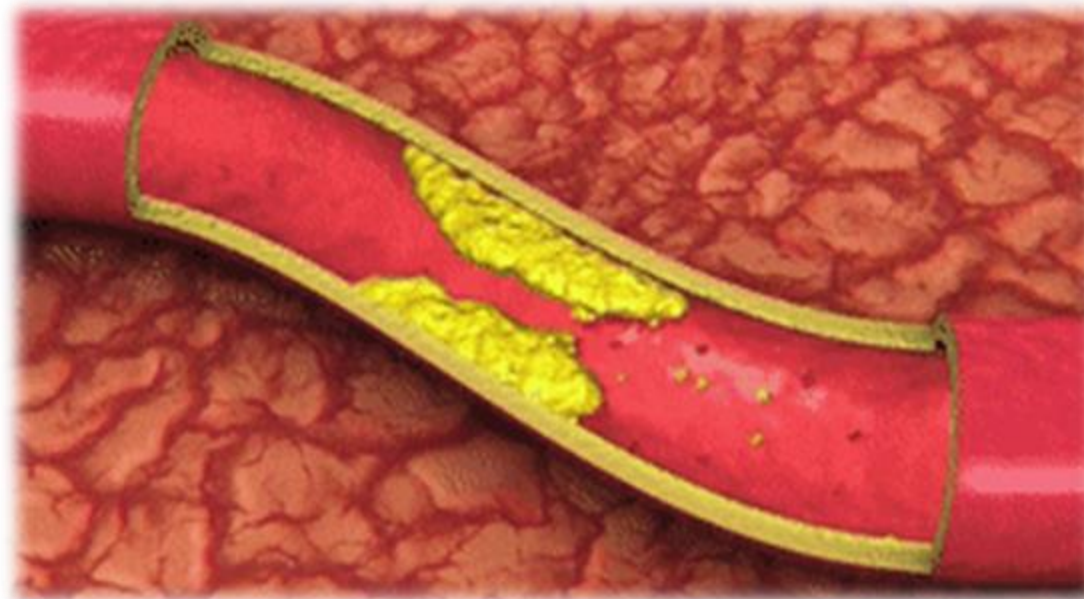
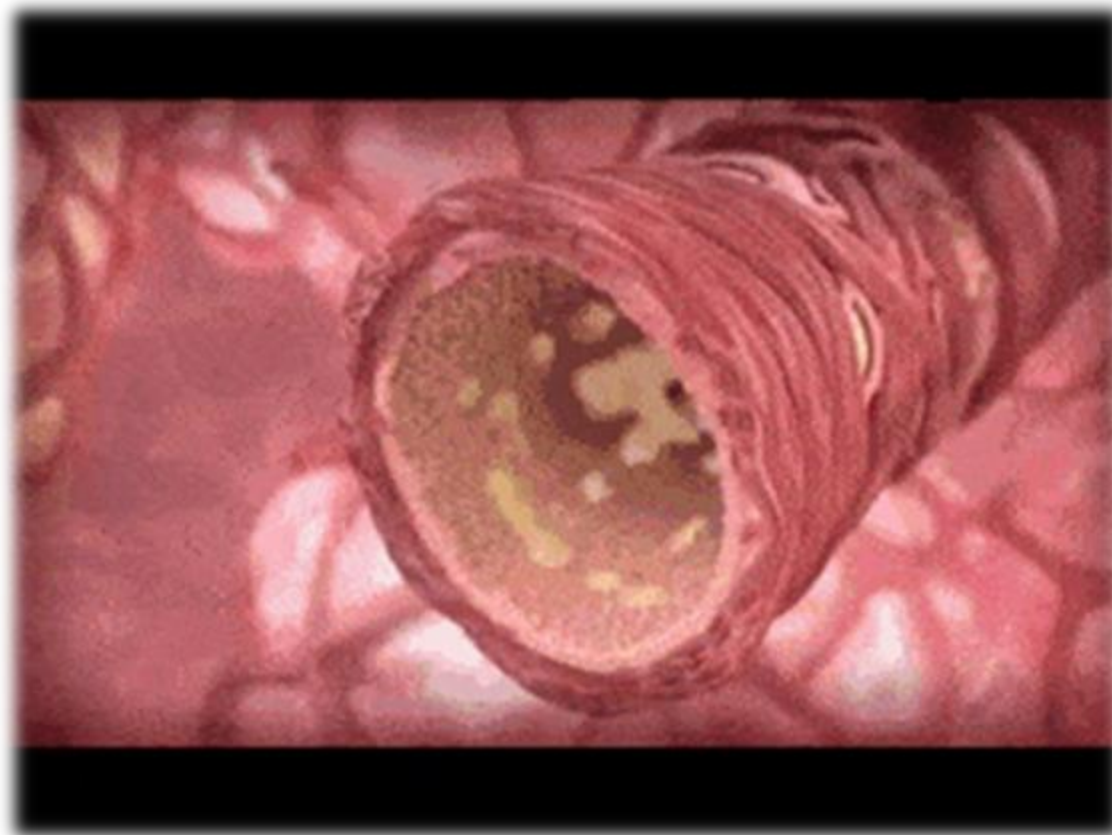




# MAPPING THE CELLULAR ORIGINS OF ATHEROSCLEROTIC PLAQUE

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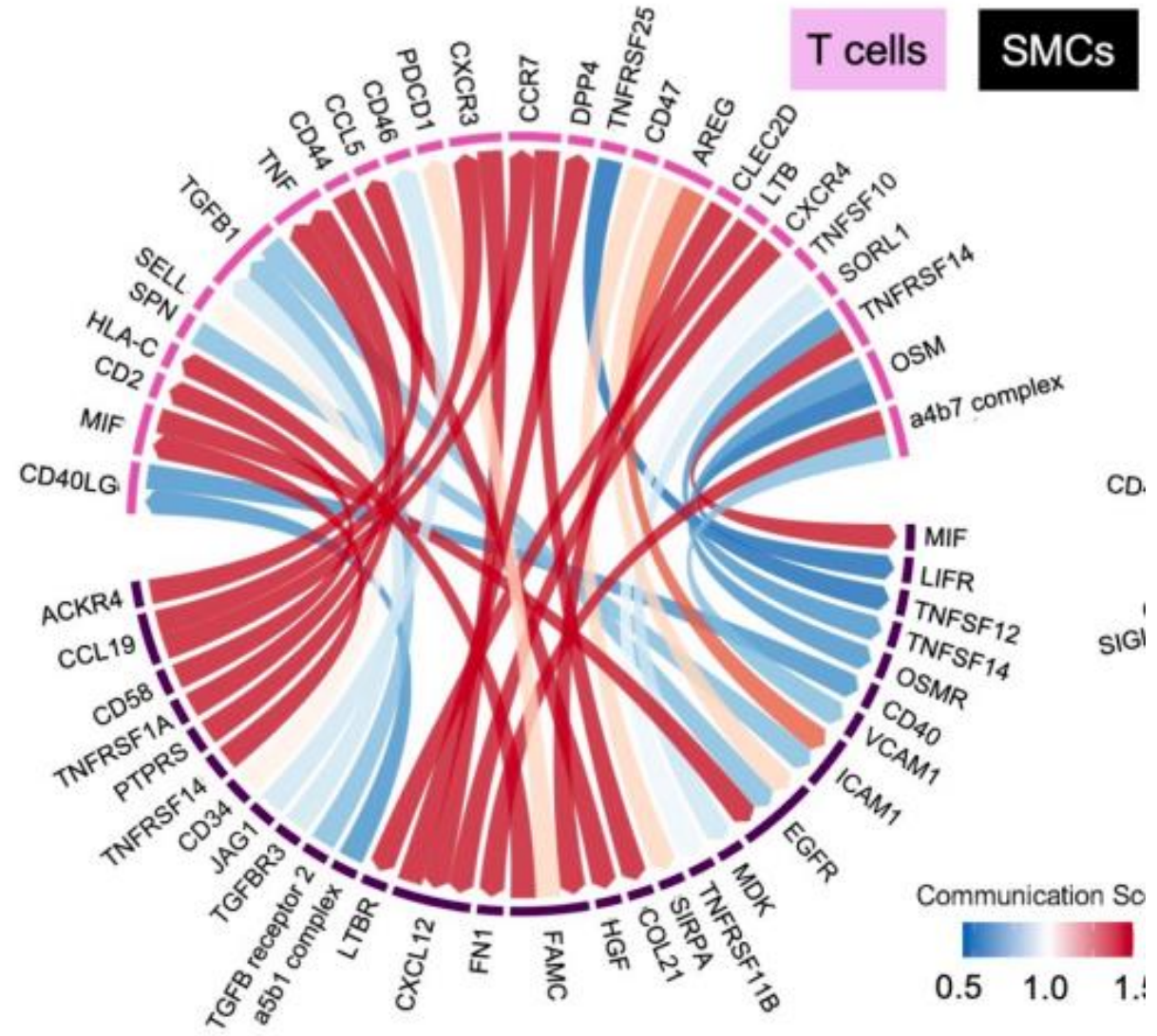
# ATHEROSCLEROSIS



# AMPHIREGULIN

AREG: secretion factor involved in fibrotic development

- possible inducer of plaque development
- chosen factor of interest



Chowdhury et al. 2022

# STATEMENT OF AIMS

**Hypothesis:** We hypothesize that immune cells play a role in the transition of smooth muscle cells to myofibroblasts and fibroblasts. We hypothesize that AREG is a prevalent secretion factor in the development of fibrosis.

**Aim 1:** To use morphology to identify the plaque structure and environment

**Aim 2:** To use protein indexing technology (CODEX), spatial transcriptomic analyses (Visium), and targeted RNA visualization (RNAscope) to visualize immune cell role in various stages of plaque and the cellular interactions with AREG

**Aim 3:** To create a rainbow-mouse model and use PCSK9 gain of function to induce atherosclerotic development in a control and double AREG knockout mouse to then isolate the plaque, compare the morphology, and lineage trace plaque components

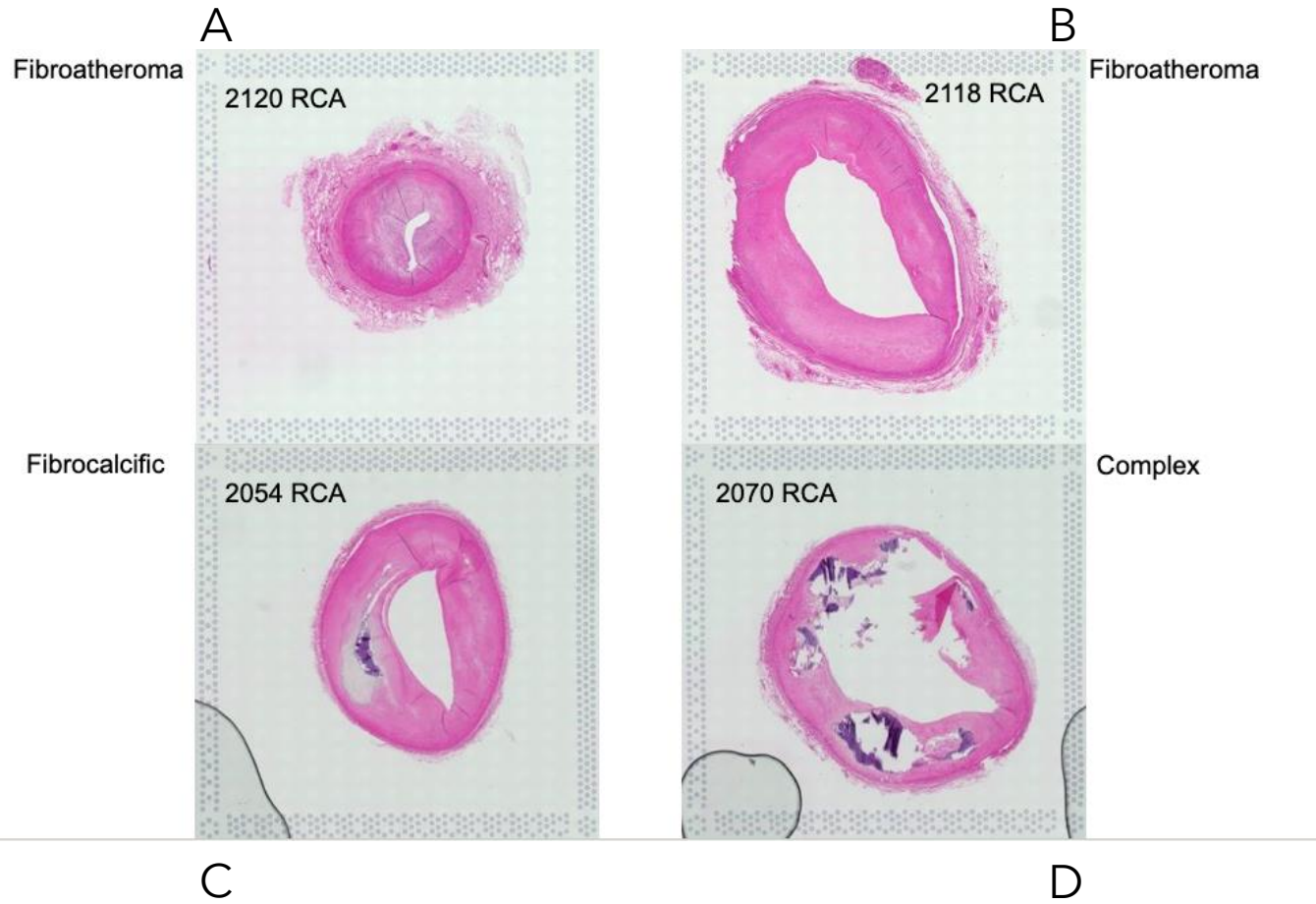
# METHODS

- Embedding:
  - Coronary arteries are received from recent heart explants and fixed in NBF or PFA for 24 hours
  - Arteries are then processed and placed in 70% EtOH or PBS (depending on embedding method)
  - Tissues are embedded and sectioned onto slides prior to histology
- Histology
  - Hematoxylin and Eosin: stains for the nuclei (H) and cytoplasm (E)
  - Movat: can visualize the components of connective tissue, including elastic fibers, collagen, proteoglycans, and fibrinoid material
  - Trichrome: can visualize connective tissue, collagen, and smooth muscle fibers
  - Dab: can visualize the location and distribution of specific antigens of interest (AREG antibody)

# METHODS

- CODEX:
  - Perform slide preparation and antigen retrieval prior to Enable Medicine's CODEX protein marking and imaging (Black *et al.* 2021 and Sanchez-Molina *et al.* 2022)
- Visium:
  - Spatial transcriptomic analysis technology that enables the profiling of gene expression in tissues with spatial resolution (Hudson & Sudmeier 2022)
  - Capture of mRNA from a tissue section on a microarray slide, followed by cDNA synthesis and sequencing
- RNAscope:
  - RNA visualization of target genes using RNAscope chromogenic assay (Zhang *et al.* 2022)
  - Hybridization of probes (positive, negative, target) and amplification of RNA signal
- Rainbow-mouse/PCSK9 gain of function:
  - Inject rainbow-mice control and rainbow-mouse AREG double knockout with PCSK9 vector via tail vein injection (Boone *et al.* 2019)
  - Sacrifice at specific week intervals and compare plaque development and lineage tracing

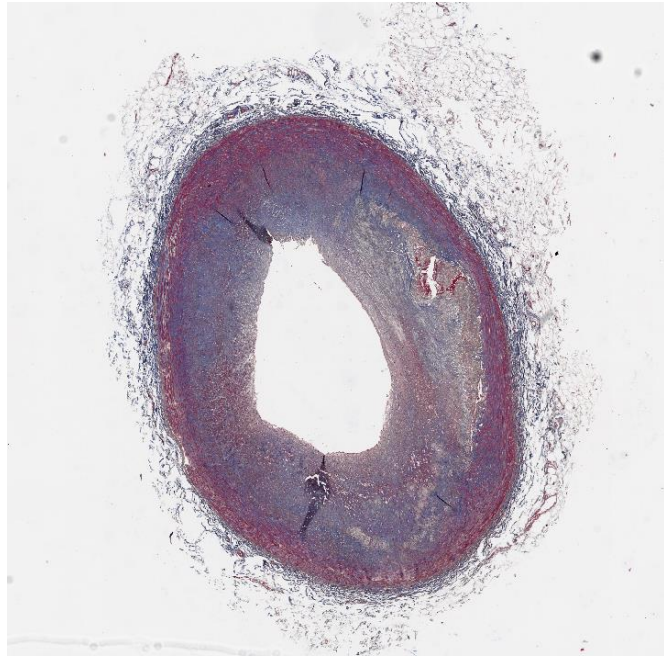
# RESULTS



H&E

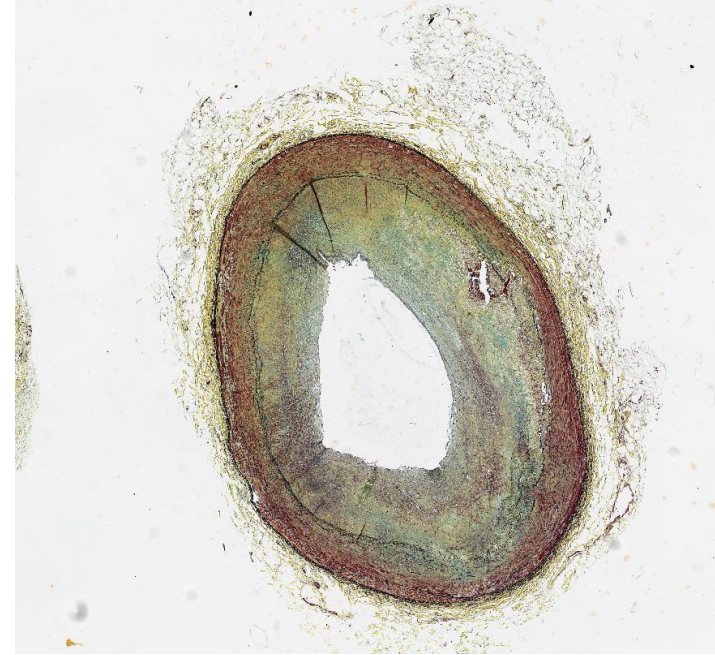
- Calcification is often seen as dark purple coloration within the tissues and causes increased fragility of artery integrity
- **A:** middle stage of plaque development
- **B:** middle stage of plaque development
- **C:** dark purple stain indicates a region of calcification within the intimal layer
- **D:** an example of the difficulties of processing calcified arteries as the calcium deposits make regions prone to tearing
- **E:** early stage of plaque development (intimal thickening, but closer to 'normal')

# RESULTS



## Masson's Trichrome

- **Red stain:** smooth muscle layer (media) composed of spindle-like smooth muscle cells
- **Blue stain:** fibrosis and collagen found in the intimal layer during late stage atherosclerosis
- **Pink stain:** cytoplasm
- **Brown stain:** nuclei

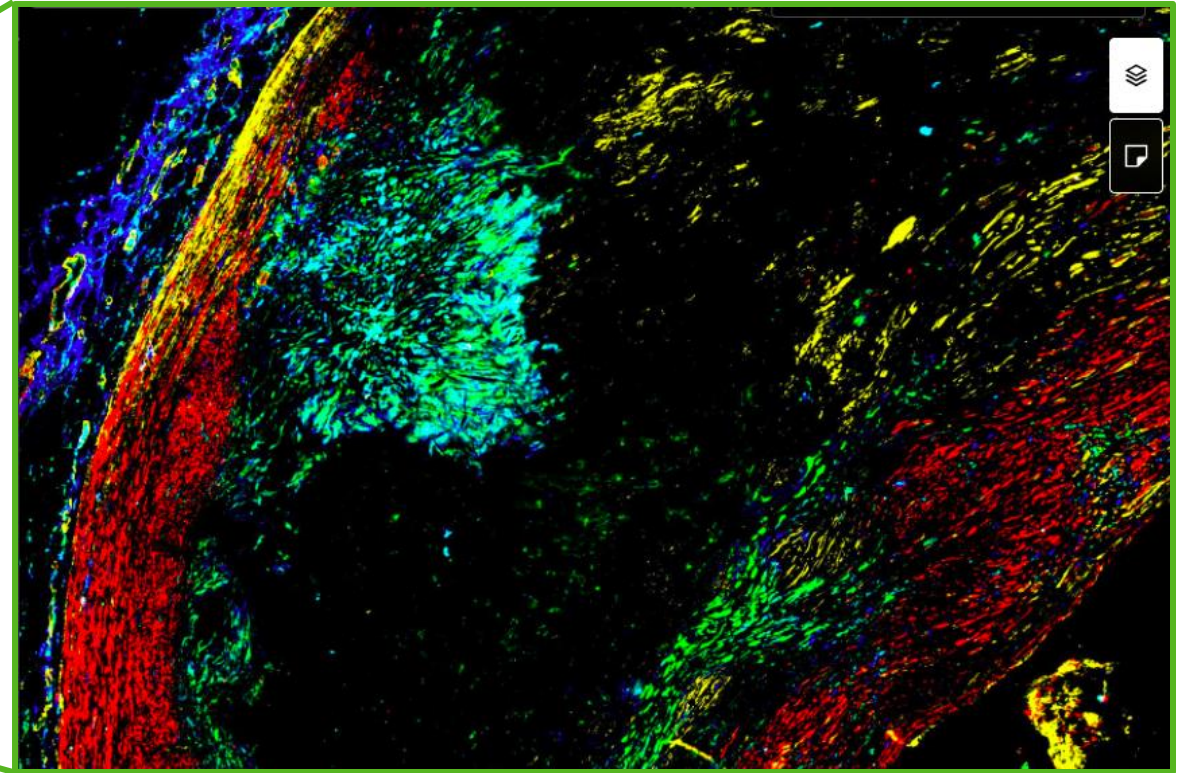
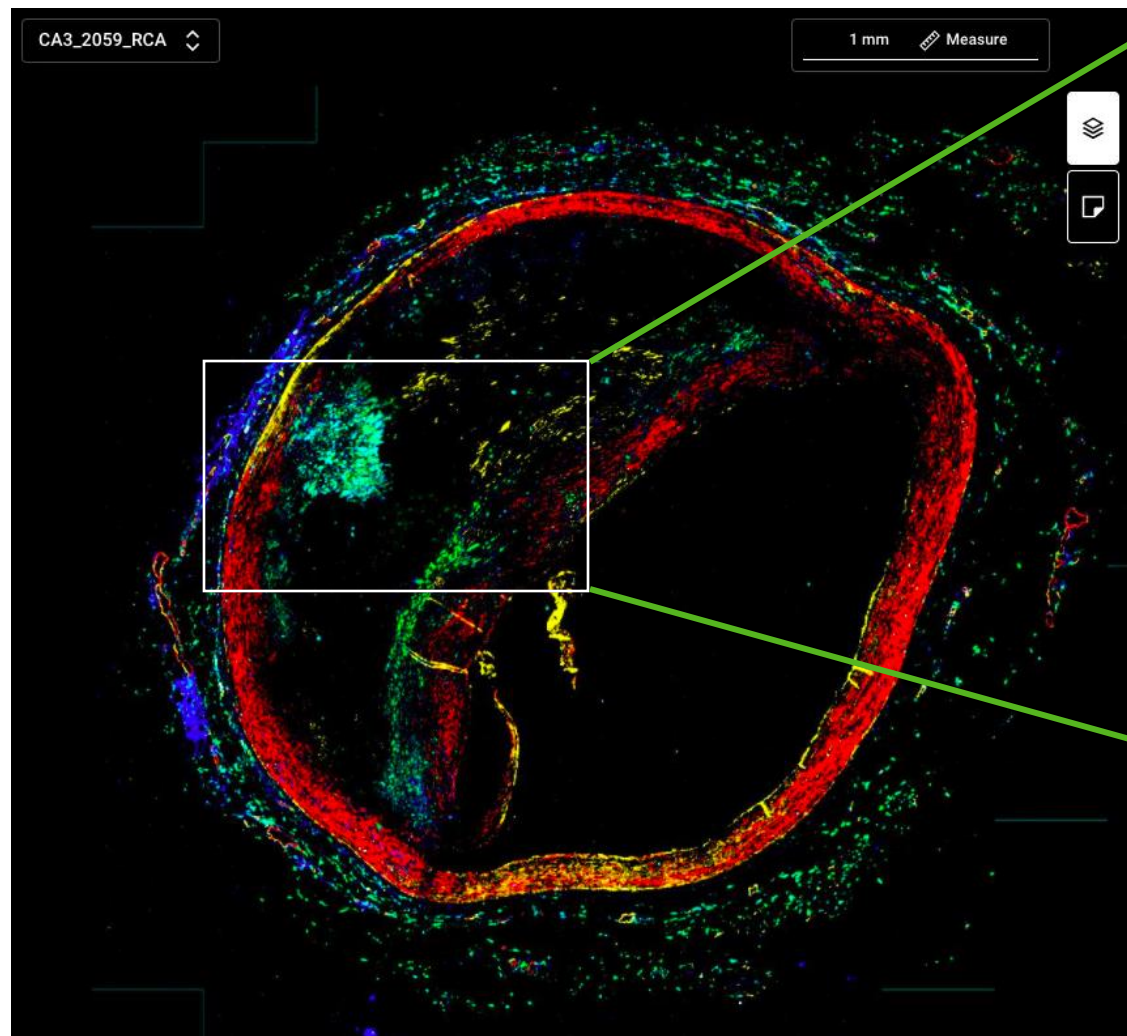


## Movat Pentachrome

- **Red stain:** smooth muscle layer (media) composed of spindle-like smooth muscle cells
- **Yellow stain:** fibrosis and collagen found in the intimal layer during late-stage atherosclerosis
- **Purple stain:** elastic fibers that define each layer of the artery
- **Black stain:** nuclei
- **Blue stain:** proteoglycans



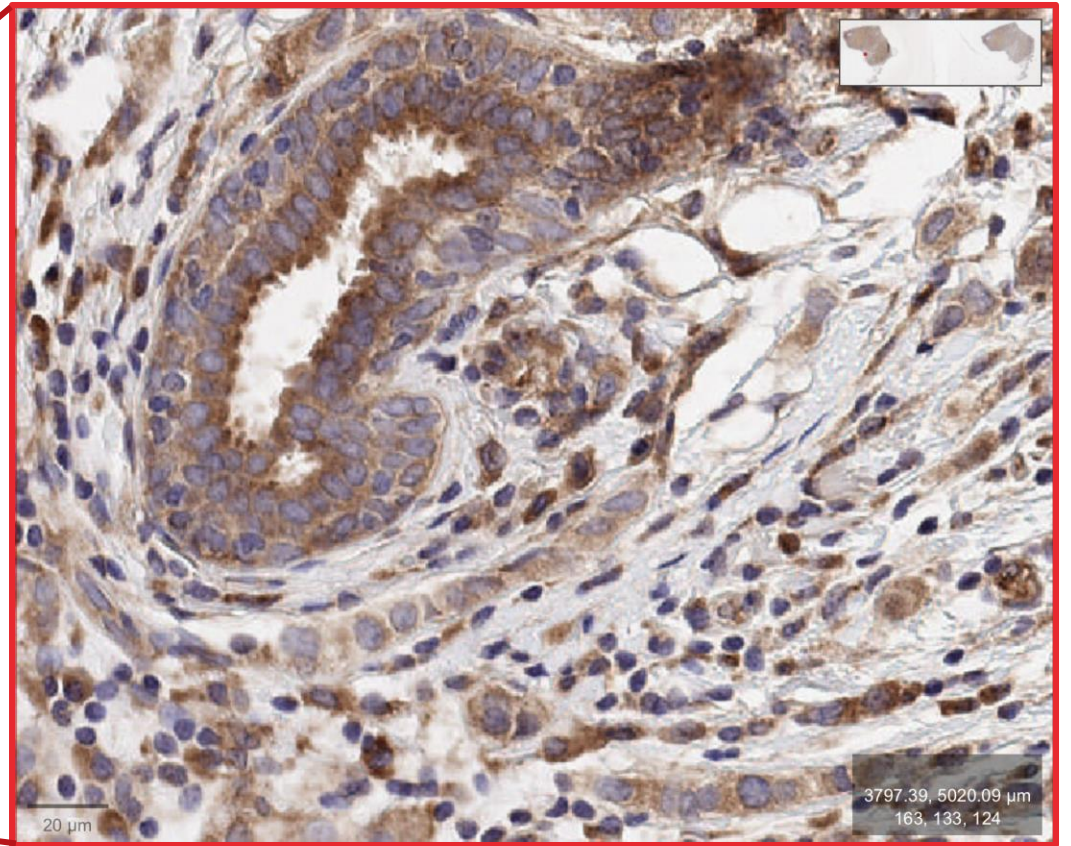
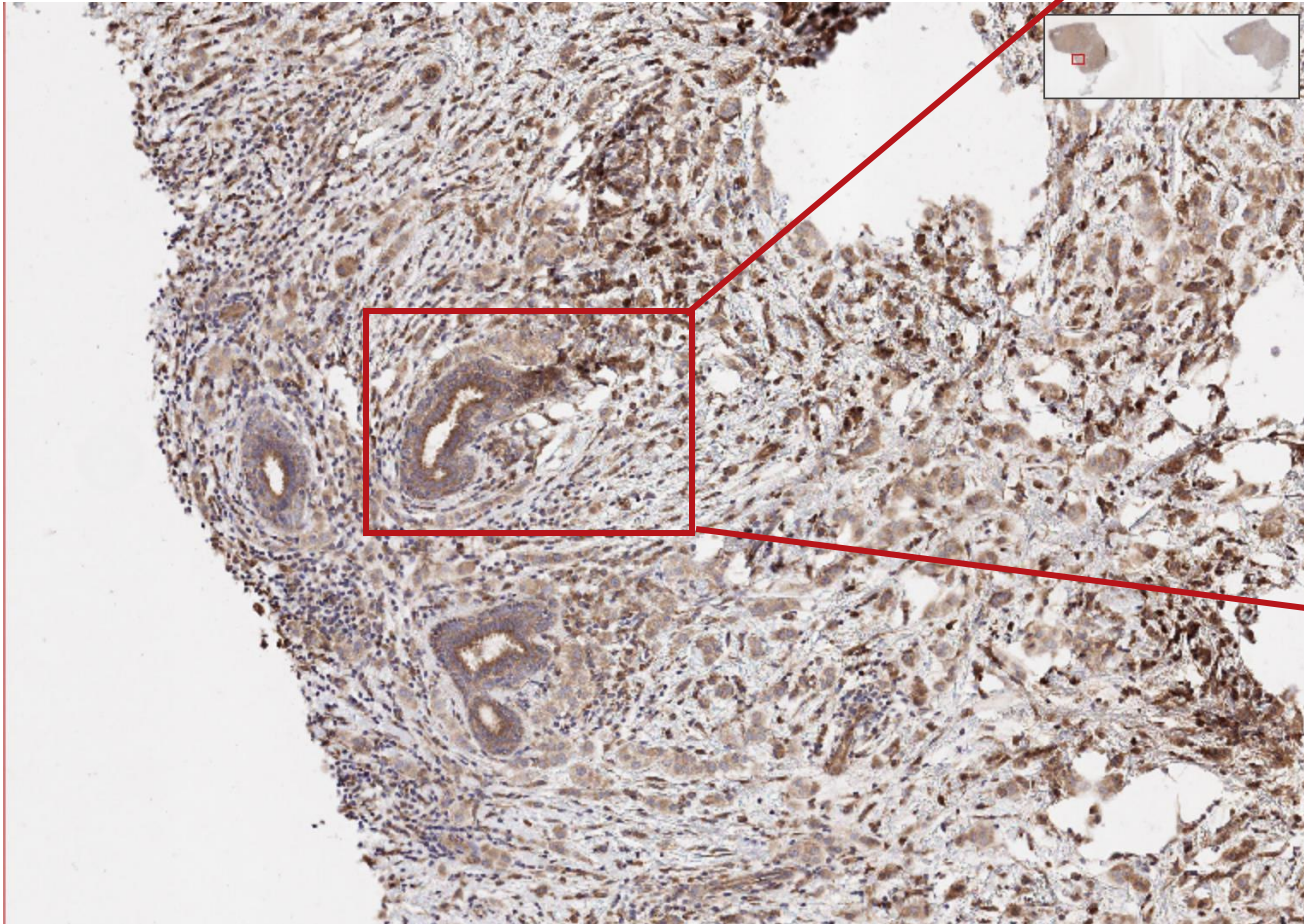
# RESULTS



## CODEX

- **CD68** (green): macrophages
- **CD8** (light blue): T cells
- **$\alpha$ SMA** (red): smooth muscle cells
- **CD45** (dark blue): leukocytes - general marker
- **CollagenIV** (yellow): collagen (indicative of fibrosis)

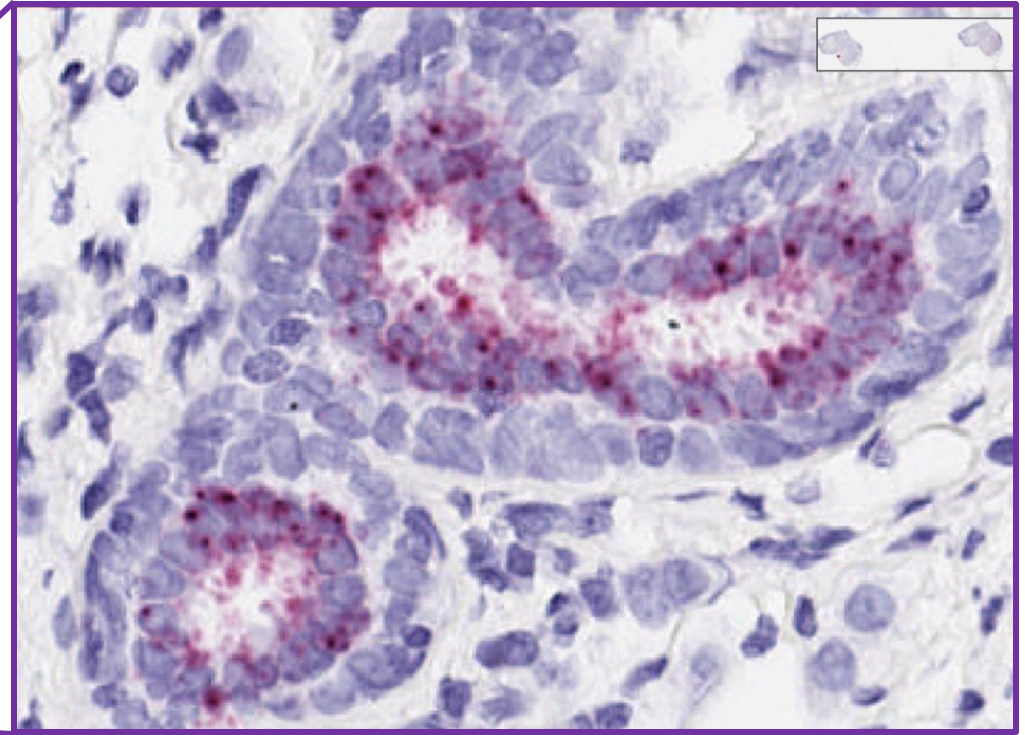
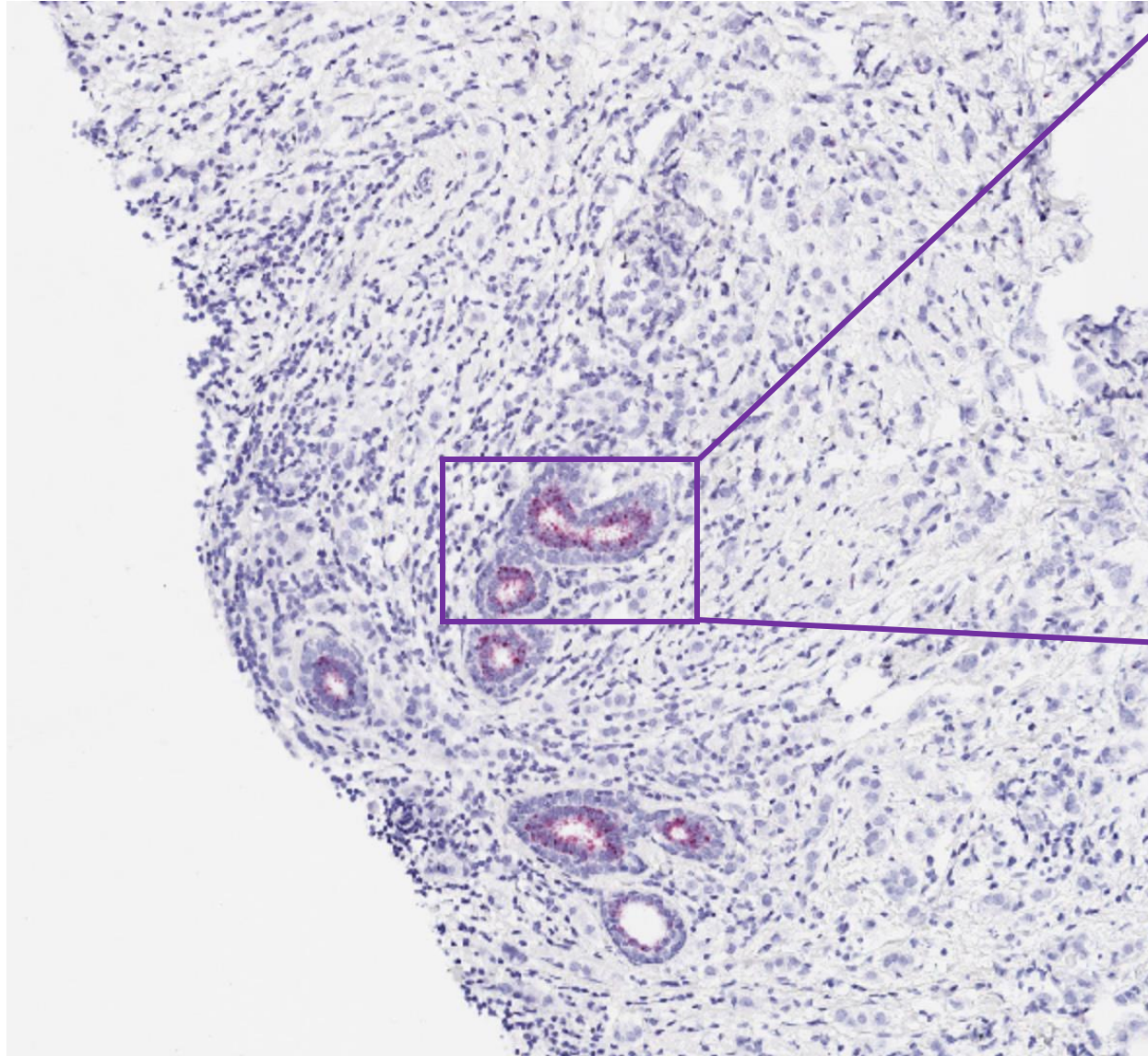
# RESULTS



## DAB - breast tissue

- **Brown:** the presence of AREG + background staining
- **Blue:** nuclei
- AREG is a cytoplasmic stain
- AREG seen in positive tissue contributing to fibrotic development, we should see in coronary arteries

# RESULTS



## RNAscope - breast tissue

- **Blue:** nuclei
- **Red dots:** indicative of target RNA presence within the tissue (nuclear)
- AREG present in breast tissue similar to Dab results
- Future experiments using diseased arteries

# CONCLUSIONS

- Continuing to identify evidence of smooth muscle cell interaction within the plaque
- Amphiregulin is found in many different fibrotic tissues (role in plaque development?)
- Aim to lineage trace rainbow-mice
- Learning more about the plaque niche and cellular components of the plaque itself
- Further experimentation to continue gaining plaque environment knowledge

# LITERATURE CITED

- Black, S., Phillips, D., Hickey, J. W., Kennedy-Darling, J., Venkatarahaman, V. G., Samusik, N., & Nolan, G. P. (2021). CODEX multiplexed tissue imaging with DNA-conjugated antibodies. *Nature protocols*, 16(8), 3802-3835.
- Boone, P. G., Rochelle, L. K., Ginzler, J. D., Lubkov, V., Roberts, W. L., Nicholls, P. J., & Snyder, J. C. (2019). A cancer rainbow mouse for visualizing the functional genomics of oncogenic clonal expansion. *Nature communications*, 10(1), 1-15.
- Chowdhury, R. R., D'Addabbo, J., Huang, X., Veizades, S., Sasagawa, K., Louis, D. M., & Nguyen, P. K. (2022). Human Coronary plaque T cells are clonal and cross-react to virus and self. *Circulation Research*, 130(10), 1510-1530.
- Hudson, W. H., & Sudmeier, L. J. (2022). Localization of T cell clonotypes using the Visium spatial transcriptomics platform. *STAR protocols*, 3(2), 101391.
- Zhang, L., Chen, D., Song, D., Liu, X., Zhang, Y., Xu, X., & Wang, X. (2022). Clinical and translational values of spatial transcriptomics. *Signal Transduction and Targeted Therapy*, 7(1), 111.
- Sanchez-Molina, P., Pratapa, A., Singh, J., Chiot, A., Bogachuk, A., Nikulina, N., & Ajami, B. (2022). Deep spatial phenotyping of microglial populations and pathological features in Alzheimer's disease brains using CODEX Multiplexed Imaging. *Alzheimer's & Dementia*, 18, e066433.