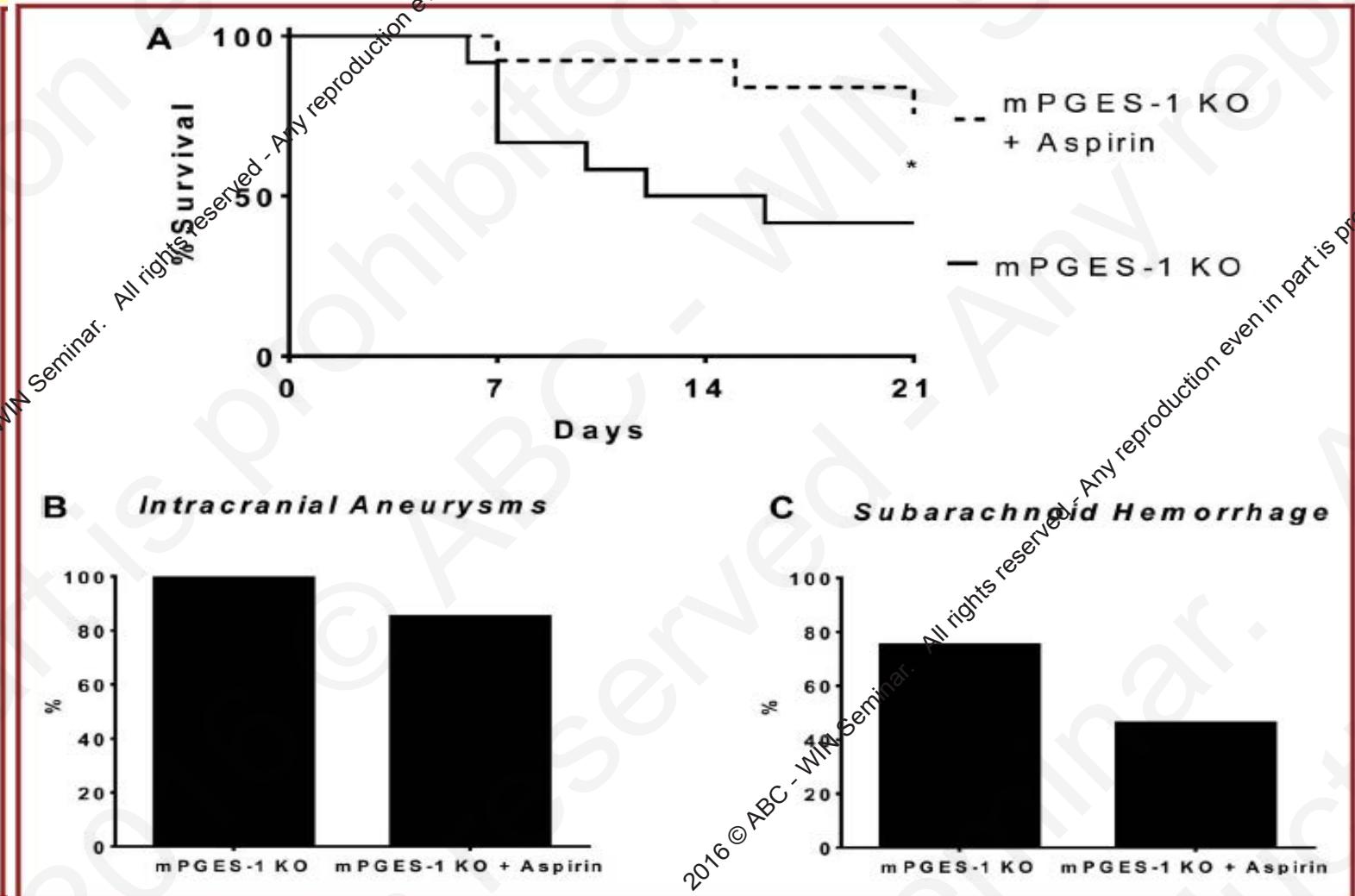
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# Blocking angiotensin II mediated signaling with Ang 1-7:

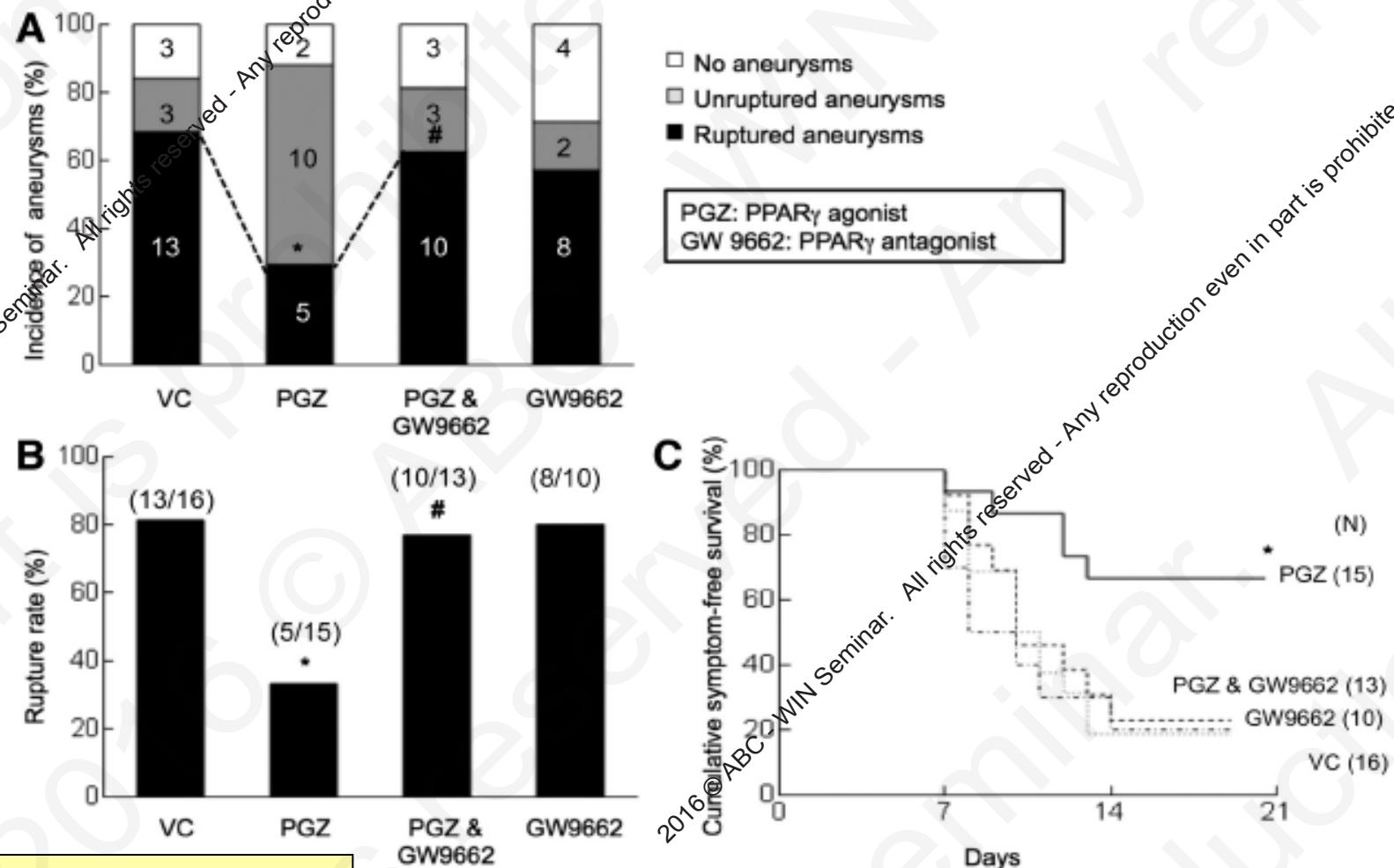
- 1) Reduced aneurysm formation
- 2) Reduced aneurysm rupture
- 3) Reduced TNFa mediated inflammation
- ...BUT
- 4) Increased COX-2 expression -?



# Blocking COX2 related mPGES-1 increased both aneurysm formation and rupture - and effect reduced by aspirin



# Activation of the immunomodulatory PPAR $\gamma$ receptor in macrophages with pioglitatzone (PGZ) reduced aneurysm formation and **rupture**



**Studies on the aneurysms induced with hypertension + elastase demonstrate:**

**Modulation of the macrophage activity DOES influence the risk of aneurysm rupture**

**... but can also increase, not just decrease rupture risk...**

**Role of the inflammatory response in aneurysm healing?**

# Do the animal models reproduce the inflammation + pathobiology of human disease?

- role of inflammation is multiplex and depends on the context
  - e.g. inflammation promotes fibrosis and wound healing
  - Inflammation promotes vessel wall thickening in vascular injury
  - Inflammation may promote plaque healing as well as plaque rupture in atherosclerosis..

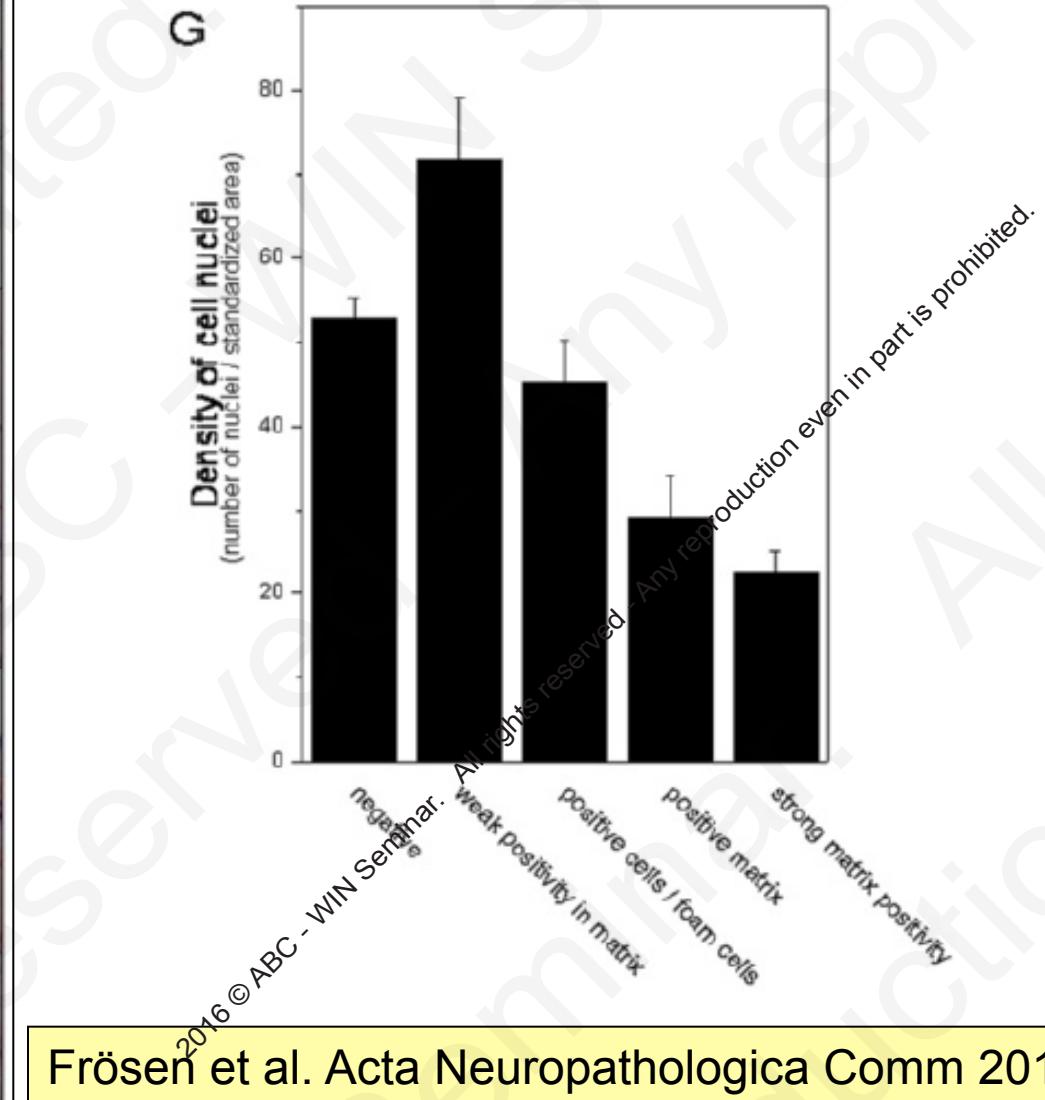
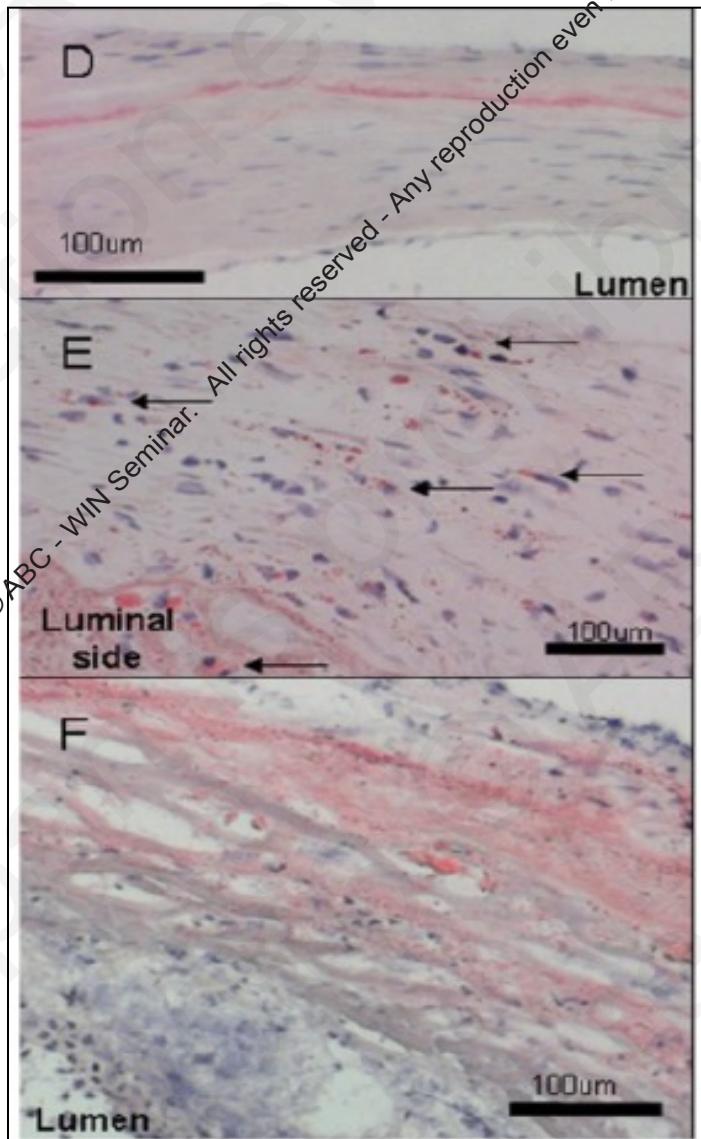
# Genome wide comparison of gene expression in ruptured and unruptured human aneurysms

**–also lipid accumulation strongly associates with rupture**

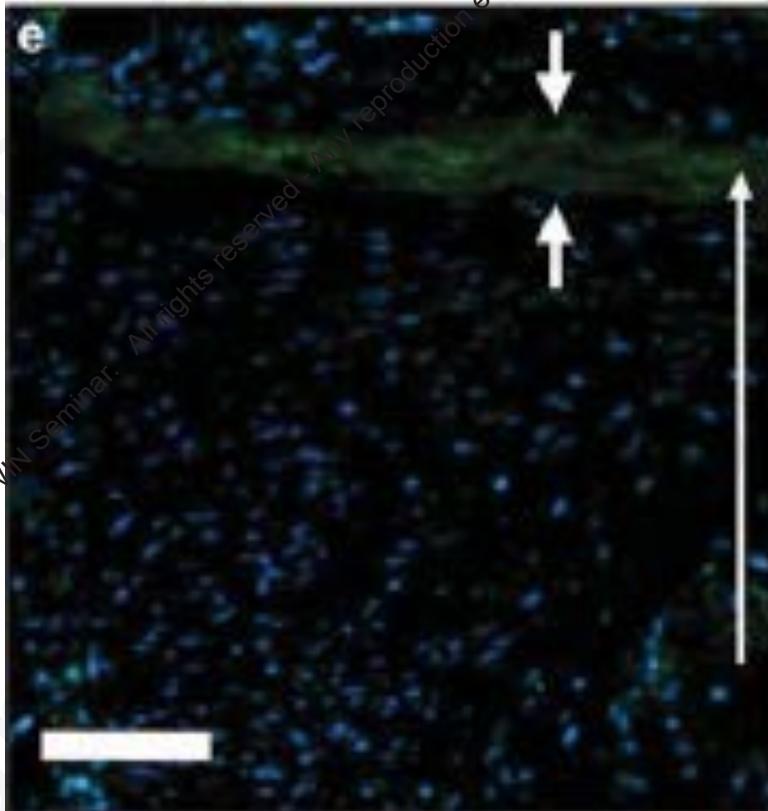
**TABLE 2. Biological Processes in Ruptured Saccular Intracranial Aneurysm Wall Samples<sup>a</sup>**

| Gene Ontology (GO) Biological Processes                  | Upregulated Genes  |                |                  |     |                    |                   |
|--|--------------------|----------------|------------------|-----|--------------------|-------------------|
|  | GO ID <sup>b</sup> | P <sup>c</sup> | FDR <sup>d</sup> | OR  | Count <sup>e</sup> | Size <sup>f</sup> |
| Chemotaxis   | GO:0006935         | 2.60E – 14     | 5.66E – 11       | 7   | 30                 | 125               |
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| Inflammatory response                                    | GO:0006954         | 1.40E – 13     | 7.52E – 11       | 4.2 | 46                 | 296               |
| Locomotory behavior                                      | GO:0007626         | 1.50E – 11     | 6.46E – 09       | 4.5 | 35                 | 208               |
| Response to stress                                       | GO:0006950         | 3.60E – 09     | 1.31E – 06       | 2.1 | 99                 | 1224              |
| Response to other organism                               | GO:0051707         | 4.70E – 08     | 1.49E – 05       | 5.2 | 18                 | 87                |
| Positive regulation of tumor necrosis factor production  | GO:0032760         | 5.80E – 08     | 1.60E – 05       | 7.7 | 6                  | 7                 |
| Locomotion   | GO:0040011         | 1.30E – 07     | 3.18E – 05       | 4.7 | 20                 | 114               |
| Cytokine production                                      | GO:0001816         | 2.50E – 05     | 5.49E – 03       | 3.8 | 16                 | 108               |
| Phosphate metabolic process                              | GO:0006796         | 5.30E – 05     | 1.06E – 02       | 1.8 | 66                 | 893               |
| Positive regulation of interleukin-6 production          | GO:0032755         | 5.90E – 05     | 1.08E – 02       | 42  | 4                  | 6                 |
| Regulation of cell proliferation                         | GO:0042127         | 1.80E – 04     | 3.10E – 02       | 1.9 | 44                 | 550               |
| Intracellular lipid transport                            | GO:0032365         | 2.60E – 04     | 3.55E – 02       | 21  | 4                  | 8                 |
| Neutrophil chemotaxis                                    | GO:0030593         | 2.70E – 04     | 3.55E – 02       | 12  | 5                  | 14                |
| Regulated secretory pathway                              | GO:0045055         | 2.70E – 04     | 3.55E – 02       | 12  | 5                  | 14                |
| Protein amino acid phosphorylation                       | GO:0006468         | 2.70E – 04     | 3.55E – 02       | 1.8 | 47                 | 613               |
| Regulation of cytokine biosynthetic process              | GO:0042035         | 3.90E – 04     | 4.55E – 02       | 4.2 | 10                 | 61                |
| Purine ribonucleoside monophosphate biosynthetic process | GO:0009168         | 3.90E – 04     | 4.55E – 02       | 11  | 5                  | 15                |

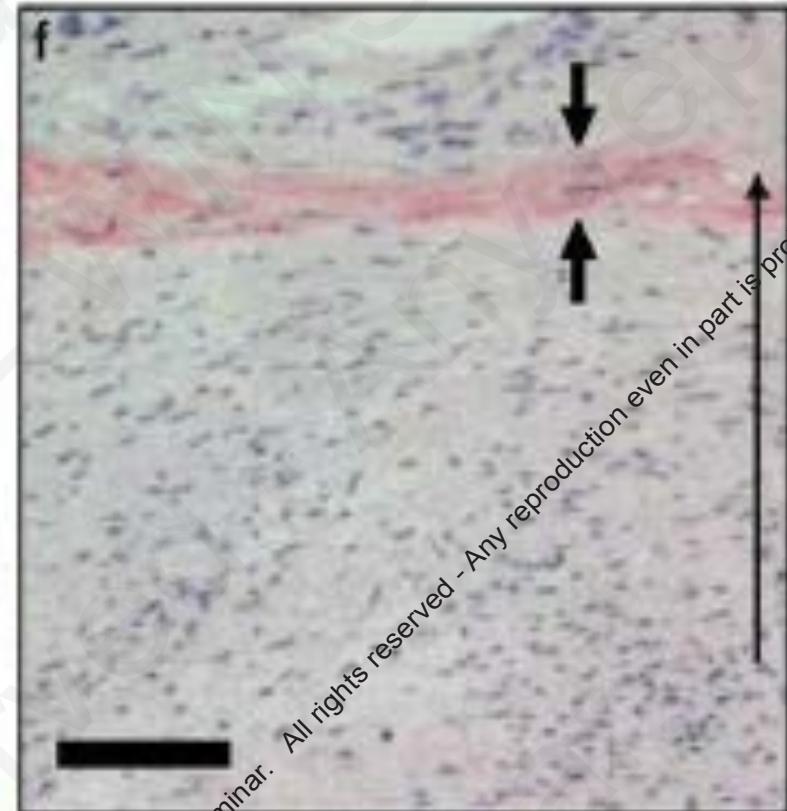
# Accumulation of lipids in the aneurysm wall associates with loss of mural cells



# Complement activation in aneurysm wall regions with lipid accumulation



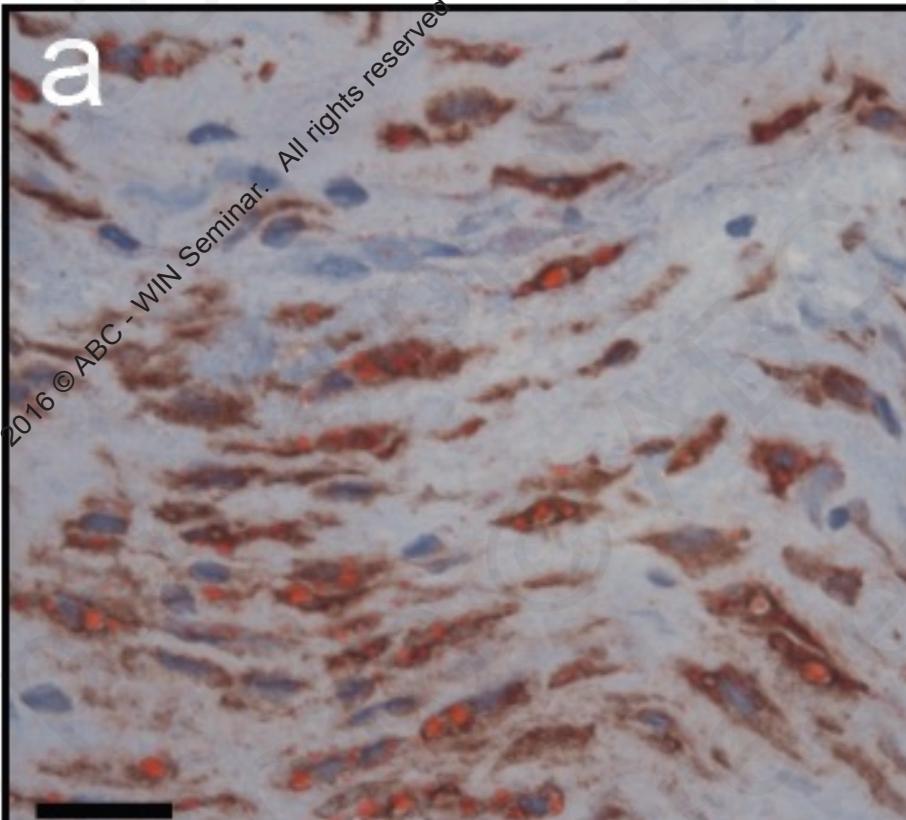
Complement in green



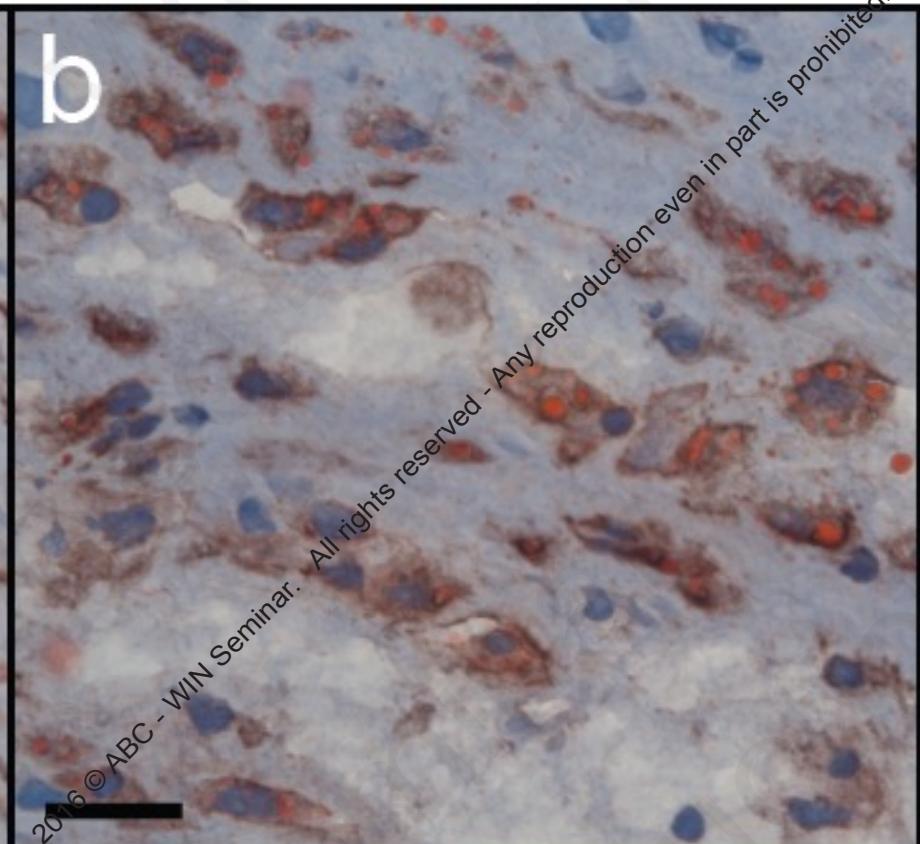
Lipids in red

# Macrophage infiltration in the aneurysm wall associates with accumulation of lipids – that the macrophages phagocytose

**CD68+ORO**



**CD163+ORO**

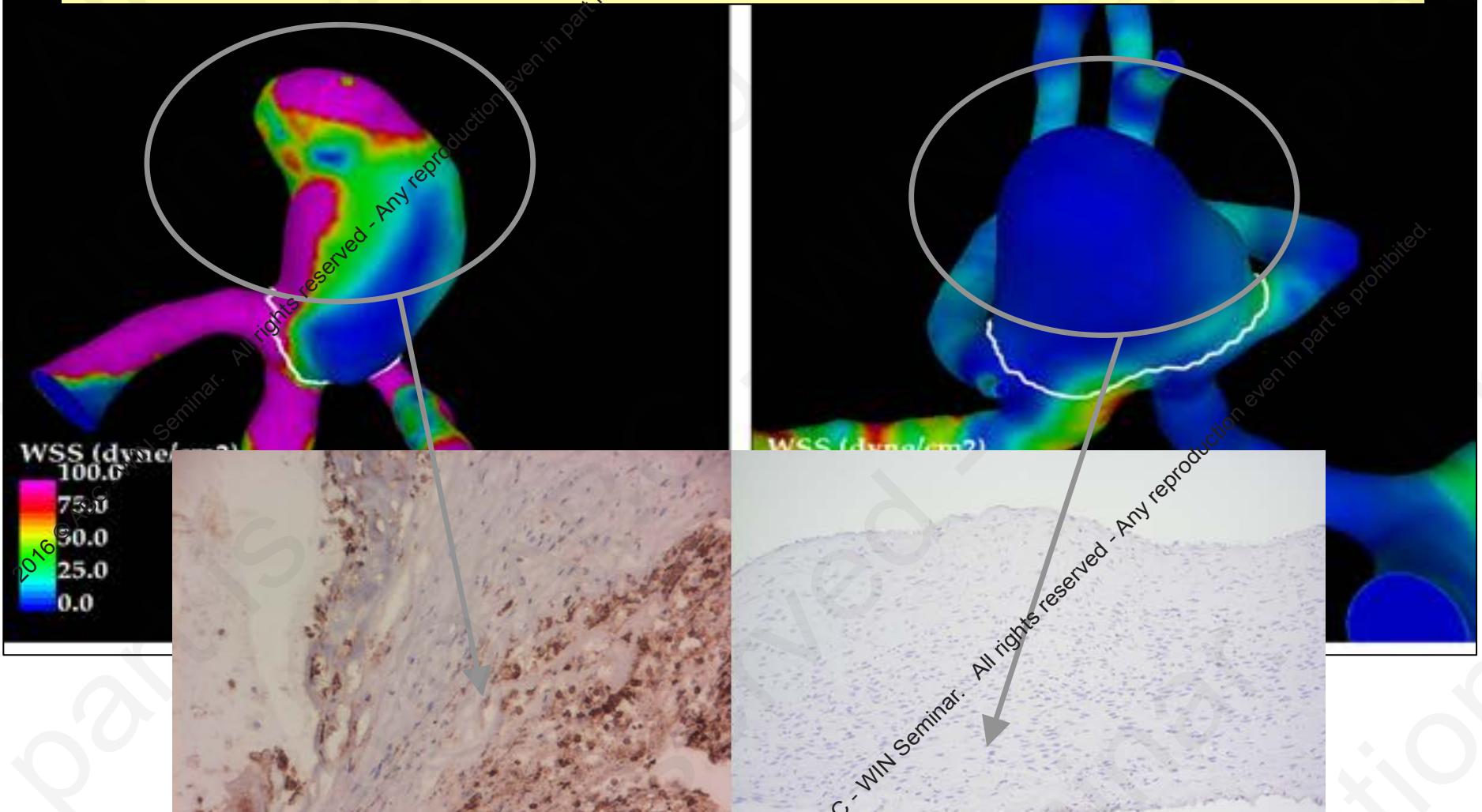


So in human intracranial aneurysms macrophages seem to remove /clear lipids that otherwise would promote death of smooth muscle cells...

-Does lipid accumulation occur in animal models of induced aneurysms?

- How does that change the effects of anti-inflammatory therapy?

# Flow dynamics in human intracranial aneurysms associate with inflammation in the wall – different flow dynamics in mouse models?



Inflammatory cells (in brown) are found especially in the walls of aneurysms with HIGH WSS - (Cebral et al. submitted)

Moreover...

## Odontogenic bacteria in human aneurysm wall

- what is their role?
- how will anti-inflammatory therapy affect?

**Table 2** The prevalence of bacterial DNA in the samples

|                   | Negative | Positive |
|-------------------|----------|----------|
| Surgical patients | 11 (38%) | 18 (62%) |
| Autopsy group     | 4 (57%)  | 3 (43%)  |
| Total             | 15 (42%) | 21 (58%) |

# In summary -

- Inflammation mediates the formation of intracranial aneurysms
  - drug therapy to reduce the formation of aneurysms?
- Inflammation of the aneurysm wall associates with degenerated wall and rupture
  - marker of rupture risk?
- Inflammation is likely not the only cause of the degenerative remodeling but seems to modulate it
  - possible target for drug therapy?

# Thank you.

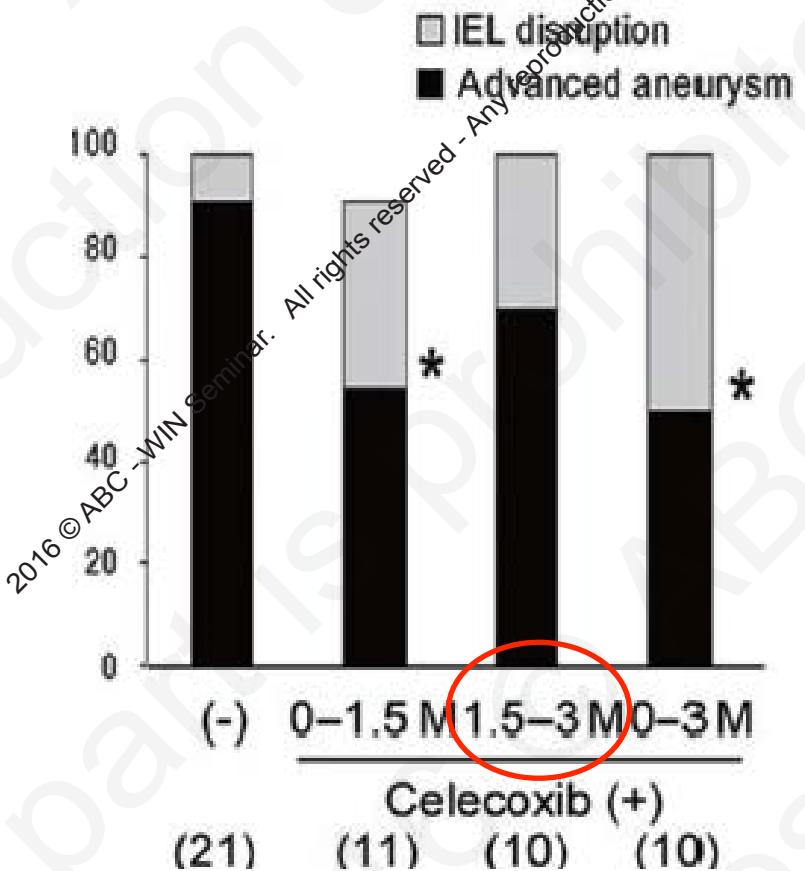
- Acknowledgements:
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- **Professor Mika Niemelä**
- **Dr. Riikka Tulamo**
- **Dr. Eliisa Ollikainen**

KUOPIO

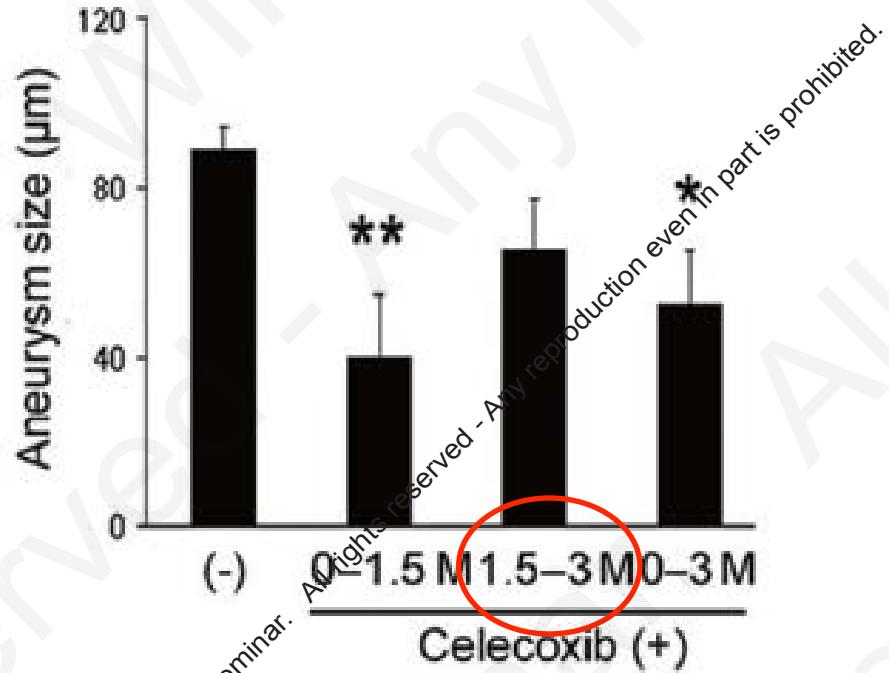


Biomedicum Helsinki

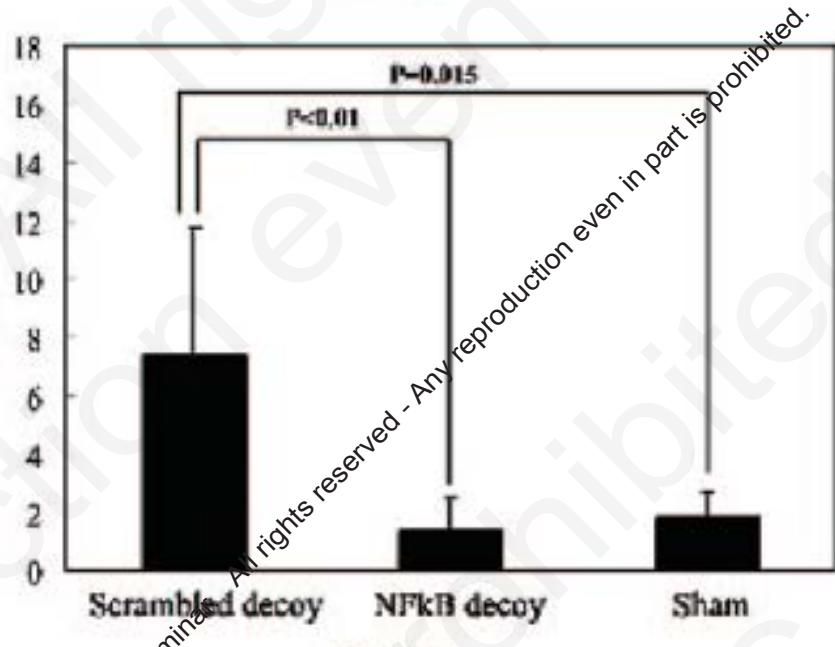
# Blocking the COX2 –EP2 feedback loop in already formed aneurysms perhaps not enough?



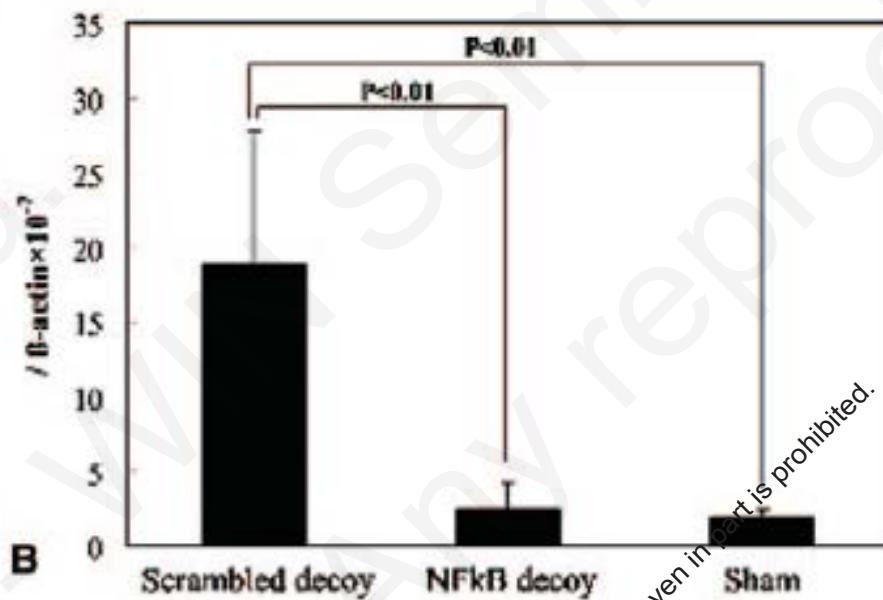
Aoki T et al. Br J Pharmacology 2011



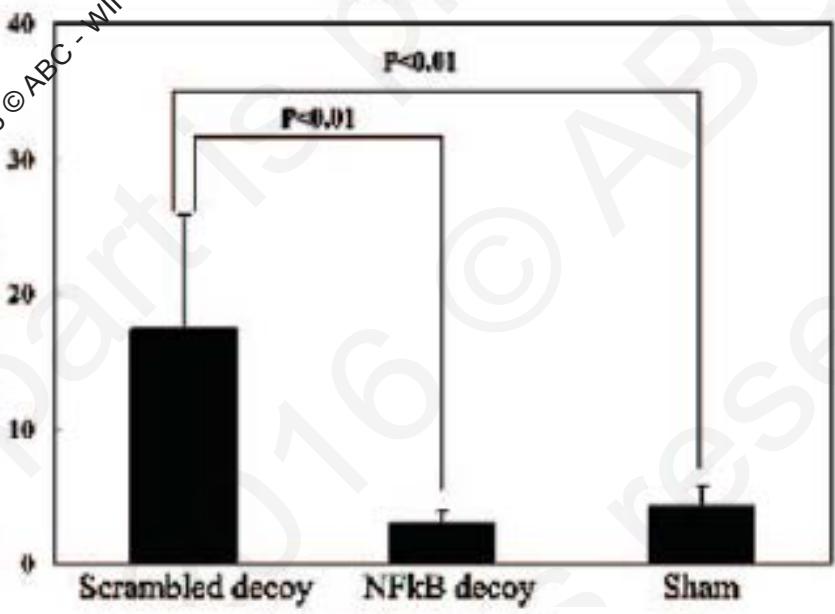
### MCP-1



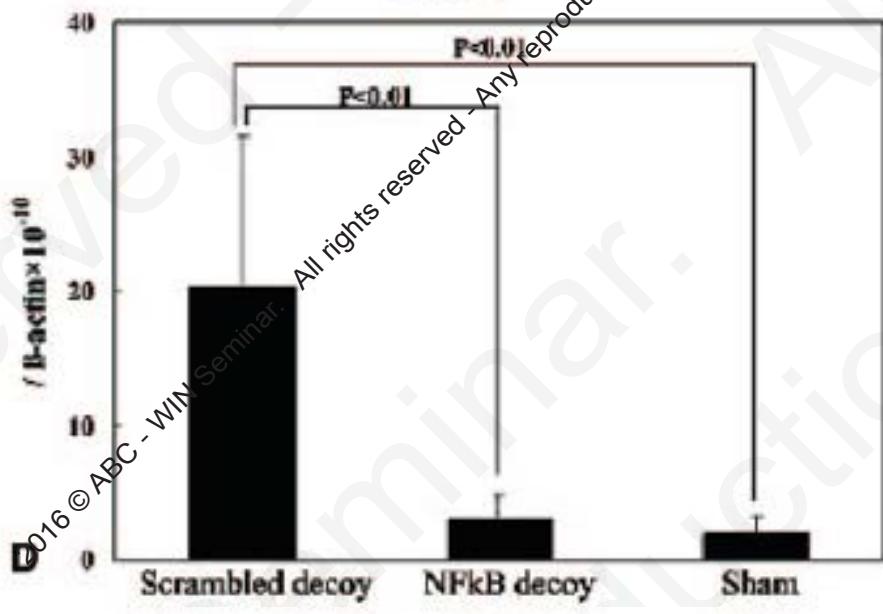
### VCAM-1



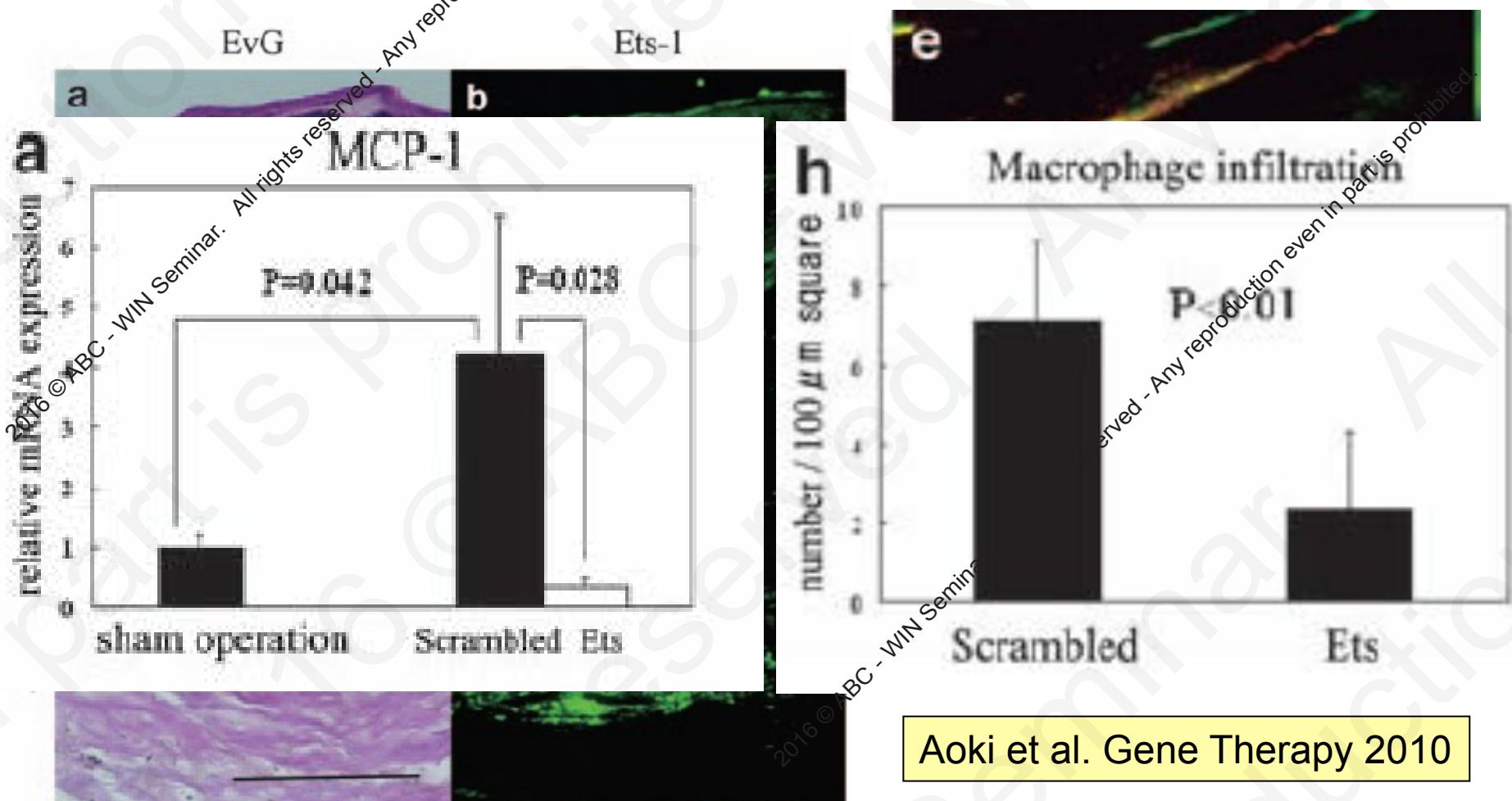
### MMP-2



### MMP-9



# Also activation of Ets-1 transcription factor in SMOOTH MUSCLE CELLS mediates in the carotid ligation + hypertension model the inflammation causing aneurysm formation



## Genome wide gene expression study:

- Inflammation mediated by activation of NFkB and ETS transcription factors in the human aneurysm wall similar to induced aneurysms in animal models

**TABLE 4.** In Silico Analysis of Transcription Factor Binding Sites in the –5-kb Promoter Regions of the 686 Upregulated Genes in the Ruptured Saccular Intracranial Aneurysm Walls <sup>a</sup>

| TFs With Increased Binding Sites<br>TRANSFAC <sup>b</sup> | HGNC <sup>c</sup> | Number of Binding Sites |                           | P           |                      | TF Family   |
|---|-------------------|-------------------------|---------------------------|-------------|----------------------|-------------|
|   |                   | 686 Upregulated Genes   | Human Genome <sup>d</sup> | Uncorrected | Bonferroni corrected |             |
| ELF1  | ELF1              | 411                     | 11 549                    | 1.66E – 08  | 3.32E – 06           | ETS         |
| PEA3  | ETV4              | 714                     | 21 887                    | 1.66E – 07  | 3.32E – 05           | ETS         |
| ETS2  | ETS2              | 413                     | 11 968                    | 3.64E – 07  | 7.28E – 05           | ETS         |
| ETS1  | ETS1              | 728                     | 22 501                    | 3.87E – 07  | 7.74E – 05           | ETS         |
| HIF1  | HIF1A             | 304                     | 8663                      | 3.12E – 06  | 6.26E – 04           | ...         |
| NFKAPPAB65  | RELA              | 96                      | 2232                      | 3.99E – 06  | 1.60E – 03           | Rel/ankyrin |