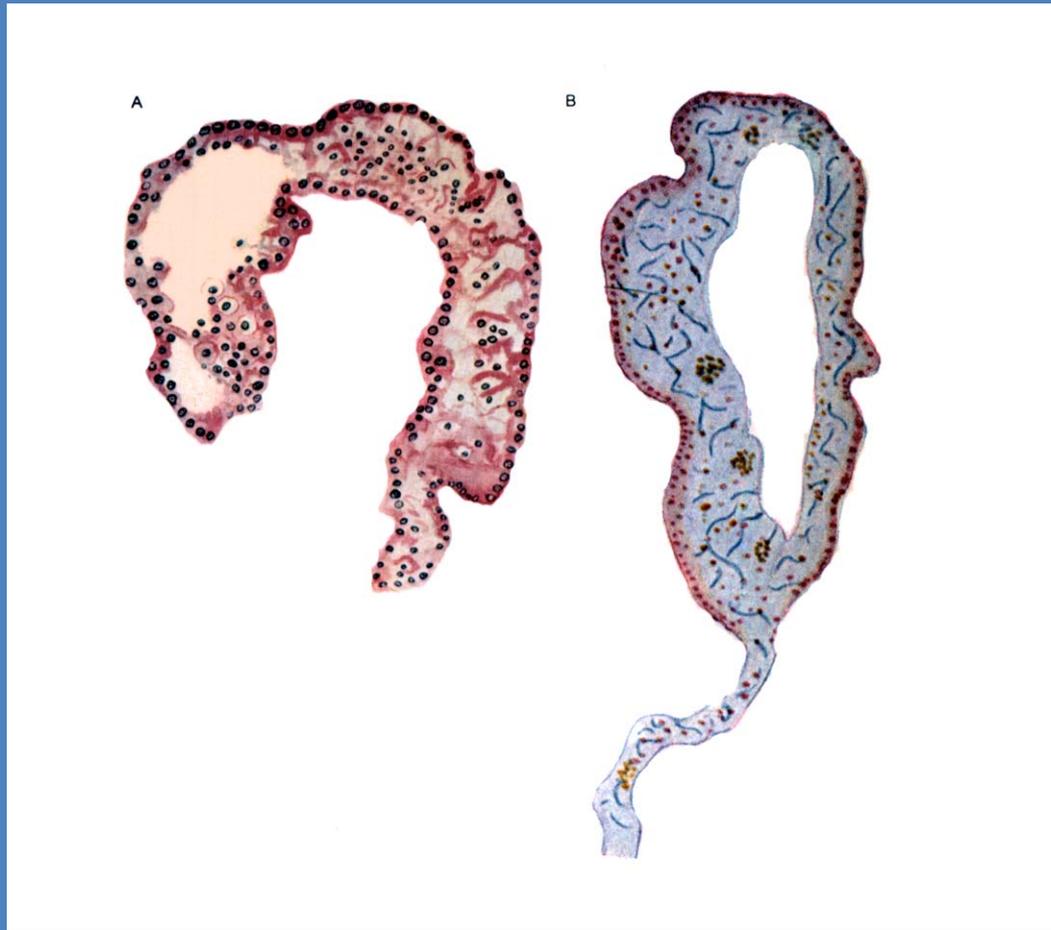


# Intestinal regeneration in the echinoderm *Holothuria glaberrima*



**José E. García-Arrarás, Ph.D.**  
Biology Department  
University of Puerto Rico  
Río Piedras Campus

# 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?

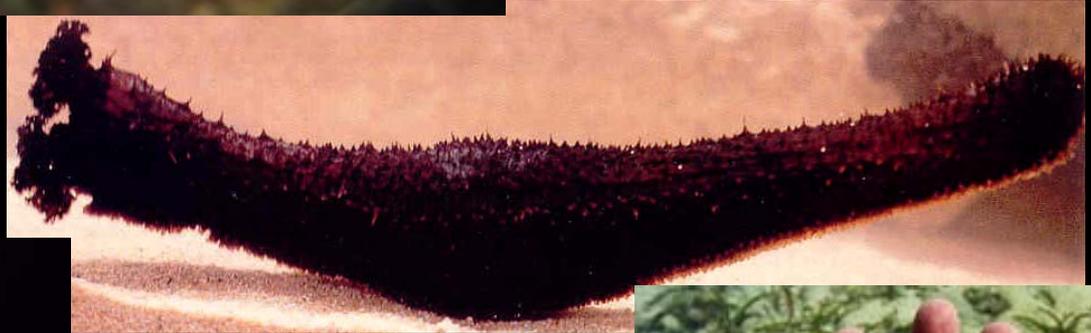


# What are the **ADVANTAGES** and **CHALLENGES** of the sea cucumber model system?



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

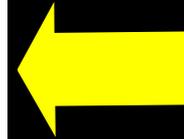
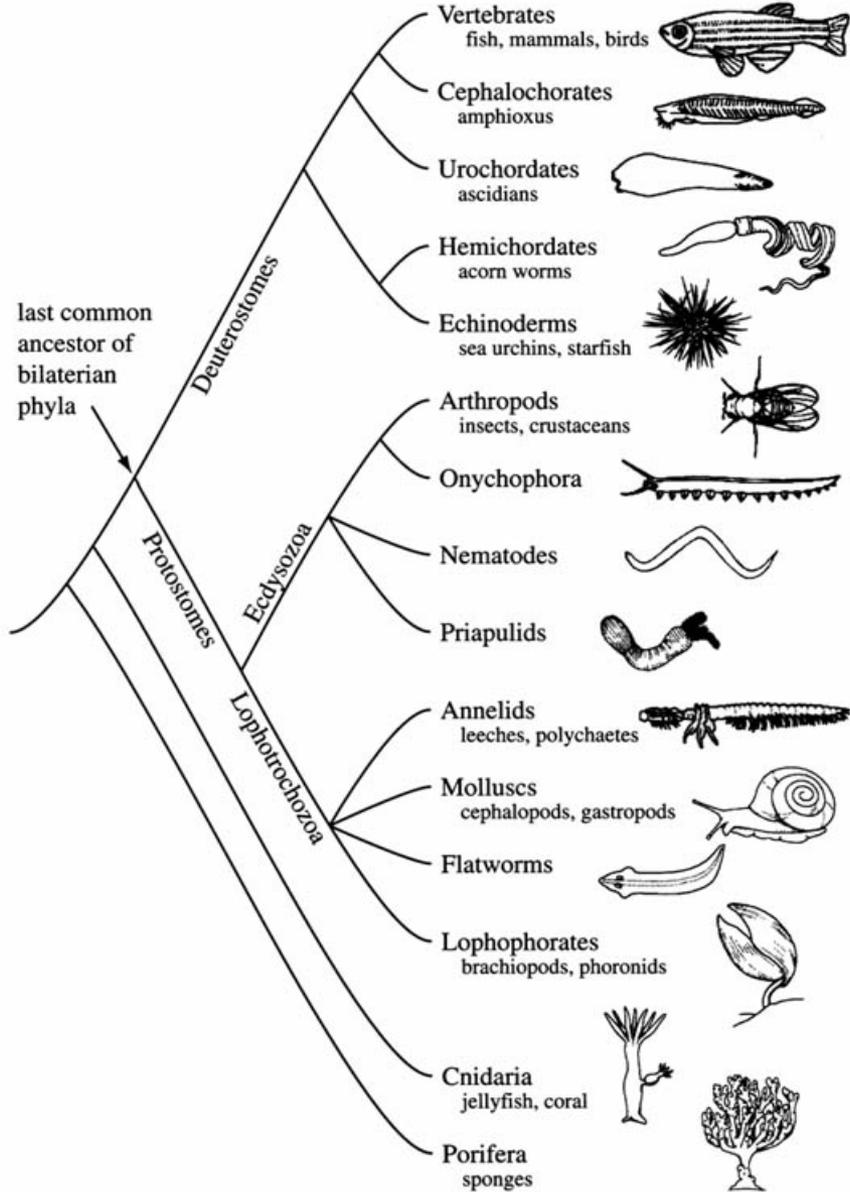
the sea cucumber, *Holothuria glaberrima*



## *Athyonidium chilensis*

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

# Advantage #2- Key phylogenetic position



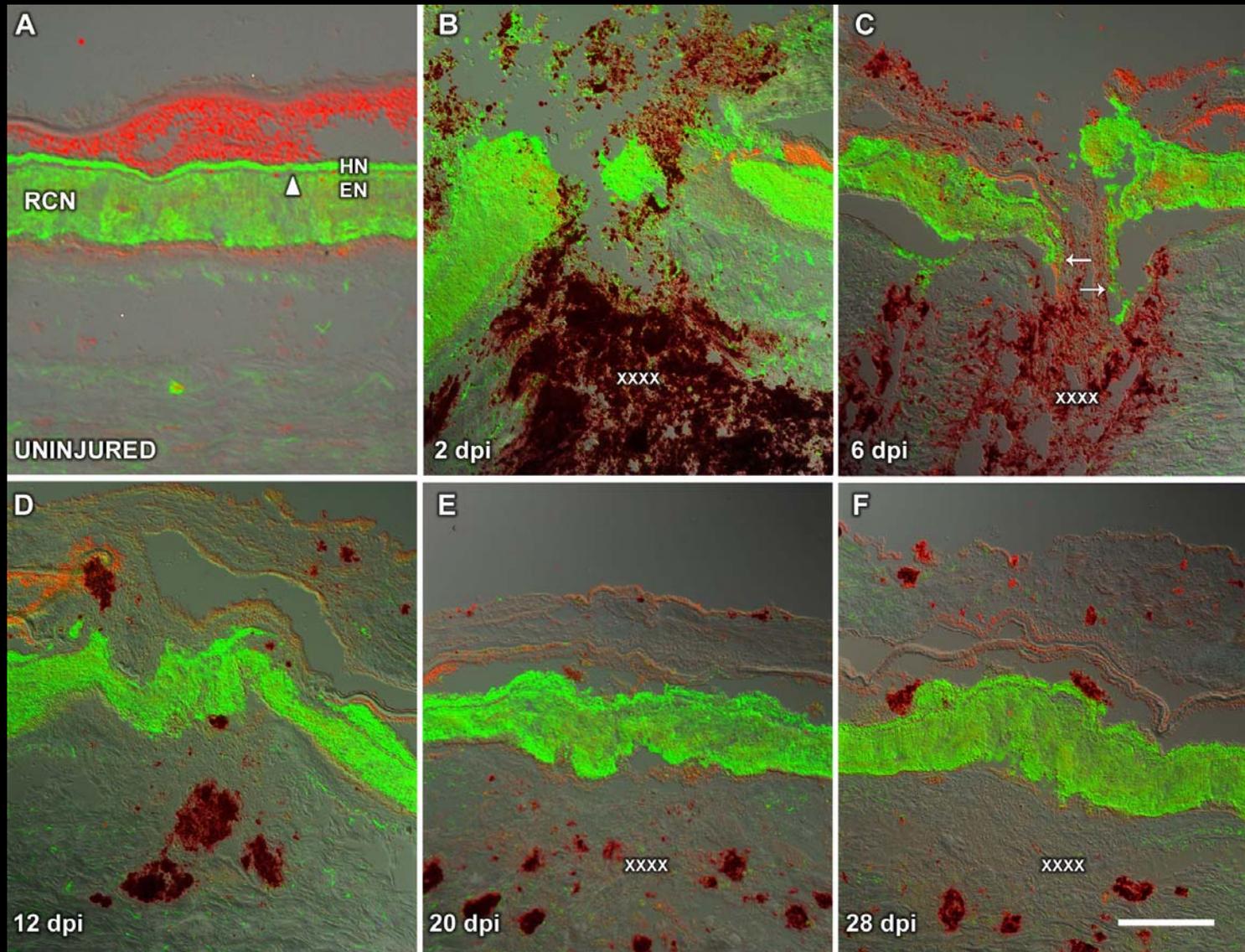
**Echinoderms**

**Sea cucumbers are deuterostomes**

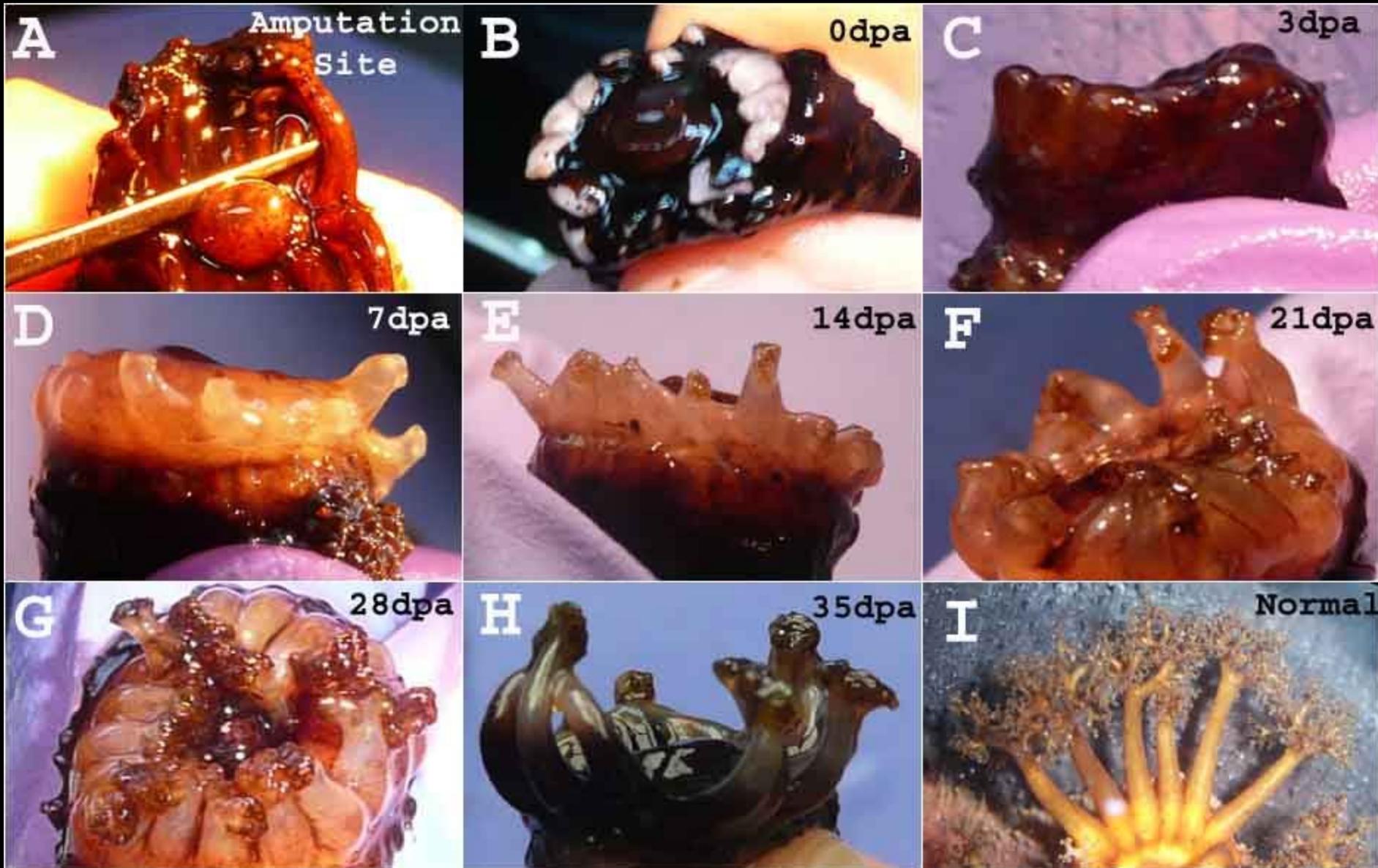
# Advantage #3 - Extraordinary regenerative properties



# Nerve fiber regeneration



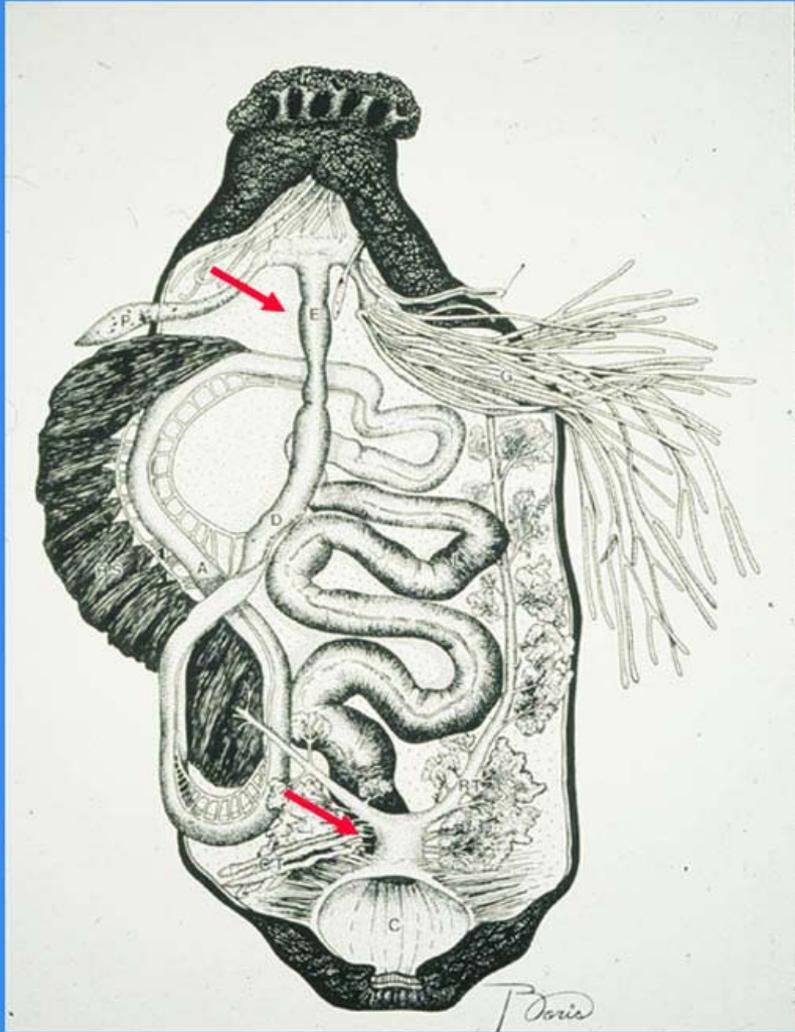
# Tentacle regeneration



## Advantage #4 - Evisceration is easily induced in lab

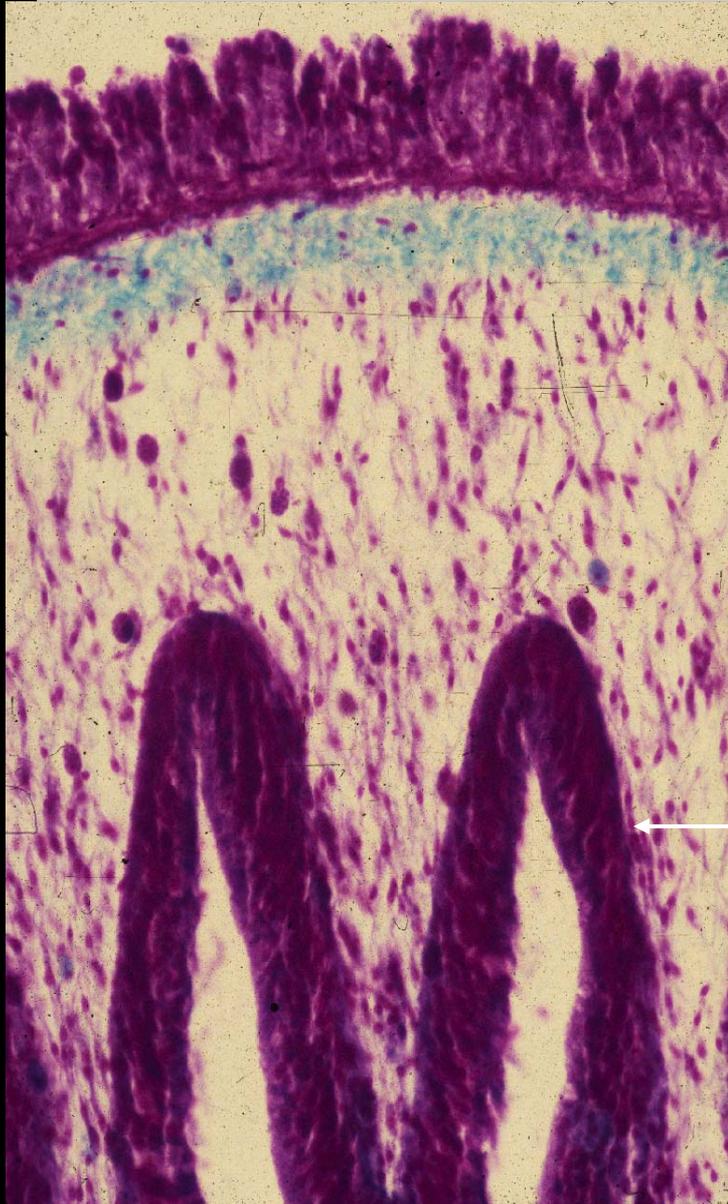


## Advantage #5 - Evisceration follows a fixed pattern, reducing variability due to surgical manipulations

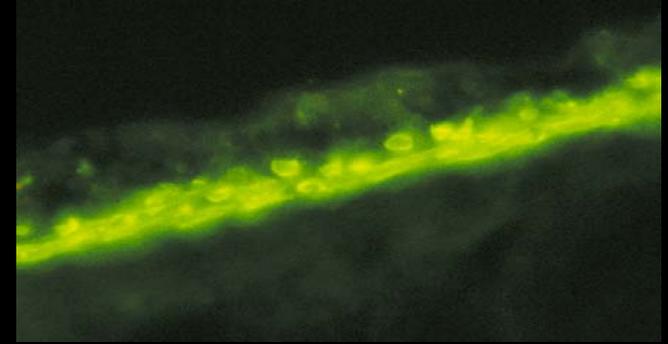


Evisceration eliminates most of the organs of the sea cucumber. In *H. glaberrima* only the left respiratory tree remains.

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



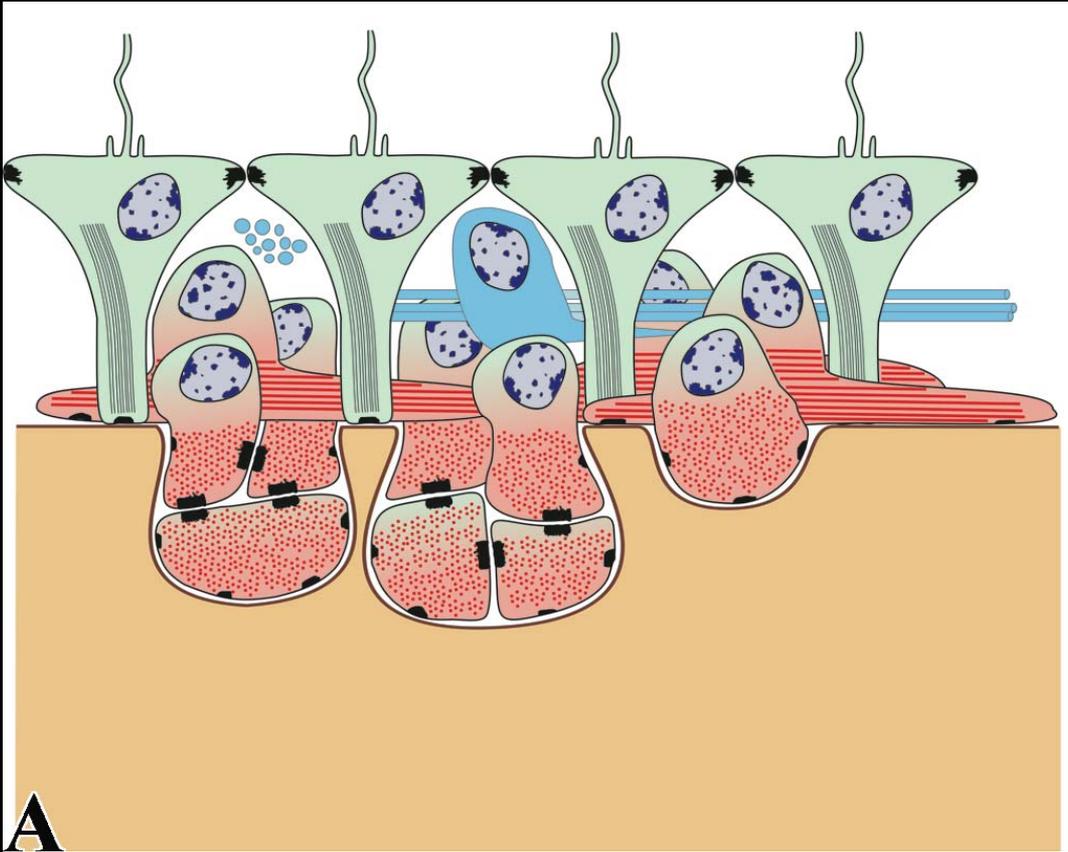
Mesothelium (includes coelomic epithelium or serosa and muscle layer)



Connective tissue layer (submucosa)

Luminal epithelium (mucosa)

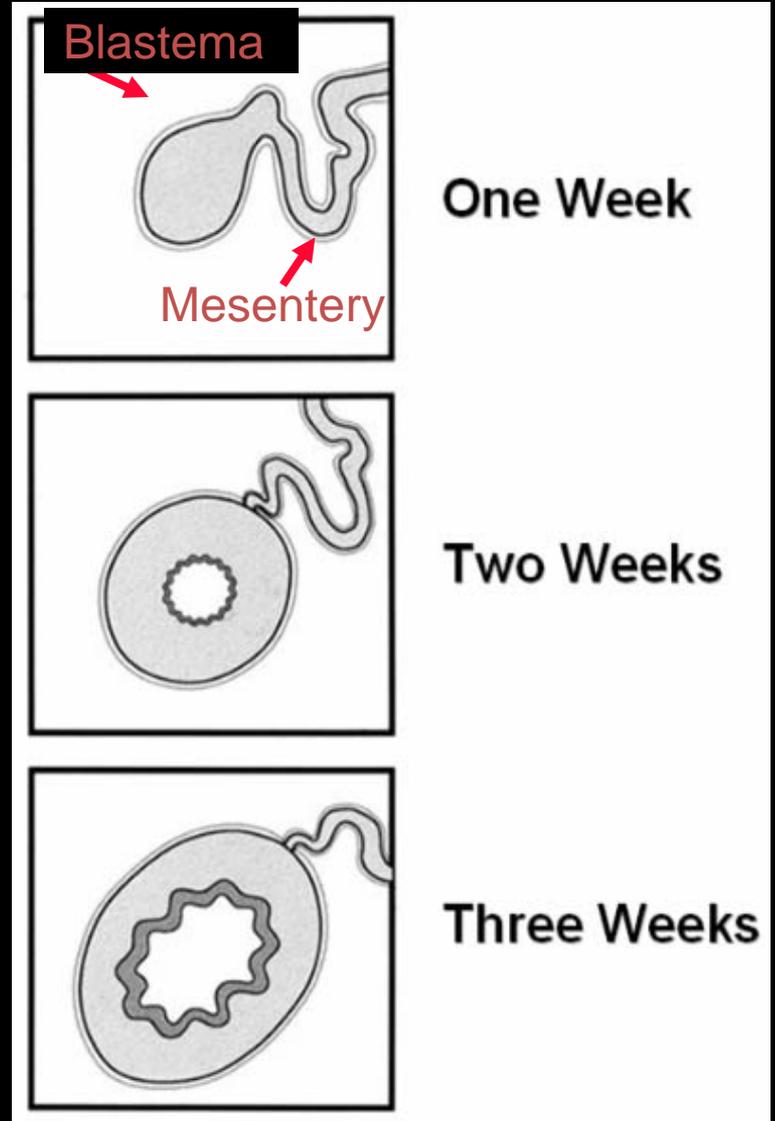
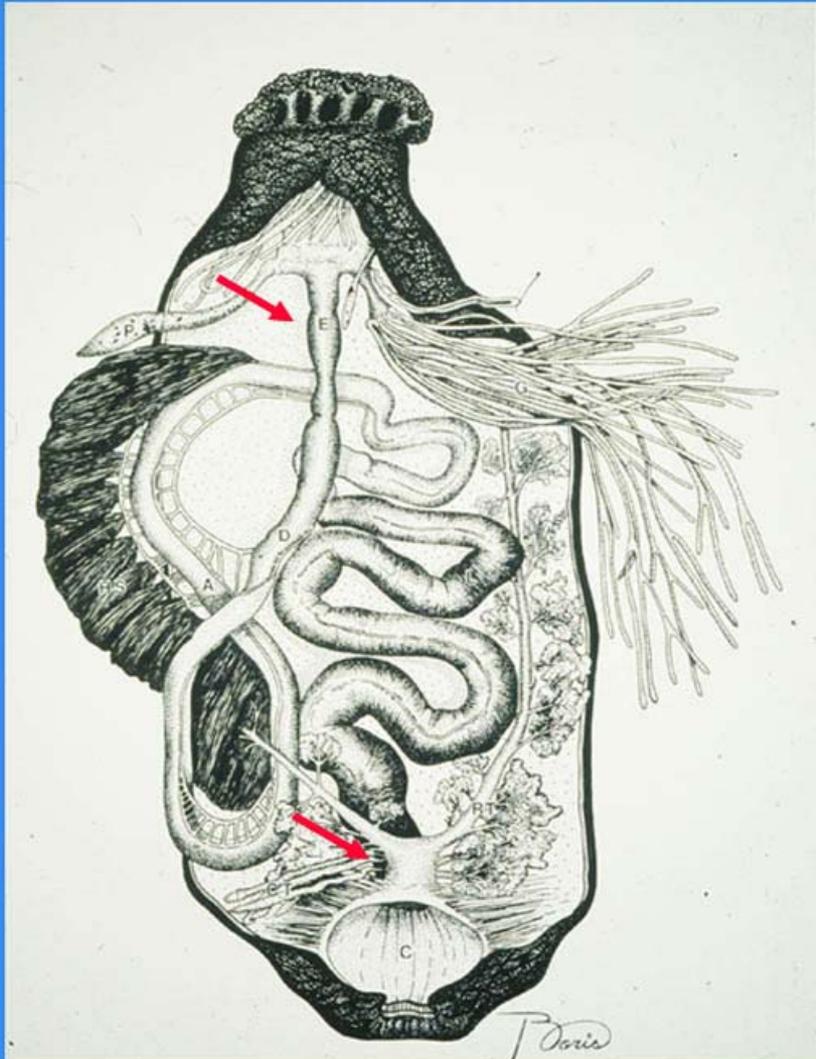
# Echinoderm mesothelium



Peritoneocytes

Myocytes

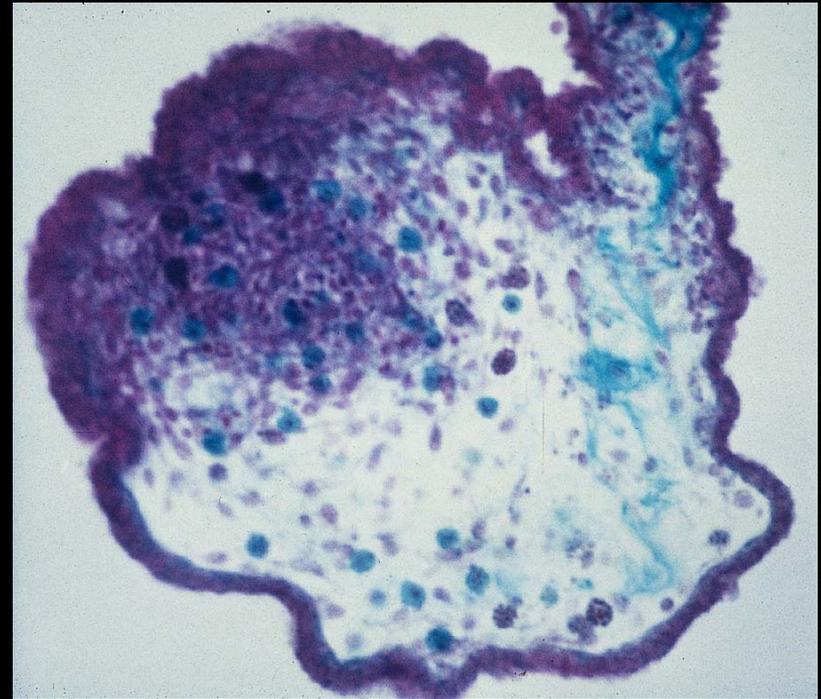
# Advantage #7 - Regeneration of a functional organ occurs within a month



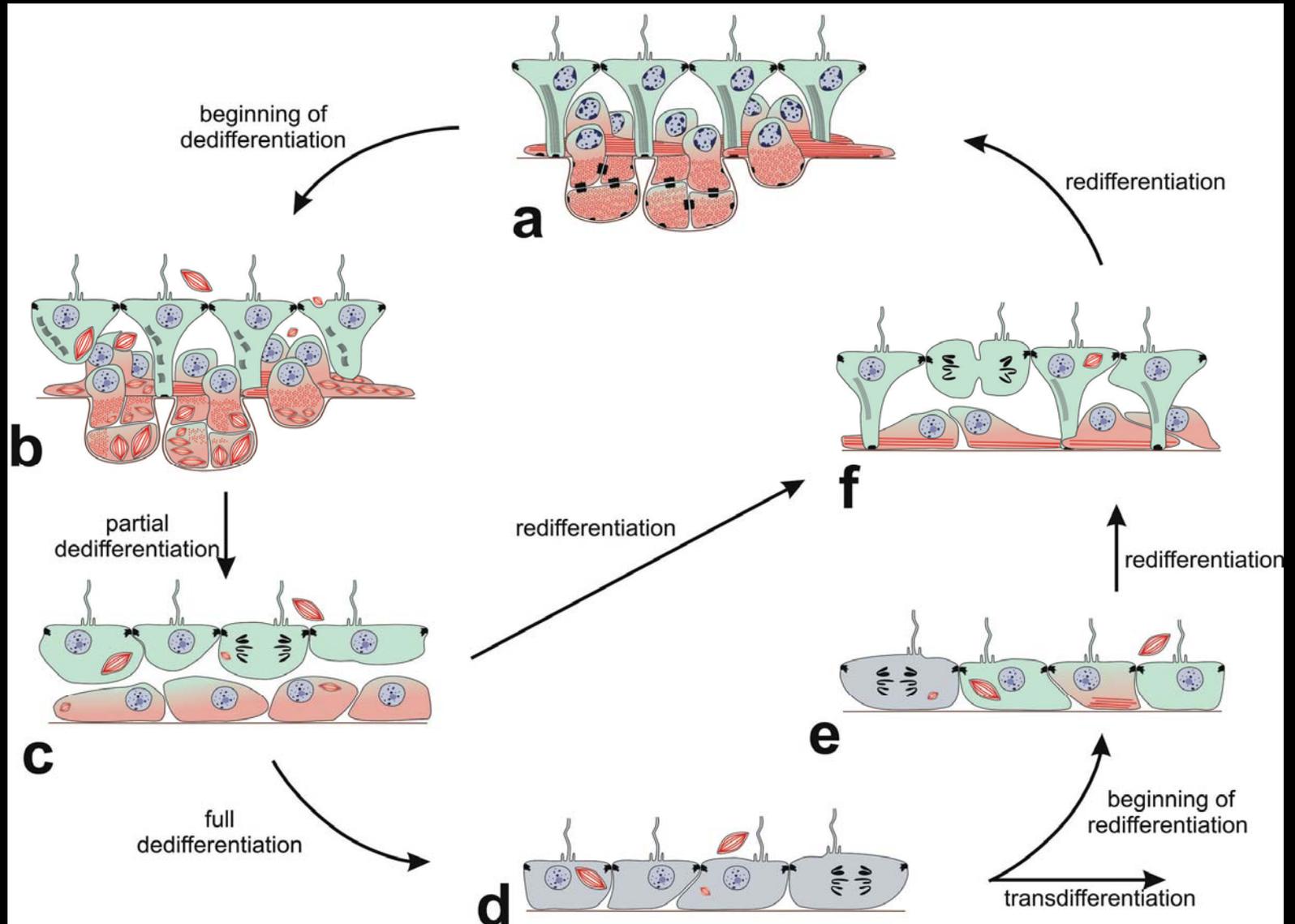
## Advantage #8 - Multiple cellular events can be studied

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

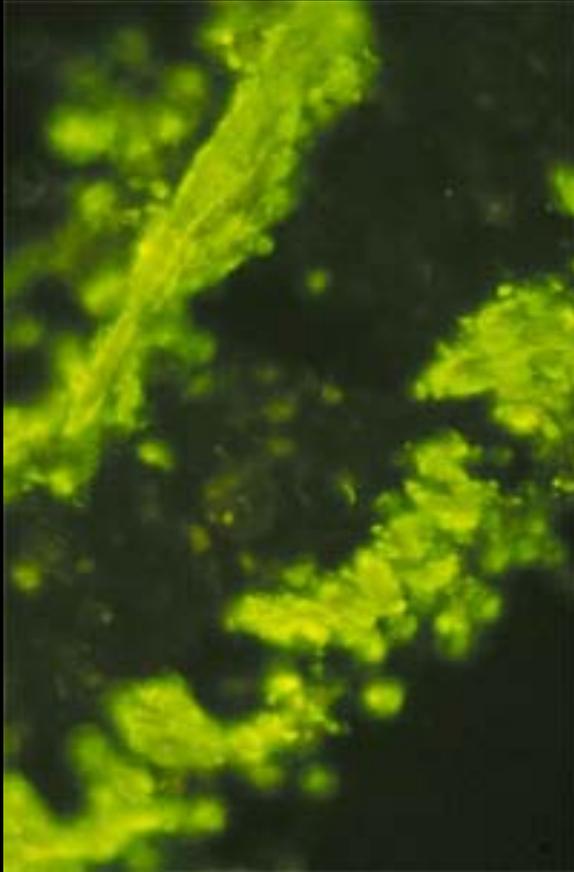
1. Cell dedifferentiation
2. Cell proliferation
3. Apoptosis
4. Epithelial to mesenchymal transition.
5. ECM remodeling
6. Cell migration
7. Cell differentiation
8. Cell-cell interactions to form a new organ



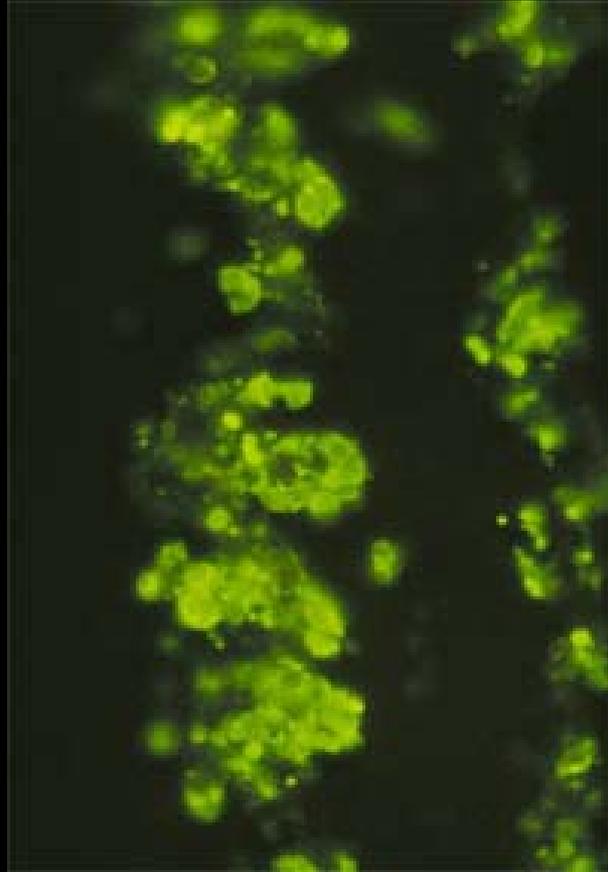
# Cell Dedifferentiation



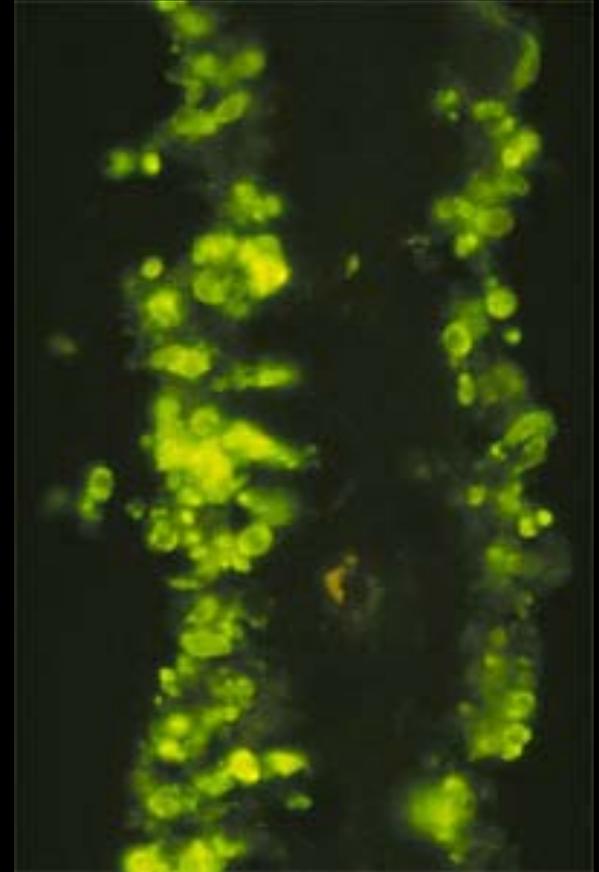
A gradient of muscle dedifferentiation can be found in the mesentery during regeneration



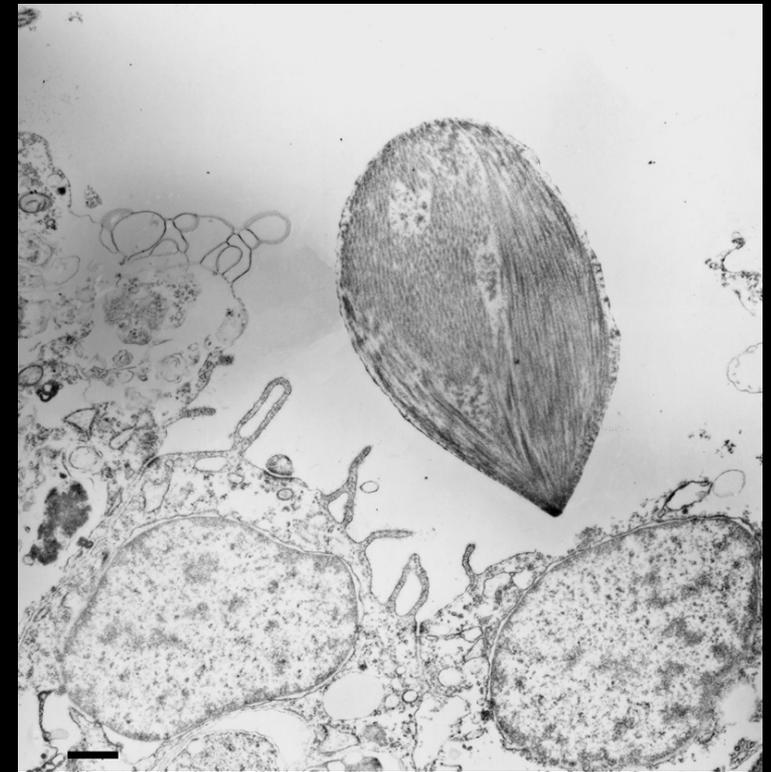
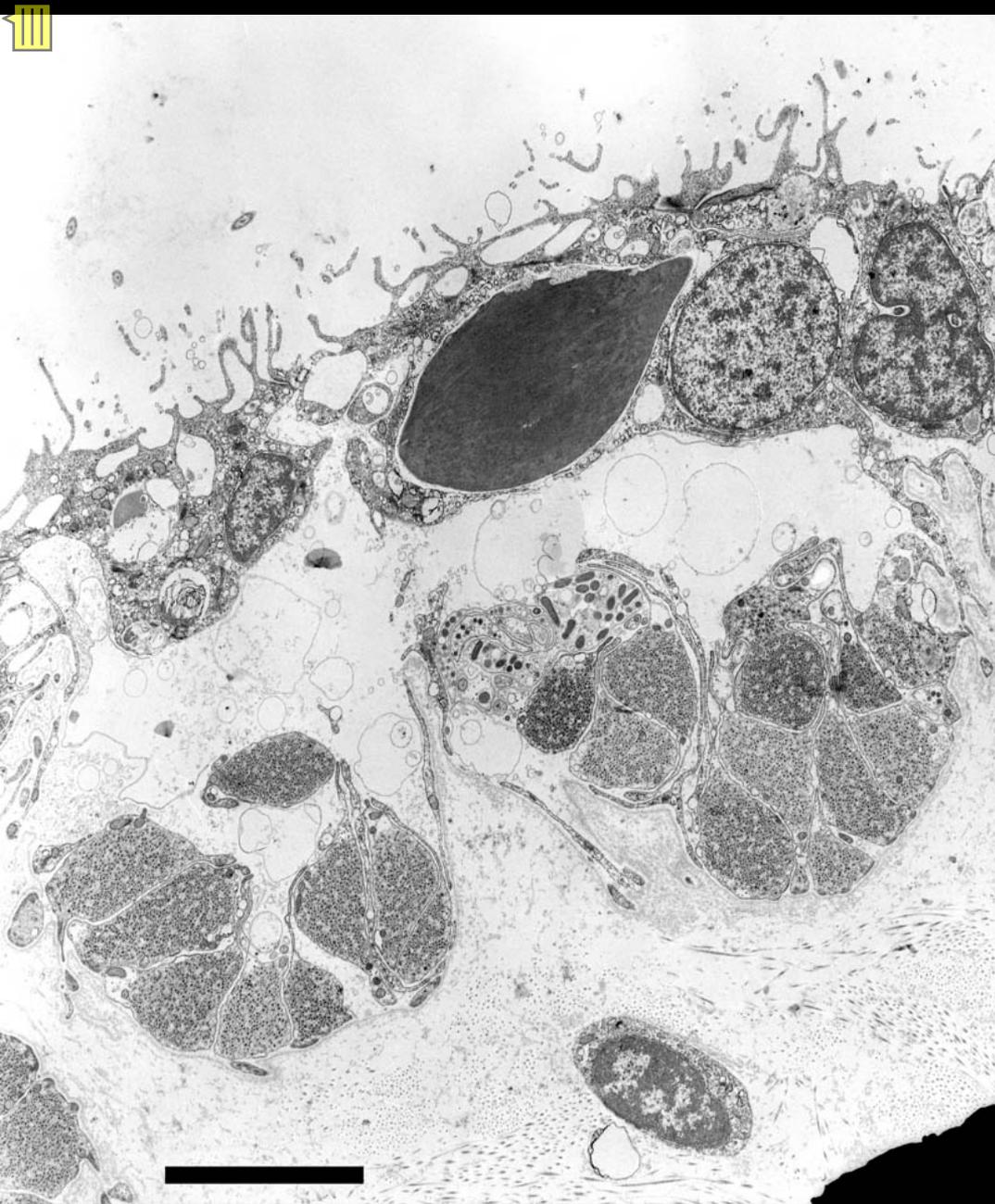
Near body wall



Medial



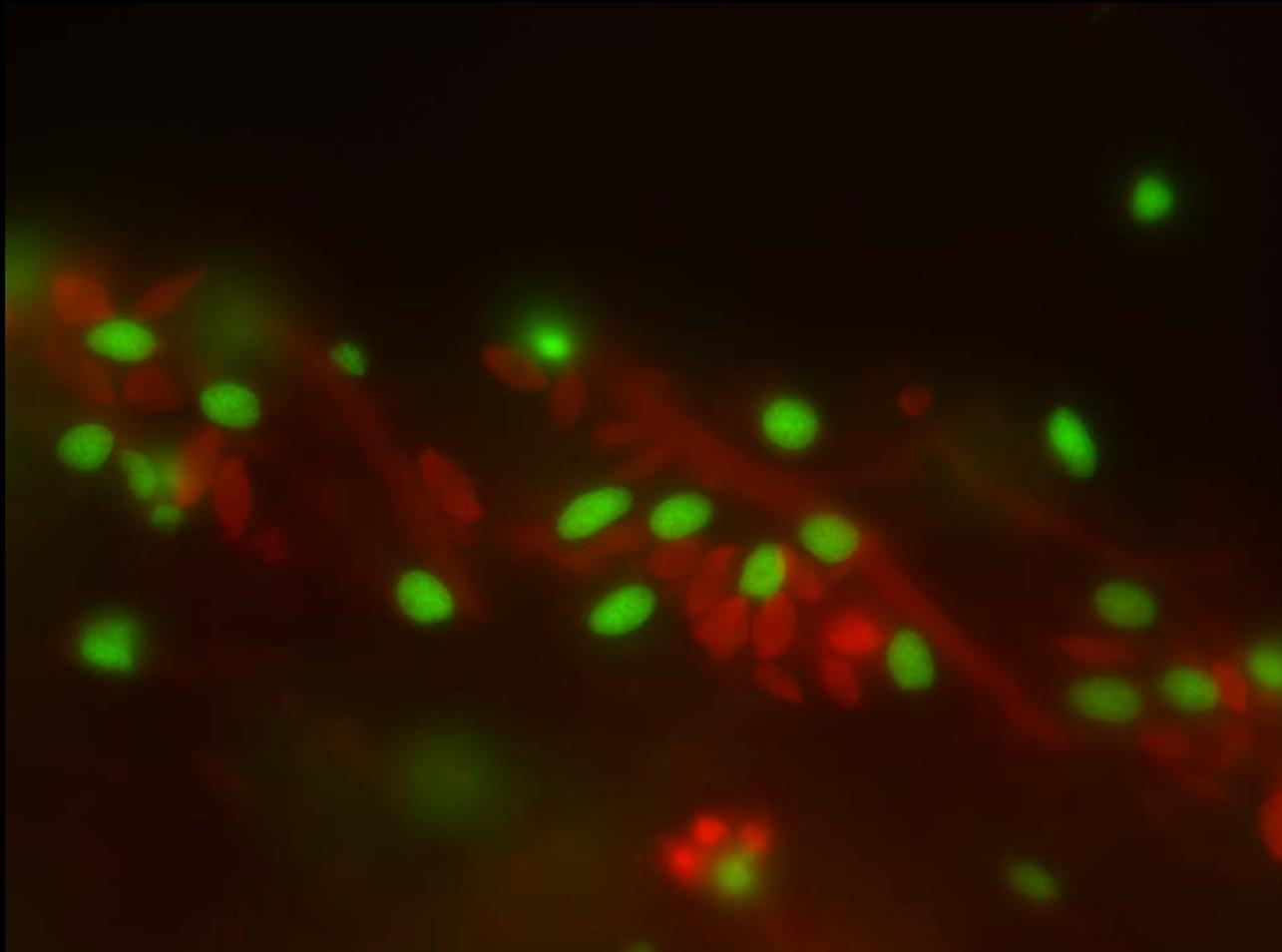
Near blastema-like structure



Muscle cell  
dedifferentiation involves  
the formation of spindle  
like structures

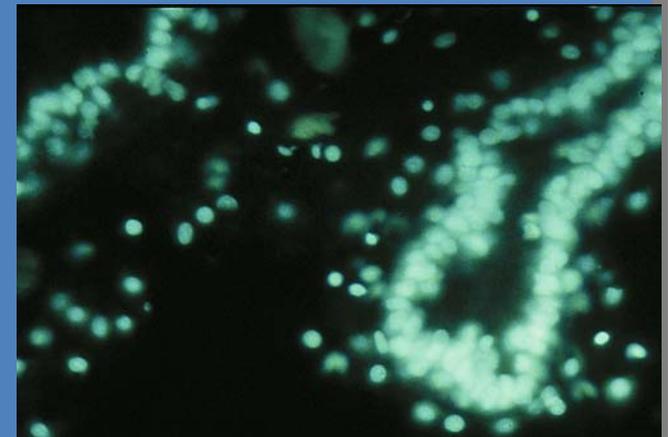
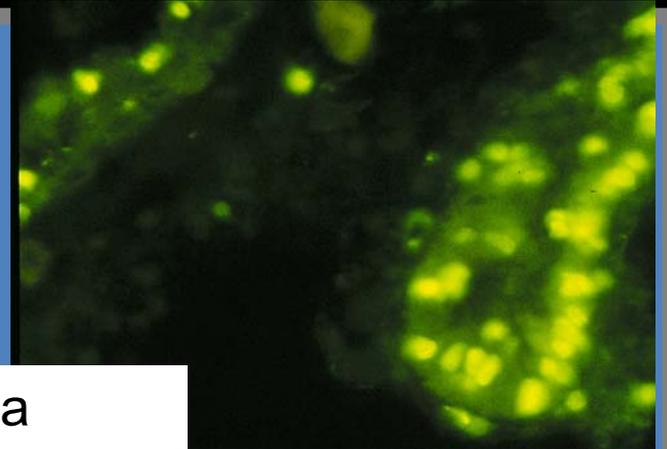
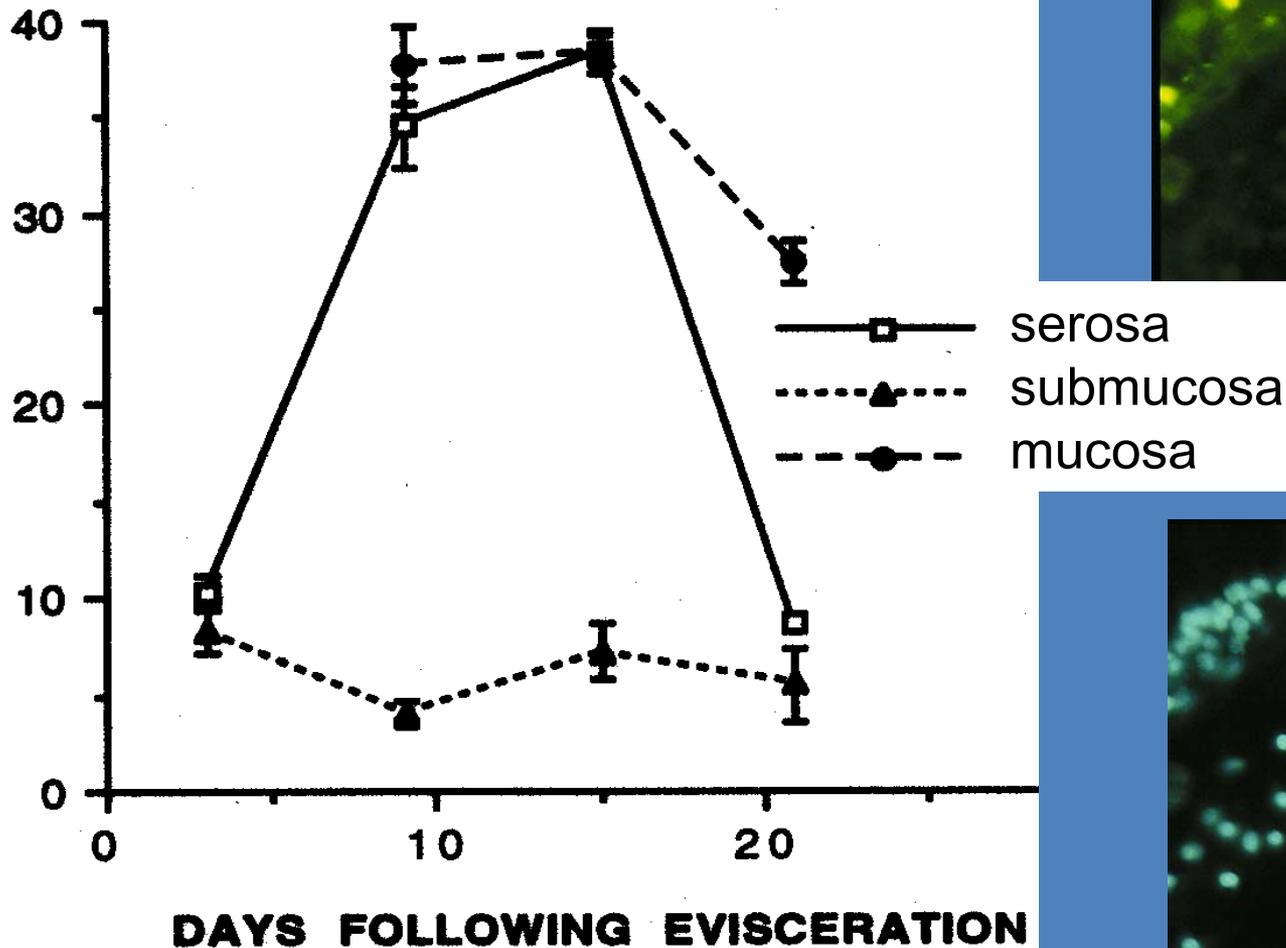


SLSs (red) are formed as cells dedifferentiate. These SLSs are not associated with cell nuclei (green)

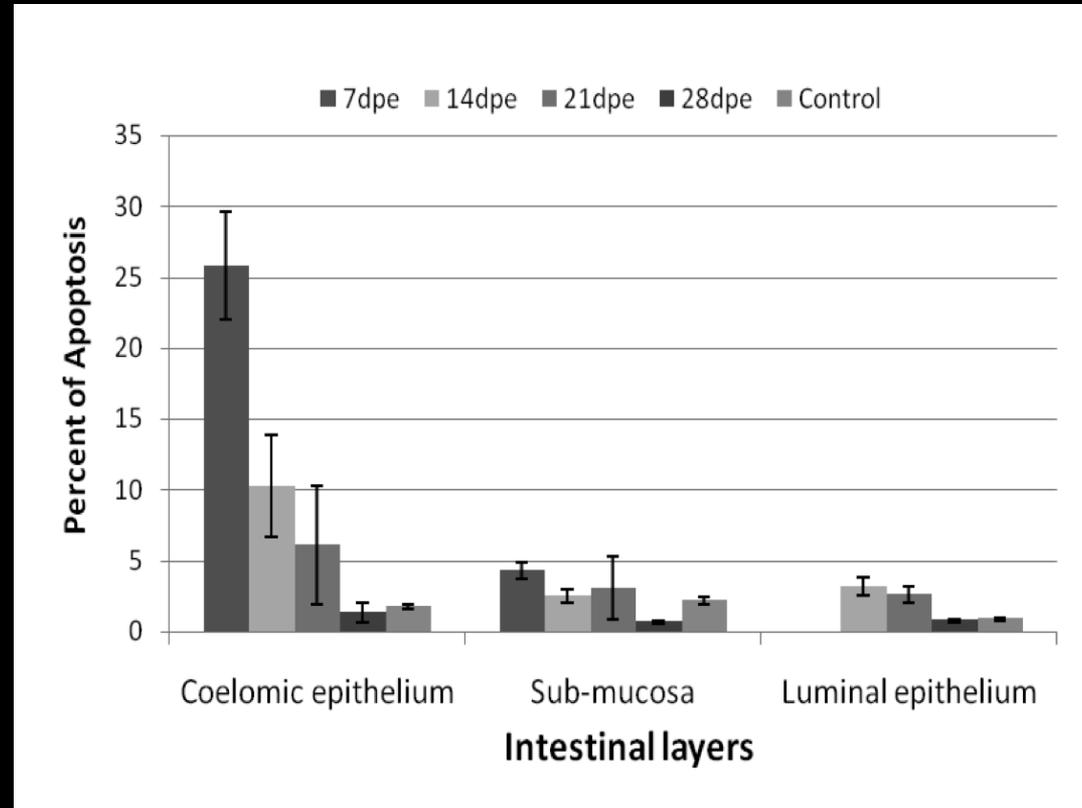
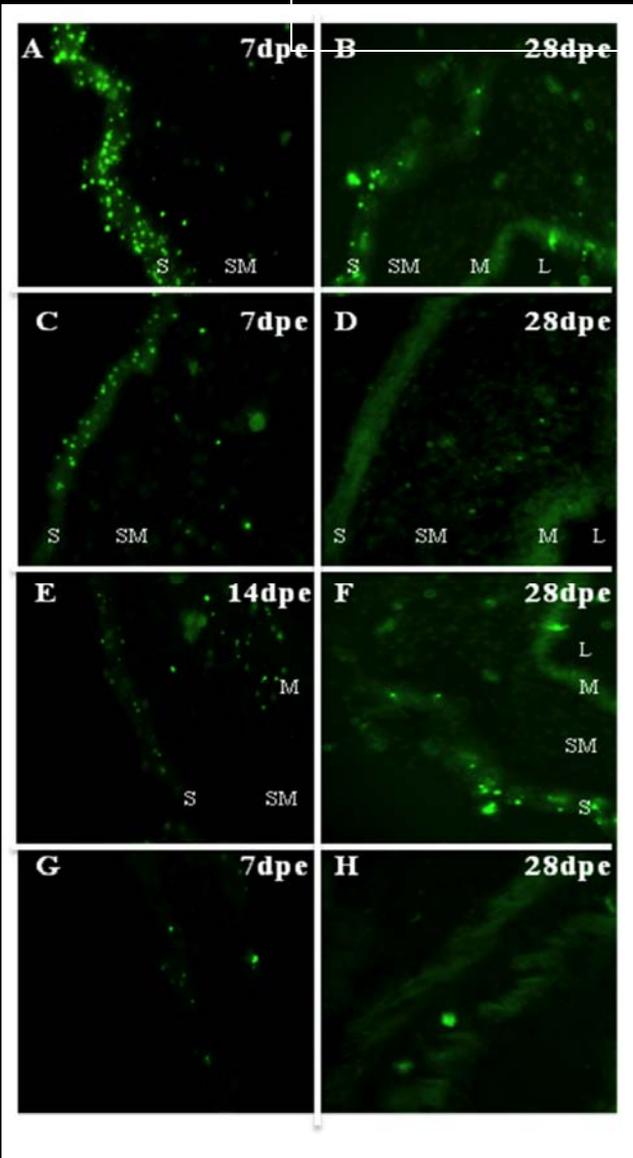


# Cell Proliferation

Percentage cell division in specimens sacrificed 24 hrs following BrdU injection



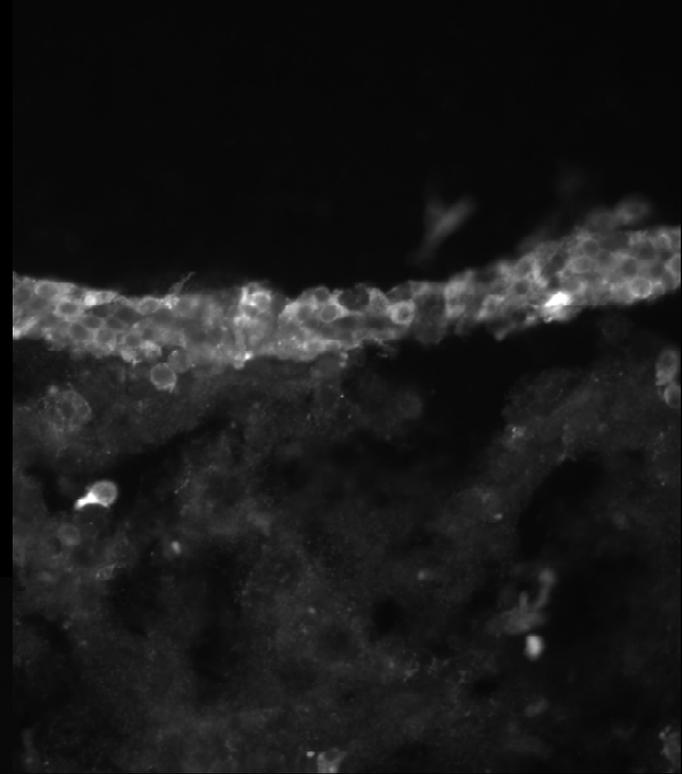
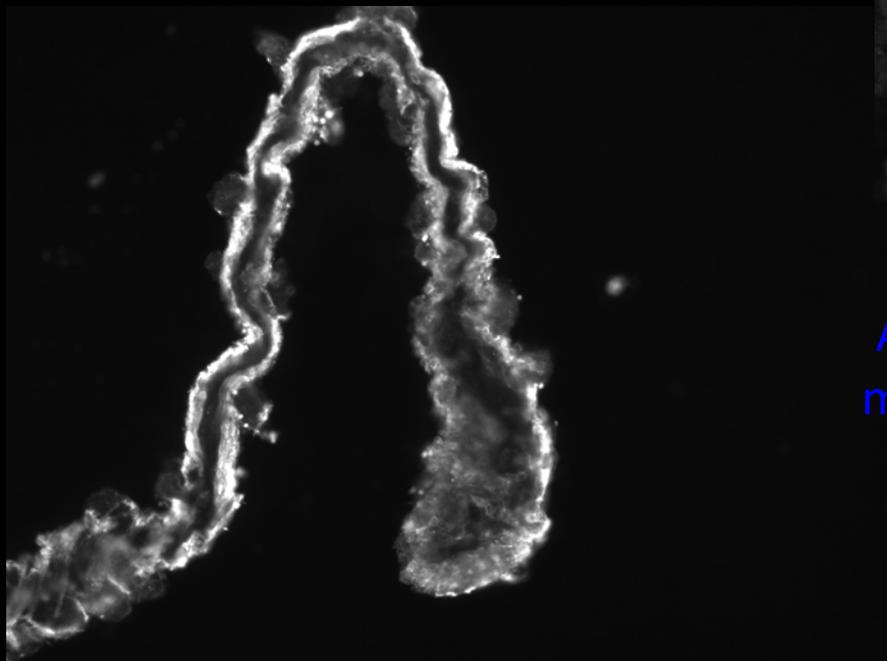
# Apoptosis



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



# Epithelial to mesenchymal transition



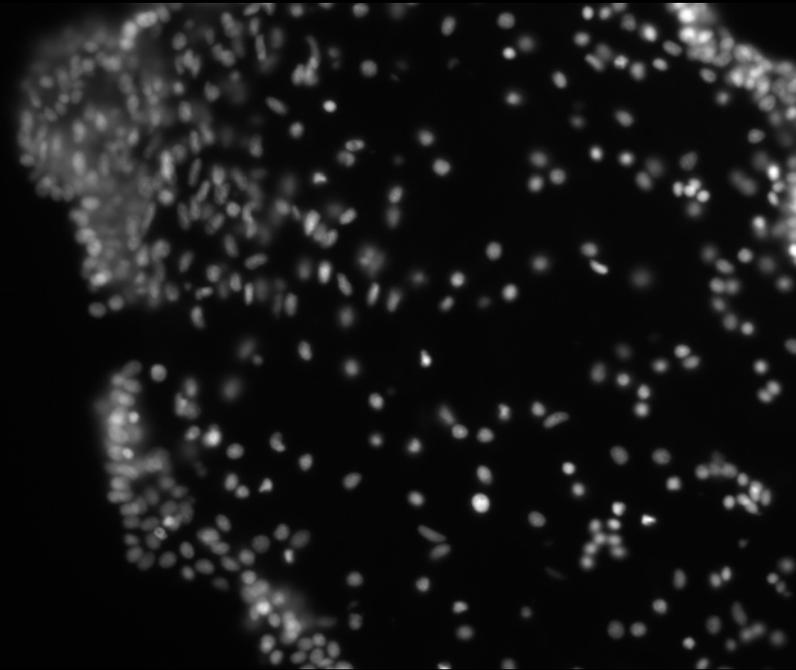
Antibody MES-1 recognizes the mesothelium and epithelial cells of the blastema.



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

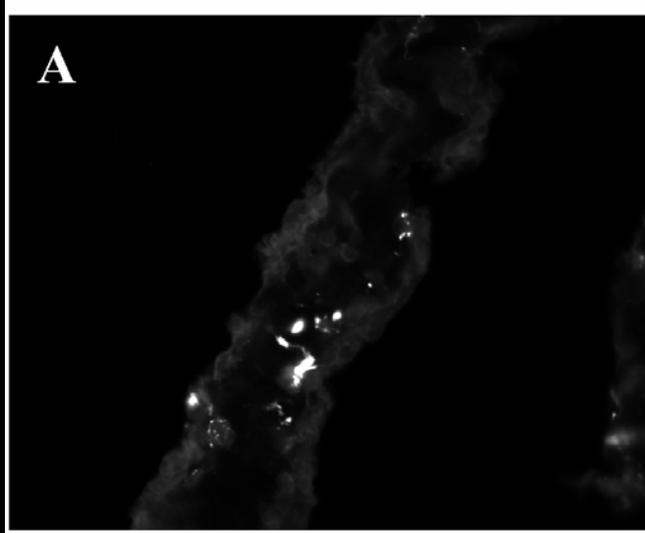


MES-1 labeling

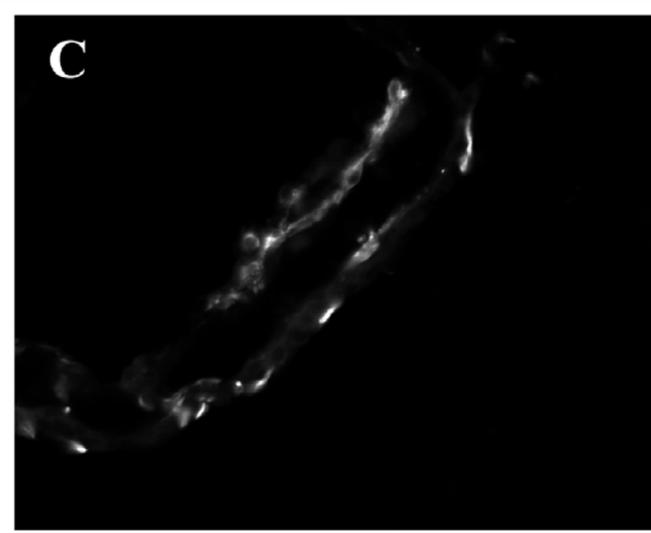


DAPI- nuclei staining

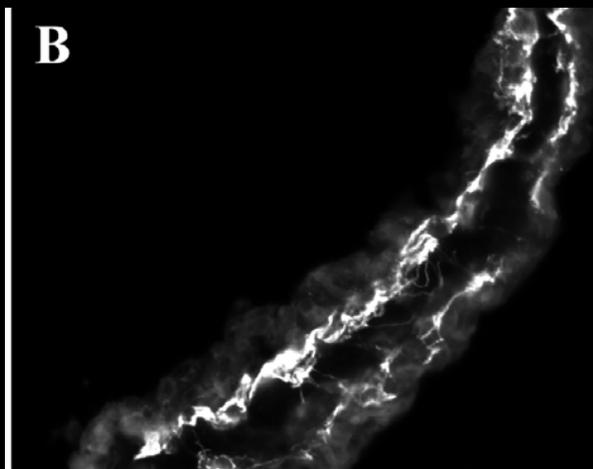
# Extracellular matrix (ECM) remodeling



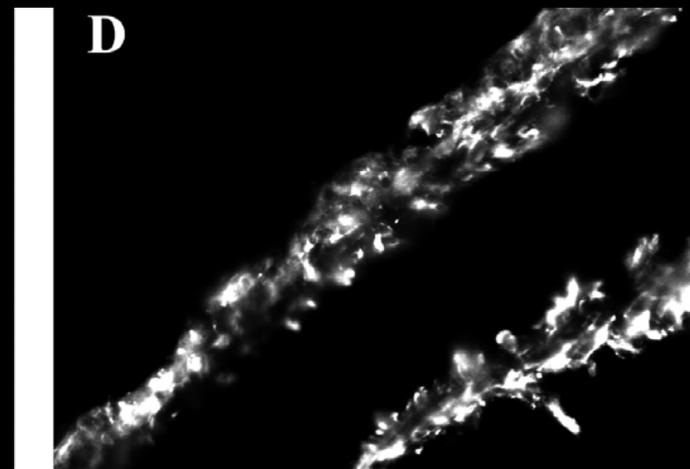
Collagen 7-d reg.



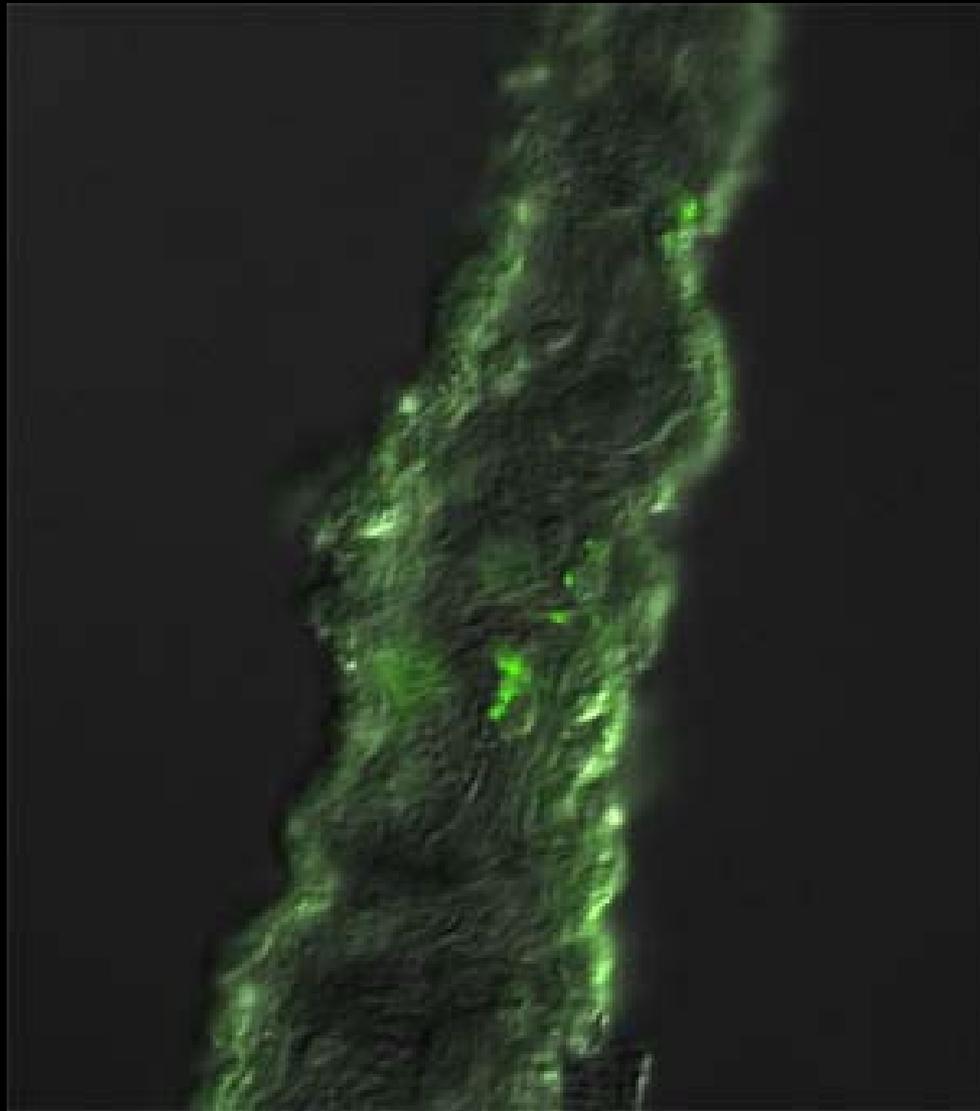
Muscle cells 7-d reg.



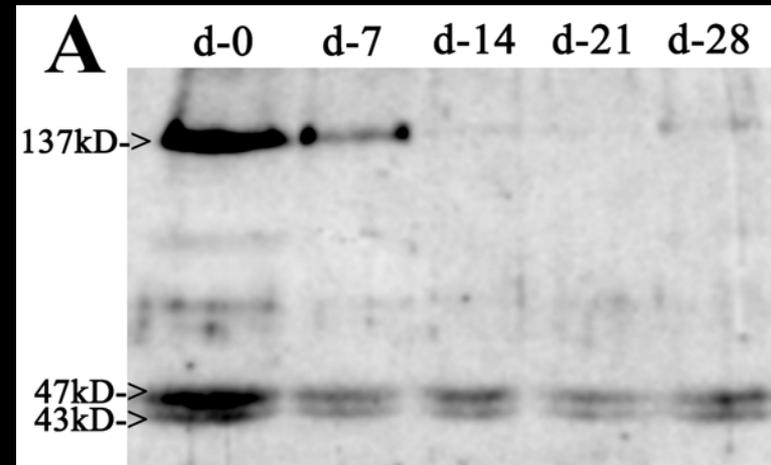
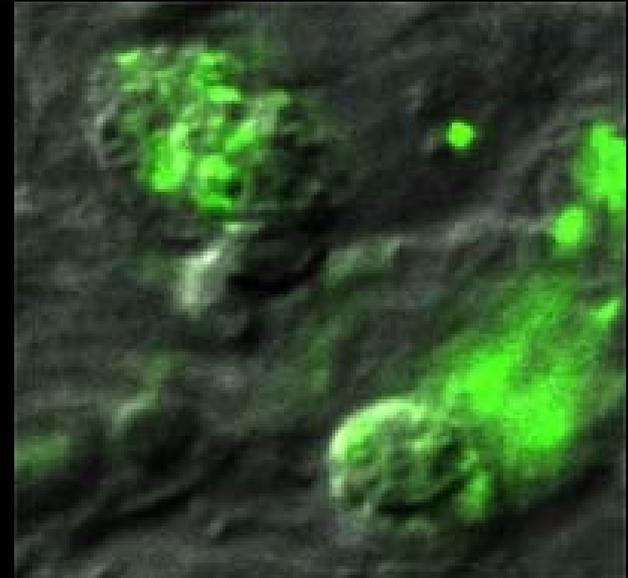
Collagen- Control



Muscle cells- Control

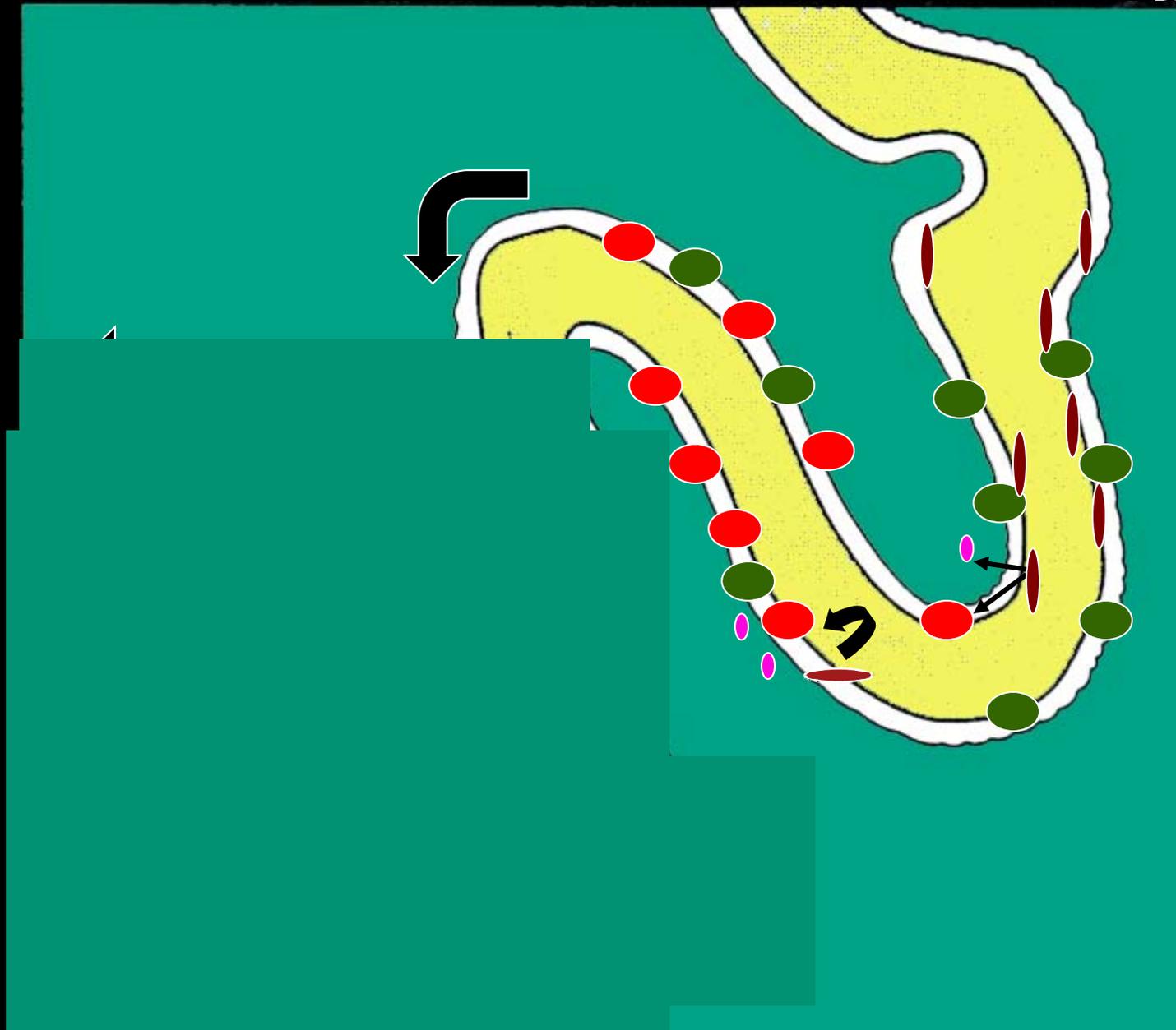


Collagen is degraded by  
phagocytic amoebocytes



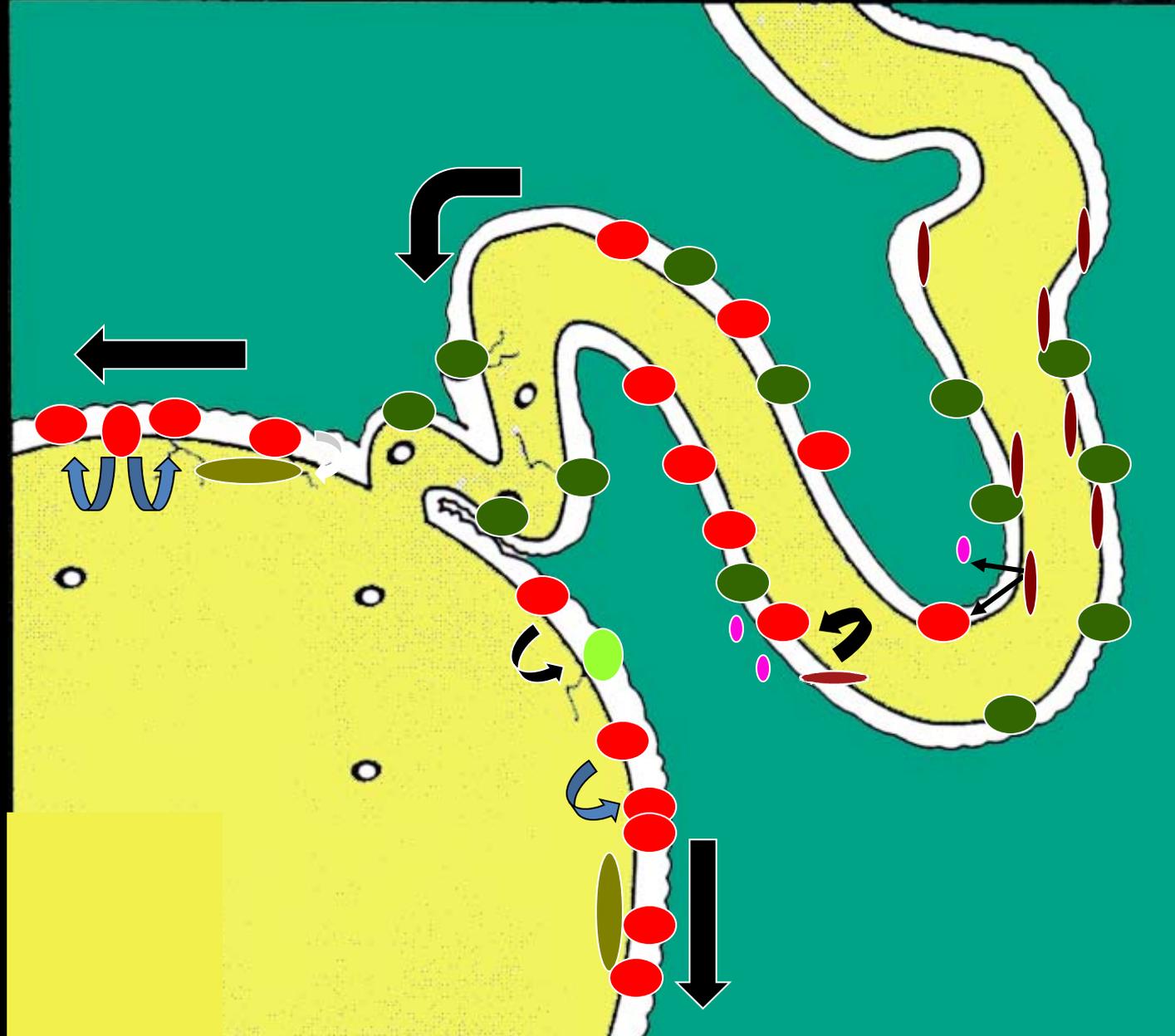


# REGENERATION MODEL- cellular origins



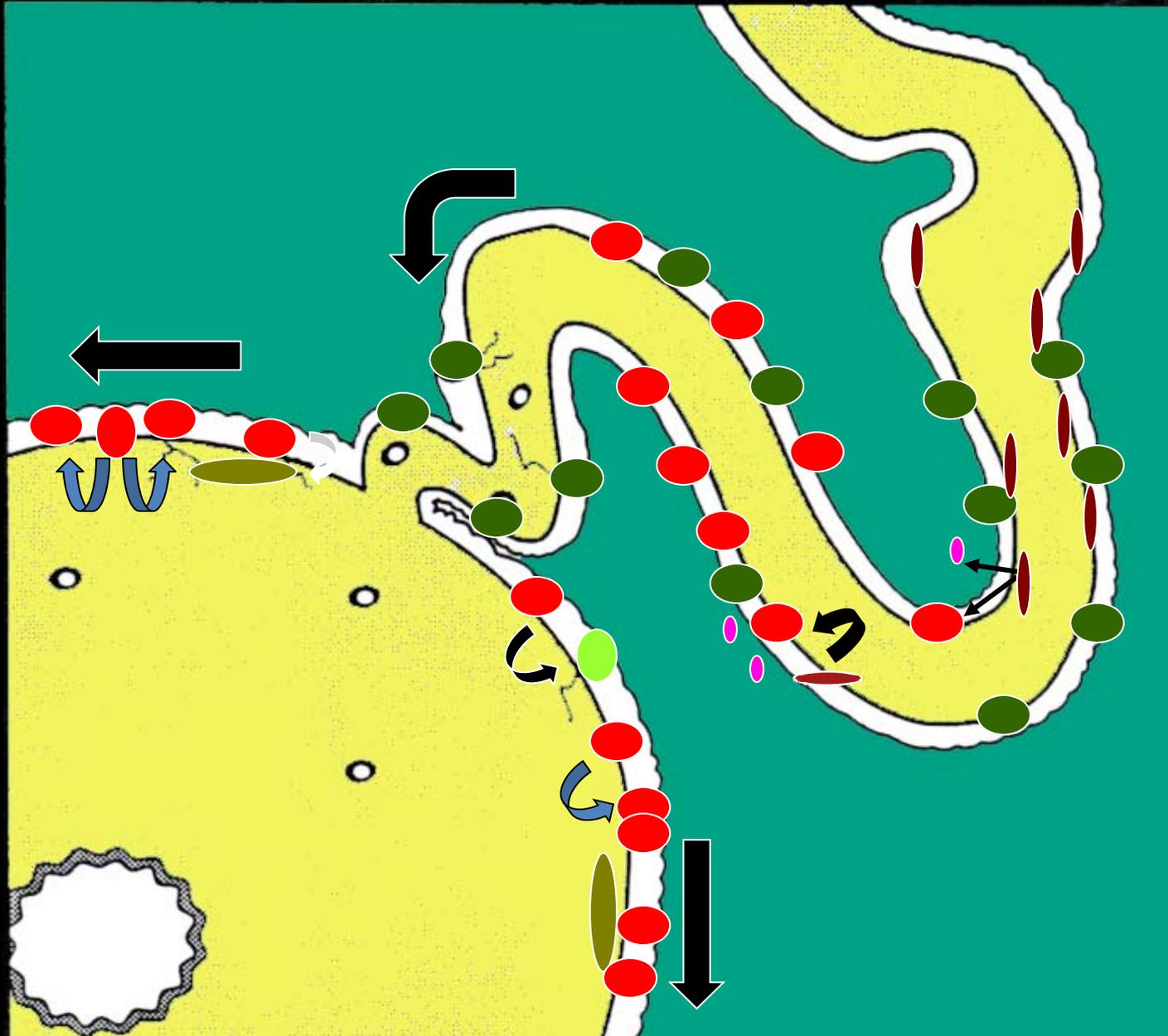


# REGENERATION MODEL- cellular origins

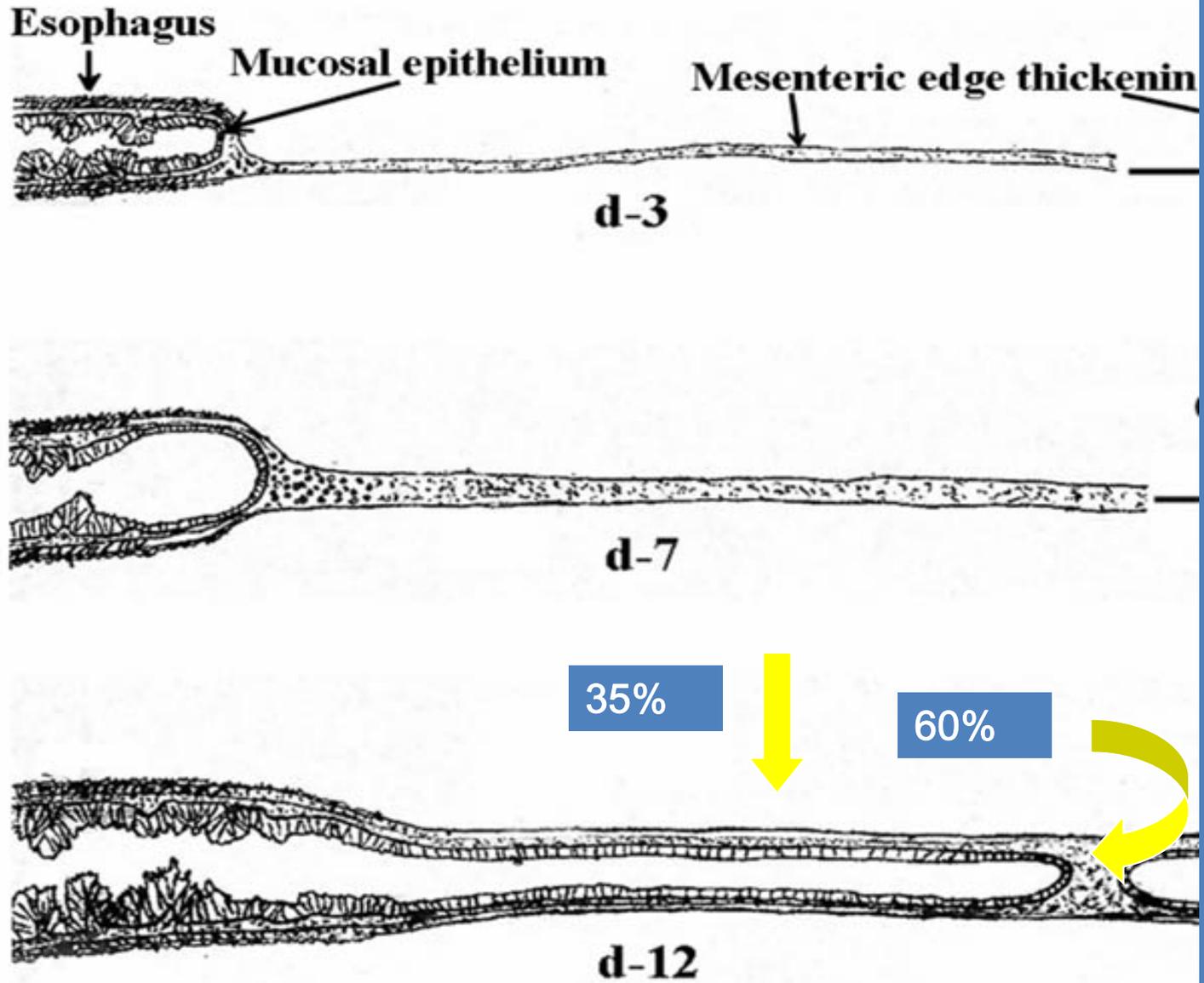




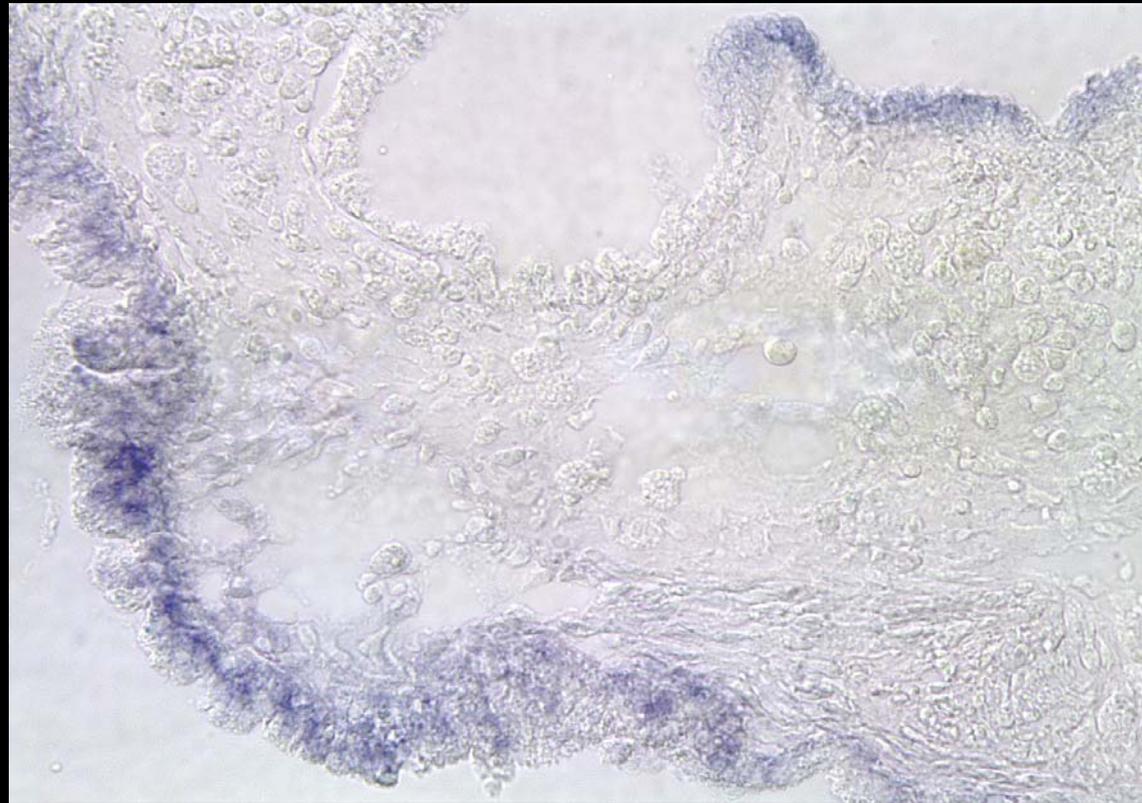
# REGENERATION MODEL- cellular origins



# Cell Migration



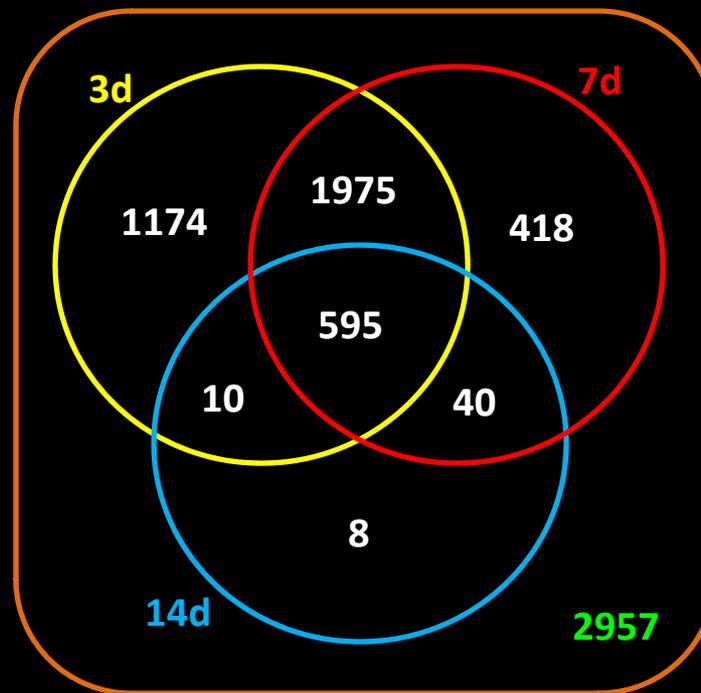
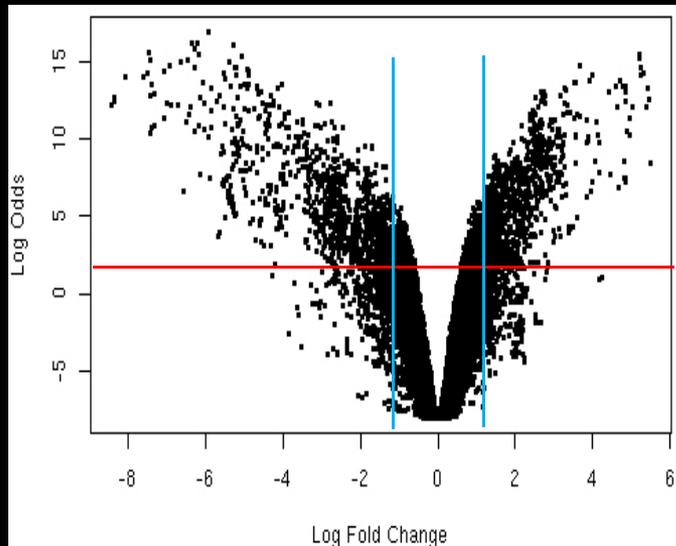
# Challenge #1 - Limited molecular tools



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

## Challenge #2 - Genomics

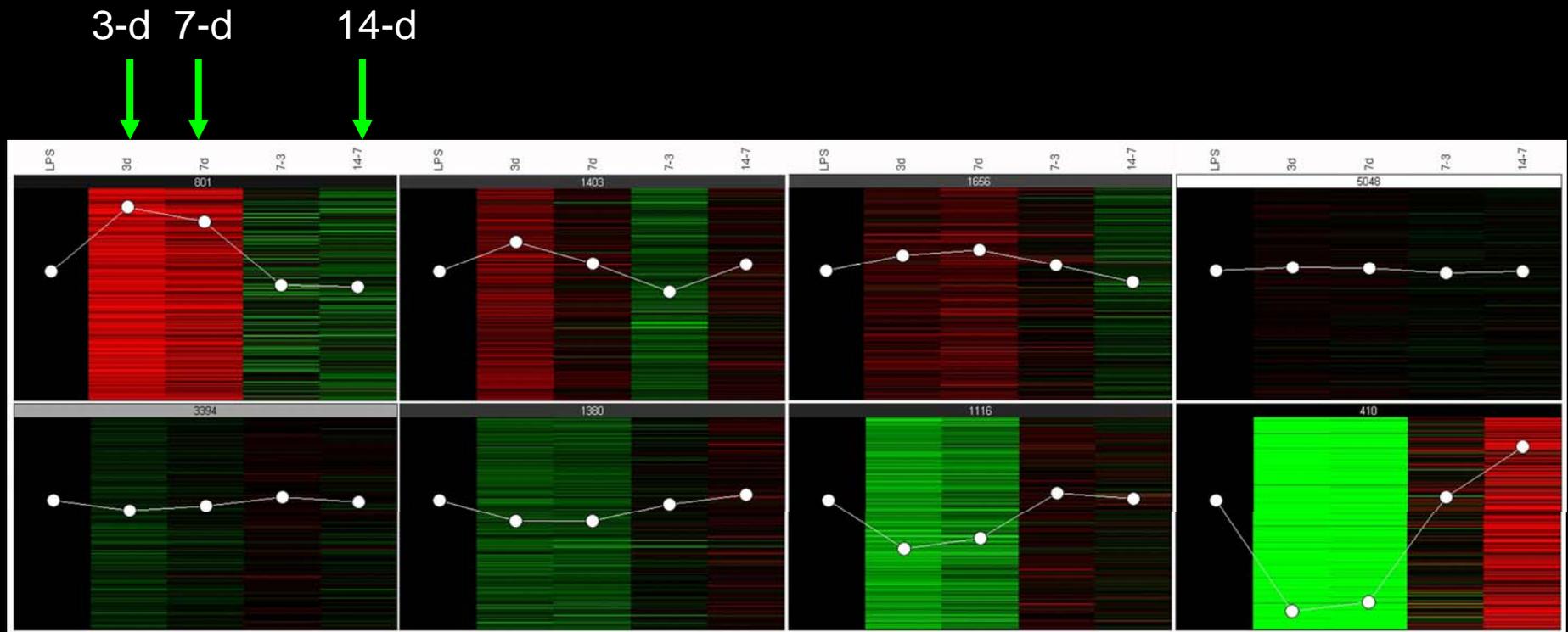
- We have a databank with over 7000 ESTs from 3 cDNA libraries of normal and regenerating intestines
  - Microarrays were done with custom made microchips with over 7000 *H. glaberrima* ESTs.
  - A large number of ESTs are differentially expressed in regenerating animals



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

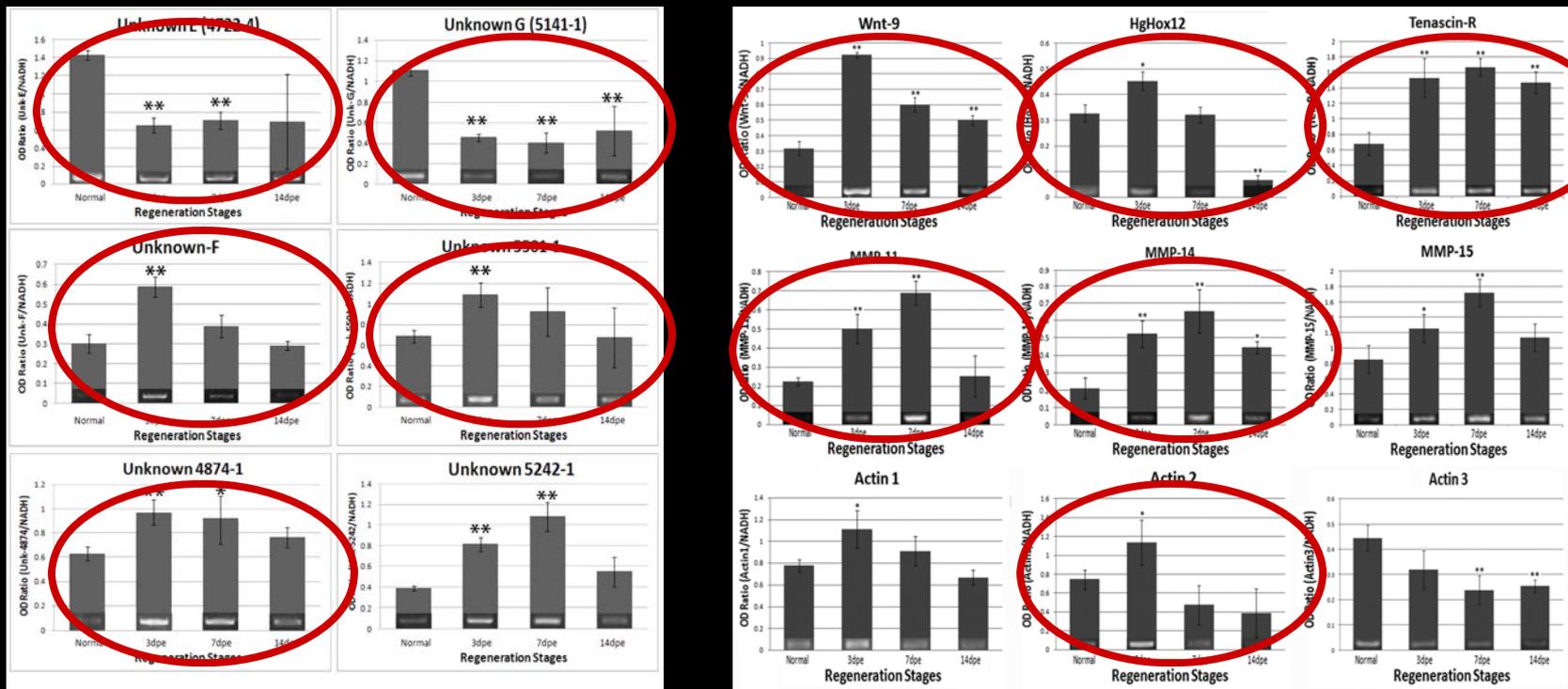
Reference = Normal (non-eviscerated)  
 $n = 7166$   $p < 0.01$

# Clusters of gene expression



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

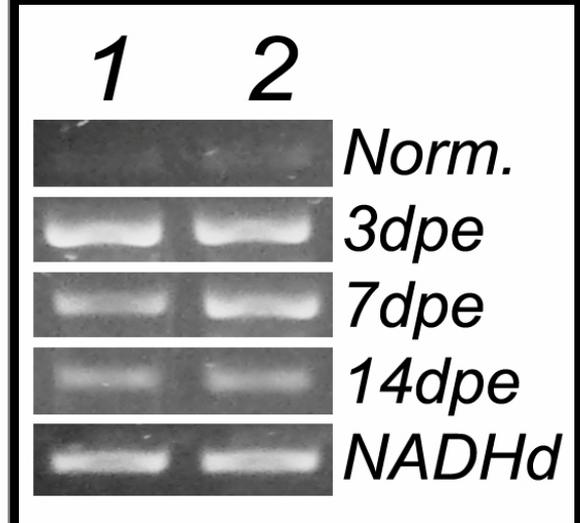
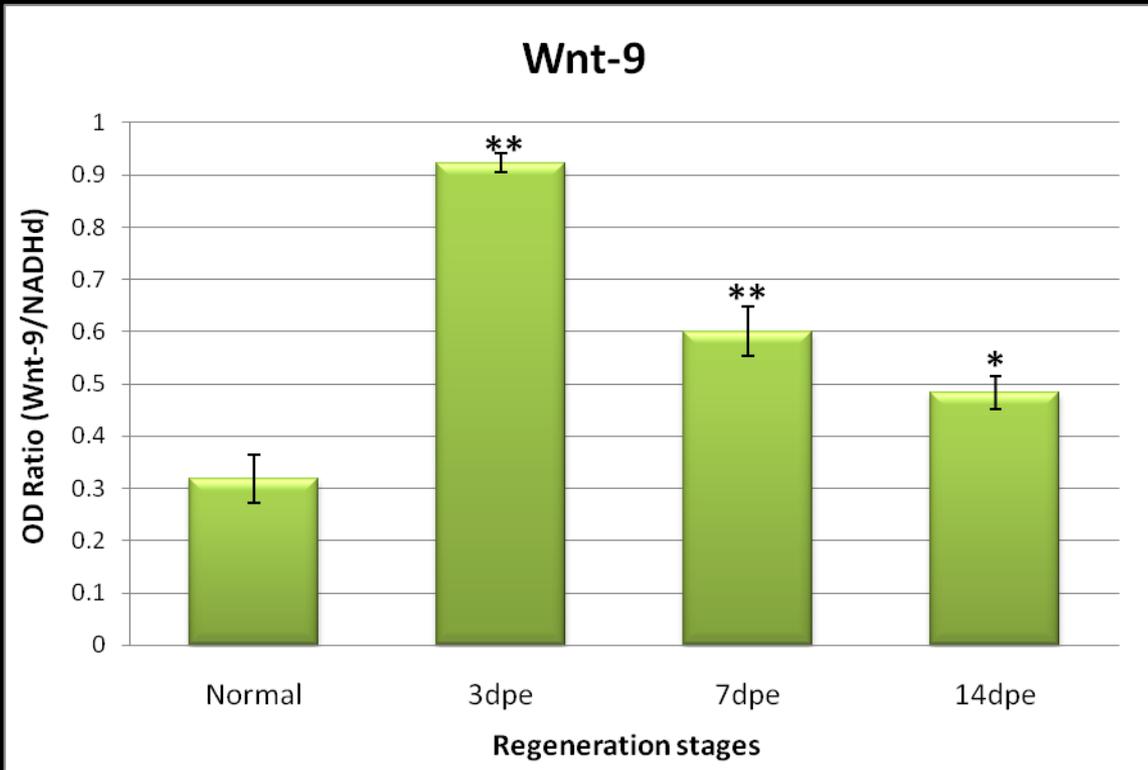
Studies can focus on selected candidates or novel genes.



# **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

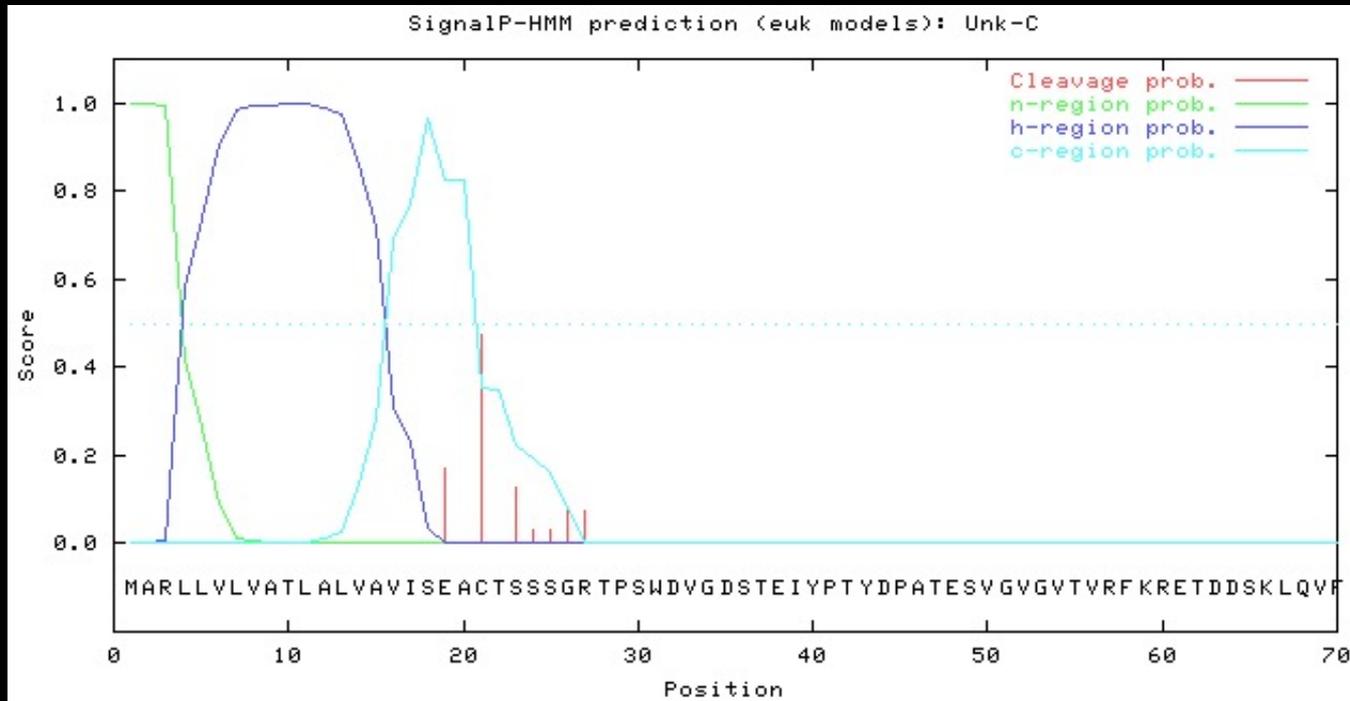


# **Orpin-** novel gene

- **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**

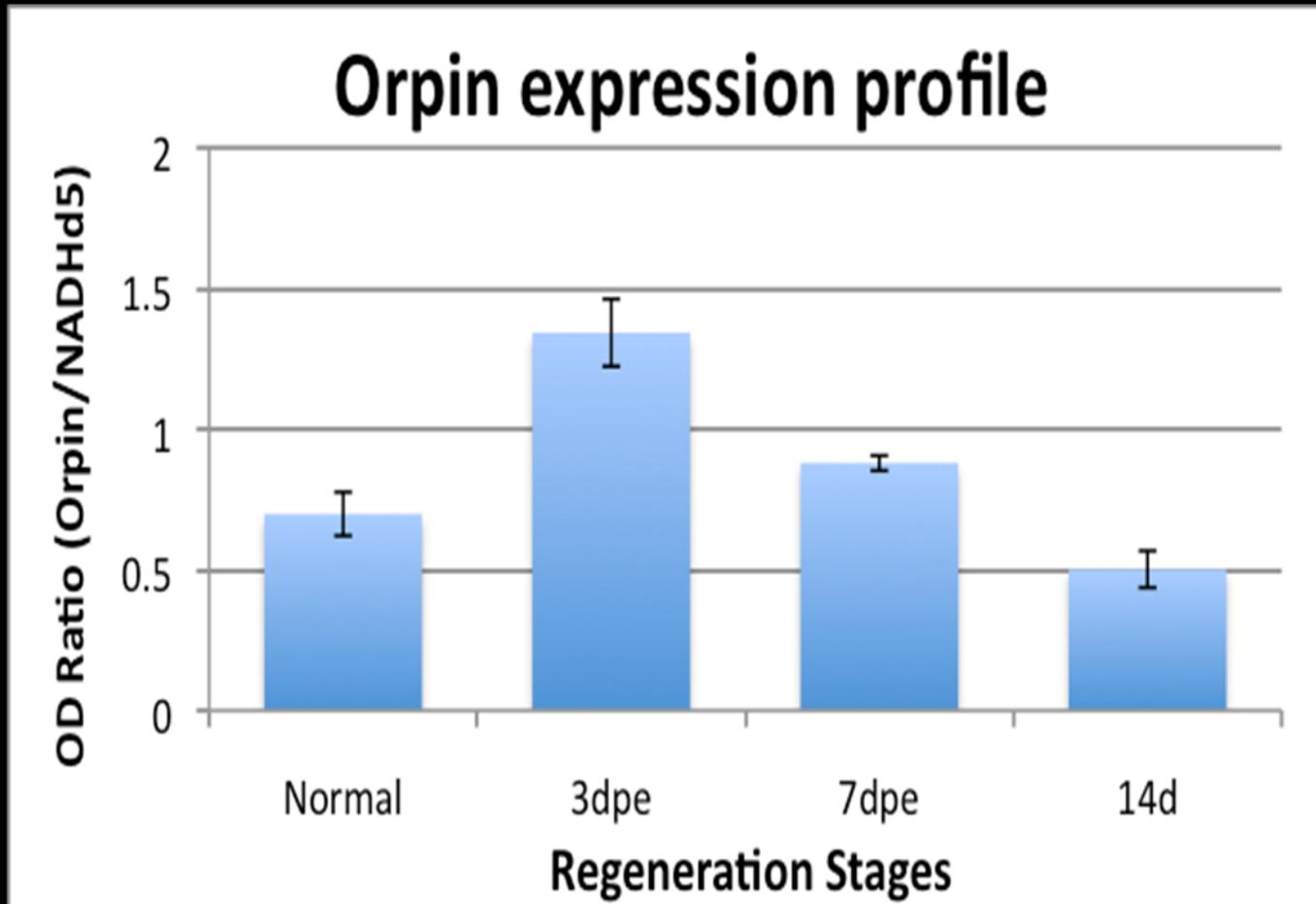
# Orpin

```
1  aaaactttctgcagaatcagttgttaagaattgtacacaggccgaaactagtcgttctaaca
62  tcgcaactcctccagaatcatccagaacaagcatttacatcaatcatgggctagacttctt
                                     M A R L L
122 gtactcgtcgctacacttgctttggctcgctgtcattagtgaaagcgtgcacctcatcatct
    V L V A T L A L V A V I S E A C T S S S
182 gggcgcacgccatcttgggacgttggcgactccactgaaatctaccccacttacgatcct
    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 gatgggtgatgagaagatcttgggaatgagttccagaaagtgtcagtcgcatagtagata
    D G D E K I S W N E F Q K V S V A *
482 gaaatagcgcggccagttccagttatTTAATATAGTCTATTTTAGAGACCTGTAGGGTAA
542 ggaacatttacttcaaagatgacagtccttccaaacctttcttcaaagctcaacgaact
602 gcacctatcatgtcaccttaagcttagagtaggatctatcgacatTTTgaaacacaatat
662 tggataacgctcttgaataaccgataataccggatgcatgggtatatatgcaagcagaa
722 aataaatacattgtctctatgtgagccattgtgaaaaaccgtgaaa
```

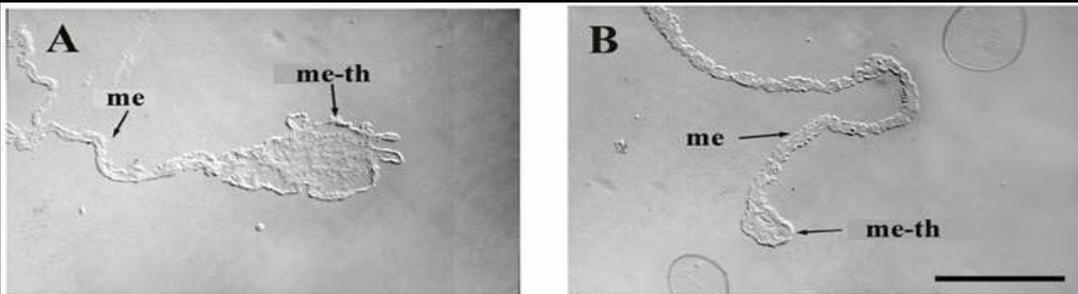




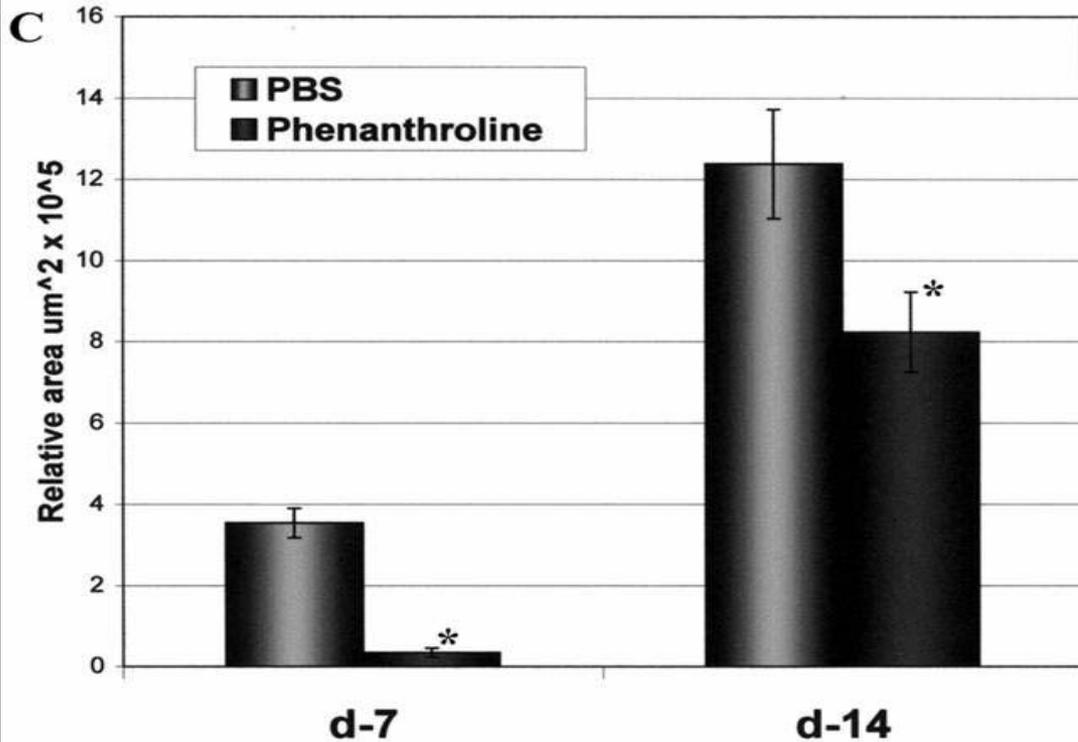
# RT-PCR validation of Orpin overexpression in the 3-day regenerating intestine.



# 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**