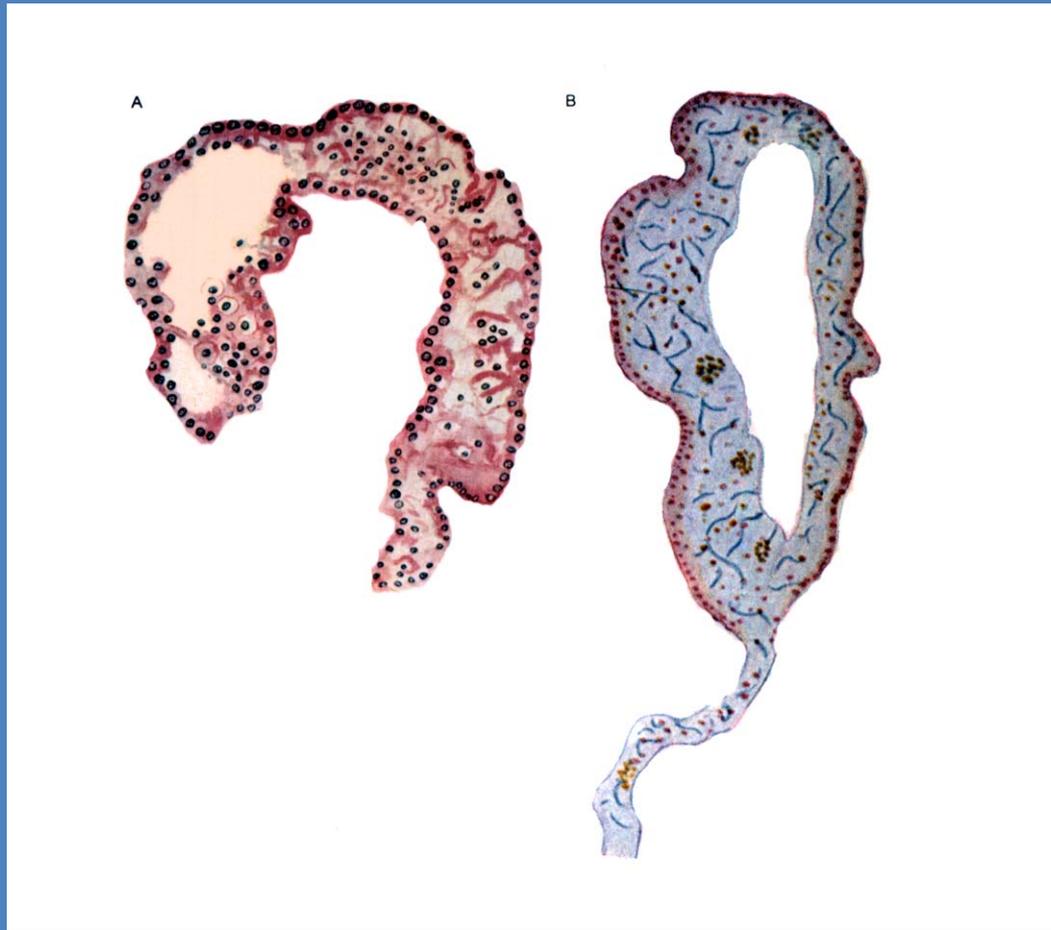


Intestinal regeneration in the echinoderm *Holothuria glaberrima*



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What are the ADVANTAGES and PITFALLS of the sea cucumber model system?



The first studies on sea cucumber gut regeneration were done in the early 1900s by Fausta Bertolini at the Stazione Zoologica Anton Dohrn in Naples

What are the **ADVANTAGES** and **PITFALLS** of the sea cucumber model system?

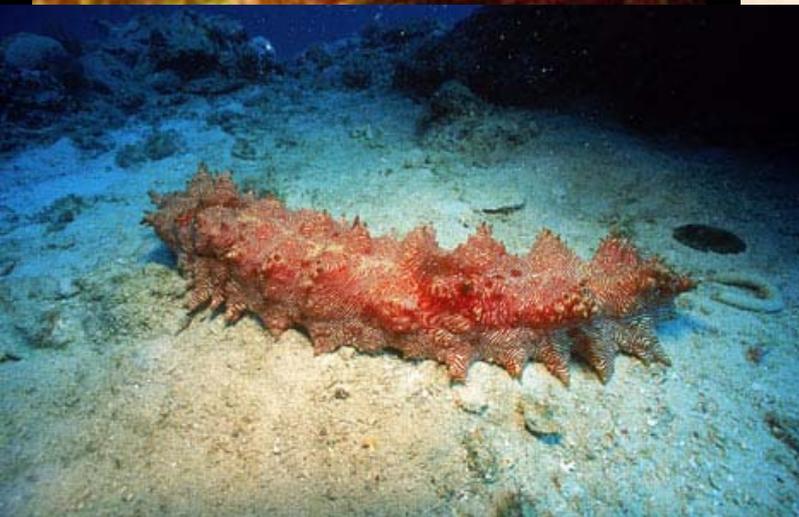
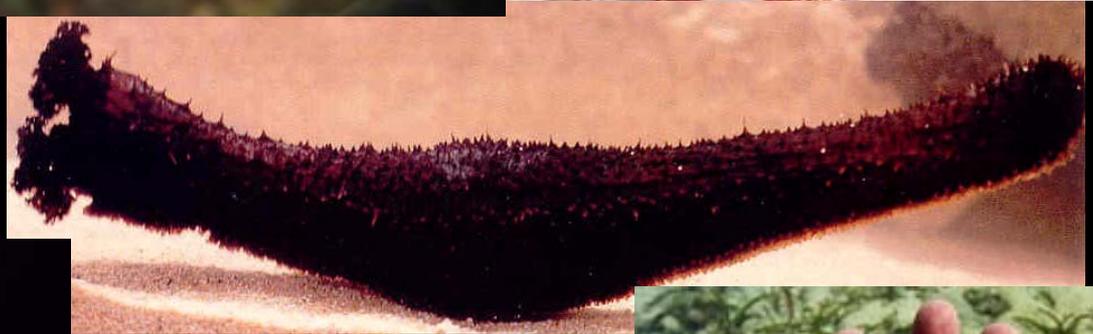


What are the **ADVANTAGES** and **CHALLENGES** of the sea cucumber model system?



Funded by- NSF-IBN, NIH- NIGMS and NINDS, Whitehall Foundation and the University of Puerto Rico

Advantage #1- Hundreds of species available for studies





Experimental Model

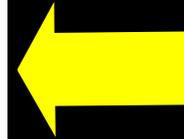
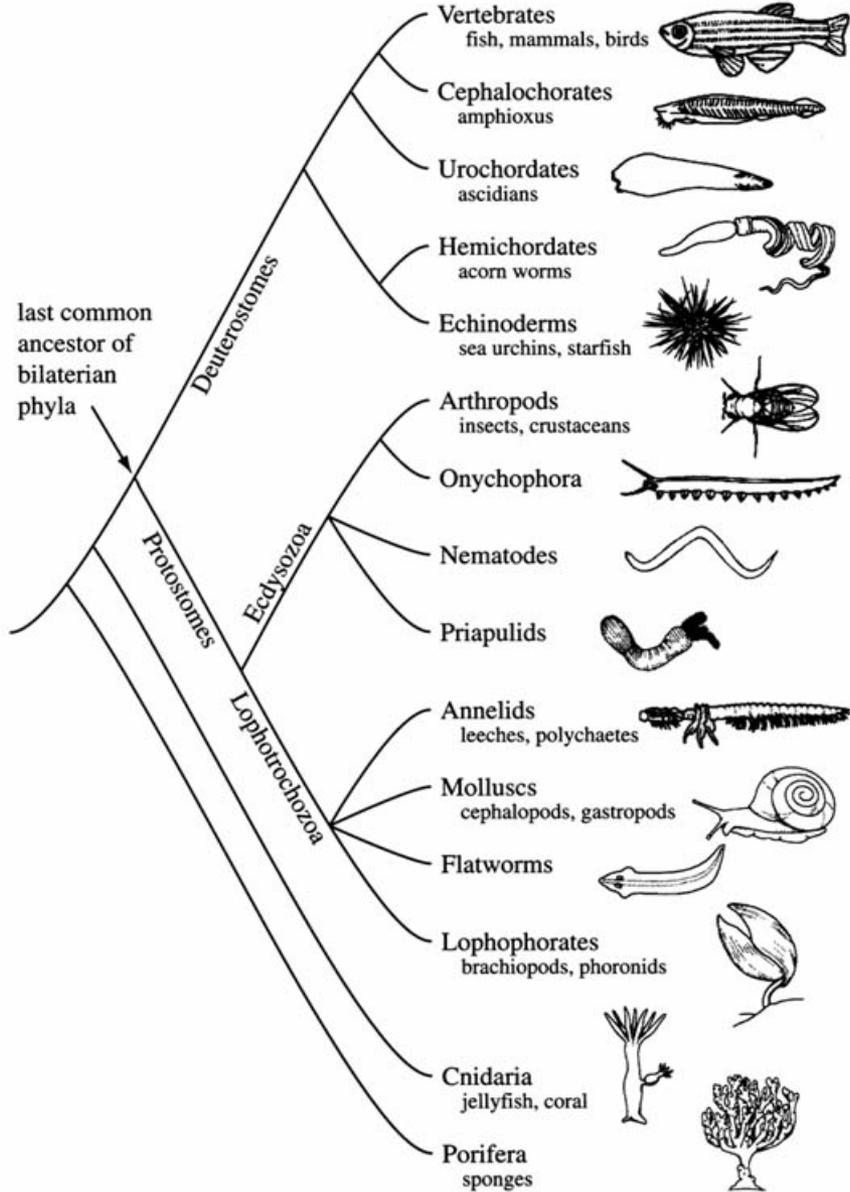
the sea cucumber, *Holothuria glaberrima*



Athyonidium chilensis

QuickTime™ and a
TIFF (Uncompressed) decompressor
are needed to see this picture.

Advantage #2- Key phylogenetic position



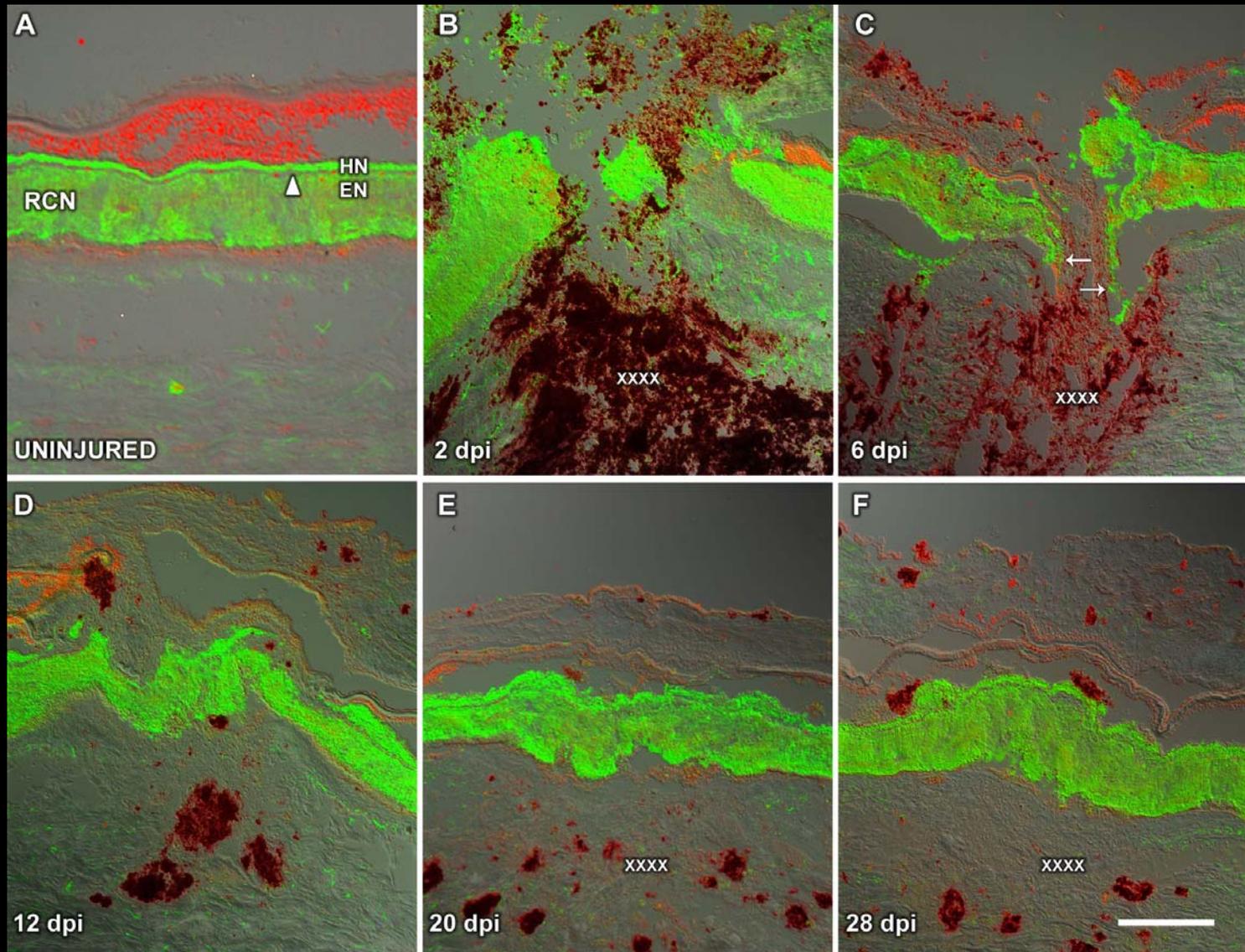
Echinoderms

Sea cucumbers are deuterostomes

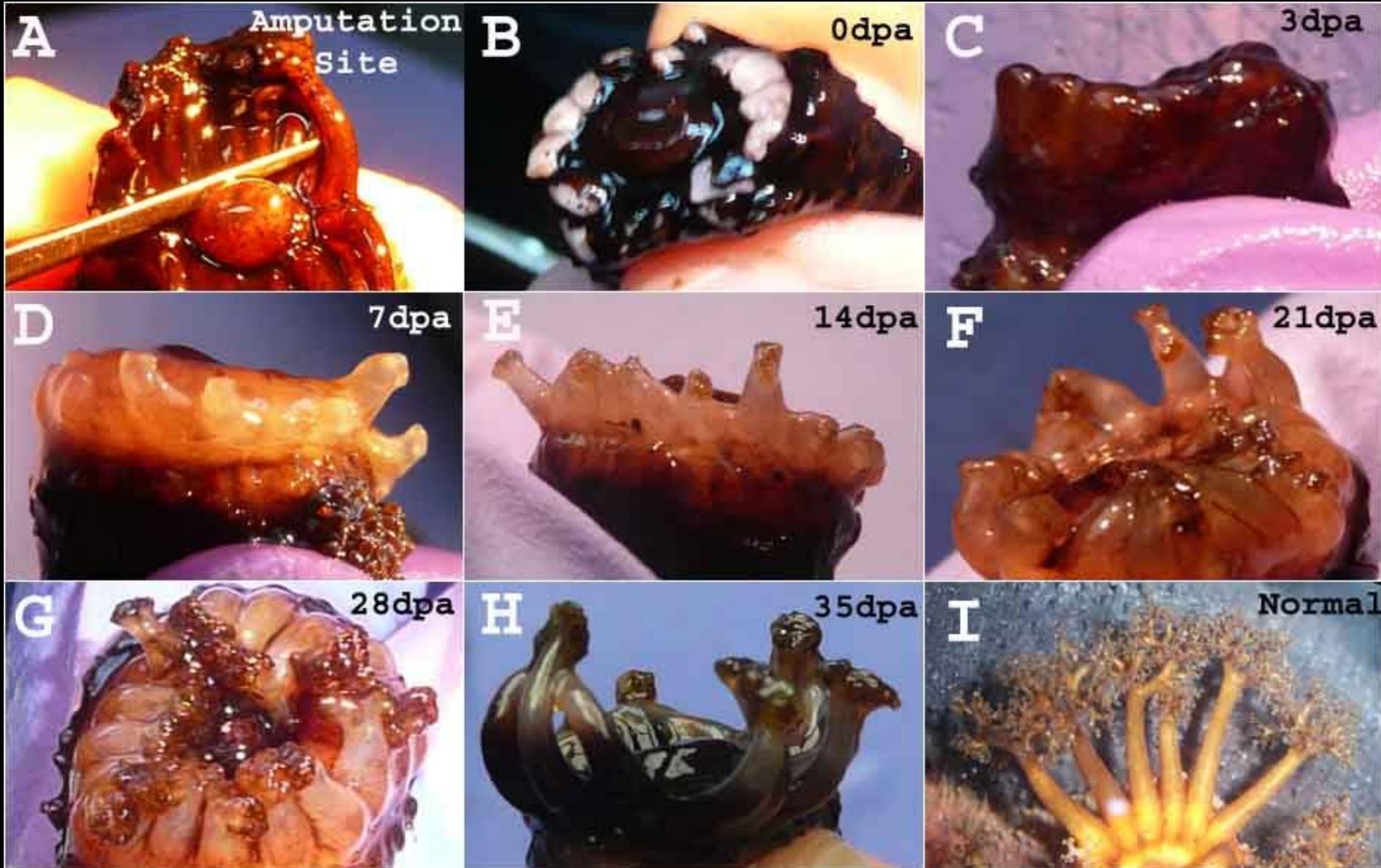
Advantage #3 - Extraordinary regenerative properties



Nerve fiber regeneration



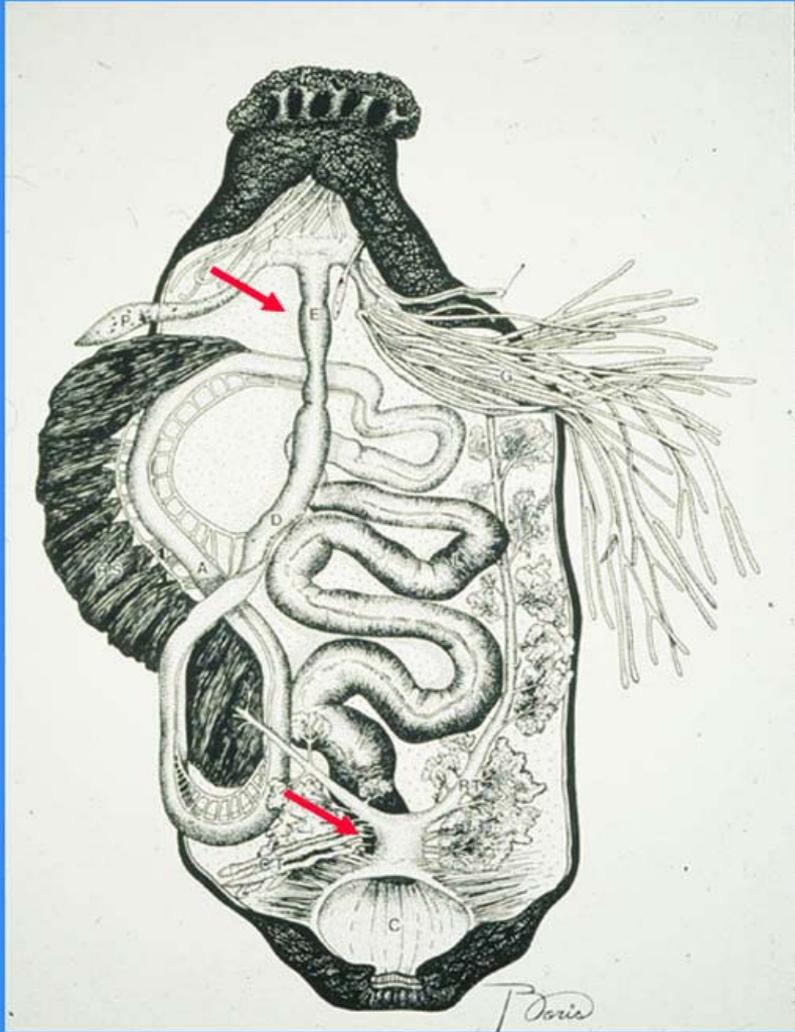
Tentacle regeneration



Advantage #4 - Evisceration is easily induced in lab

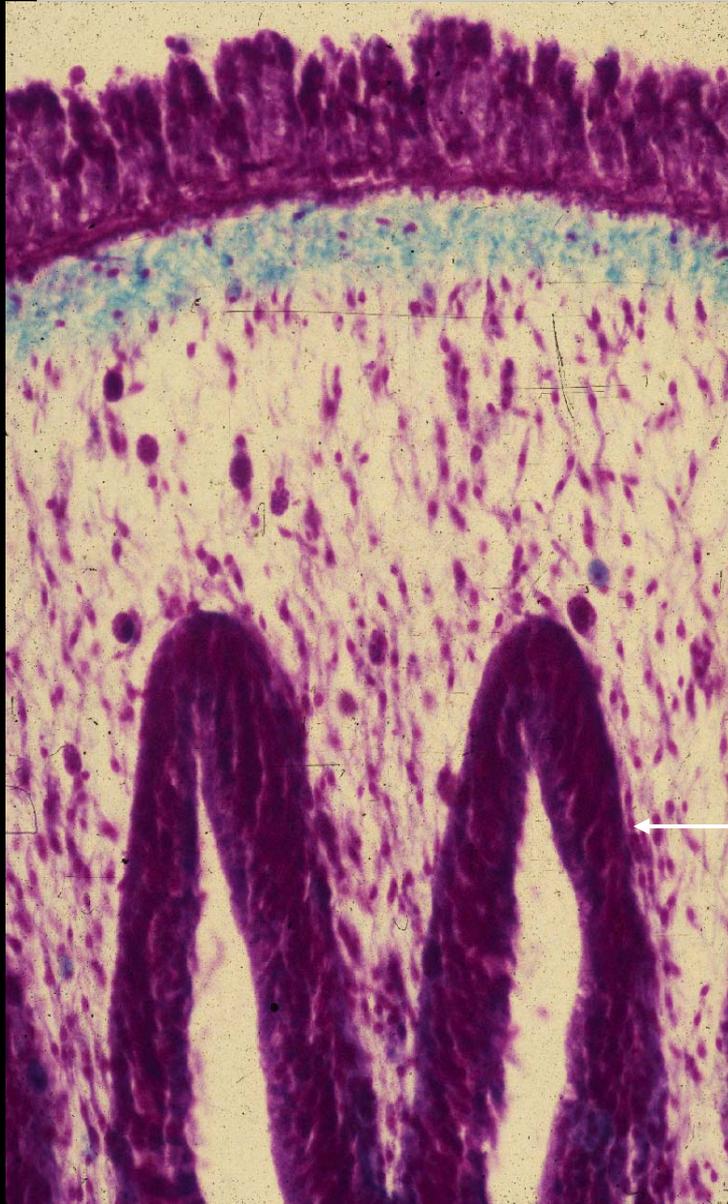


Advantage #5 - Evisceration follows a fixed pattern, reducing variability due to surgical manipulations

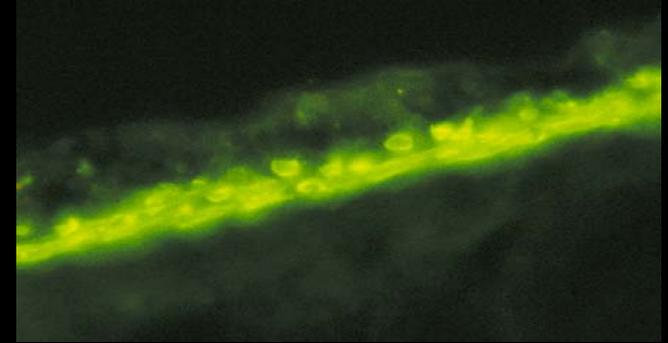


Evisceration eliminates most of the organs of the sea cucumber. In *H. glaberrima* only the left respiratory tree remains.

Advantage #6 - The digestive tract is well conserved among animal groups, particularly in deuterostomes



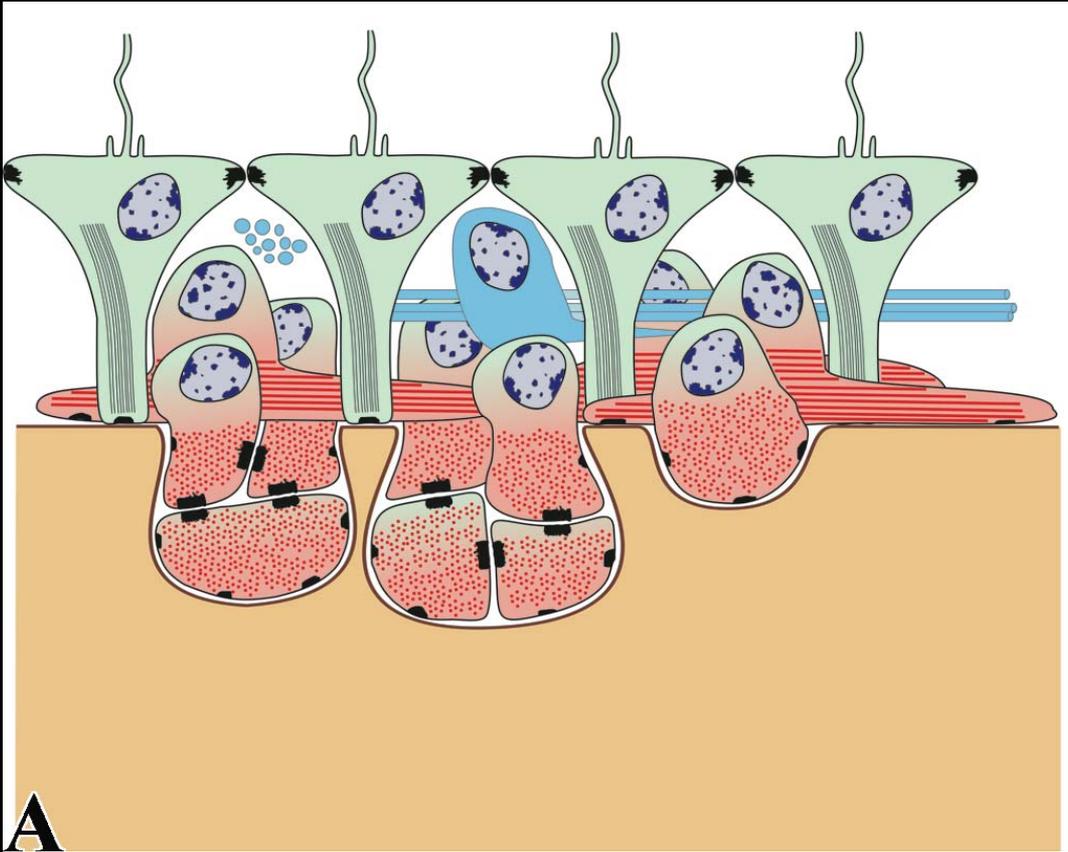
Mesothelium (includes coelomic epithelium or serosa and muscle layer)



Connective tissue layer (submucosa)

Luminal epithelium (mucosa)

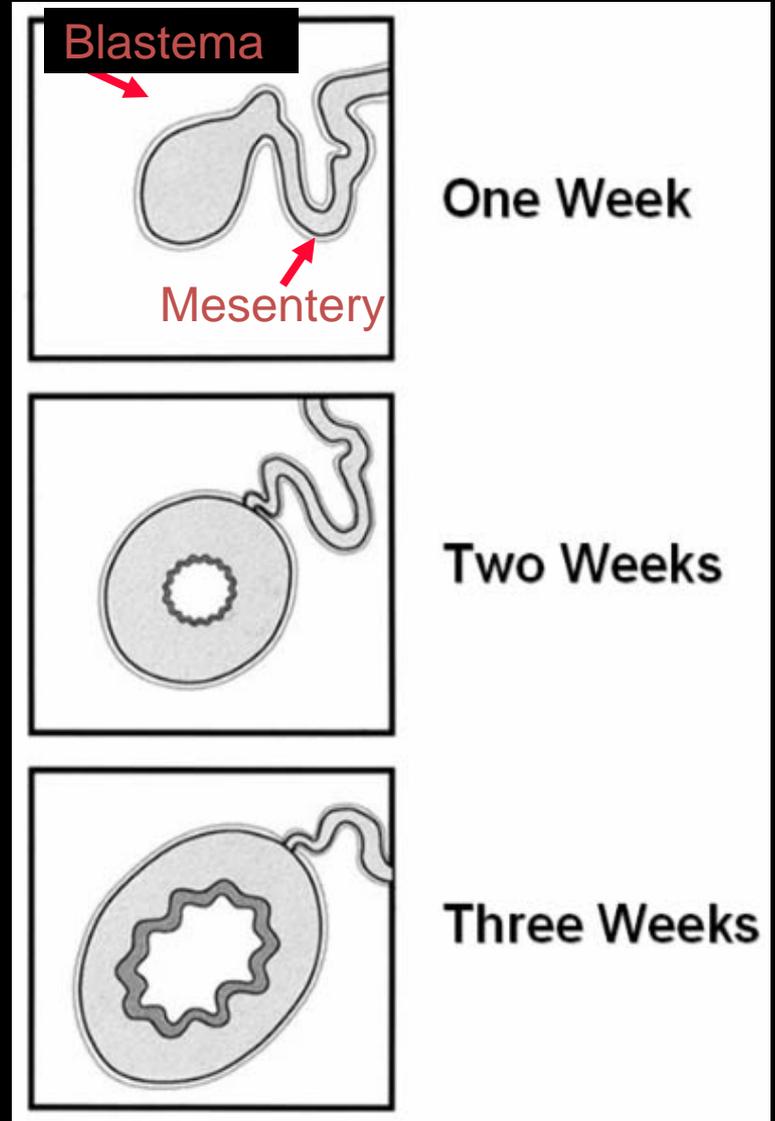
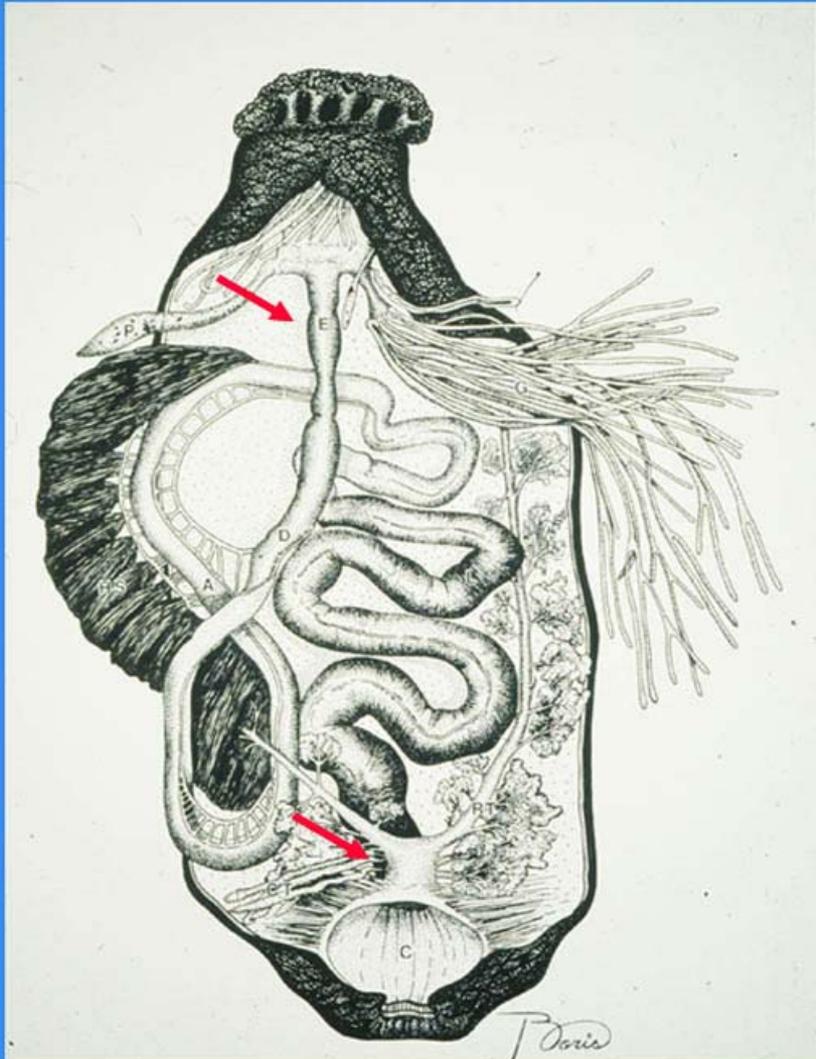
Echinoderm mesothelium



Peritoneocytes

Myocytes

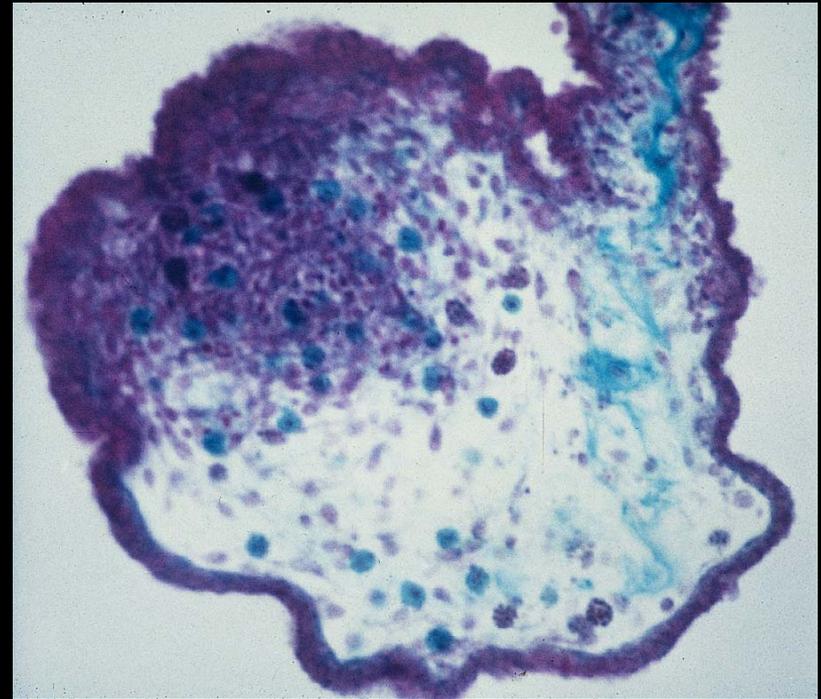
Advantage #7 - Regeneration of a functional organ occurs within a month



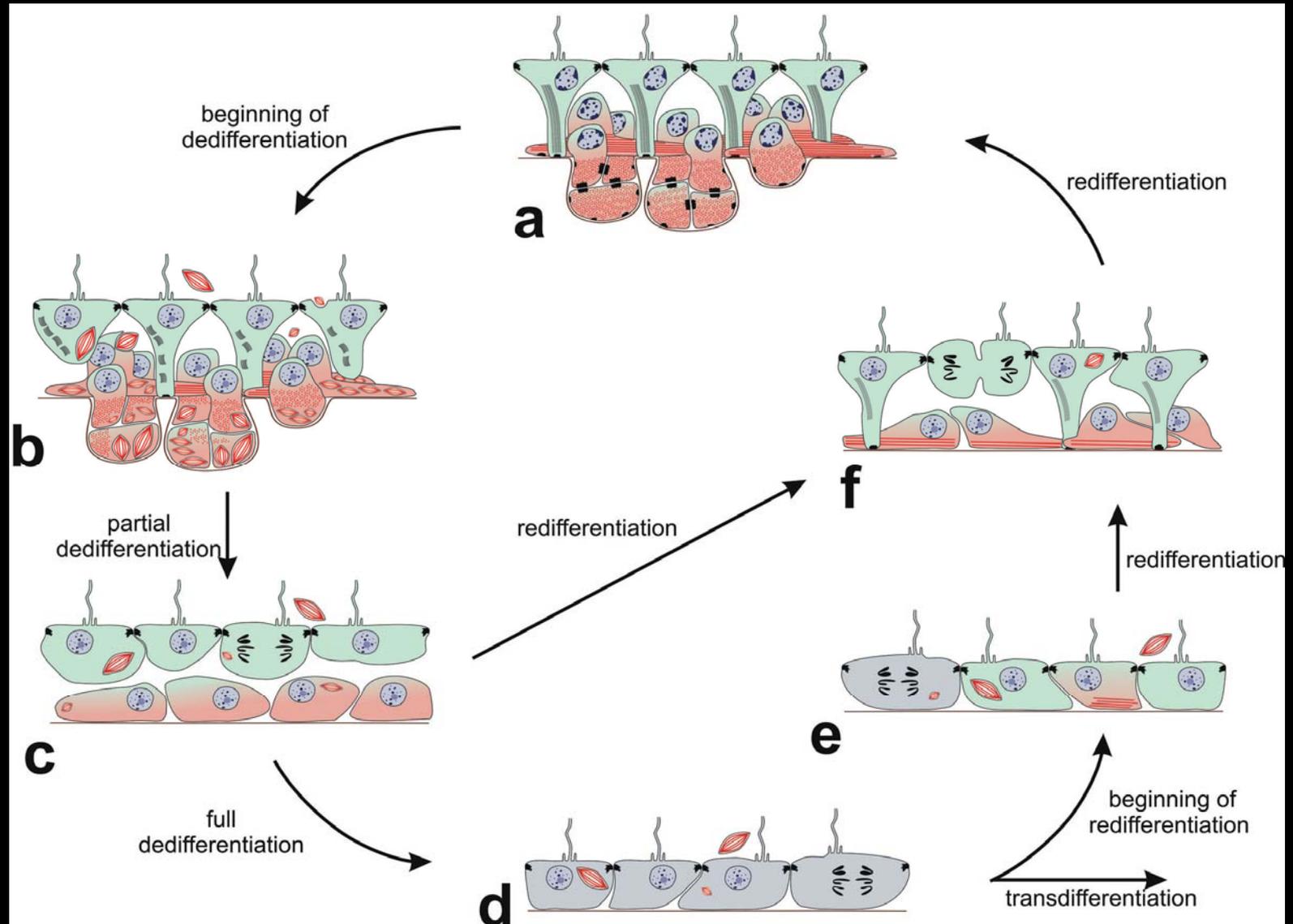
Advantage #8 - Multiple cellular events can be studied

Cellular events associated with the formation of a regeneration blastema include:

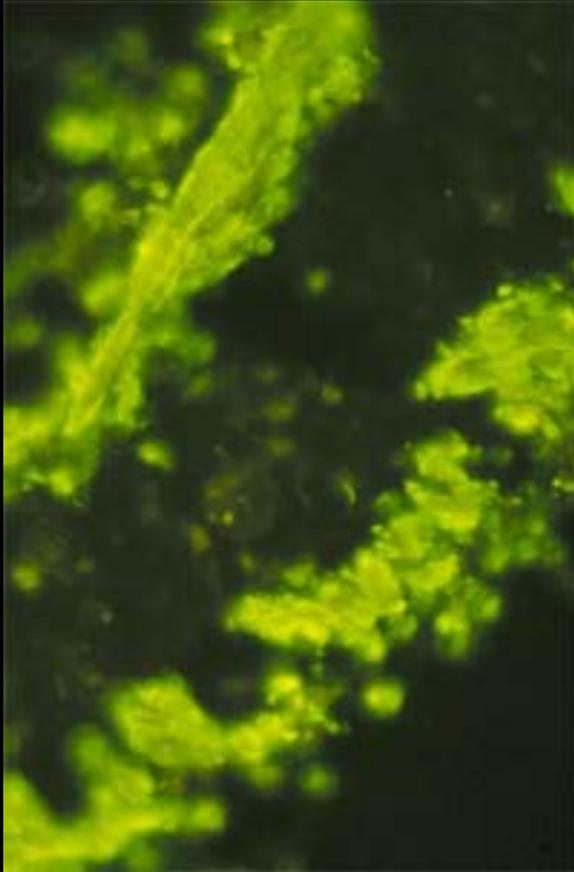
1. Cell dedifferentiation
2. Cell proliferation
3. Apoptosis
4. Epithelial to mesenchymal transition.
5. ECM remodeling
6. Cell migration
7. Cell differentiation
8. Cell-cell interactions to form a new organ



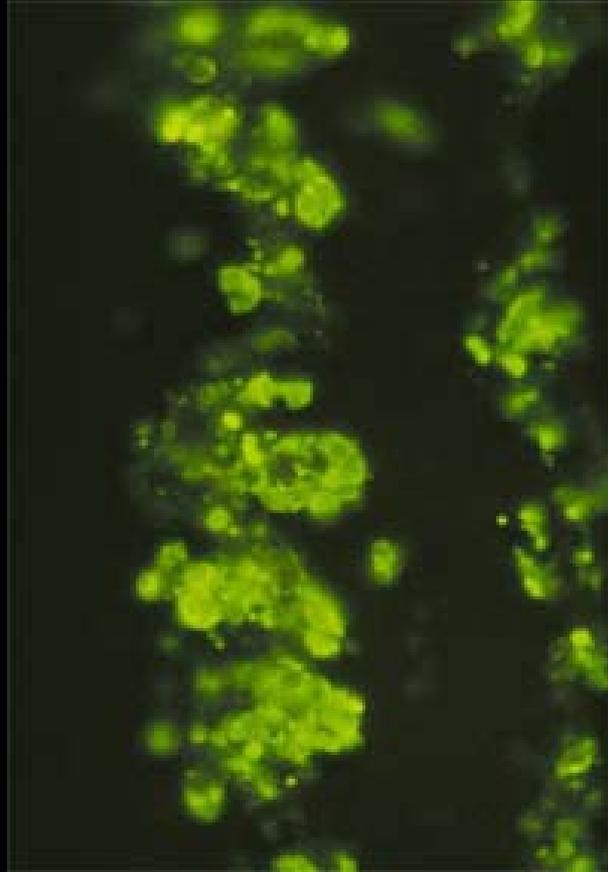
Cell Dedifferentiation



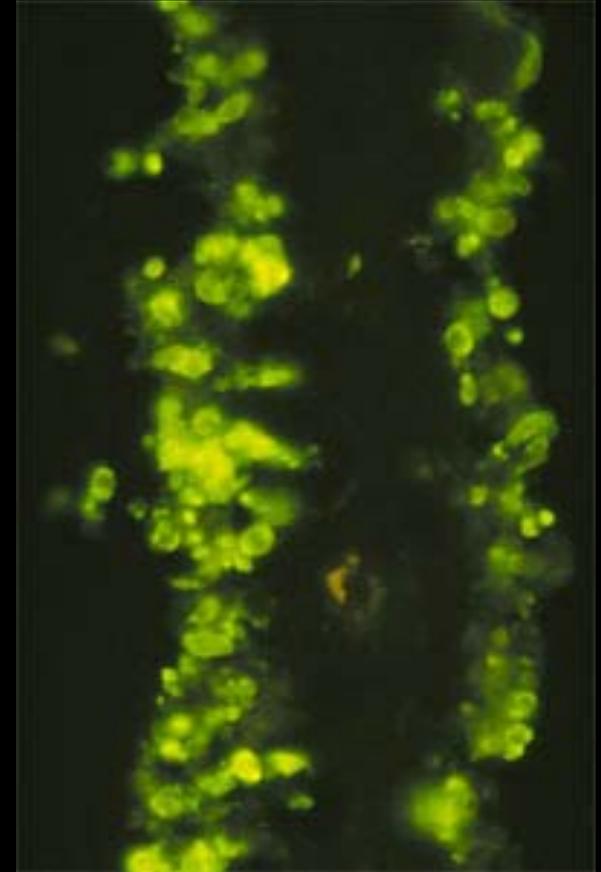
A gradient of muscle dedifferentiation can be found in the mesentery during regeneration



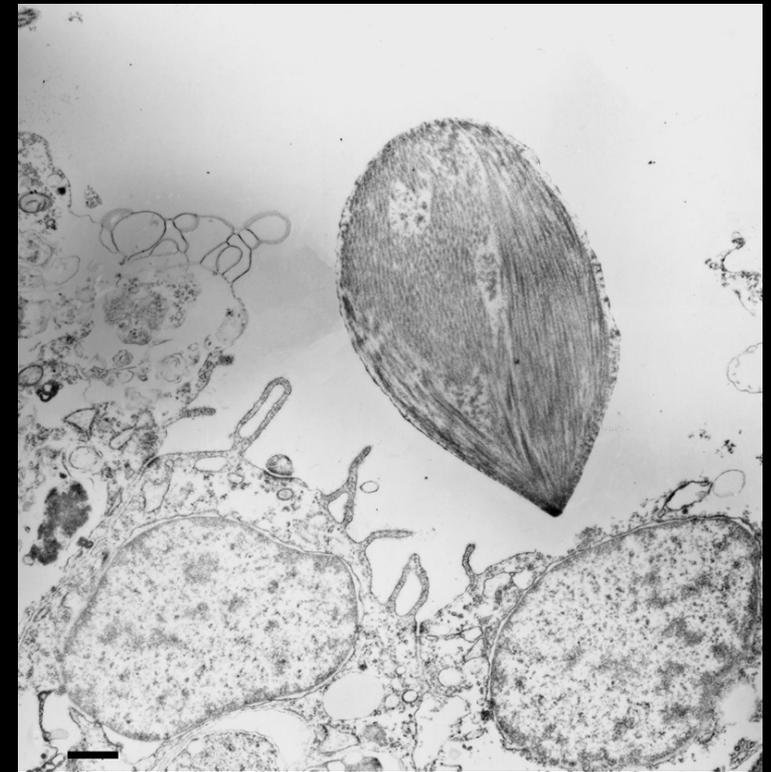
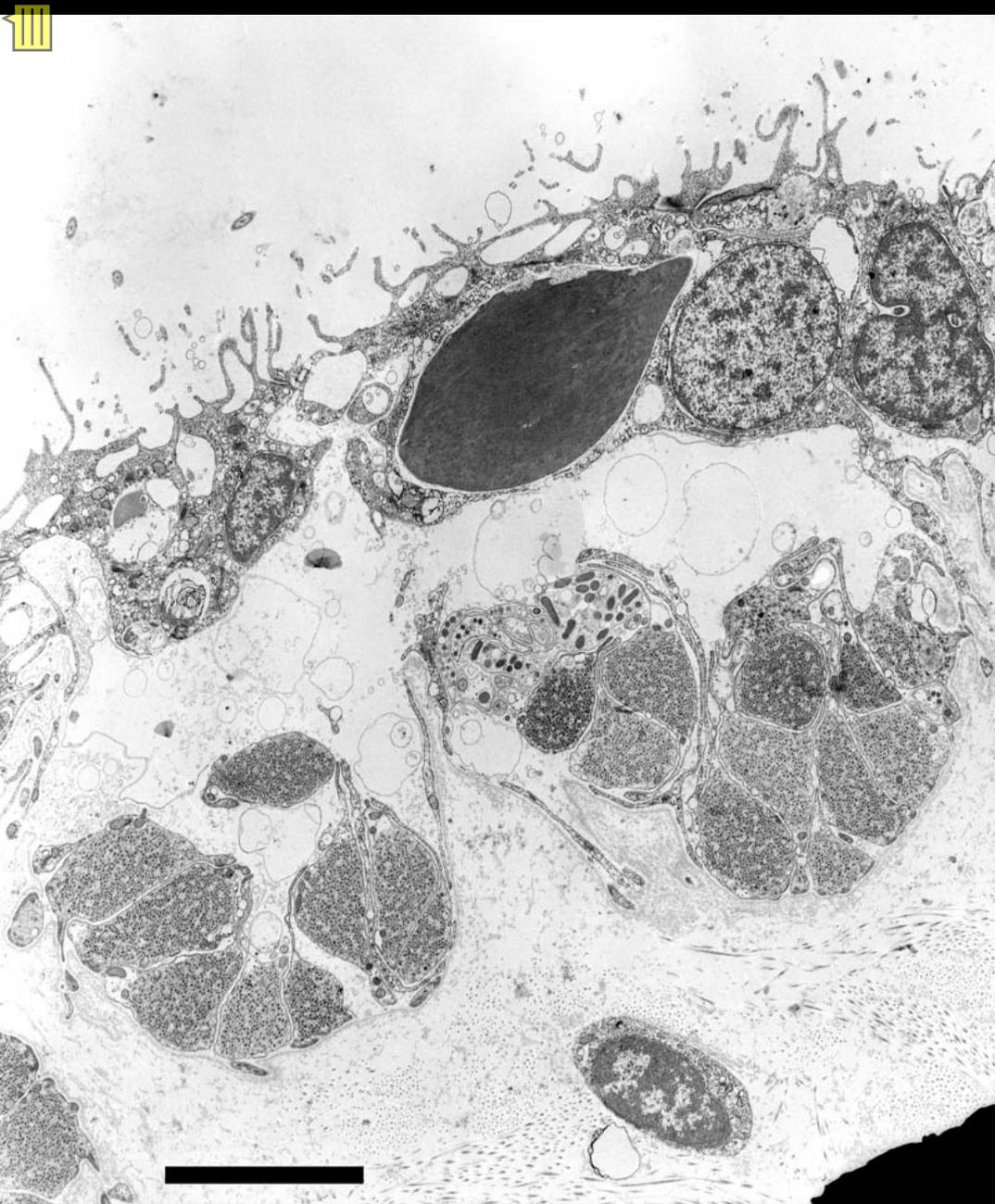
Near body wall



Medial



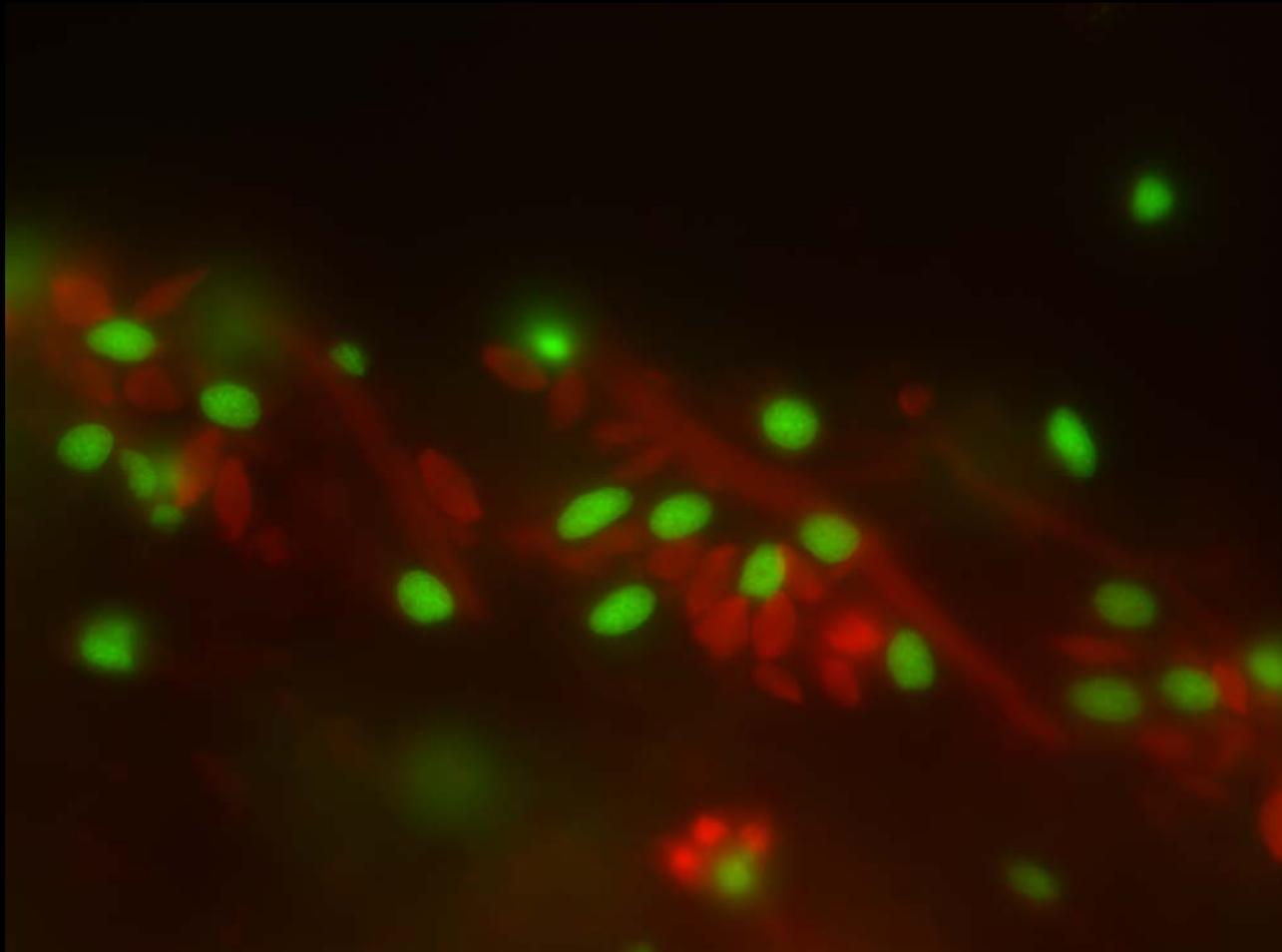
Near blastema-like structure



Muscle cell differentiation involves the formation of spindle like structures

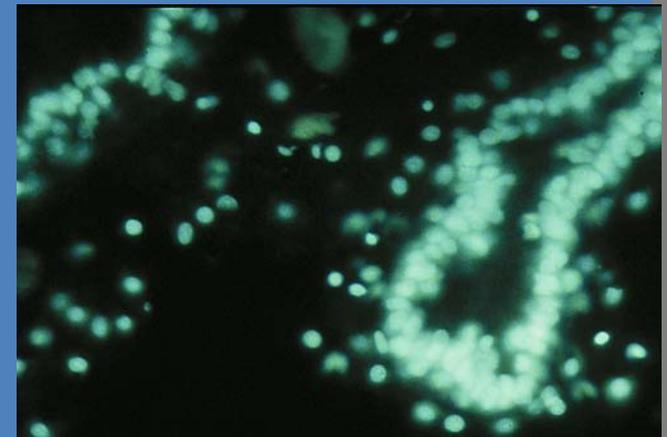
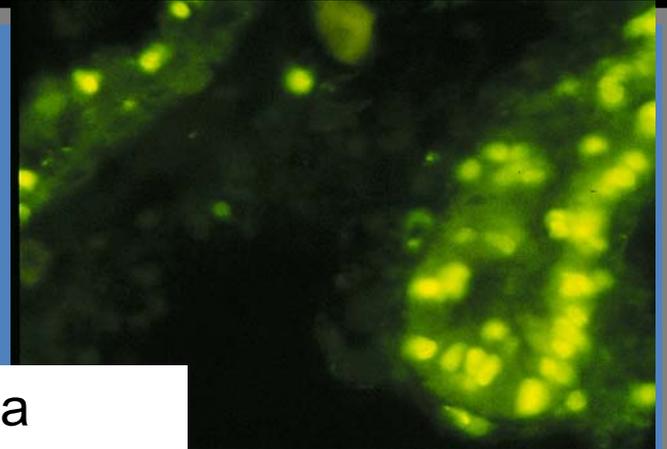
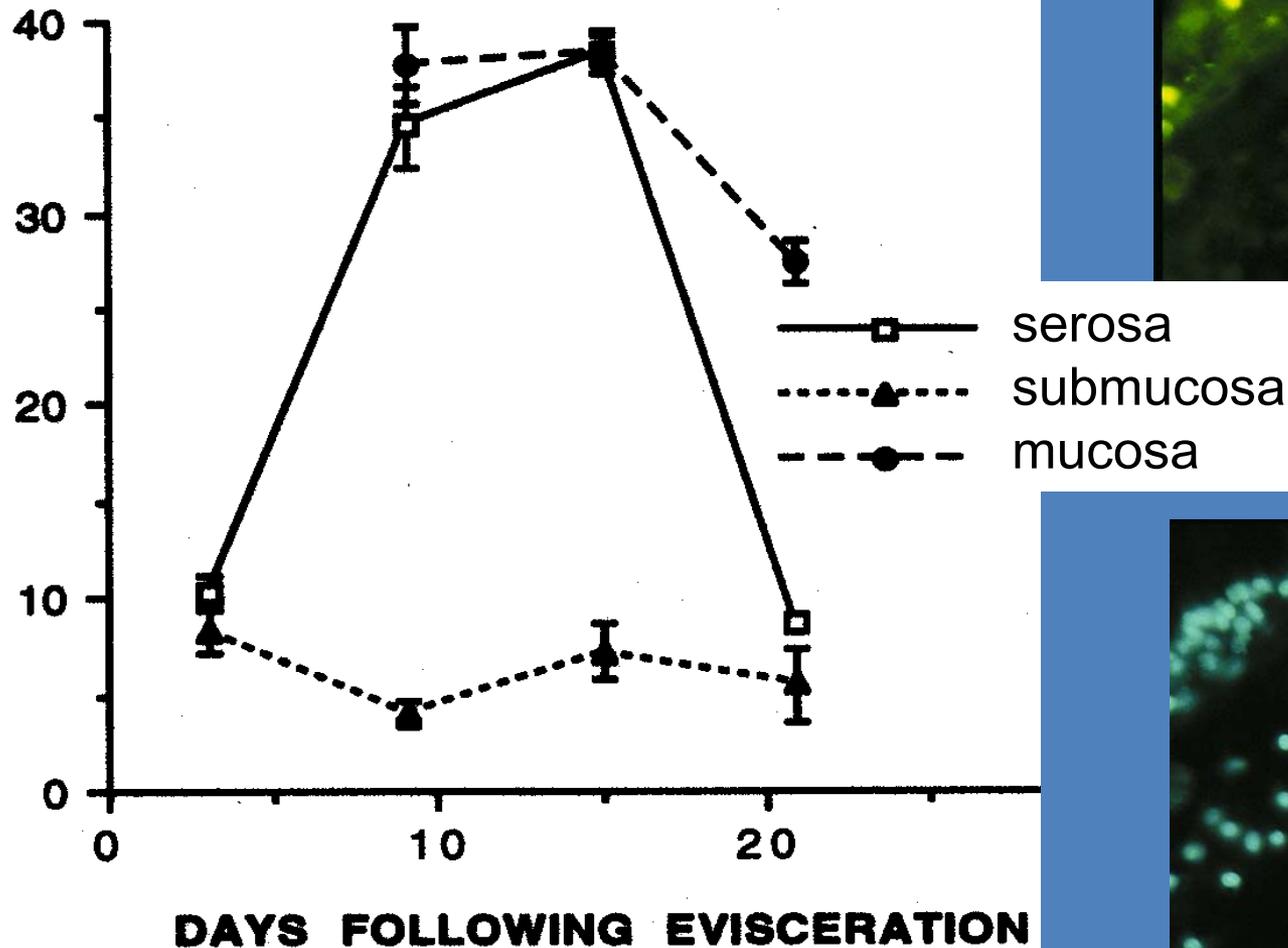


SLSs (red) are formed as cells dedifferentiate. These SLSs are not associated with cell nuclei (green)

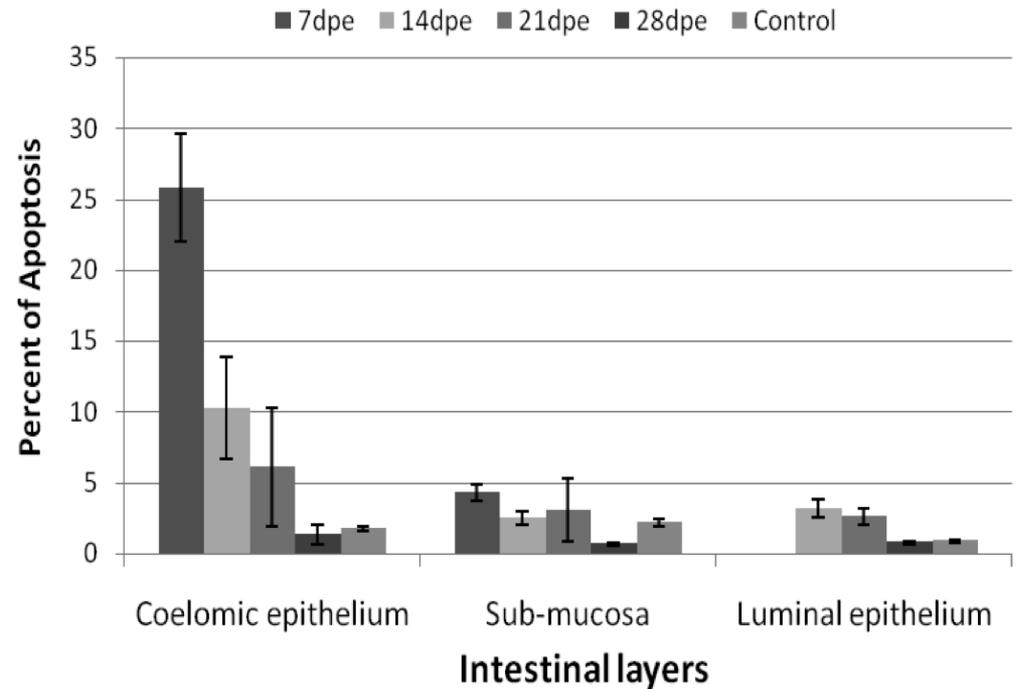
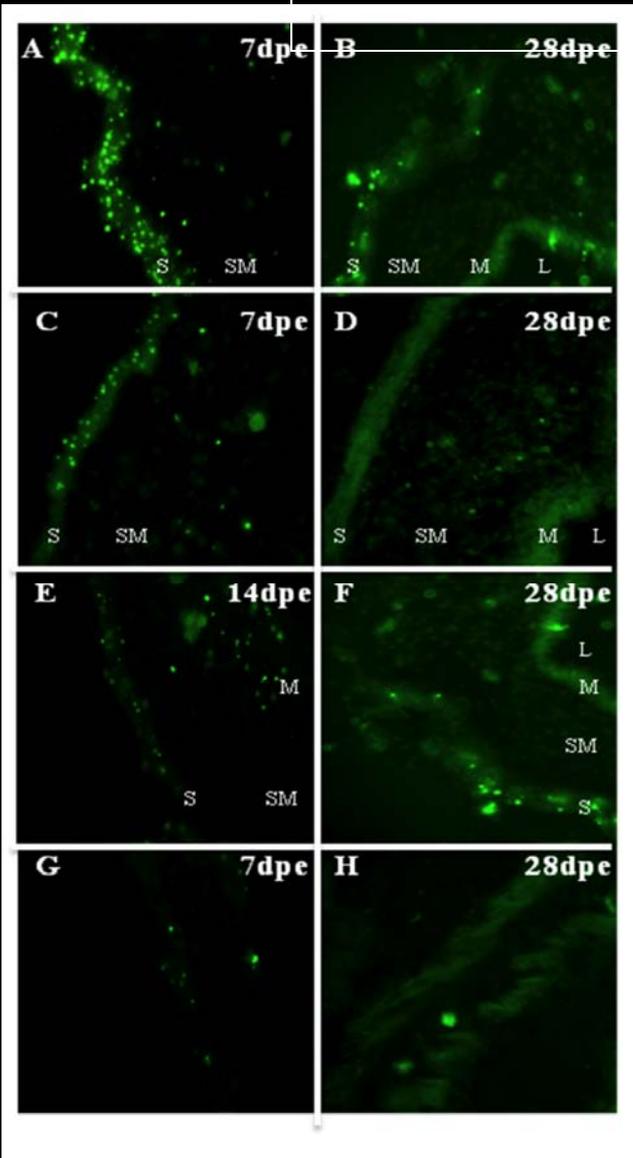


Cell Proliferation

Percentage cell division in specimens sacrificed 24 hrs following BrdU injection



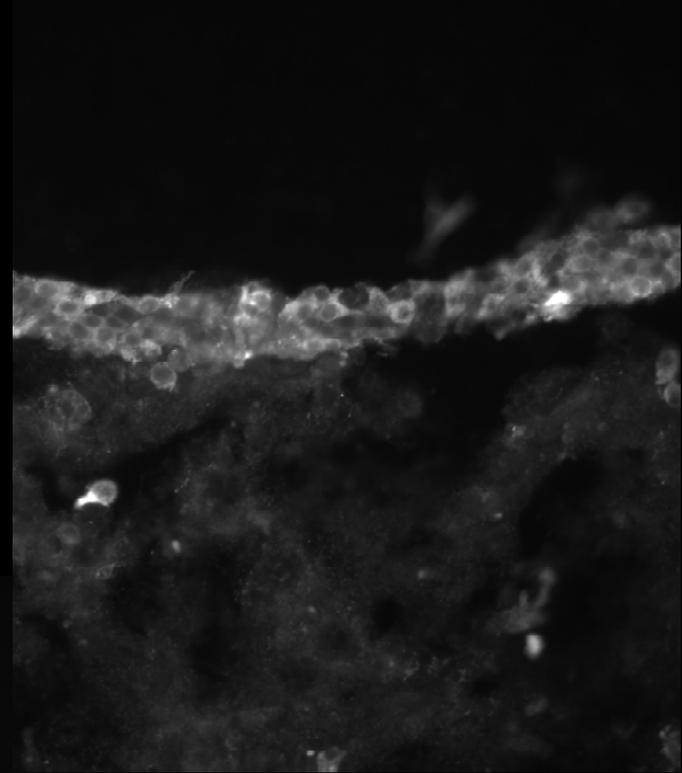
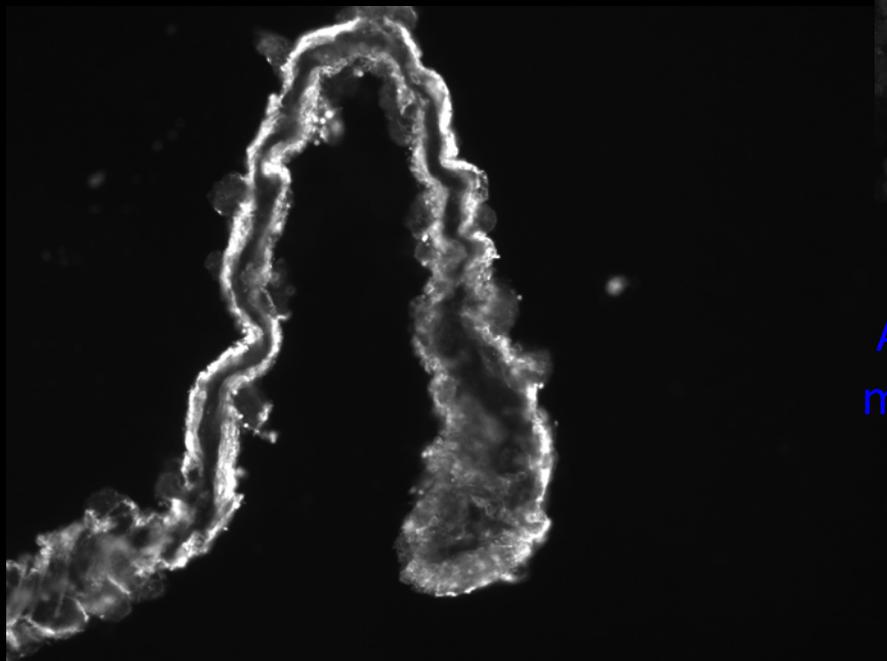
Apoptosis



TUNEL assays show large number of apoptotic cells in the regenerating epithelia.



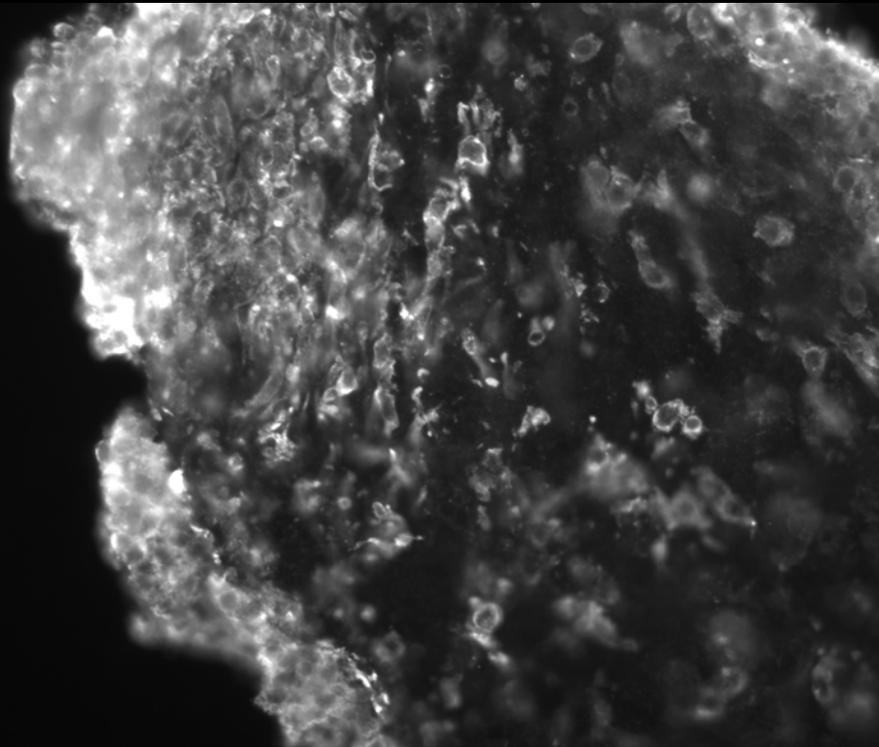
Epithelial to mesenchymal transition



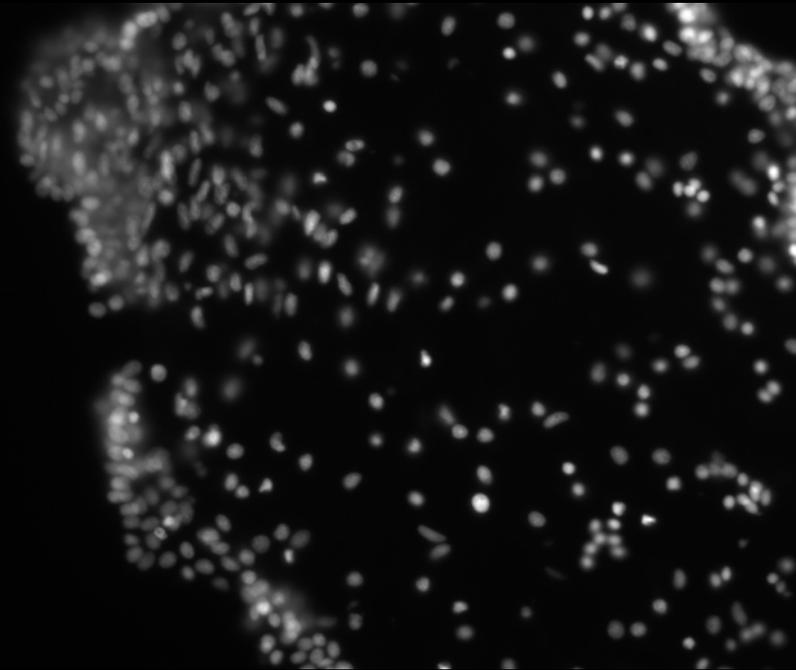
Antibody MES-1 recognizes the mesothelium and epithelial cells of the blastema.



Epithelial cells at the tip of the regenerating mesentery ingress to form the underlying mesenchyme

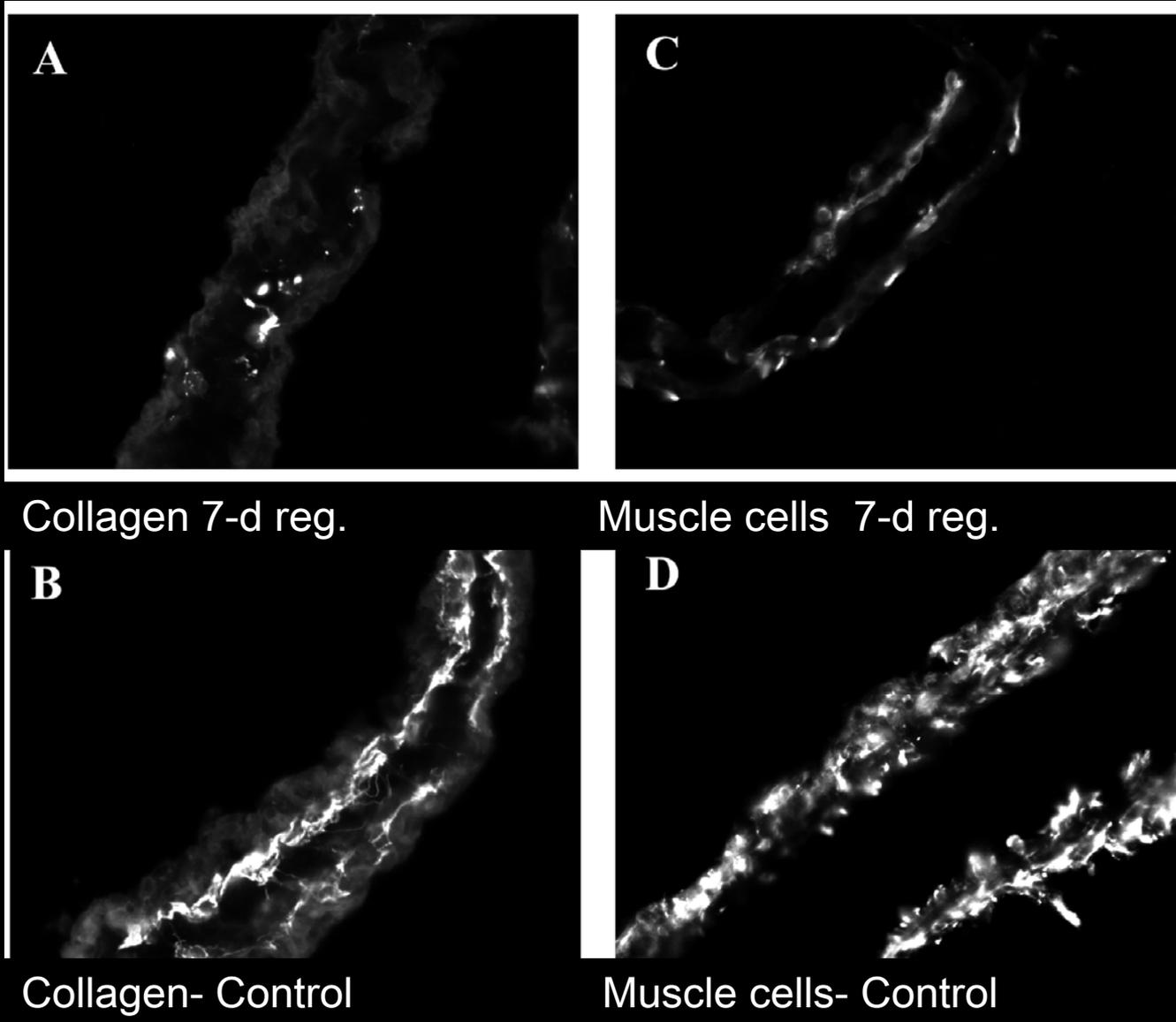


MES-1 labeling

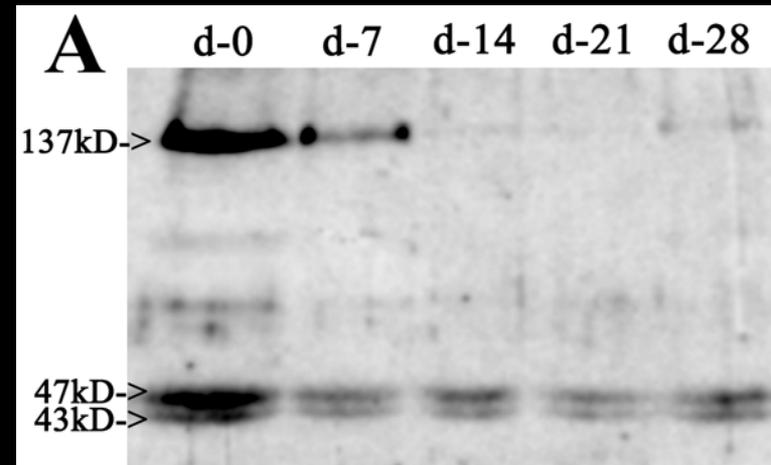
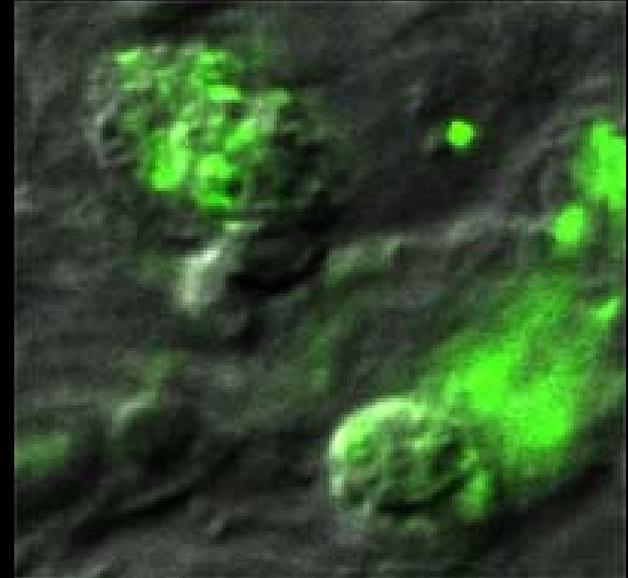
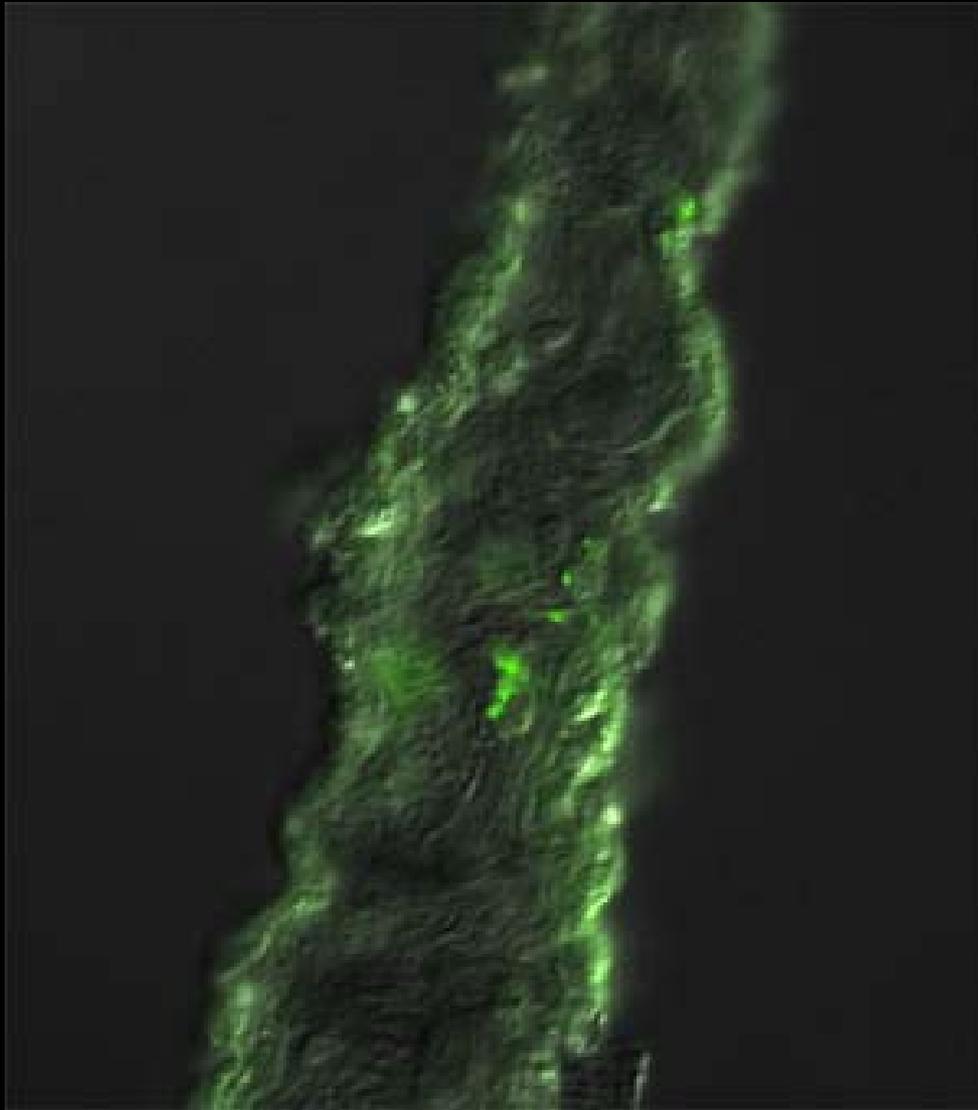


DAPI- nuclei staining

Extracellular matrix (ECM) remodeling

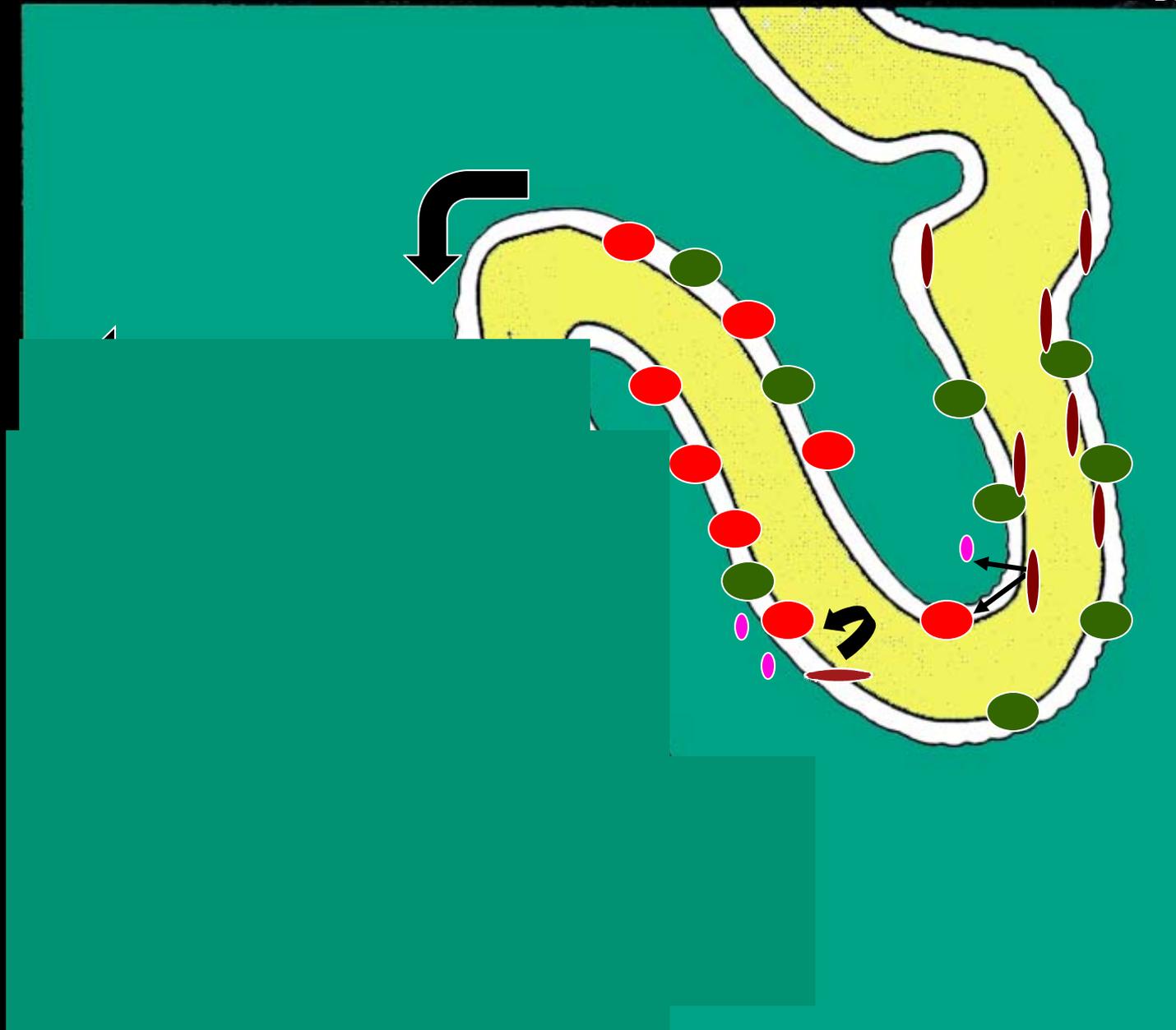


Collagen is degraded by phagocytic amoebocytes



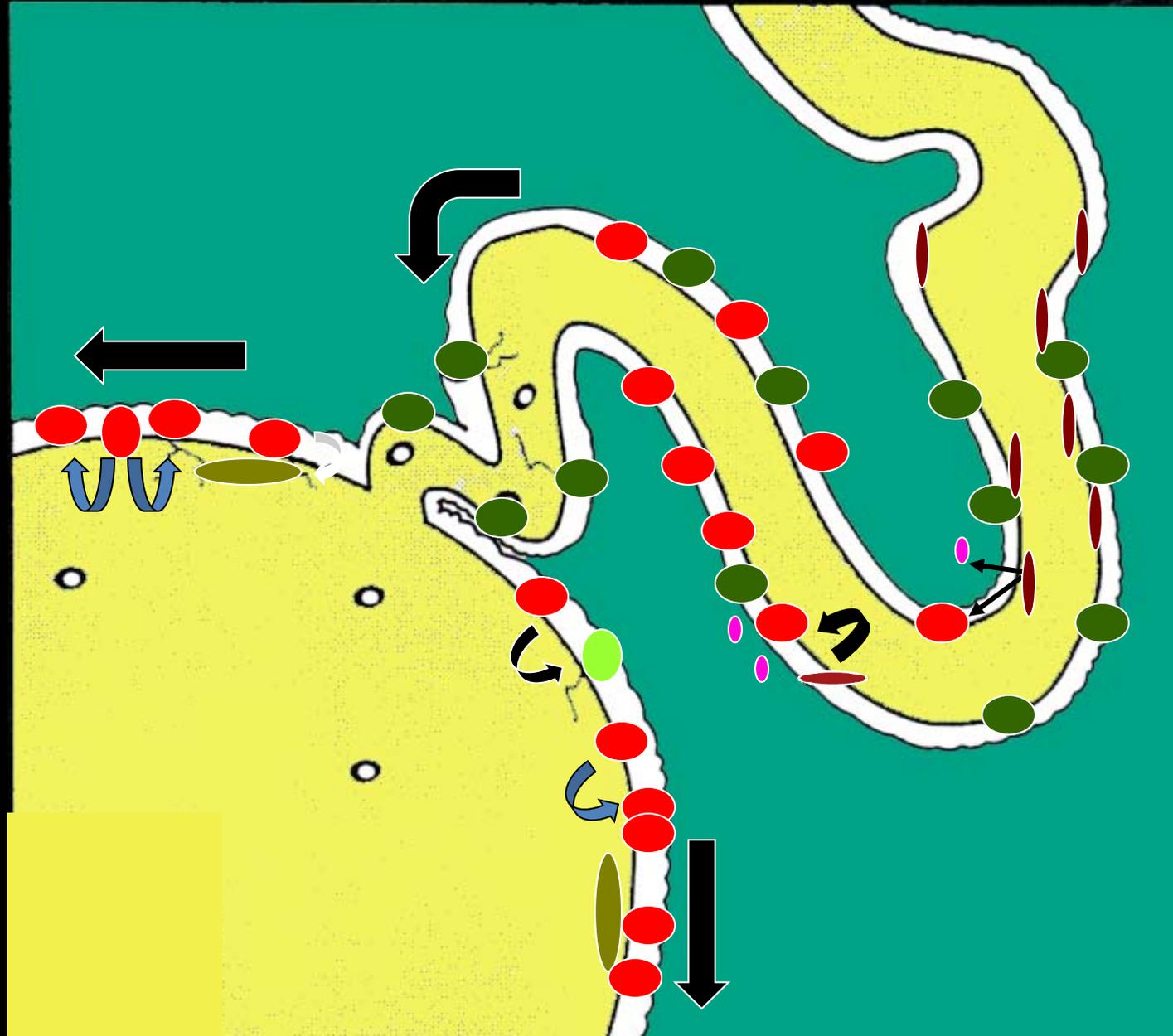


REGENERATION MODEL- cellular origins



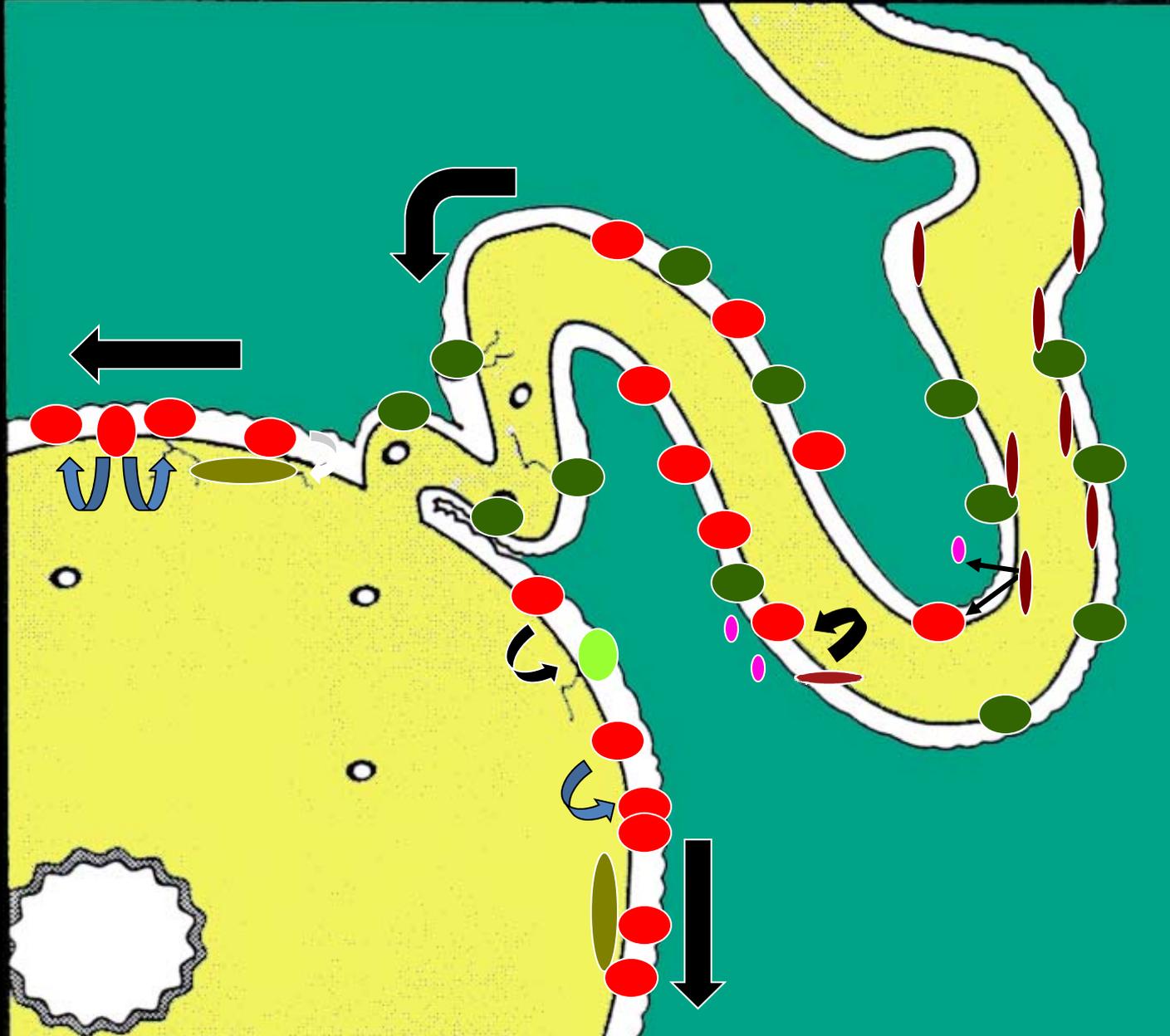


REGENERATION MODEL- cellular origins

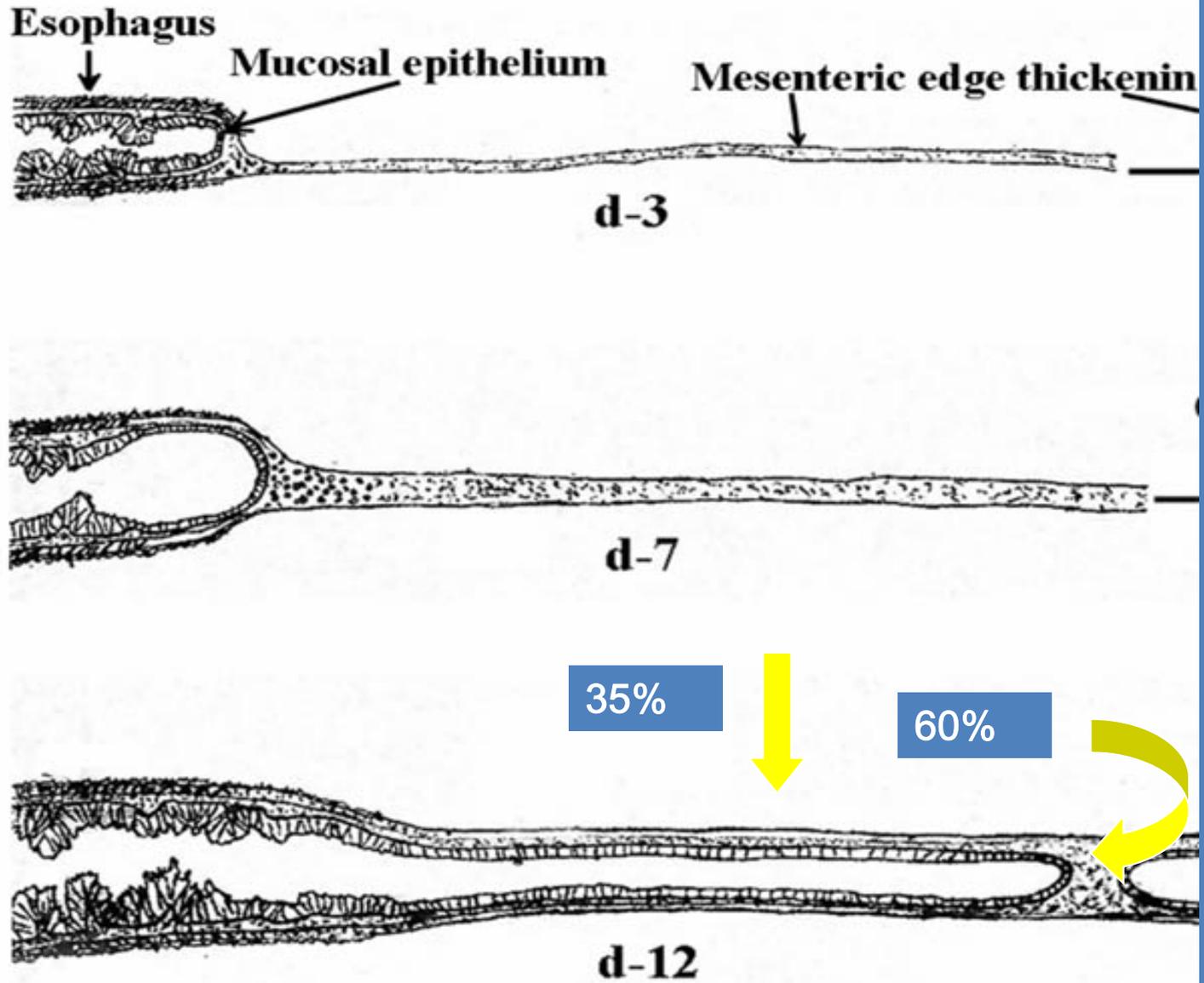




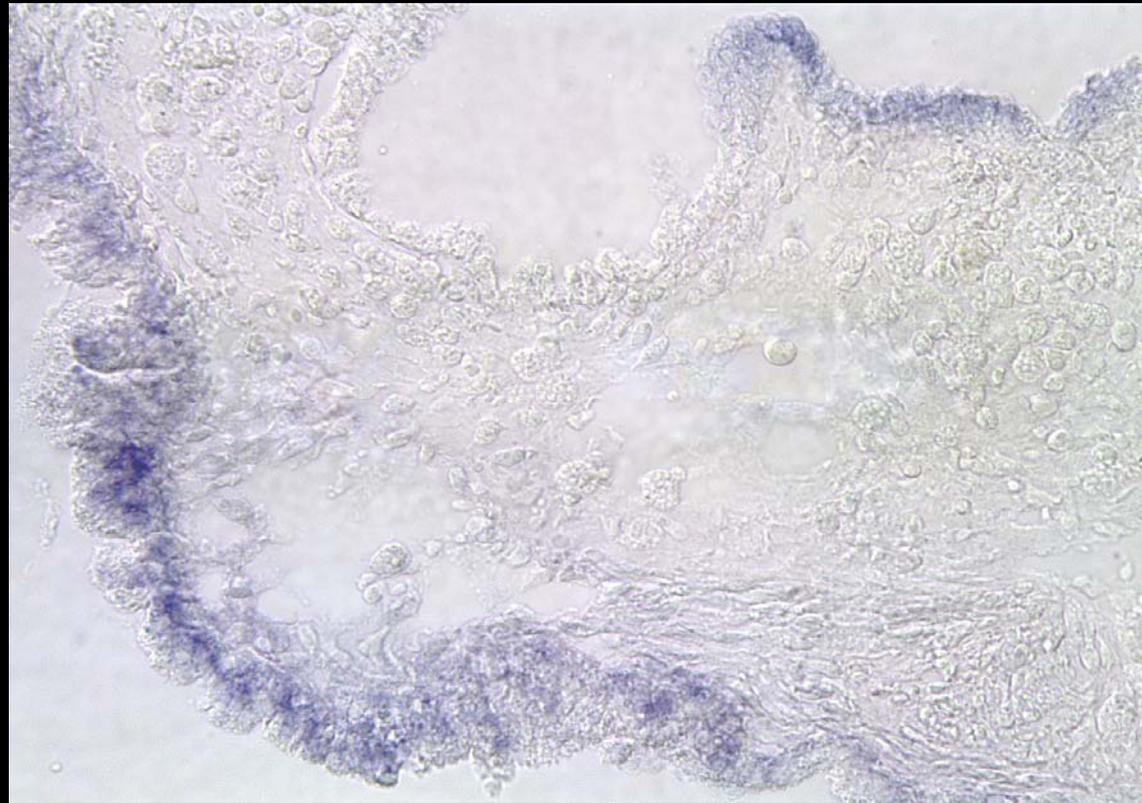
REGENERATION MODEL- cellular origins



Cell Migration



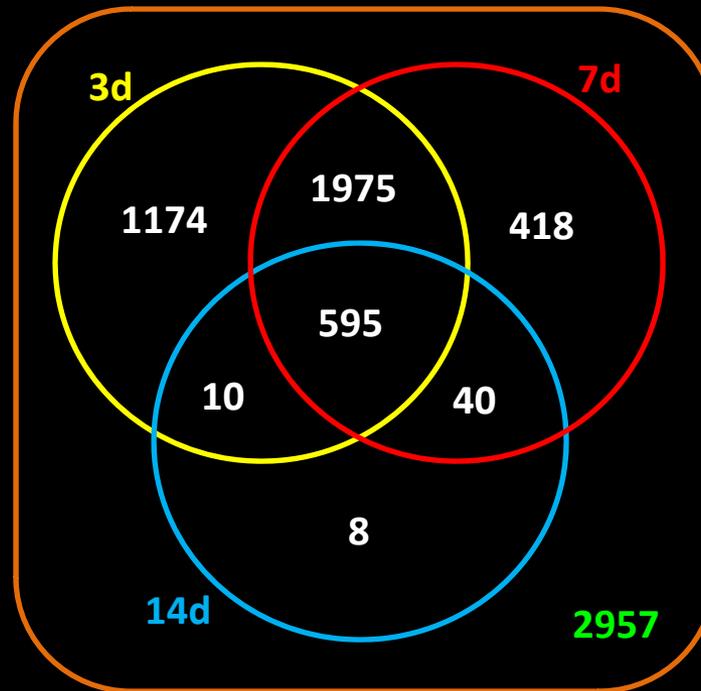
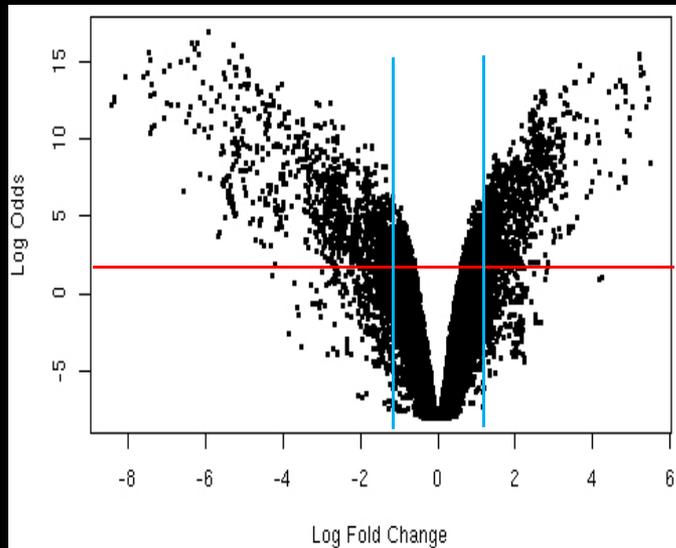
Challenge #1 - Limited molecular tools



In situ hybridization for Survivin mRNA
in 10-day regenerating animal

Challenge #2 - Genomics

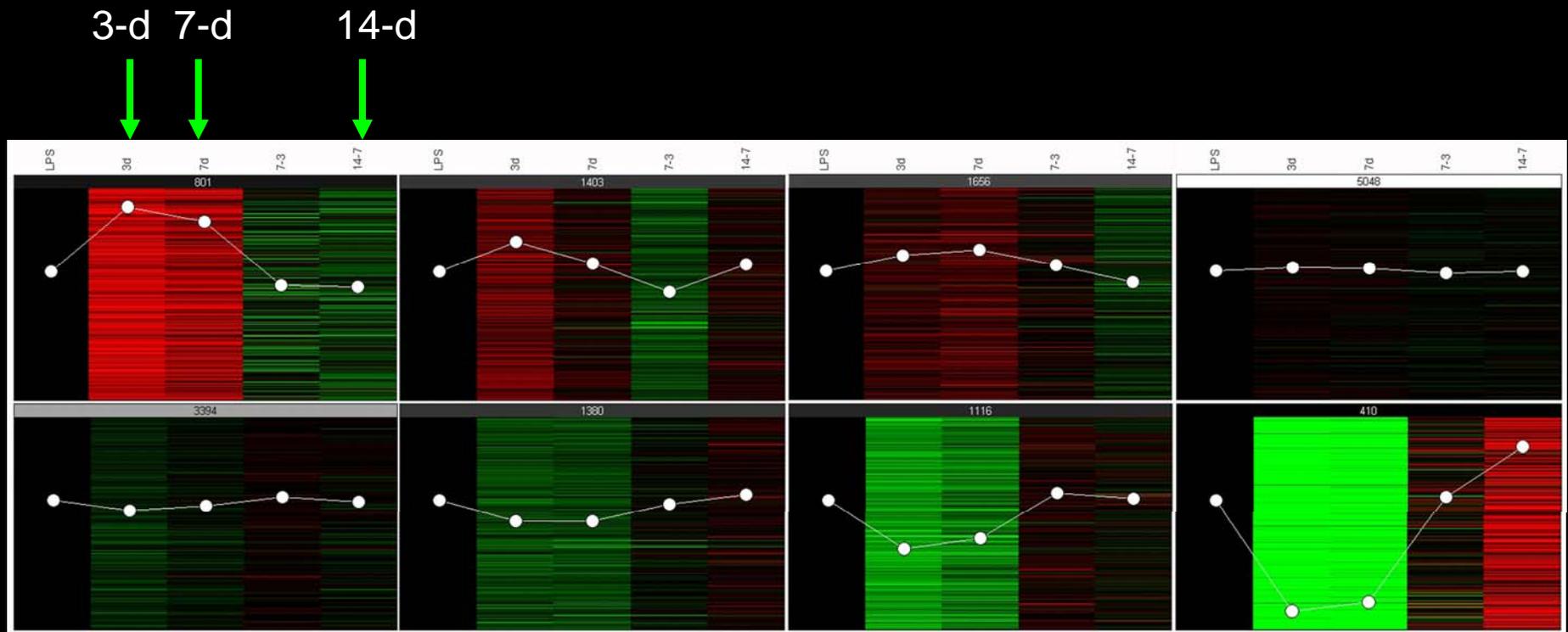
- We have a databank with over 7000 ESTs from 3 cDNA libraries of normal and regenerating intestines
 - Microarrays were done with custom made microchips with over 7000 *H. glaberrima* ESTs.
 - A large number of ESTs are differentially expressed in regenerating animals



$p < 0.05$ -- 73%
 $p < 0.01$ -- 58%
 $P < 0.001$ -- 39%

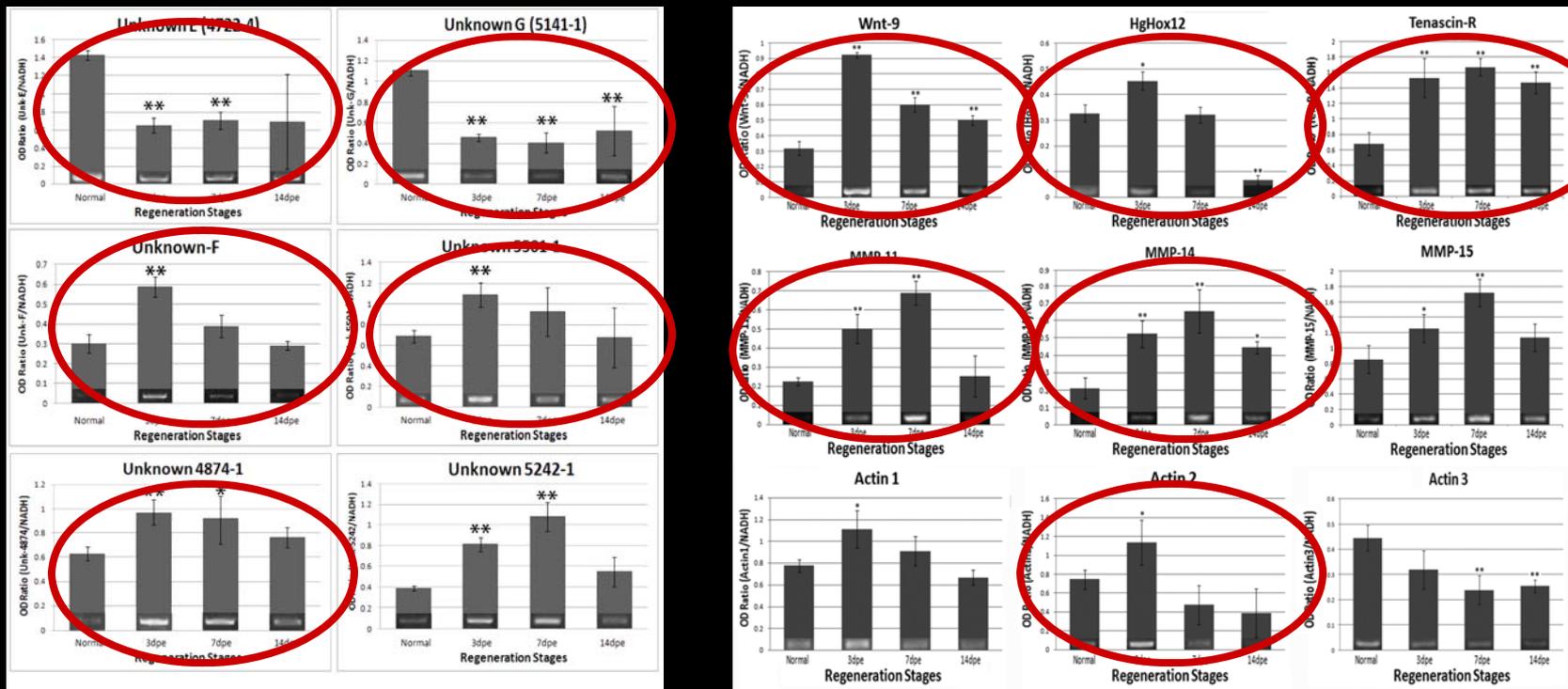
Reference = Normal (non-eviscerated)
 $n = 7166$ $p < 0.01$

Clusters of gene expression



Over 85% of genes validated with PCR showed the same level of significant differences as the microarray.

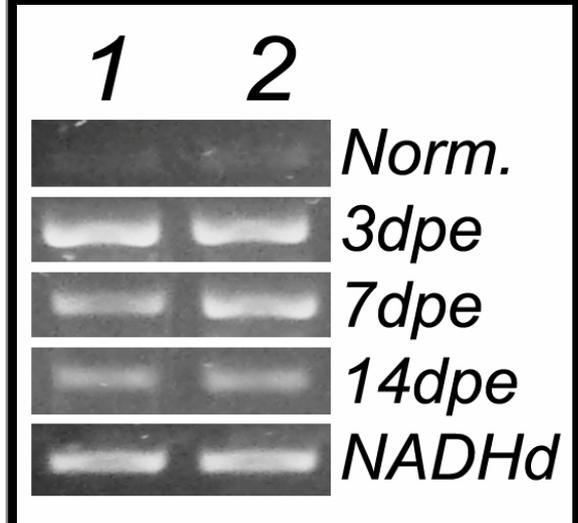
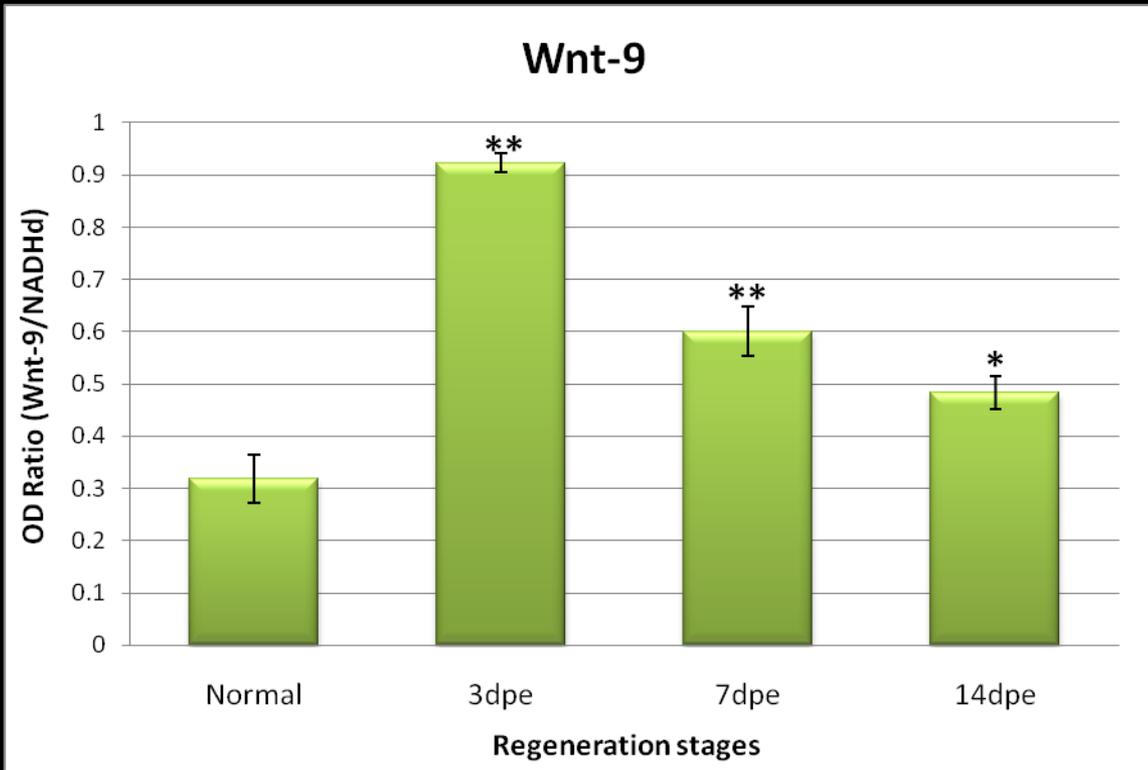
Studies can focus on selected candidates or novel genes.



Wnt- candidate gene

- **Gene family of secreted factors with important roles as regulators of embryonic development**
- **Important role in maintenance and the activation of proliferation of stem cells**
- **Associated with regeneration processes in various animal models**

Wnt-9 is over-expressed during intestinal regeneration showing the highest values in the 3-day intestine

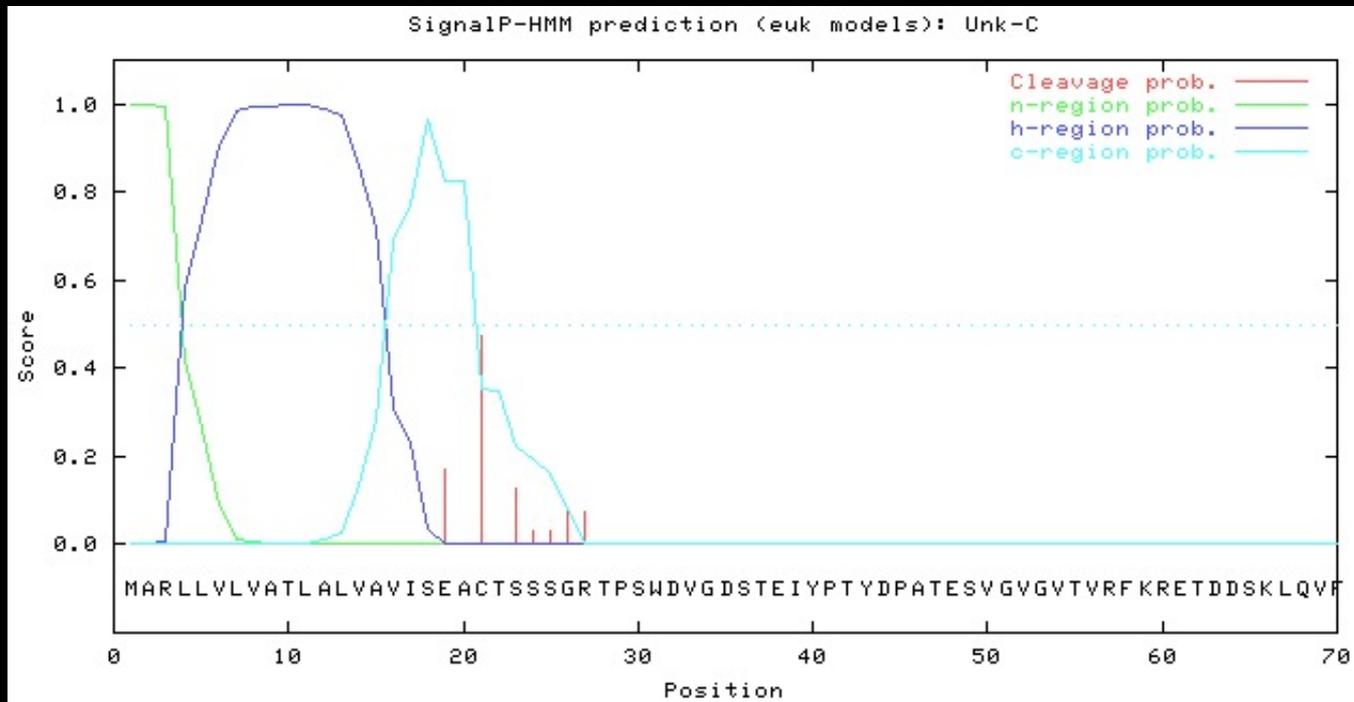


Orpin- novel gene

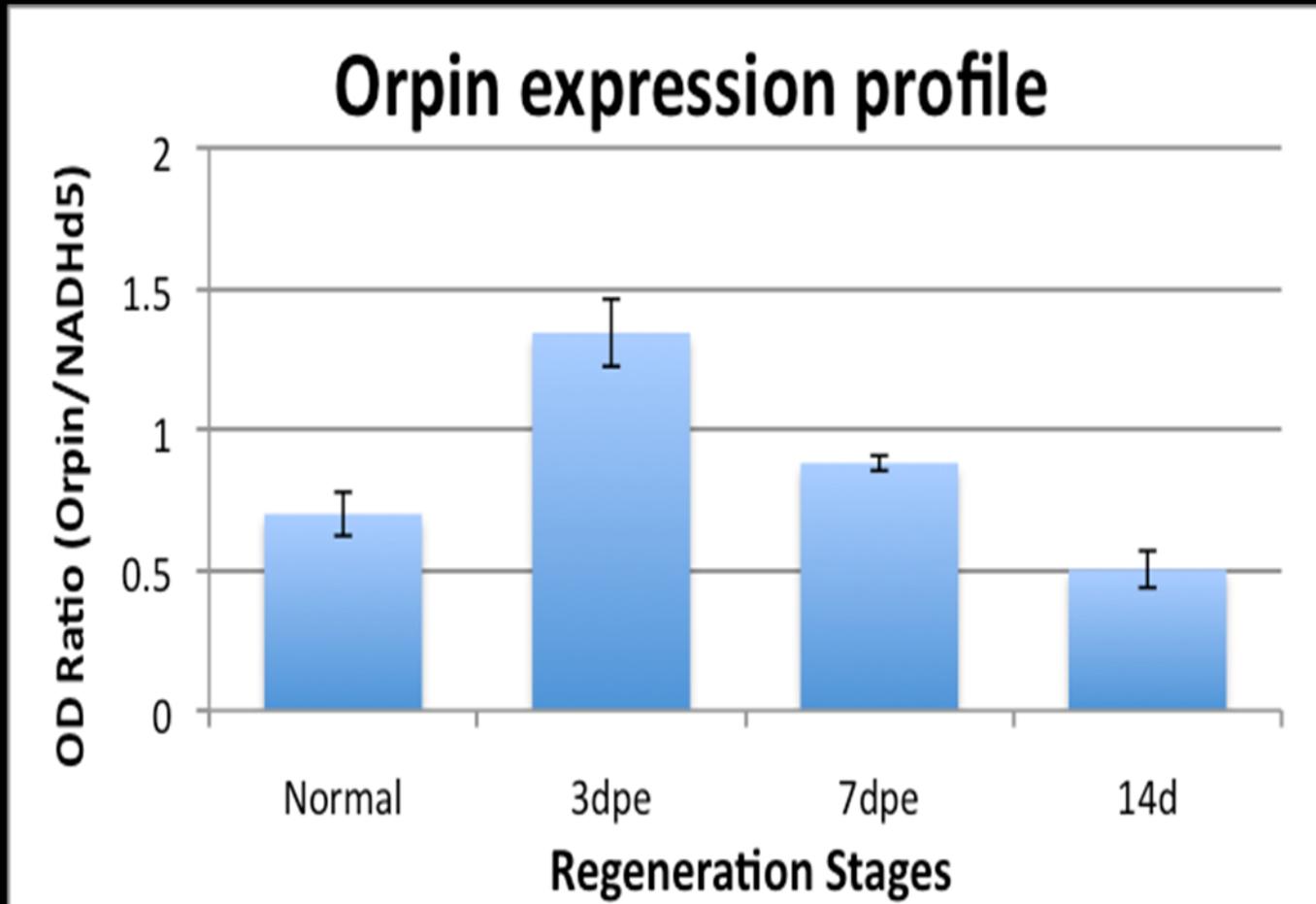
- Large number of ESTs in regenerating cDNA libraries that assemble into one contig
- Over-expressed at 3-days of regeneration in the microarray
- No similarity to genes in database

Orpin

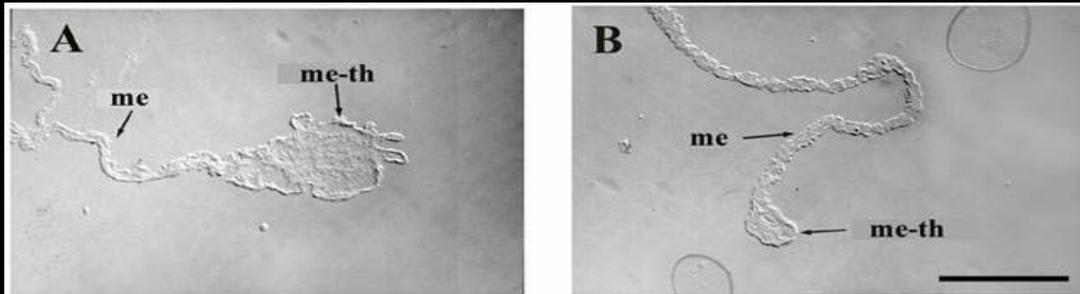
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62  tcgcaactcctccagaatcatccagaacaagcatttacatcaatcatgggctagacttctt
                                     M A R L L
122 gtactcgtcgctacacttgctttgggtcgctgtcattagtgaaagcgtgcacctcatcatct
    V L V A T L A L V A V I S E A C T S S S
182 gggcgcacgccatcttgggacggttggcgactccactgaaatctaccccacttacgatcct
    G R T P S W D V G D S T E I Y P T Y D P
242 gctaccgaatcagttggagtcggagtcactggttaggtttaaaagagaaaccgacgatagt
    A T E S V G V G V T V R F K R E T D D S
302 aaactgcaggtcttcaaattaattgacgcatctgggtgacggttacattgacgcttgcgag
    K L Q V F K L I D A S G D G Y I D A C E
362 tggttaatagagggcggtattgtcaaaaactttgtccagttcctgacagacgatgatgtc
    W L I E G G I V K N F V Q F L T D D D V
422 gatggatgatgagaagatcttggatgagttccagaaagtgtcagtcgcatagtagata
    D G D E K I S W N E F Q K V S V A *
482 gaaatagcgcggcggcagttccagttatttaatatagtcatttttagagacctgtagggtaa
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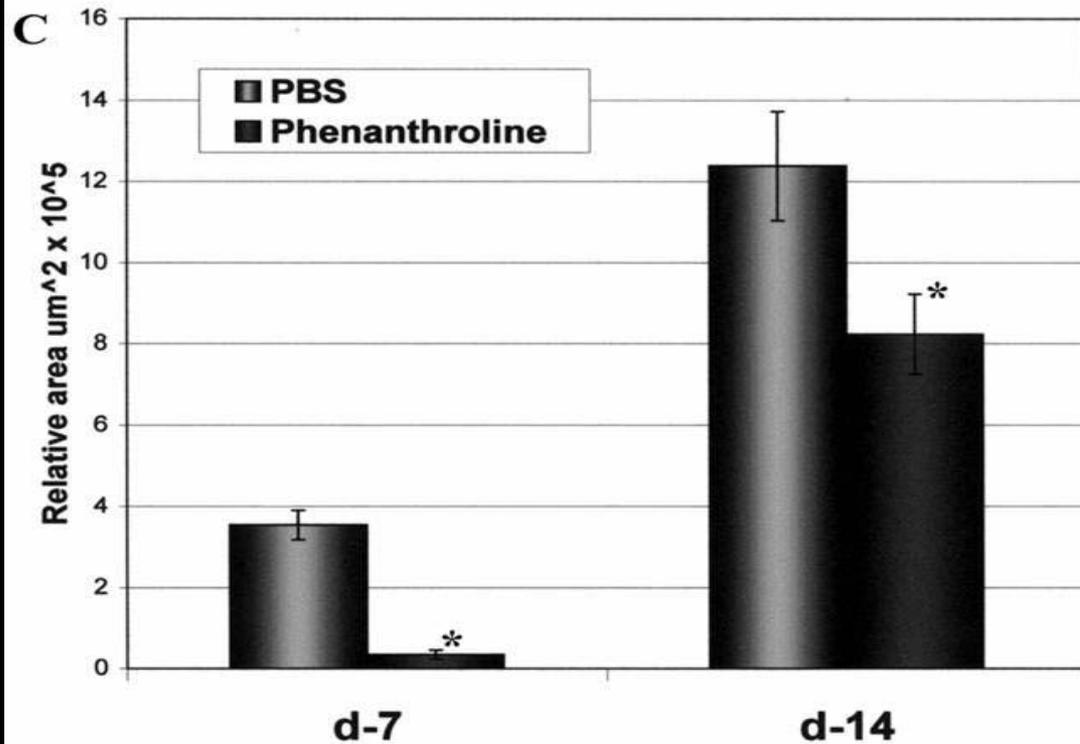
RT-PCR validation of Orpin overexpression in the 3-day regenerating intestine.



Challenge #3 - Loss of function/Transgenics



Use of pharmacological tools.
For example, MMP inhibitors,
apoptosis inhibitors, Wnt
pathway activators.



Missing
RNAi
Transfections
Genetics

Inhibitors of matrix metalloproteases inhibit
intestinal regeneration

TOP ADVANTAGE -

VERY LOW POSSIBILITY OF BEING SCOOPED

TOP ADVANTAGE -
VERY LOW POSSIBILITY OF BEING SCOOPED



THE END