

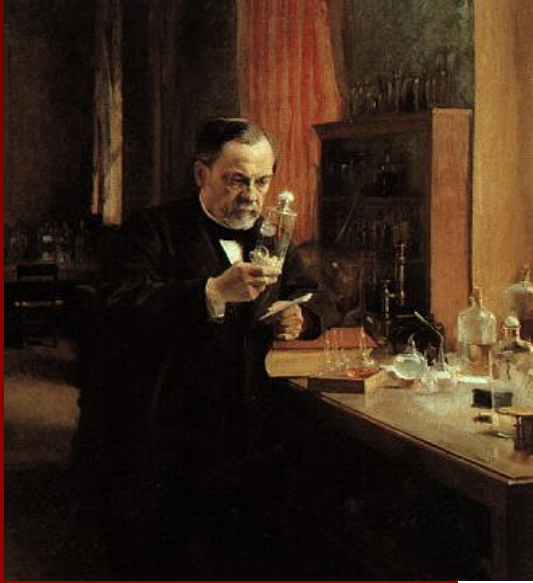
Mucosal Flora in IBD



Charité

Alexander Swidsinski

Supported by Broad Medical Research Program





Multicellular bacteria forming stromatolith
in Australian salt lakes

FISH Analysis of mucosal Flora



Eub338
Alf1b
Beta42a
Gam42a
Ebac
Ec1531
Y16s-69
Srb385
Sgd
Hpy-1
Arc1430
HGC
LGC
Sfb
Erec
Lach
Ehal
Chis150
Clit135
Lab158
Stre493
Enc131
Efaec
Ato291
Cor653
Ecy1
Phasco
Veil
Rbro, Rfla
UroA, UroB
Ser1410
Bif164
CF319a
Bac303
Bfra602
Bdis656
Fprau
Dss658
Arch915

using r-RNA probes

Analysis of mucosal biofilms using Fluorescence In-Situ Hybridization (FISH)

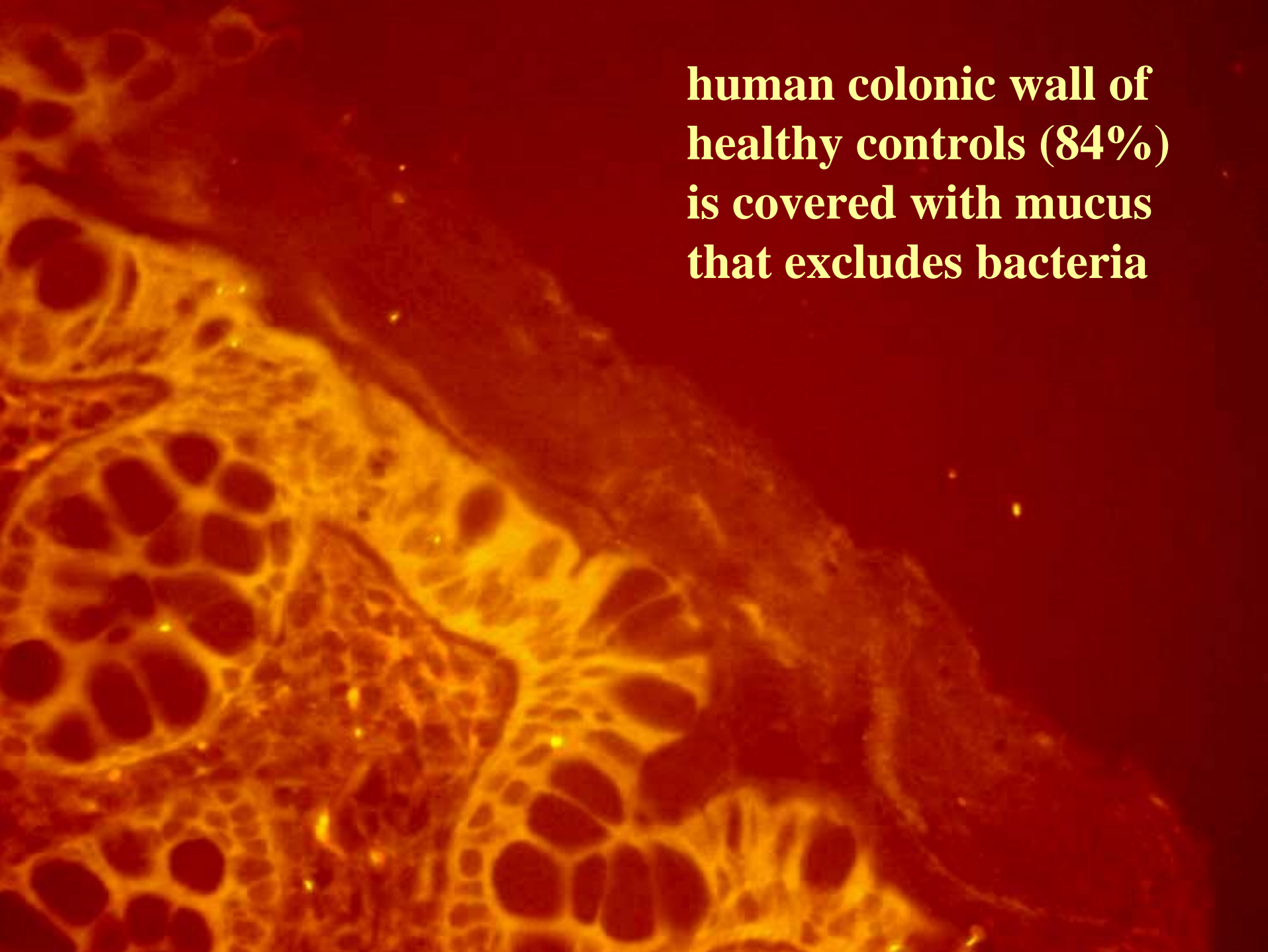
Adherent biofilms are the most prominent feature of IBD, which was not recognized due to lack of appropriate methods. Biofilms disappear after fixation in formalin – a main fixative in clinical pathology

The same patient and the same location fixed either with

Carnoy or Formalin

← →
Crohn's Disease, DAPI stain

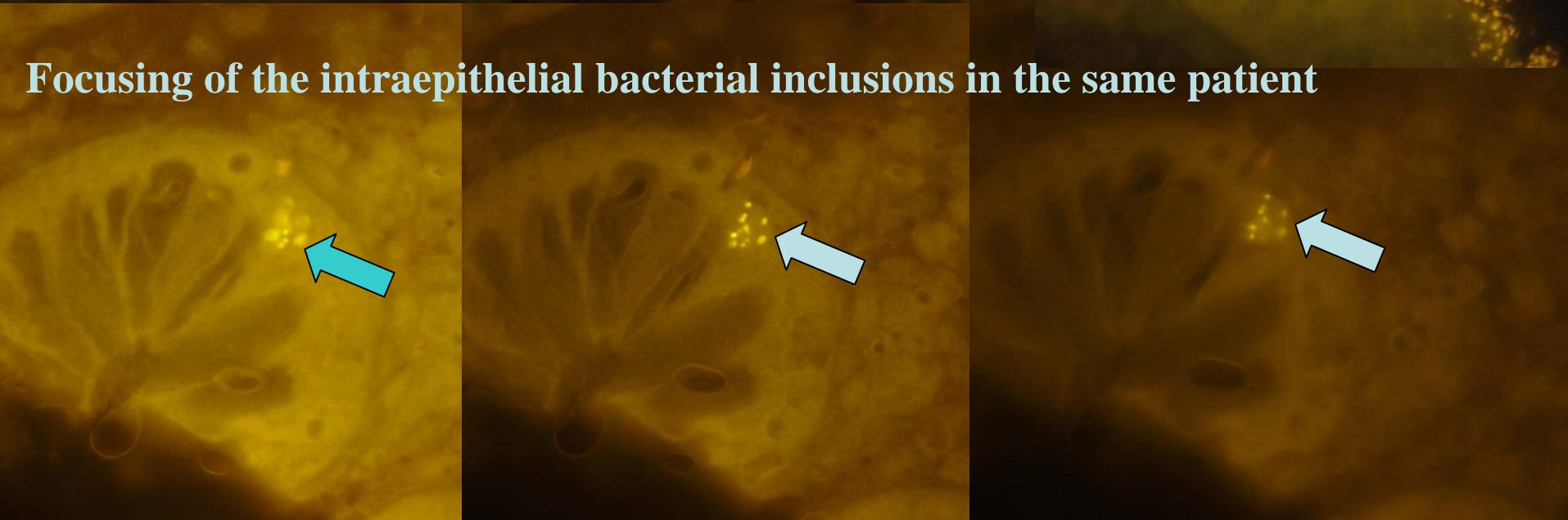
**human colonic wall of
healthy controls (84%)
is covered with mucus
that excludes bacteria**

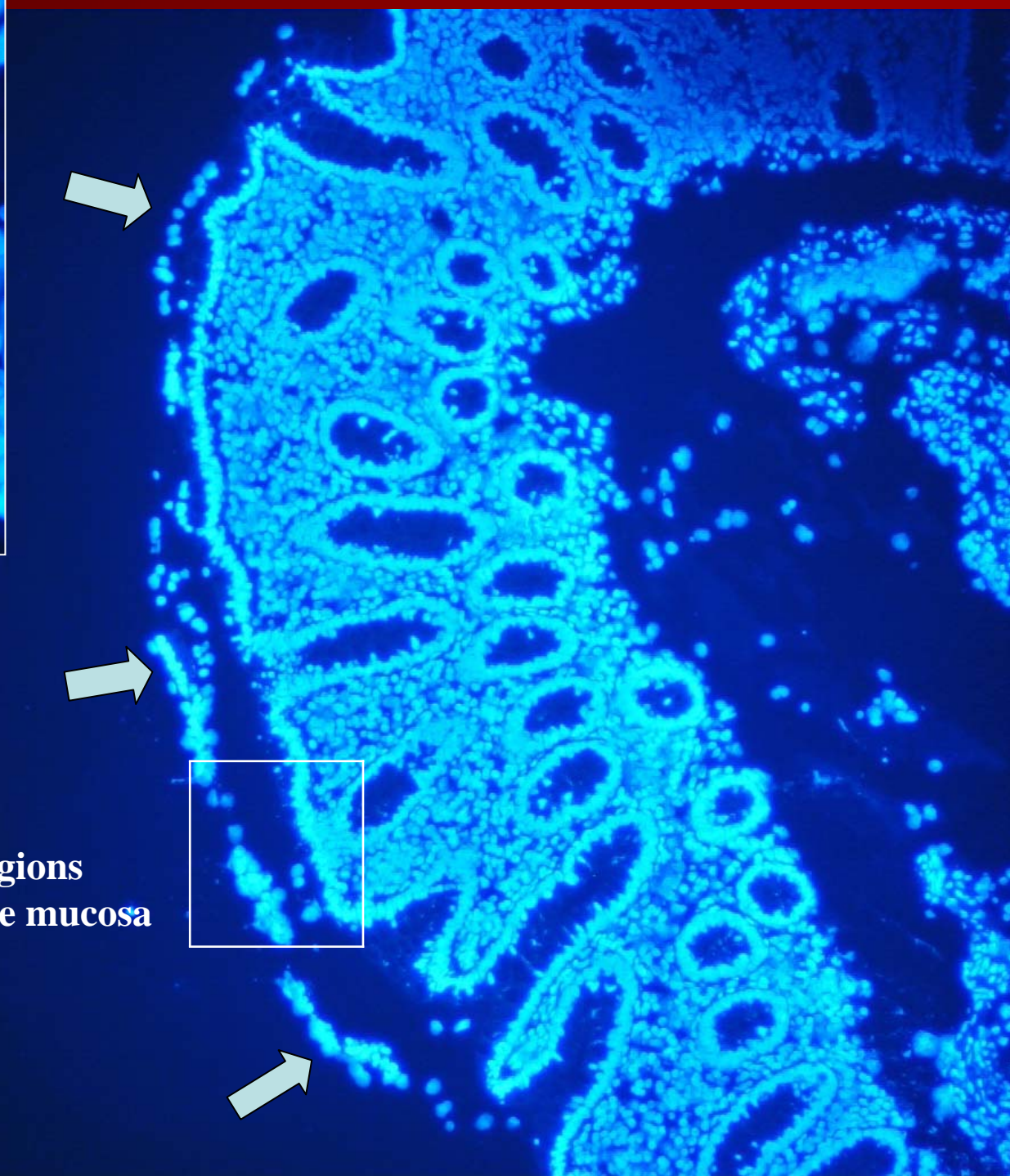
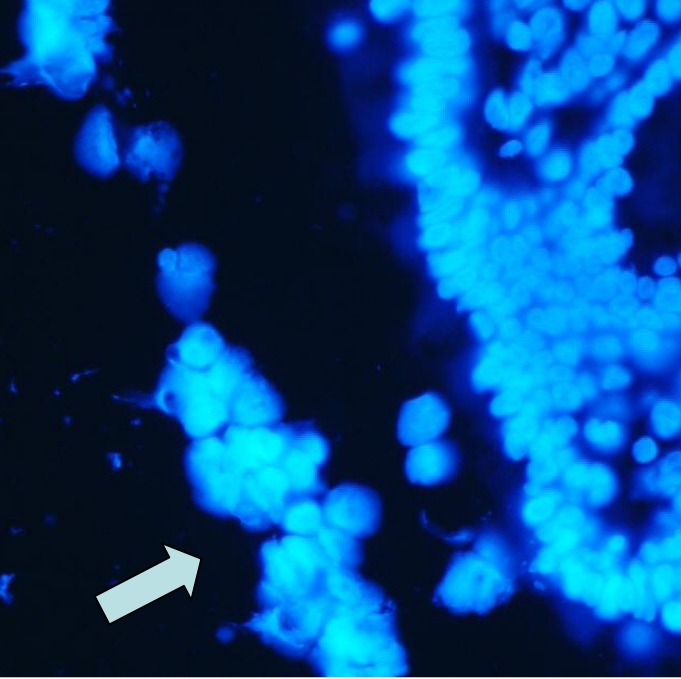


Prolific *Bacteroides fragilis* biofilm completely covers the mucosal surface and enters crypts in a CD patient



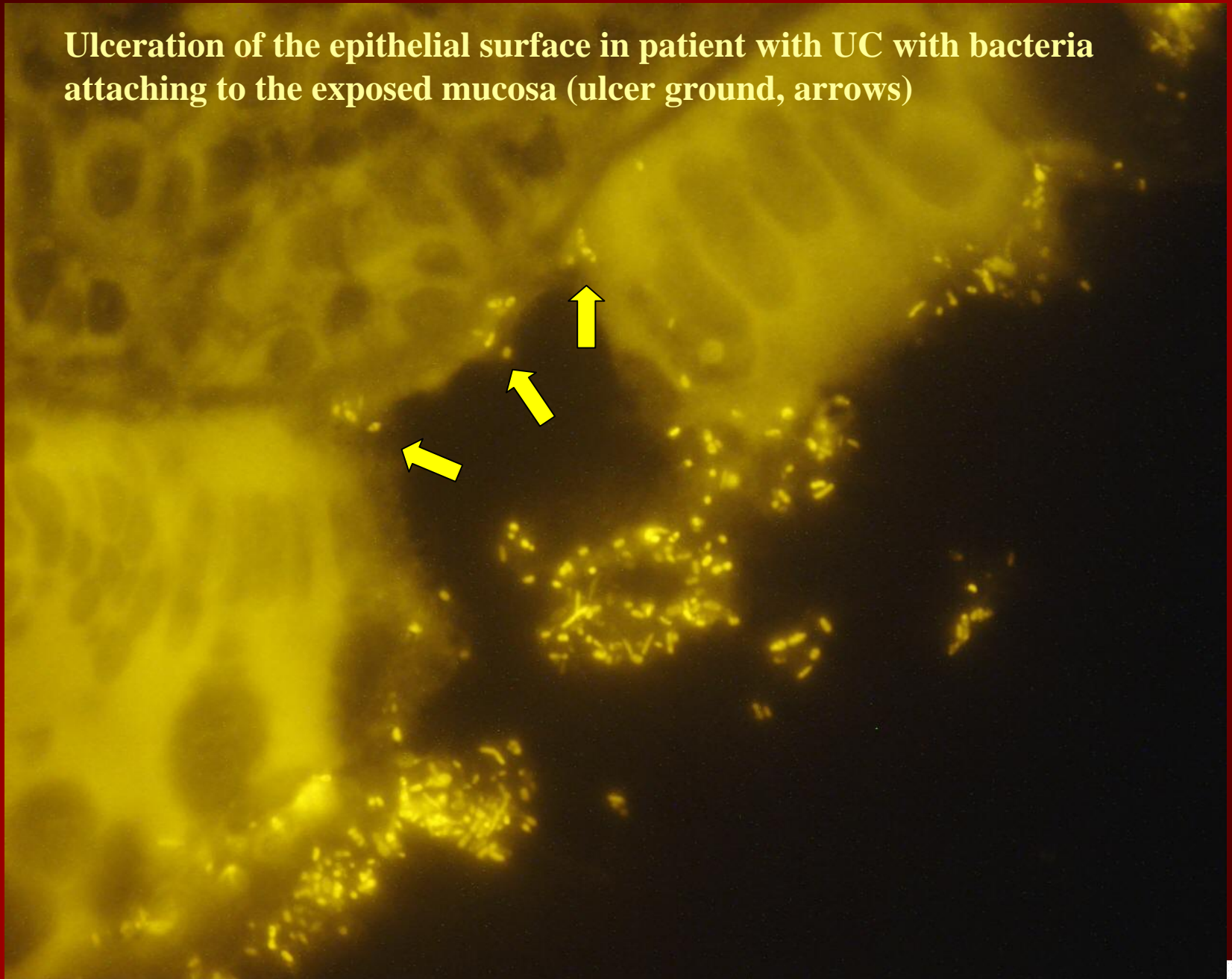
Focusing of the intraepithelial bacterial inclusions in the same patient

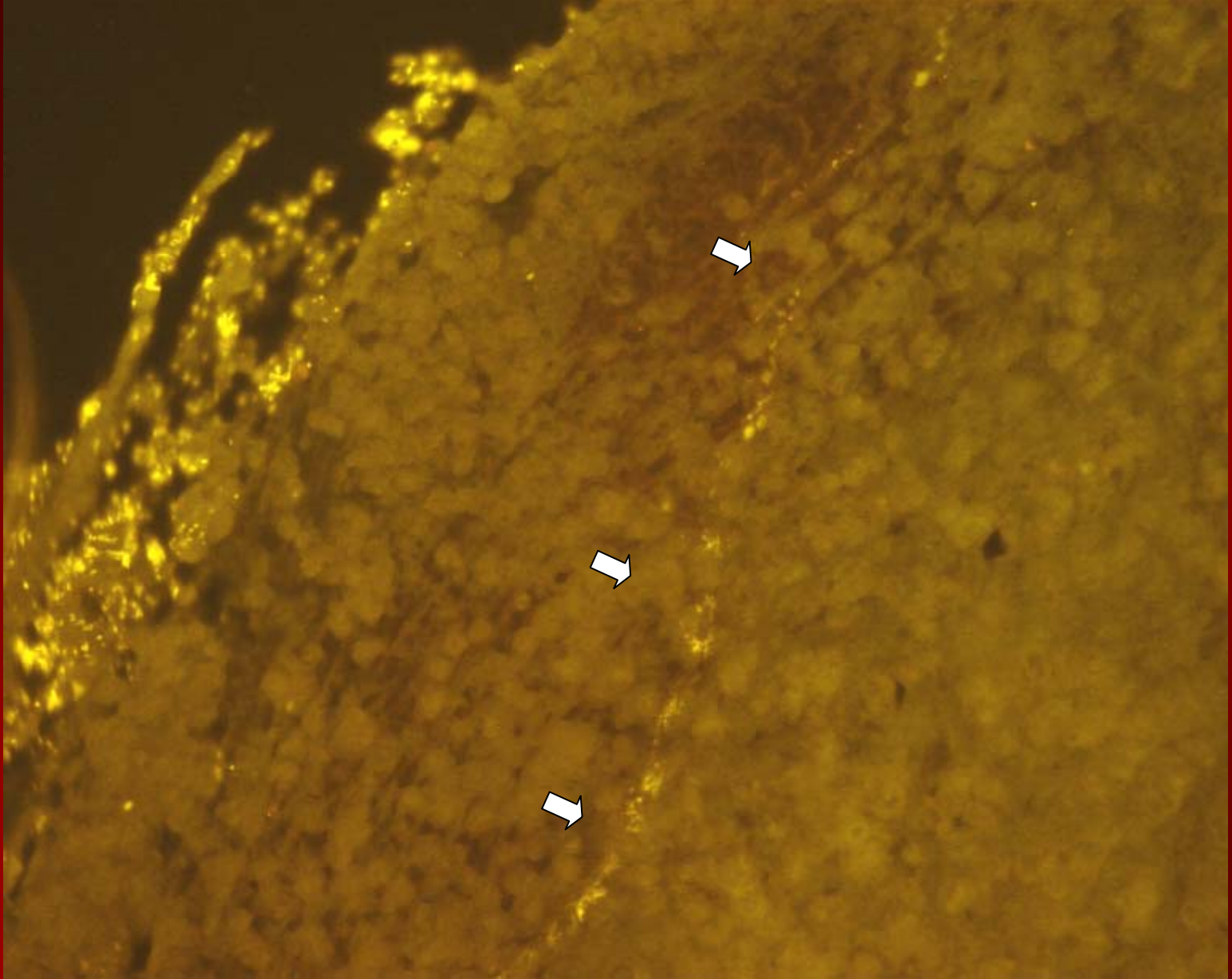




**Leukocytes migrate in
mucus, array in outer regions
and prevent access to the mucosa**

Ulceration of the epithelial surface in patient with UC with bacteria attaching to the exposed mucosa (ulcer ground, arrows)



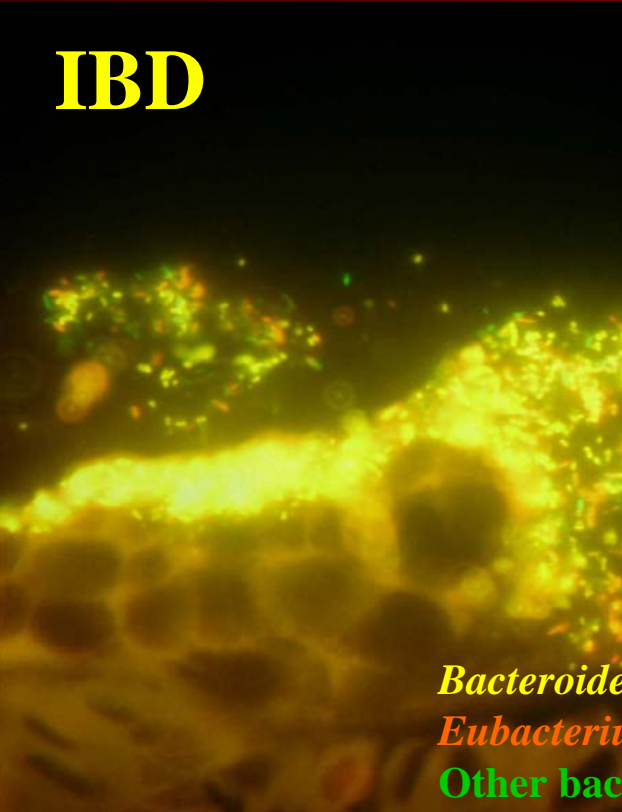


***Bacteroides* infiltration of the intestinal wall, CD**

**10 bacteria within a quadrant of this size
correspond to concentrations of 10^9 /ml**



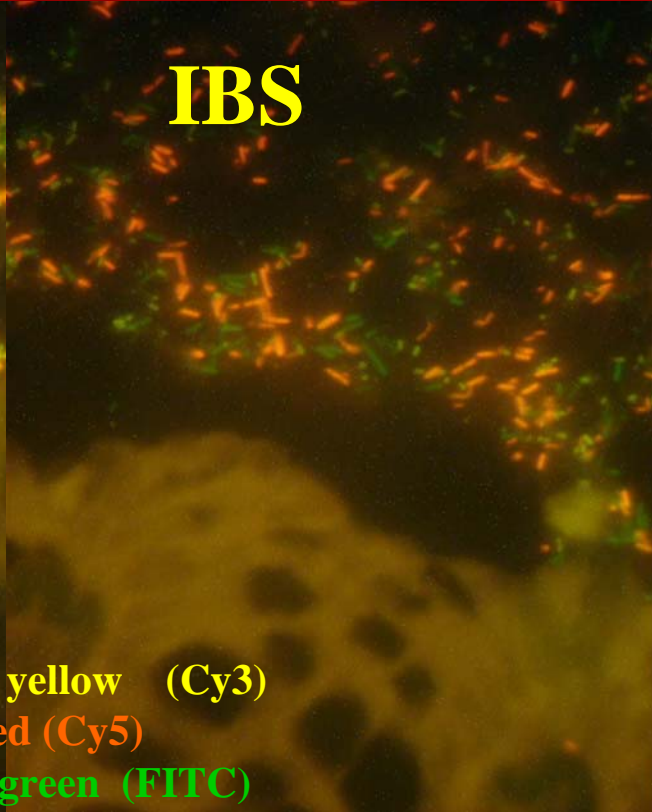
IBD



SI - colitis



IBS



Bacteroides fragilis (Bfra Probe)

Eubacterium rectale group (Erec Probe)

Other bacteria (Eub338)

yellow (Cy3)

red (Cy5)

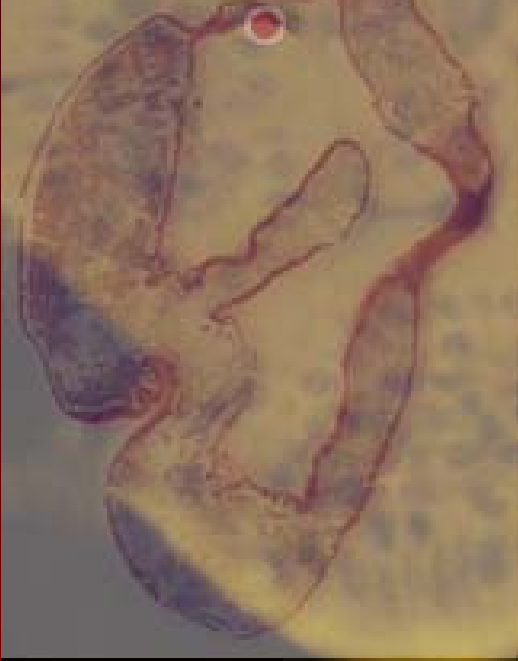
green (FITC)

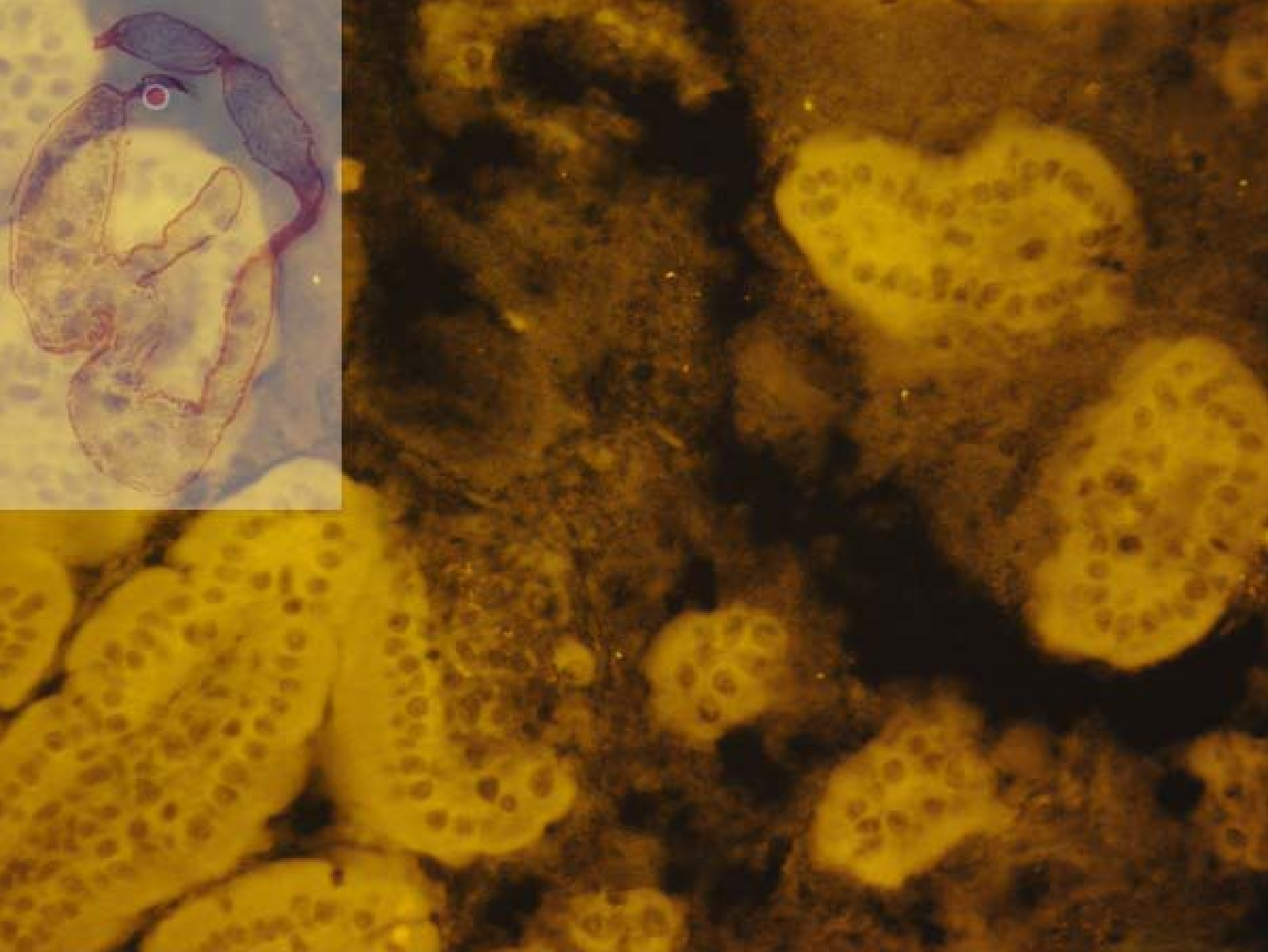
Percent of patients with 10^9 bacteria/ml

Percent of bacteria within biofilm

| | CD | UC | SIc | IBS | Contr. |
|-------------|-----|-----|-----|-----|--------|
| | 98% | 94% | 78% | 38% | 16% |
| Bfra | 60% | 30% | 31% | 14% | 16% |
| Erec | 10% | 5% | 18% | 48% | 32% |

**The number of bacteria in small intestine
of a healthy wild type mouse
is low**







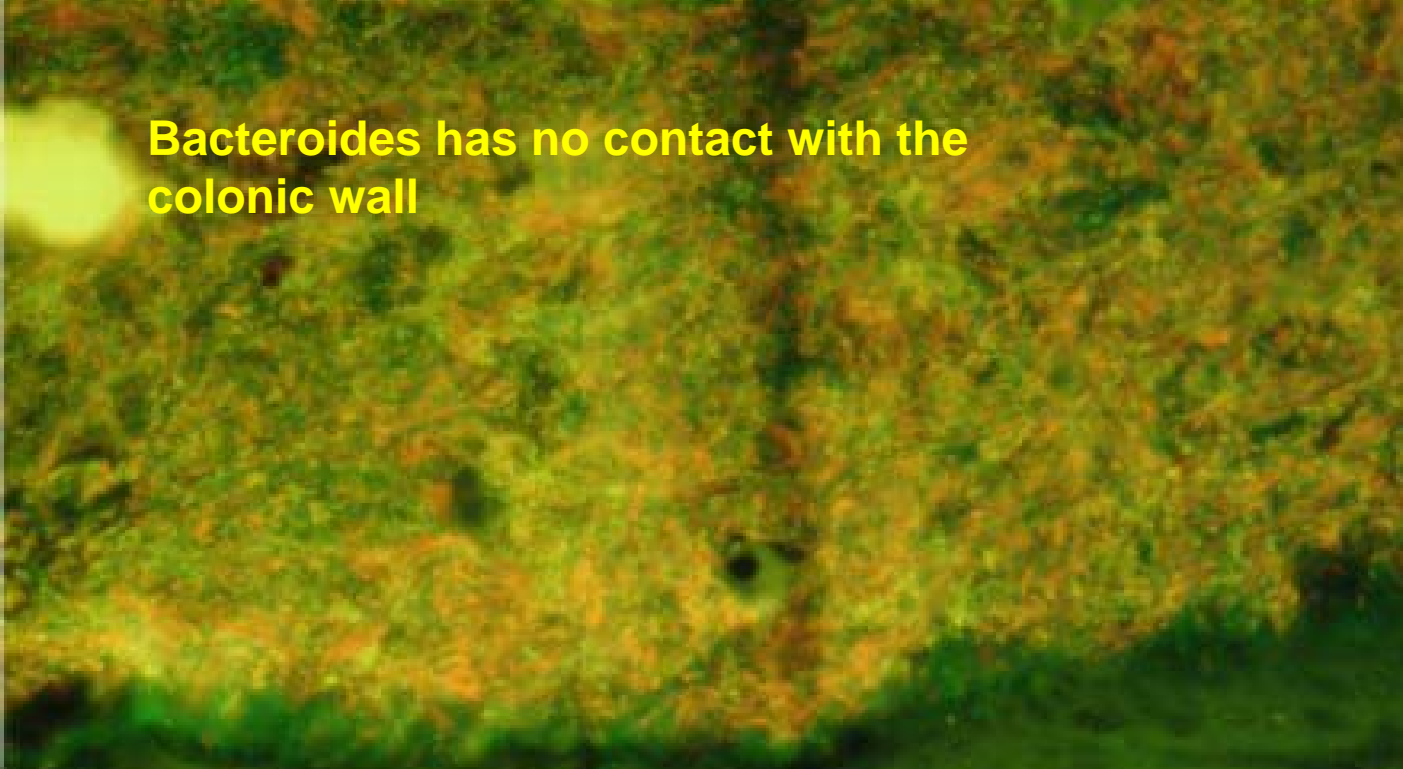
The bacterial concentrations
In the cecum of large intestine are
extremely high

Bacteria (with an exception of *Bacteroides*)
are contacting the colonic wall and entering
crypts.



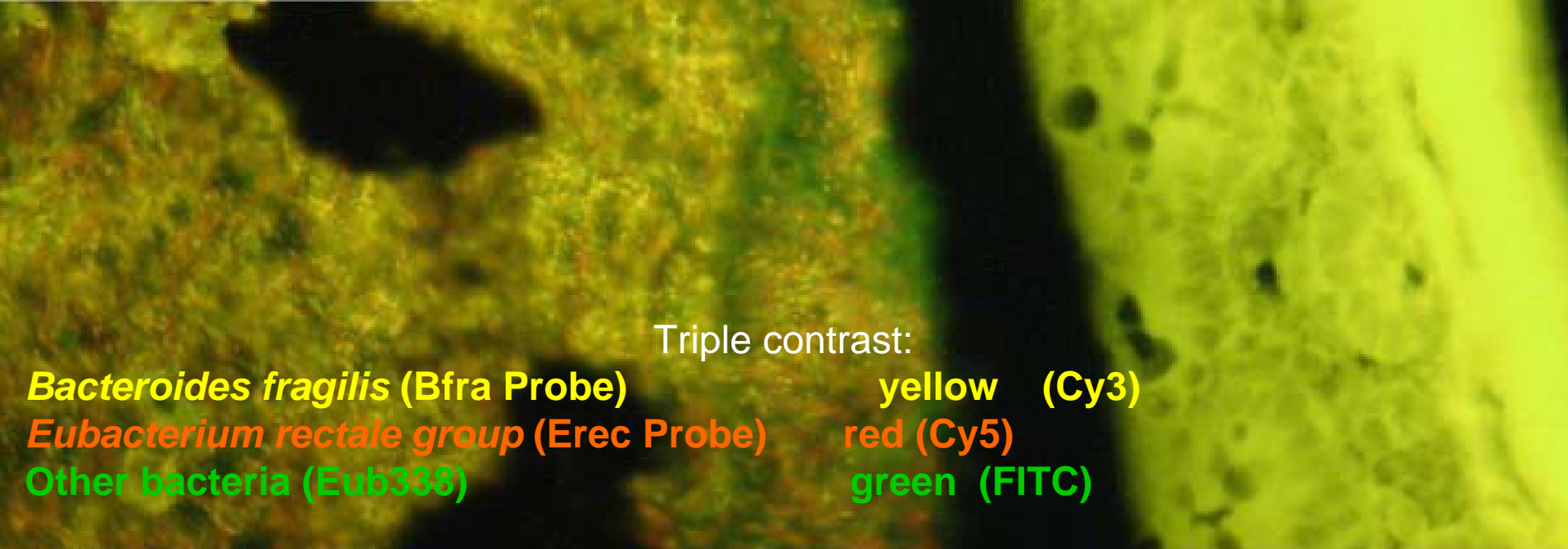


Bacteroides has no contact with the colonic wall



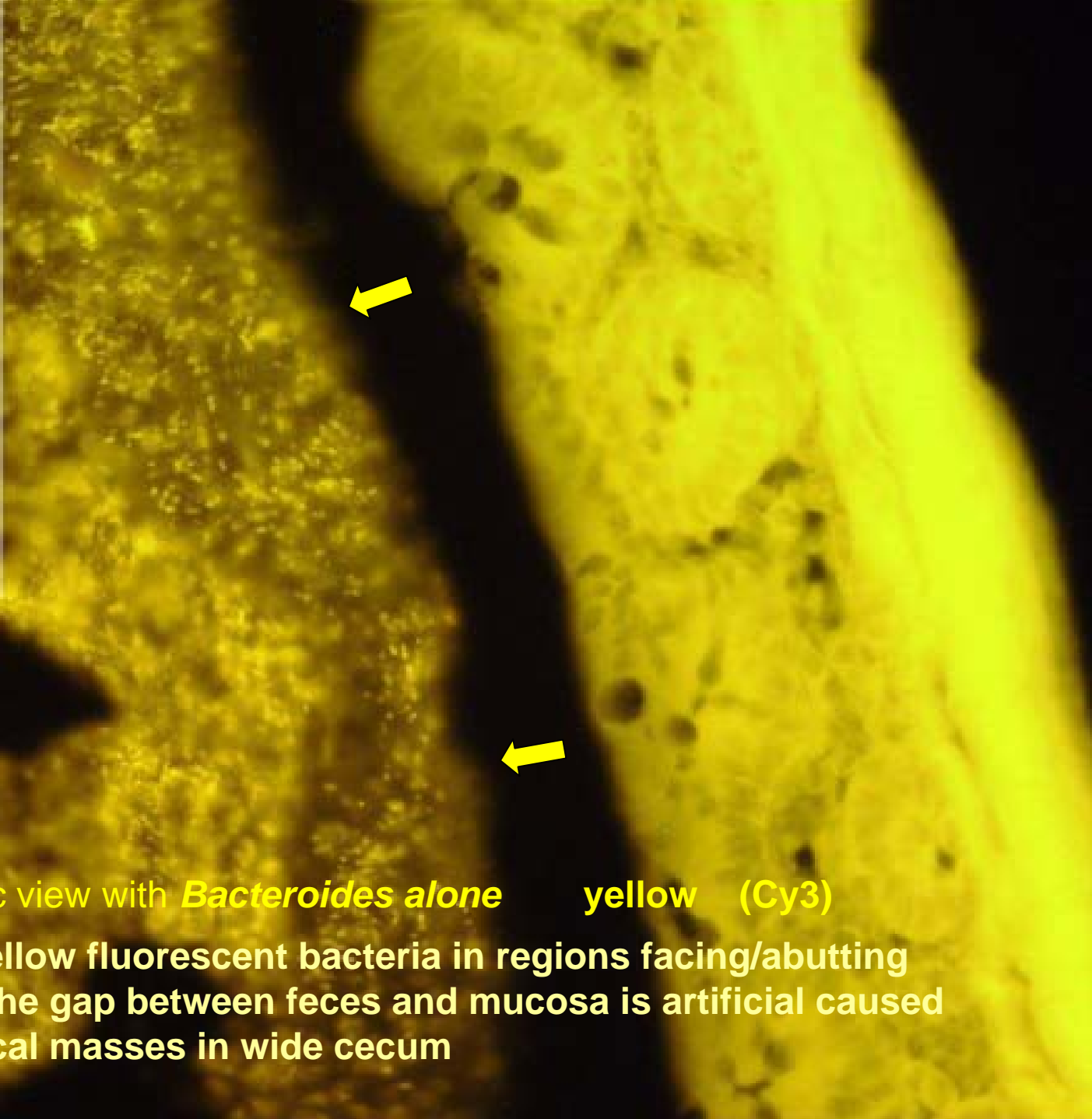
Triple contrast:

| | |
|--|---------------------|
| <i>Bacteroides fragilis</i> (Bfra Probe) | yellow (Cy3) |
| <i>Eubacterium rectale</i> group (Erec Probe) | red (Cy5) |
| Other bacteria (Eub338) | green (FITC) |



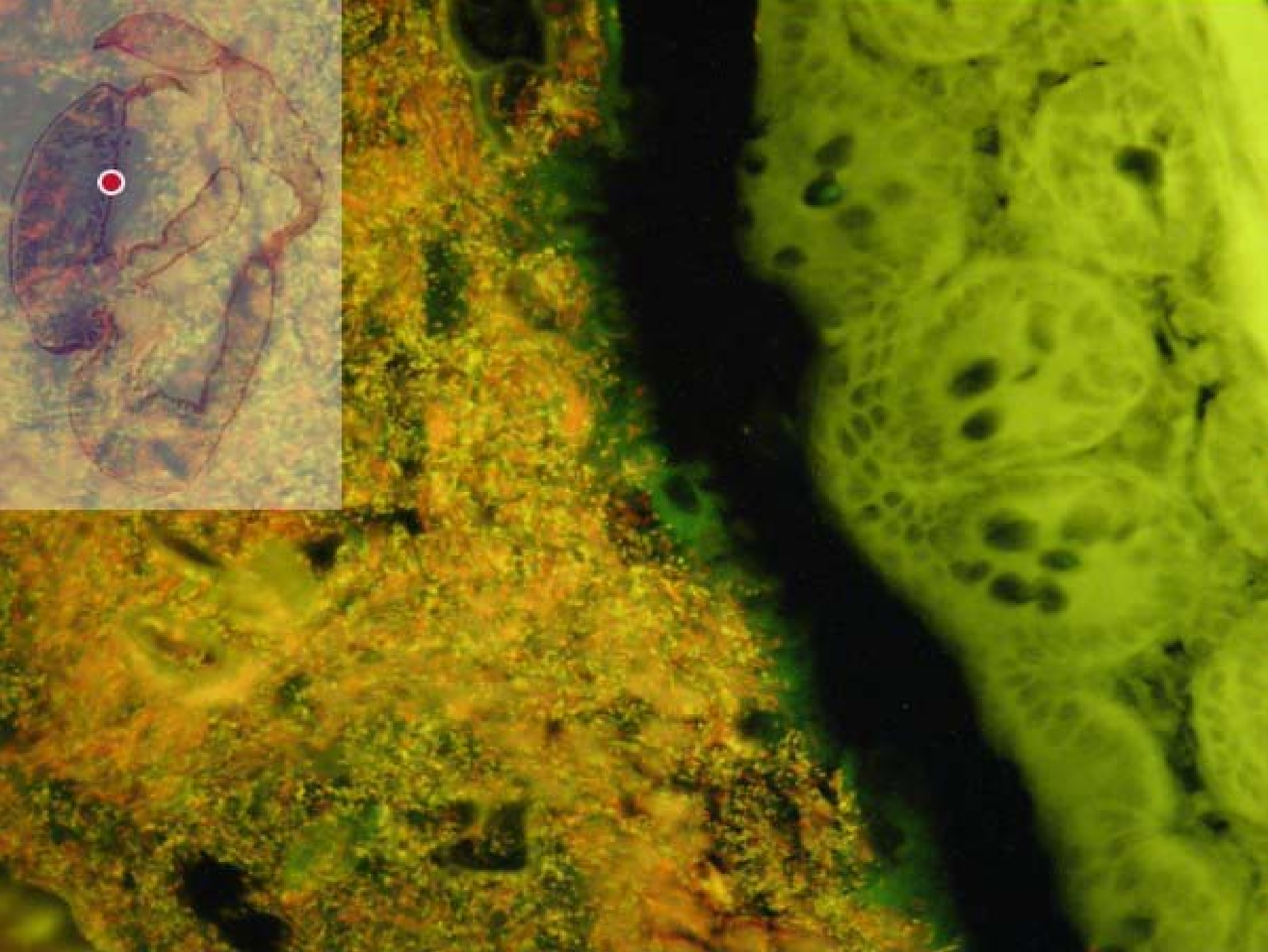
Triple contrast:

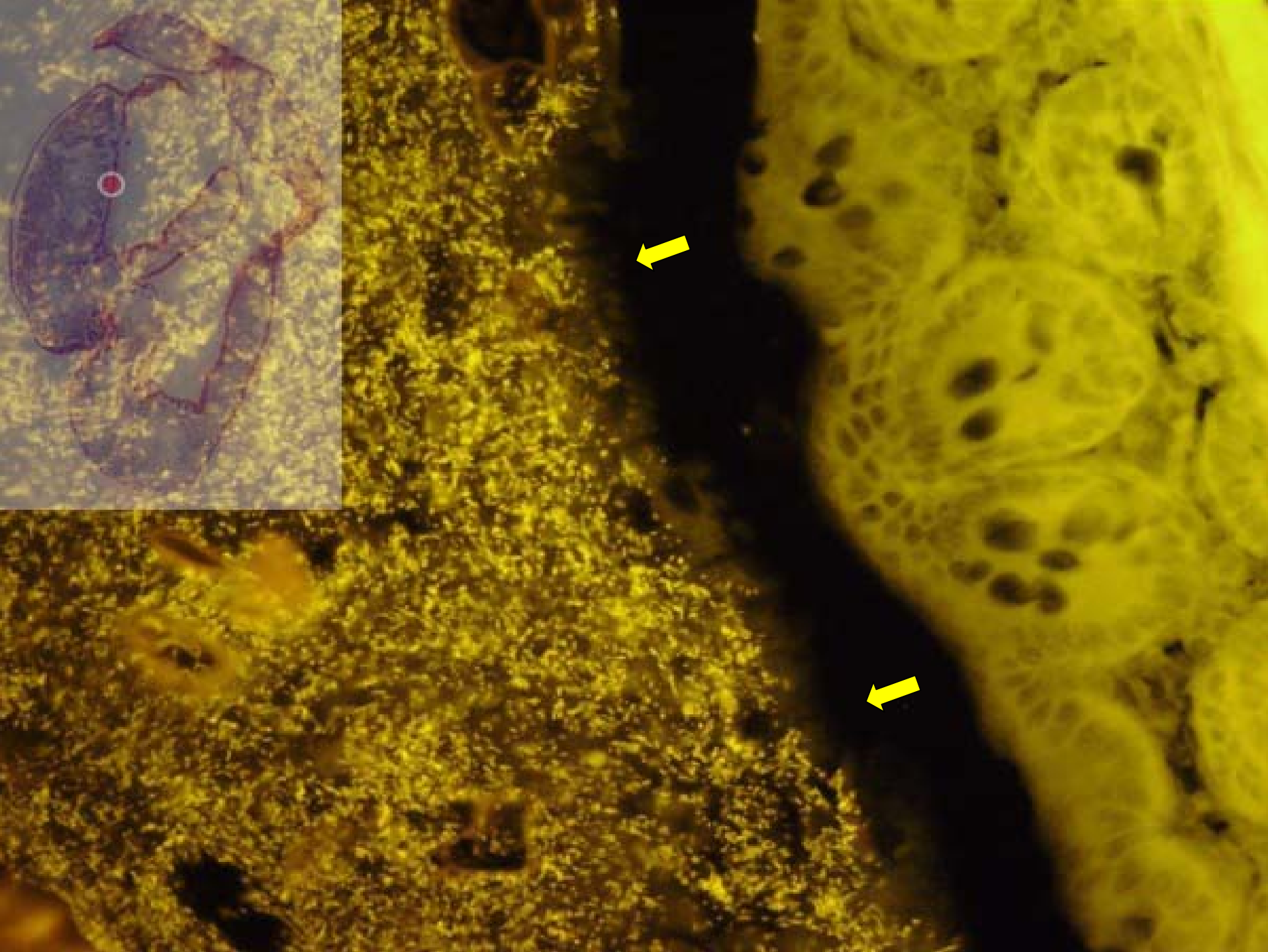
| | |
|---|--------------|
| <i>Bacteroides fragilis</i> (Bfra Probe) | yellow (Cy3) |
| <i>Eubacterium rectale</i> group (Erec Probe) | red (Cy5) |
| Other bacteria (Eub338) | green (FITC) |

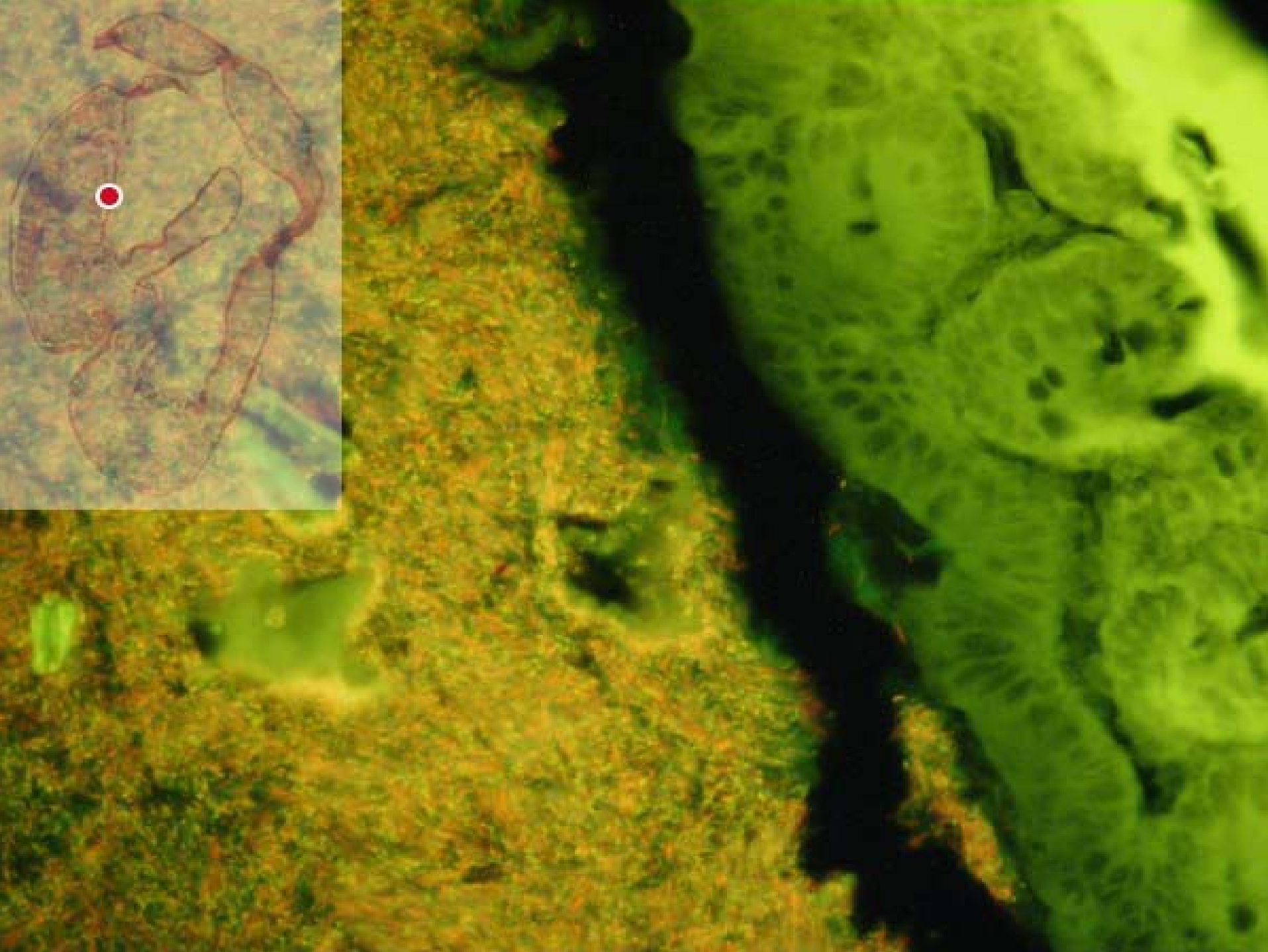


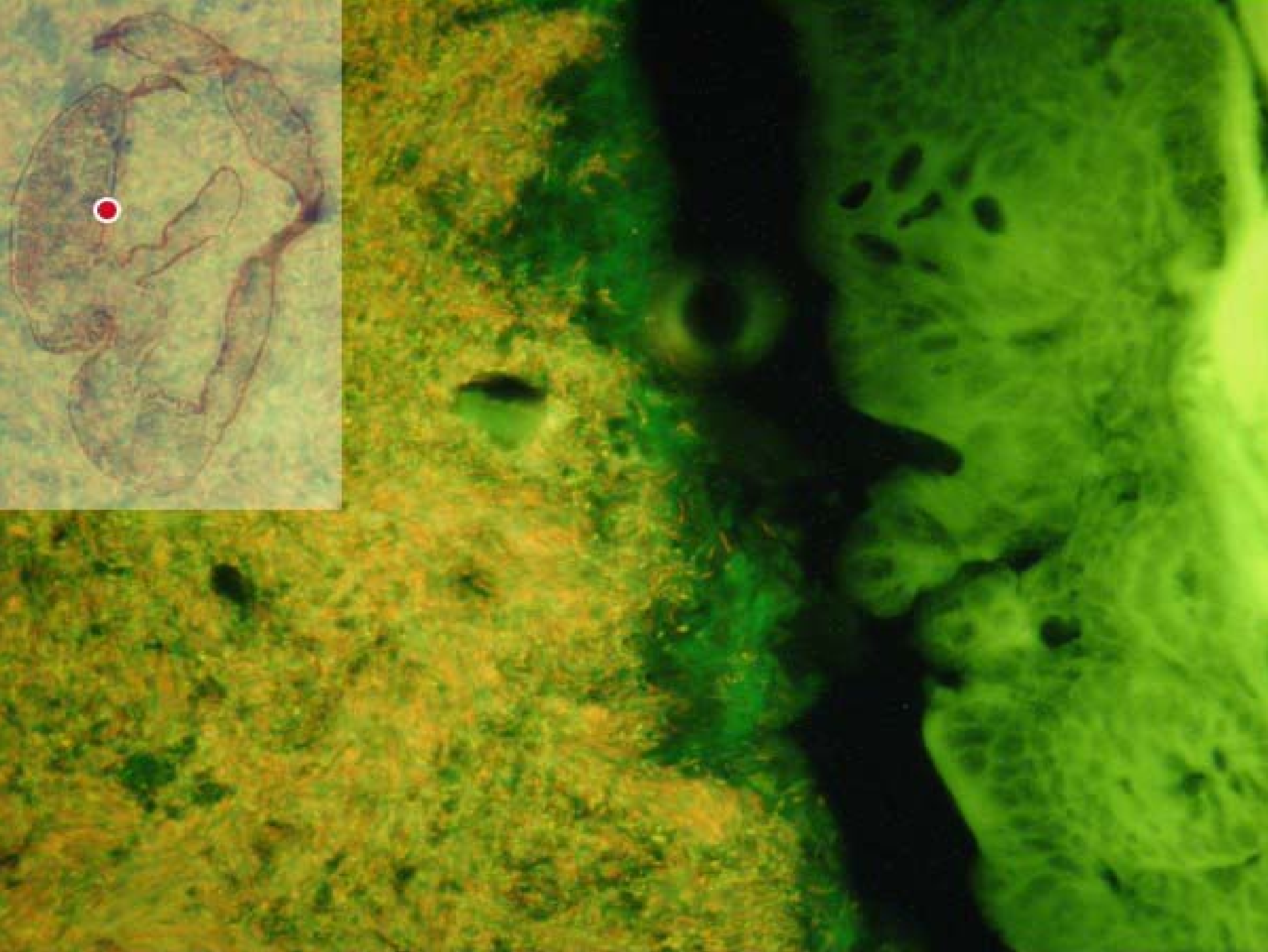
same microscopic view with *Bacteroides* alone yellow (Cy3)

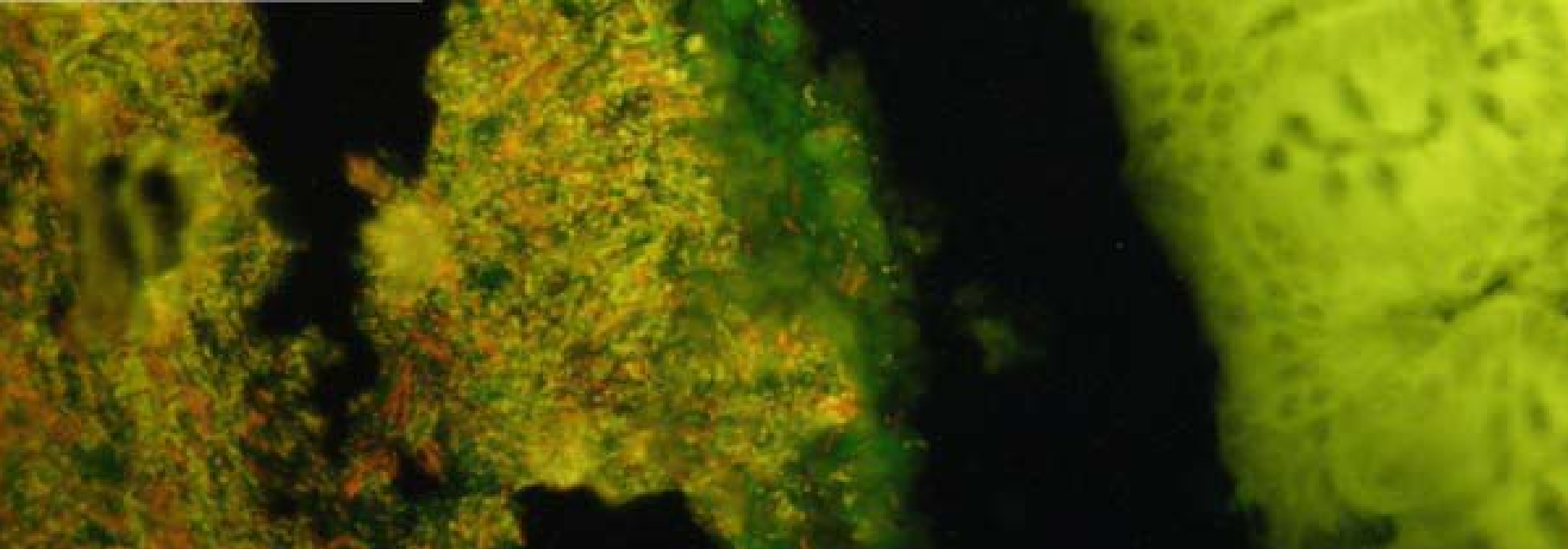
note absence of yellow fluorescent bacteria in regions facing/abutting mucosal surface, the gap between feces and mucosa is artificial caused by shrinkage of fecal masses in wide cecum

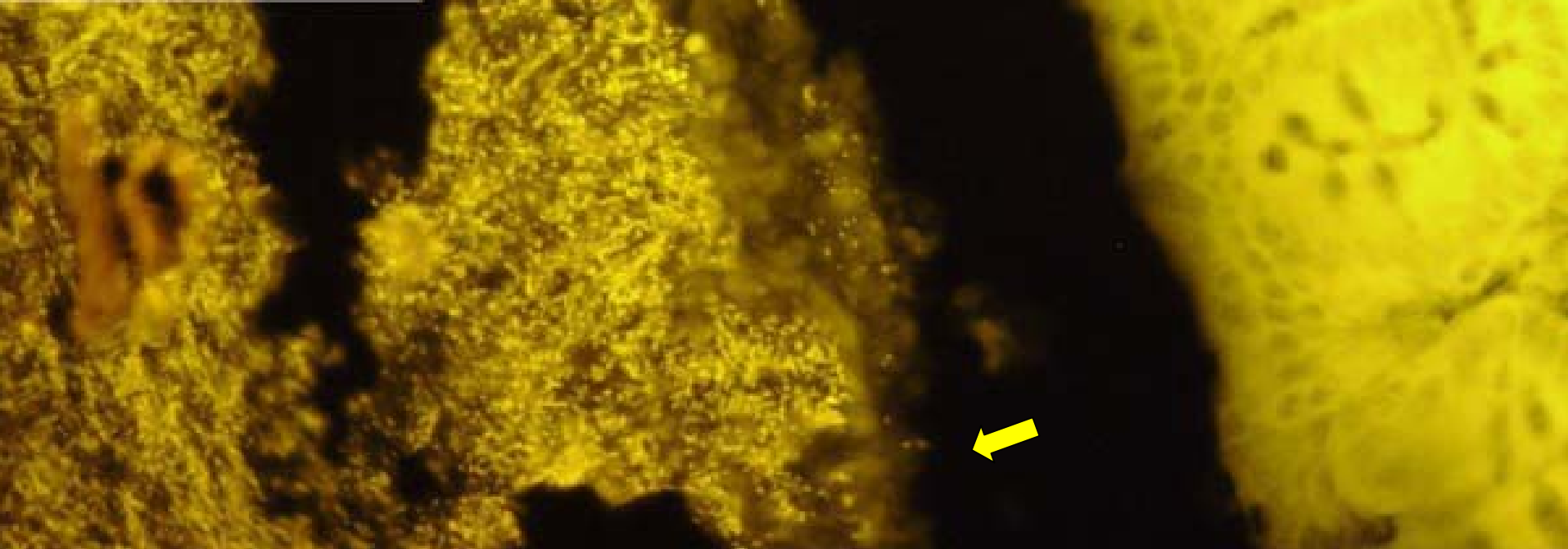


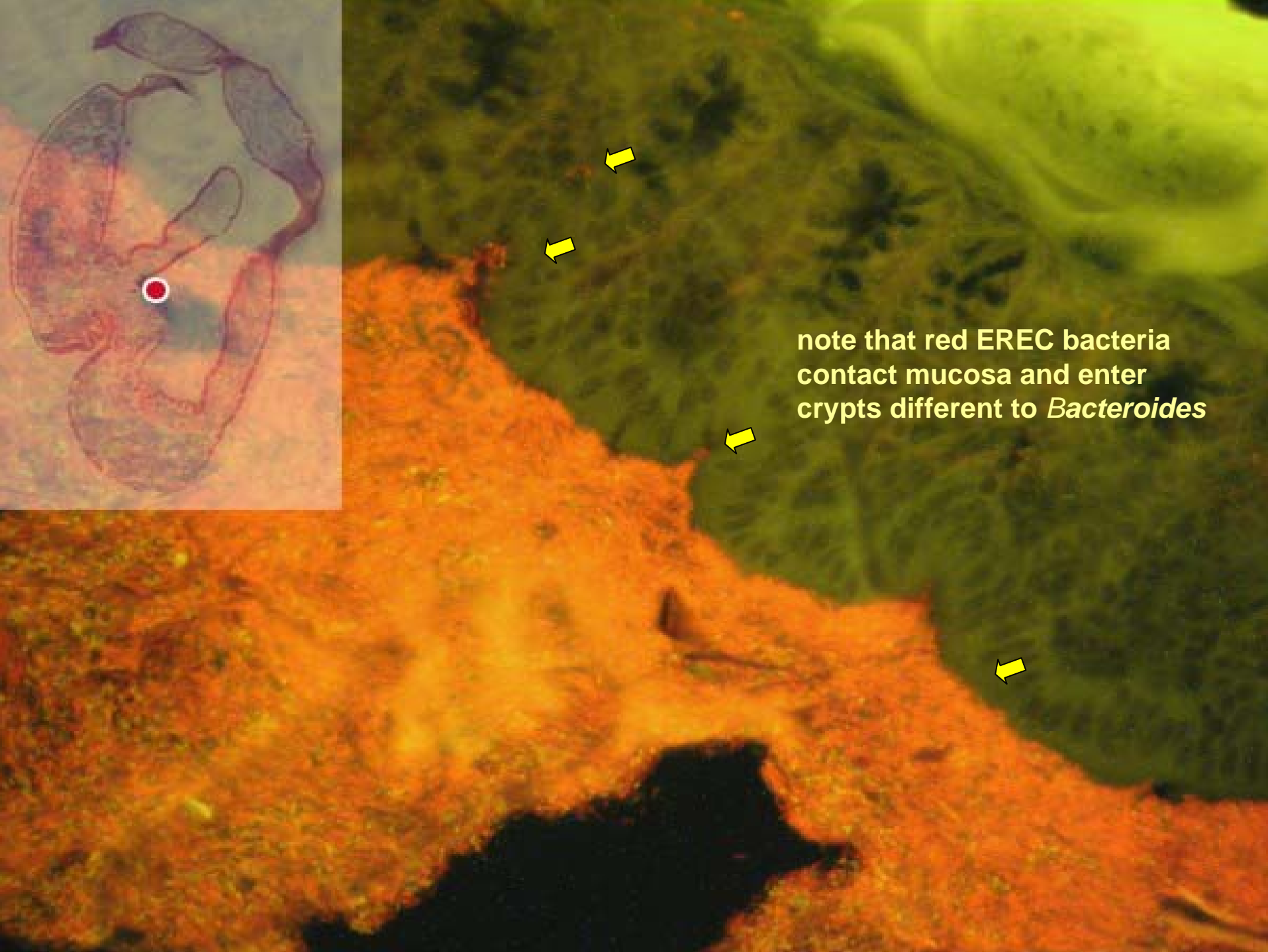








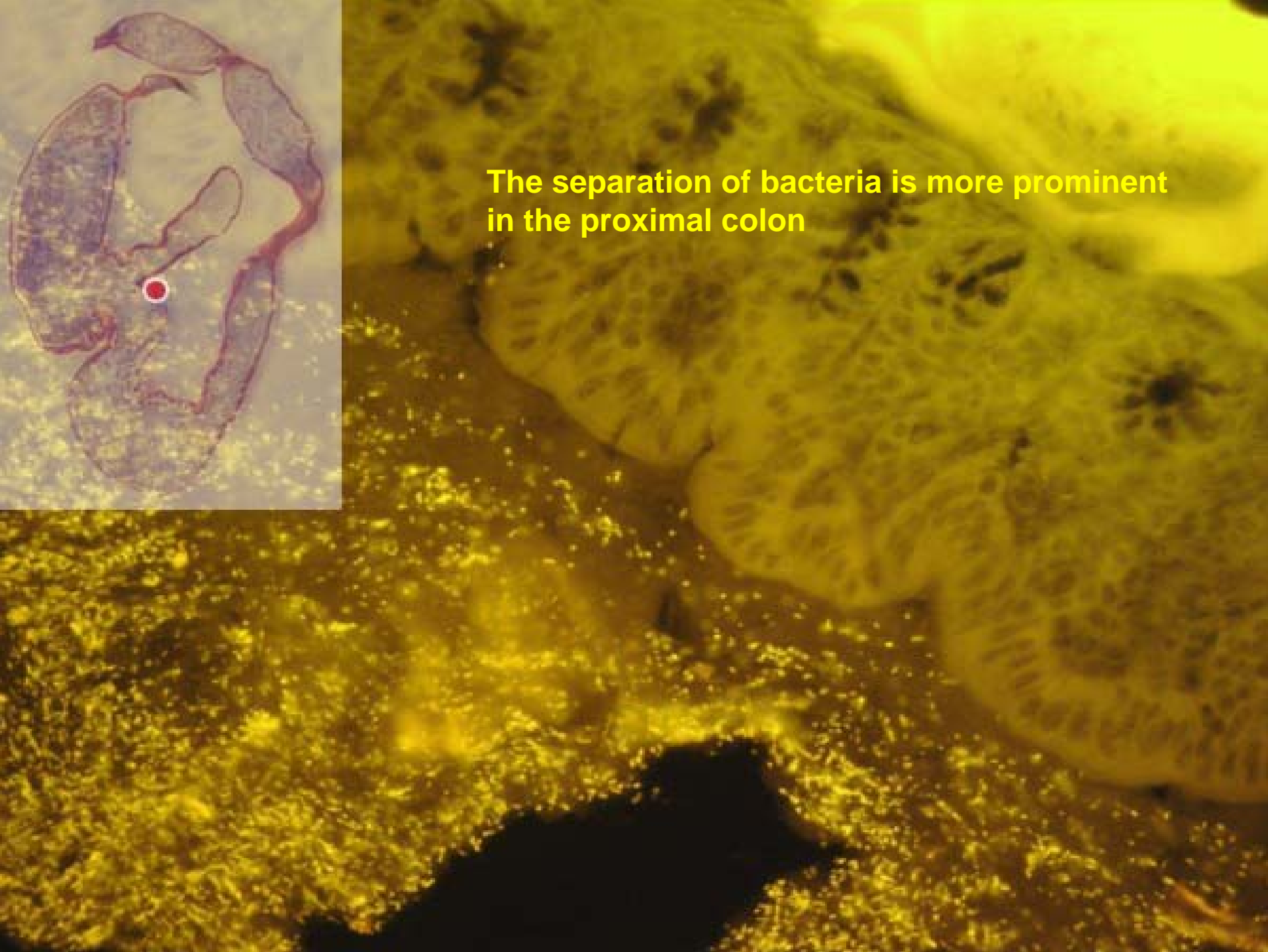


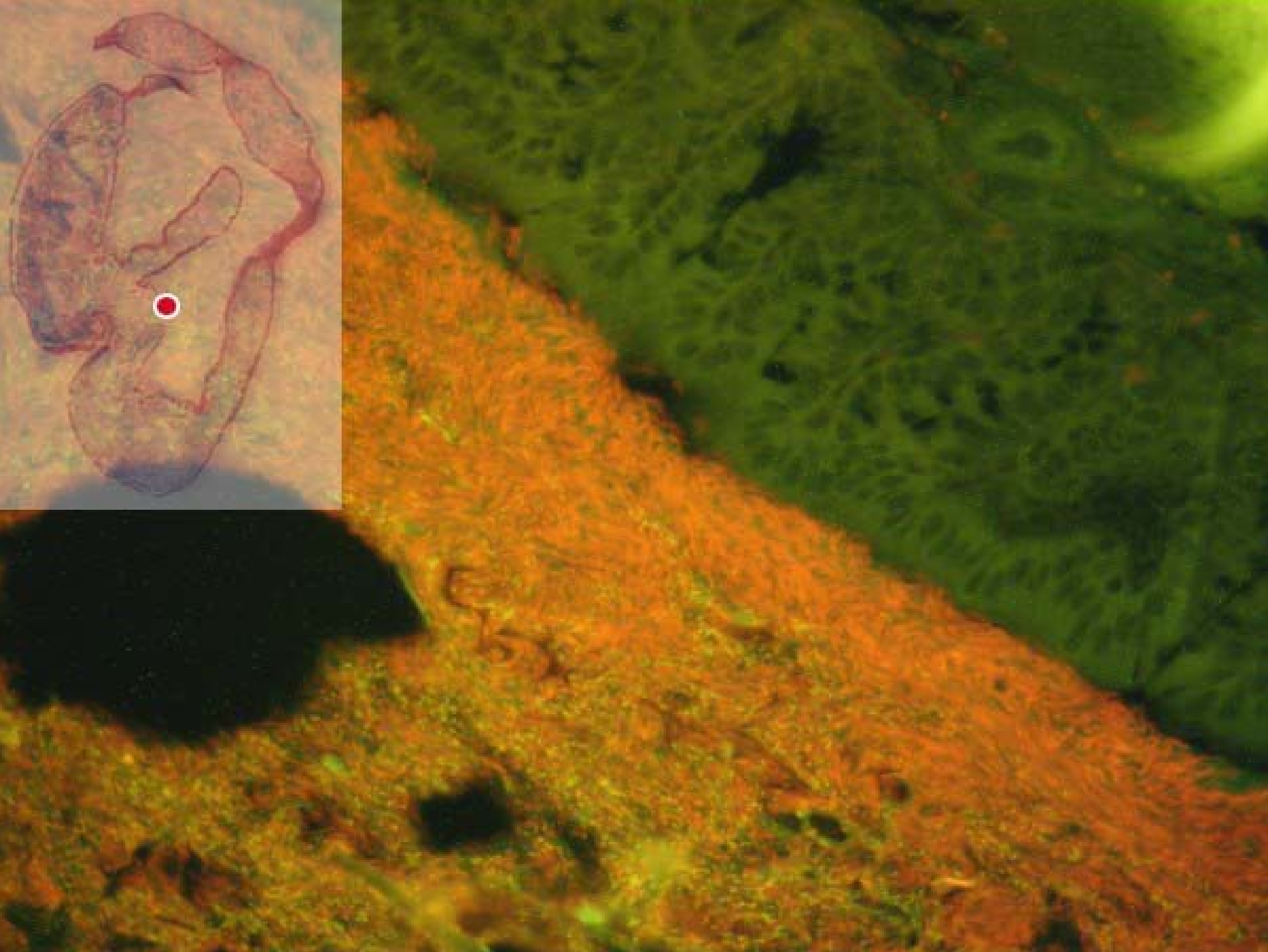


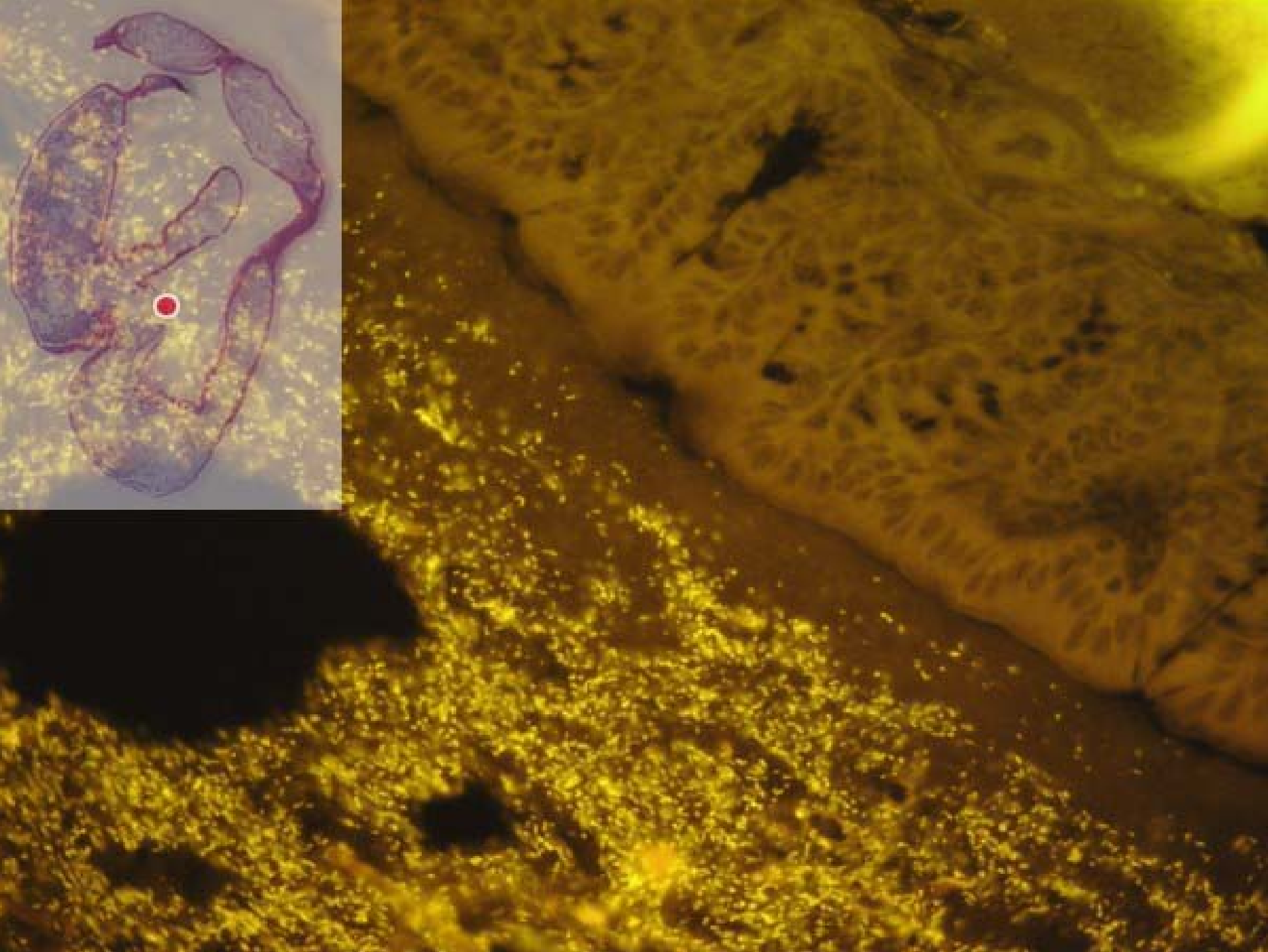
note that red EREC bacteria contact mucosa and enter crypts different to *Bacteroides*

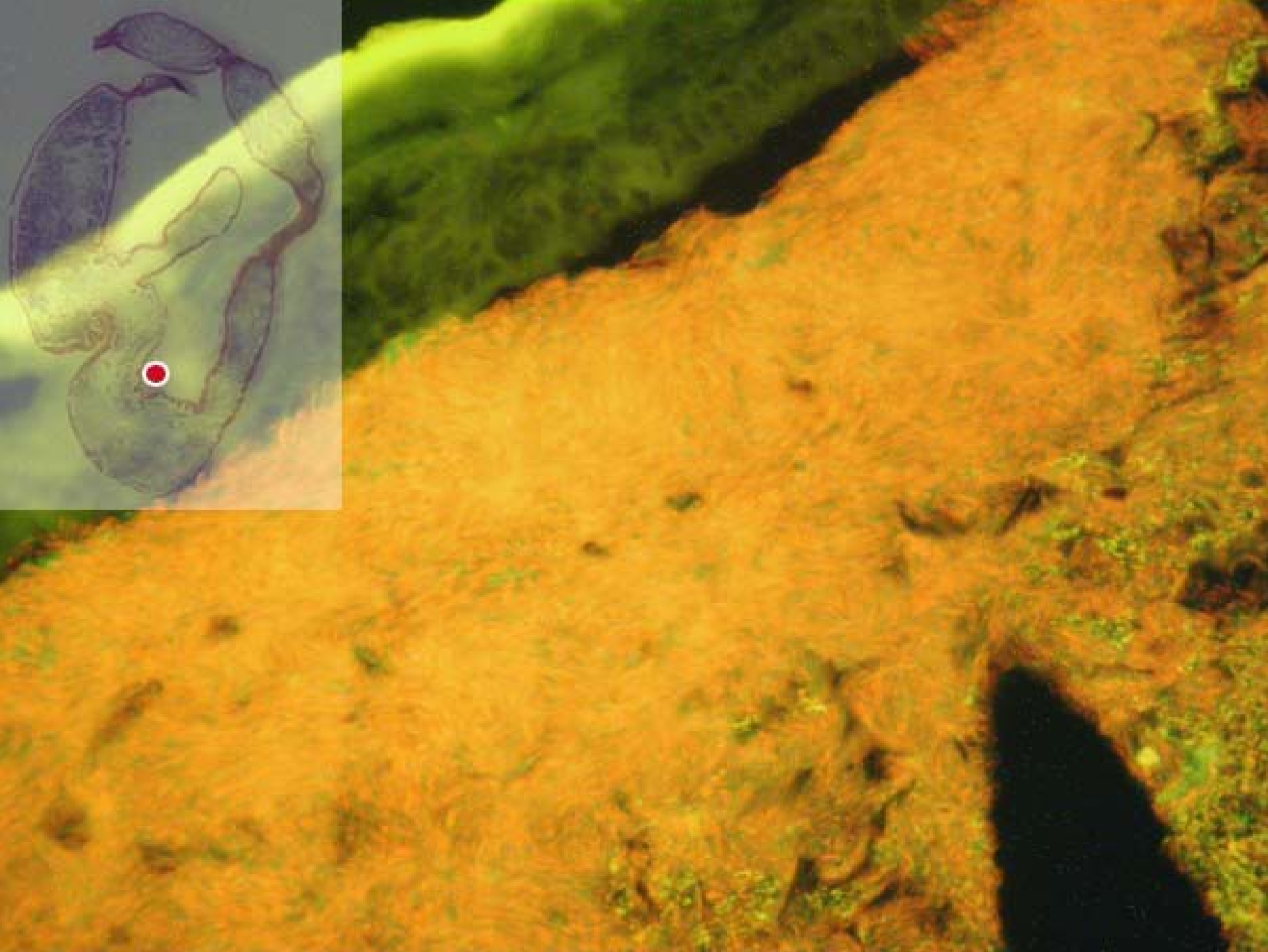


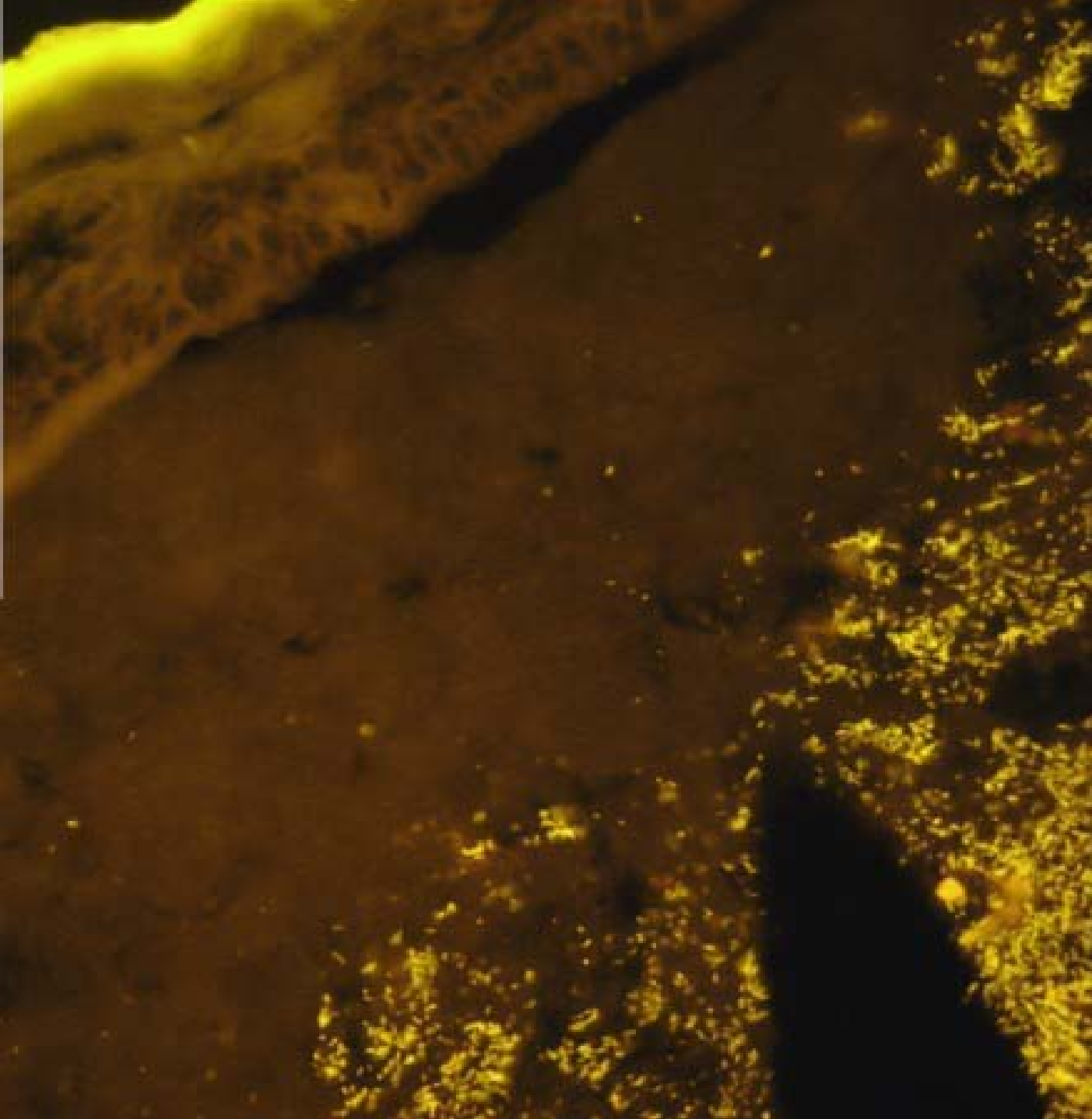
The separation of bacteria is more prominent in the proximal colon

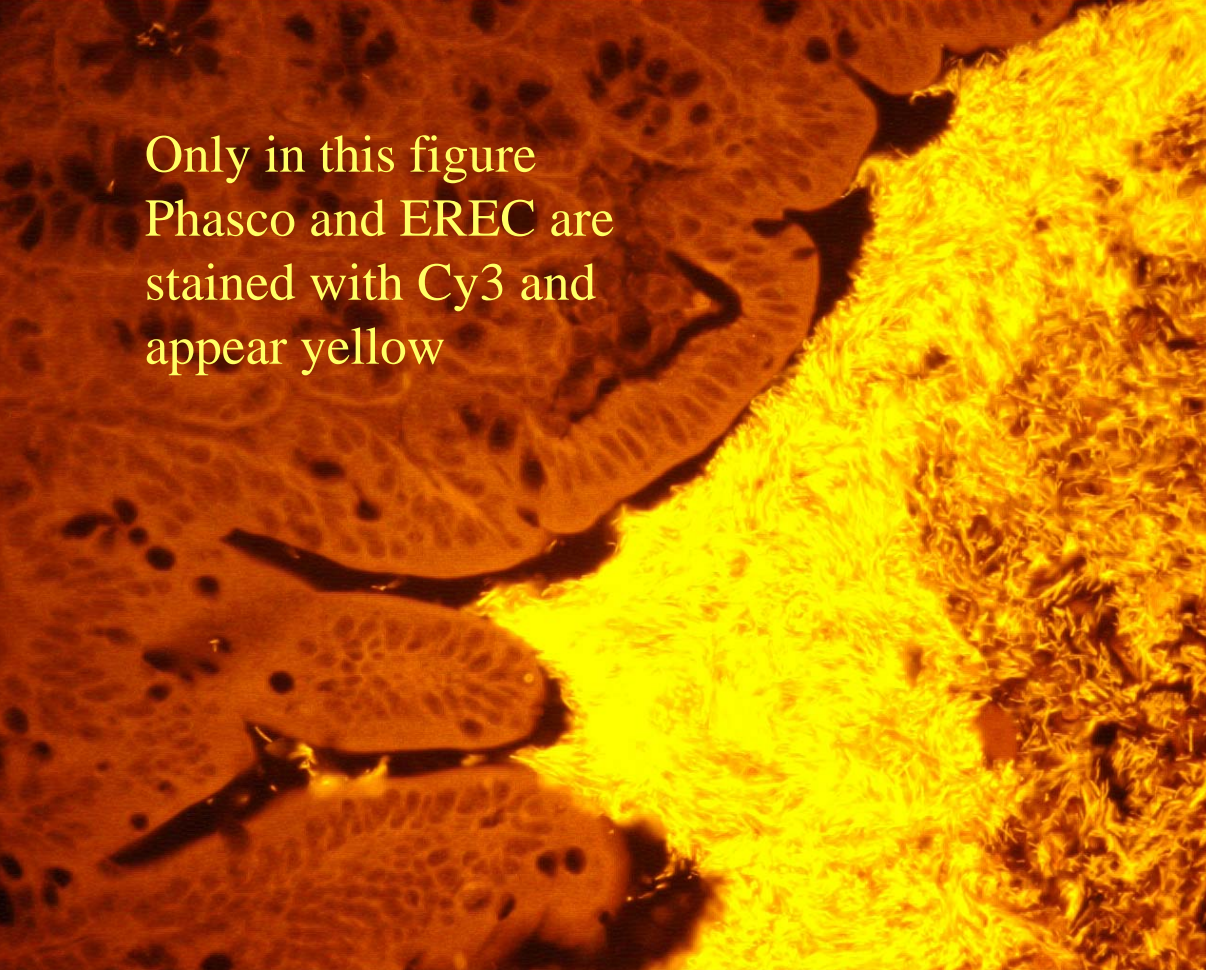












Only in this figure
Phasco and EREC
are stained with Cy3 and
appear yellow

EREC
Lab,
Bif,
Phasco
Lach



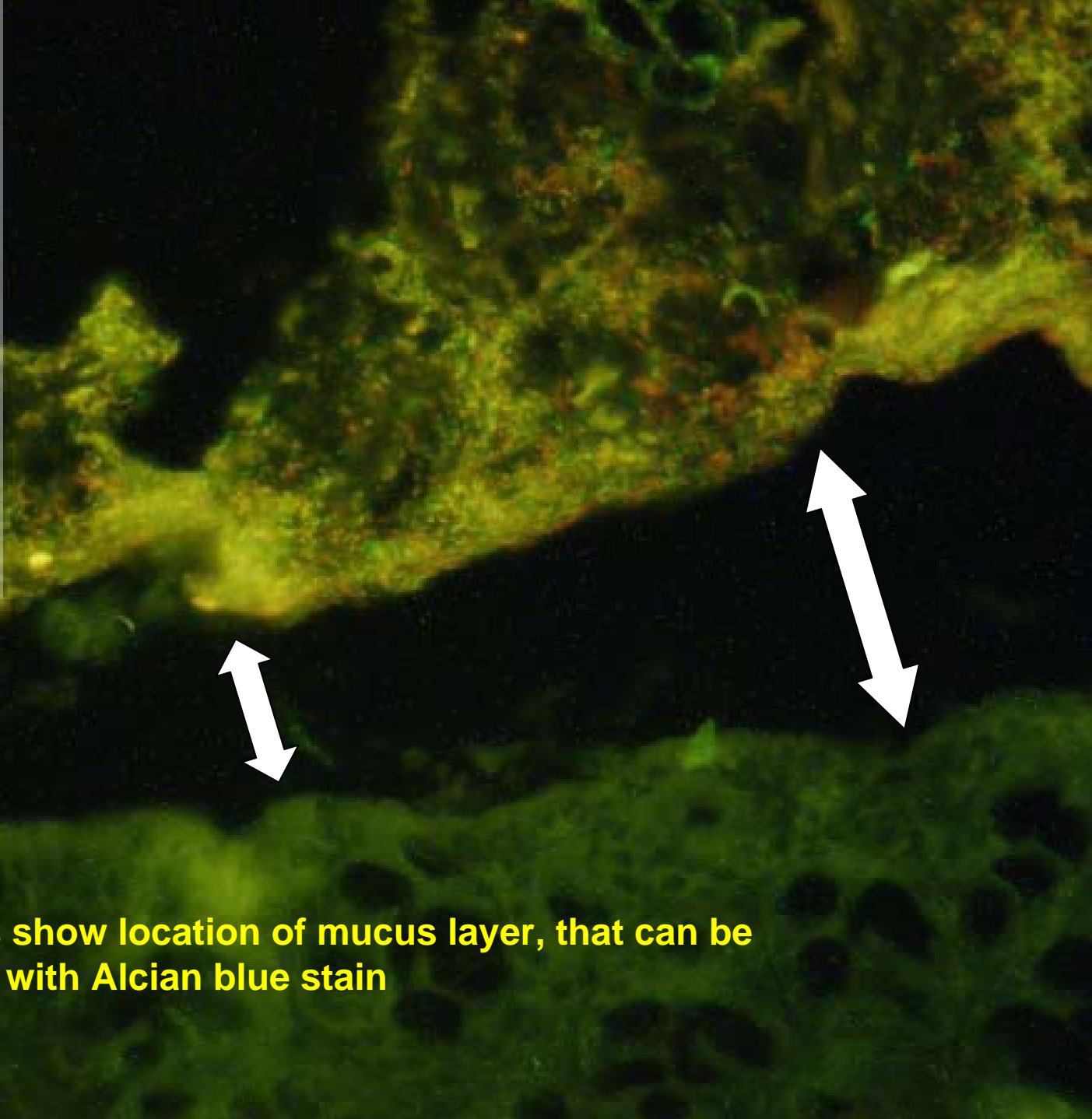
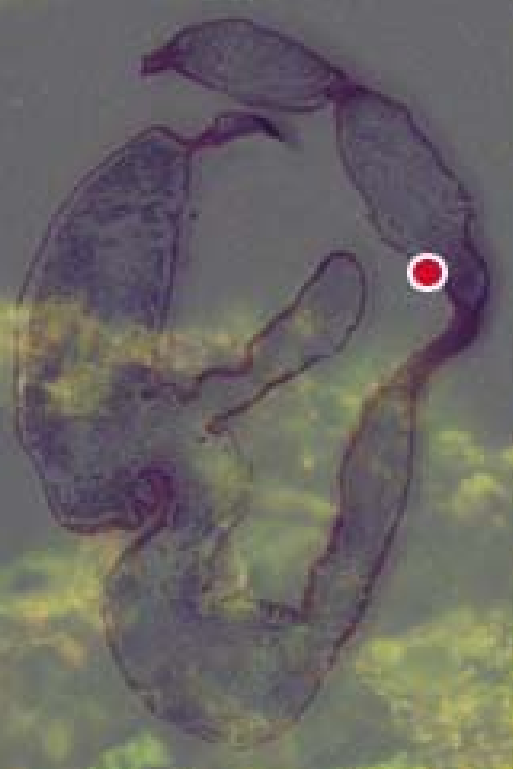
Lach is red (Cy5)

Composition of the interlaced layer

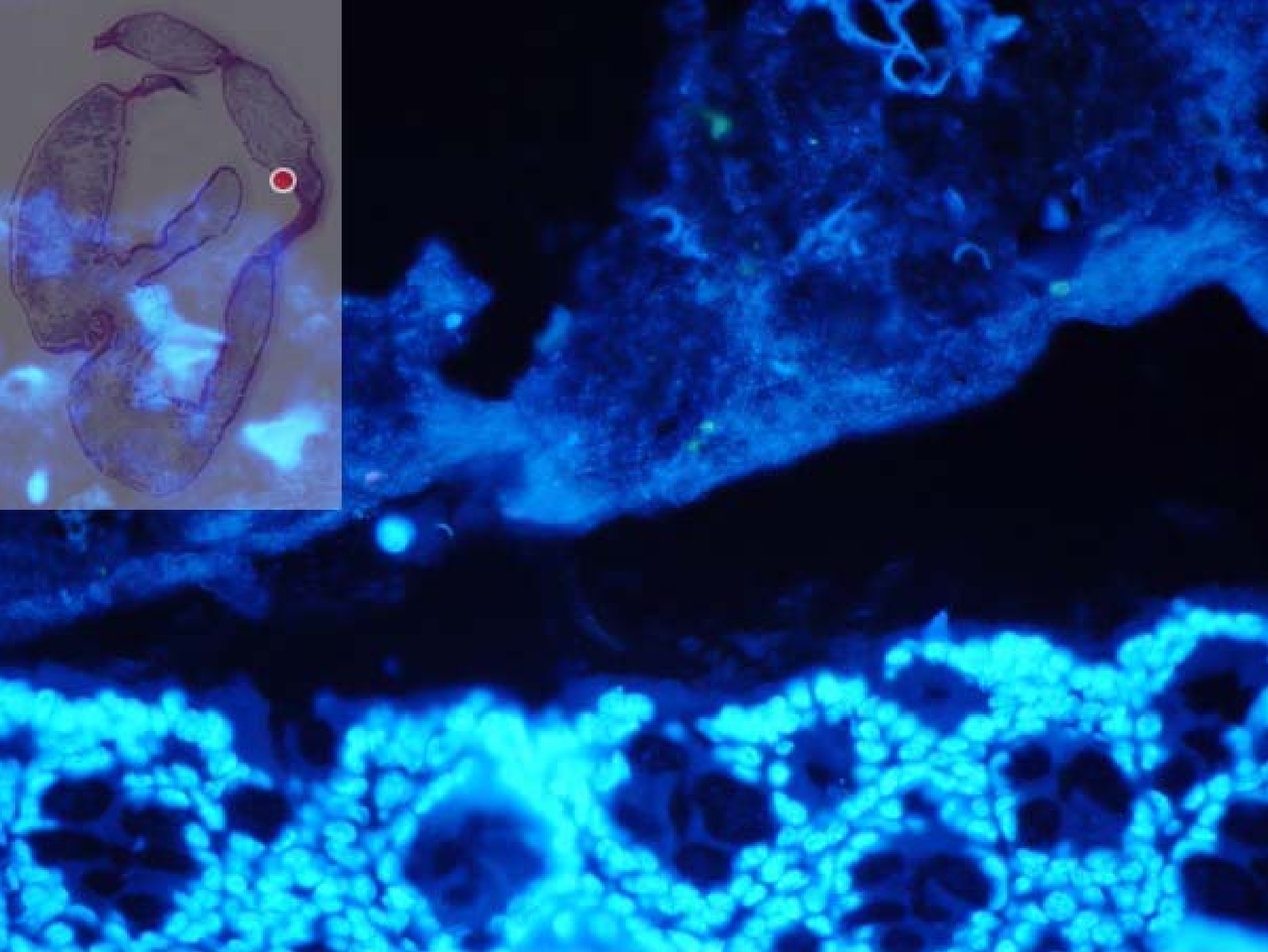
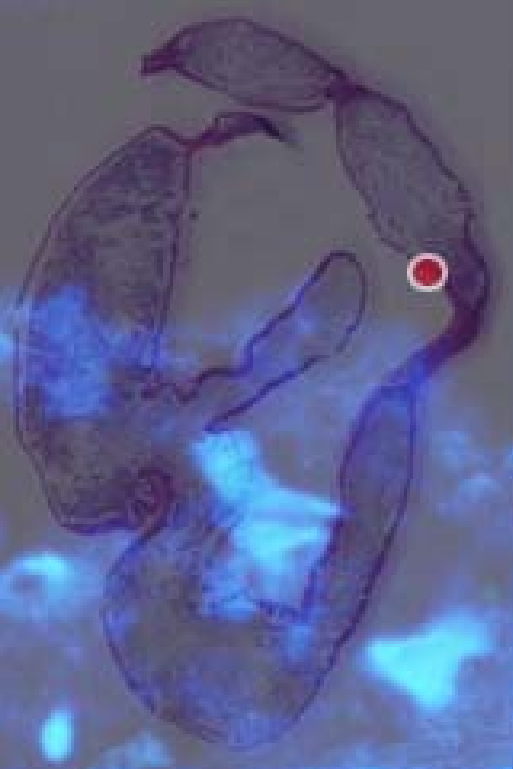


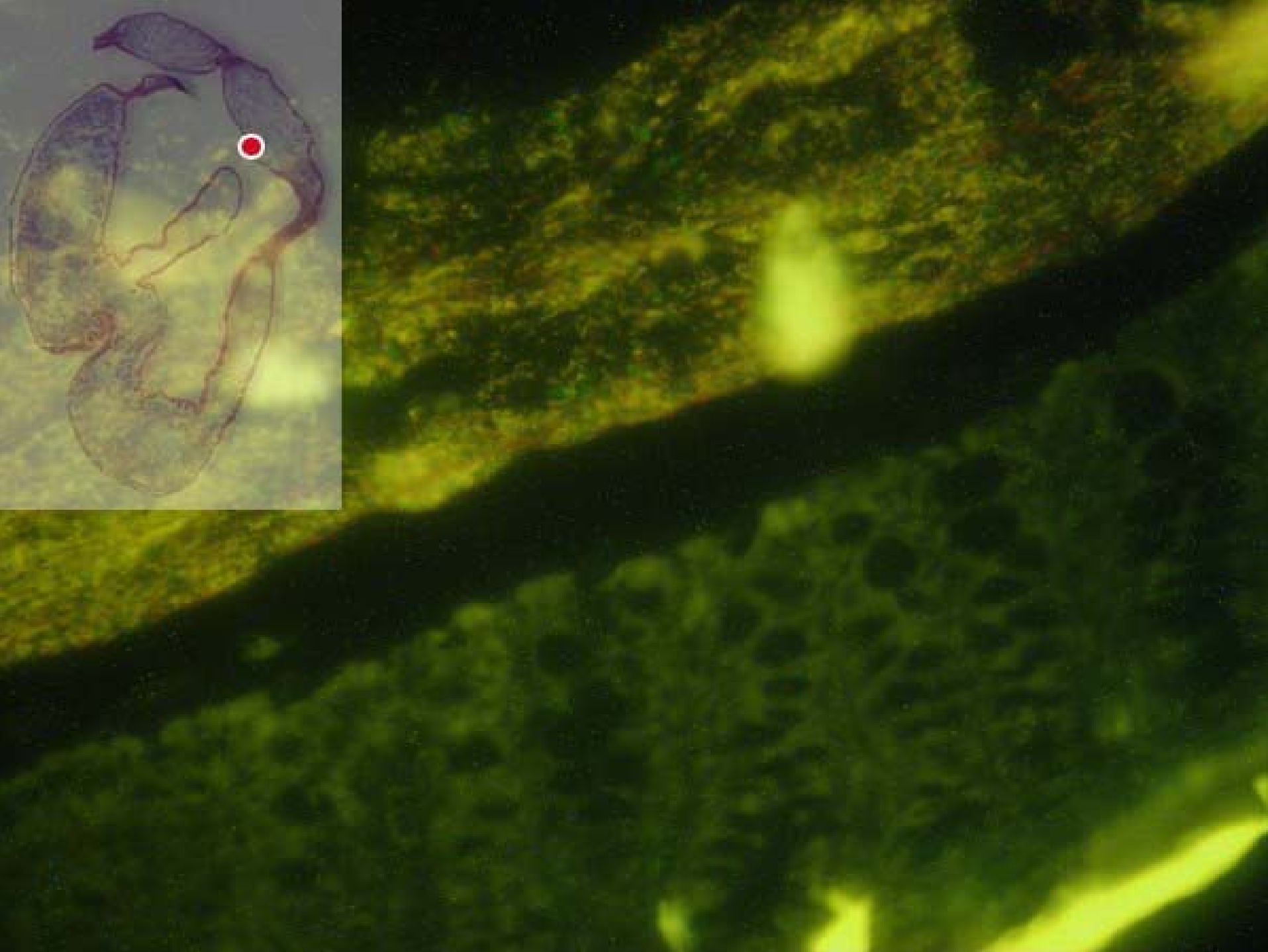
Short rods of
Bacteroides,
Enterobacteriaceae,
Clostridium difficile,
Veillonella groups
have no contact with
the colonic wall

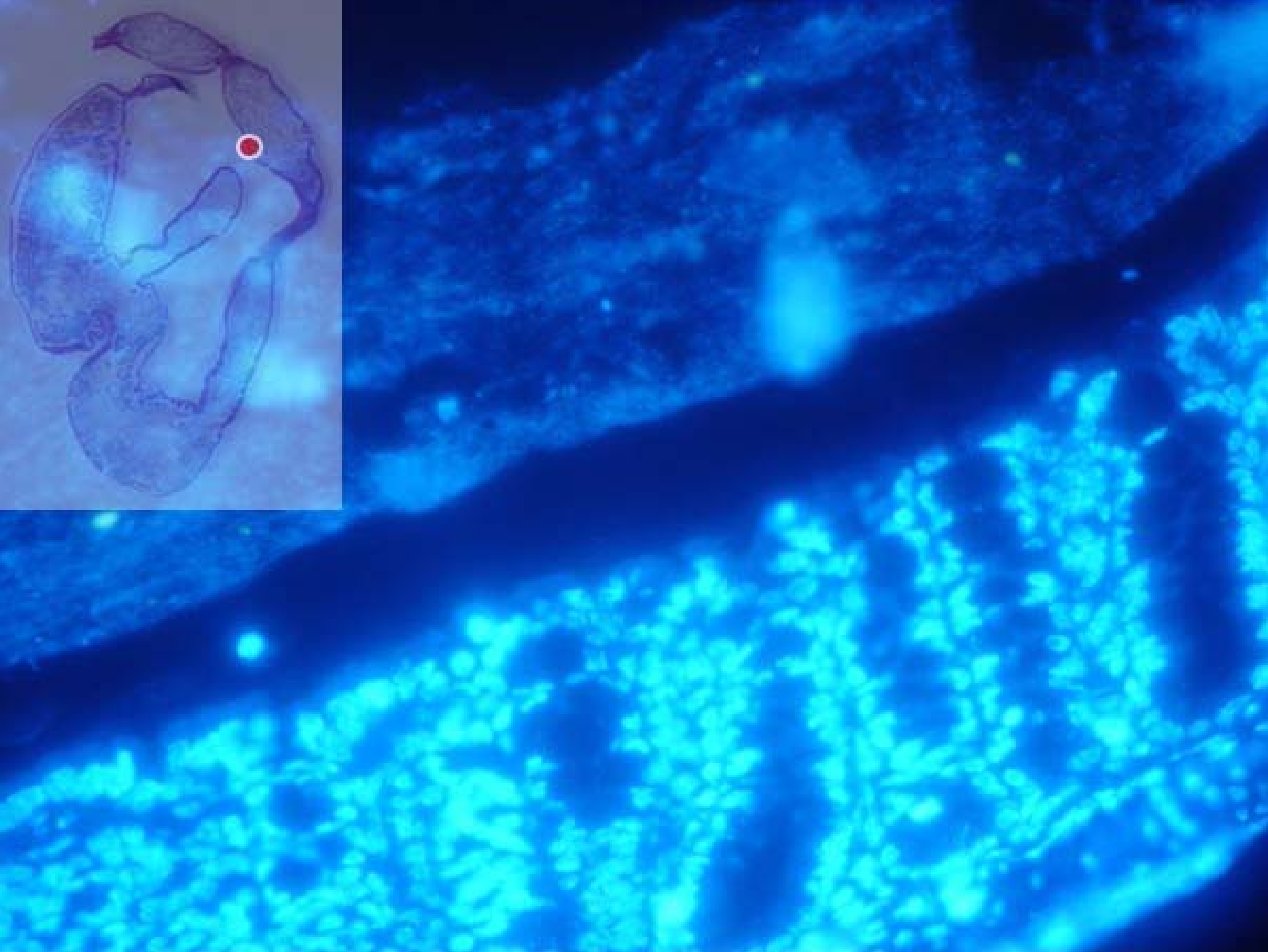


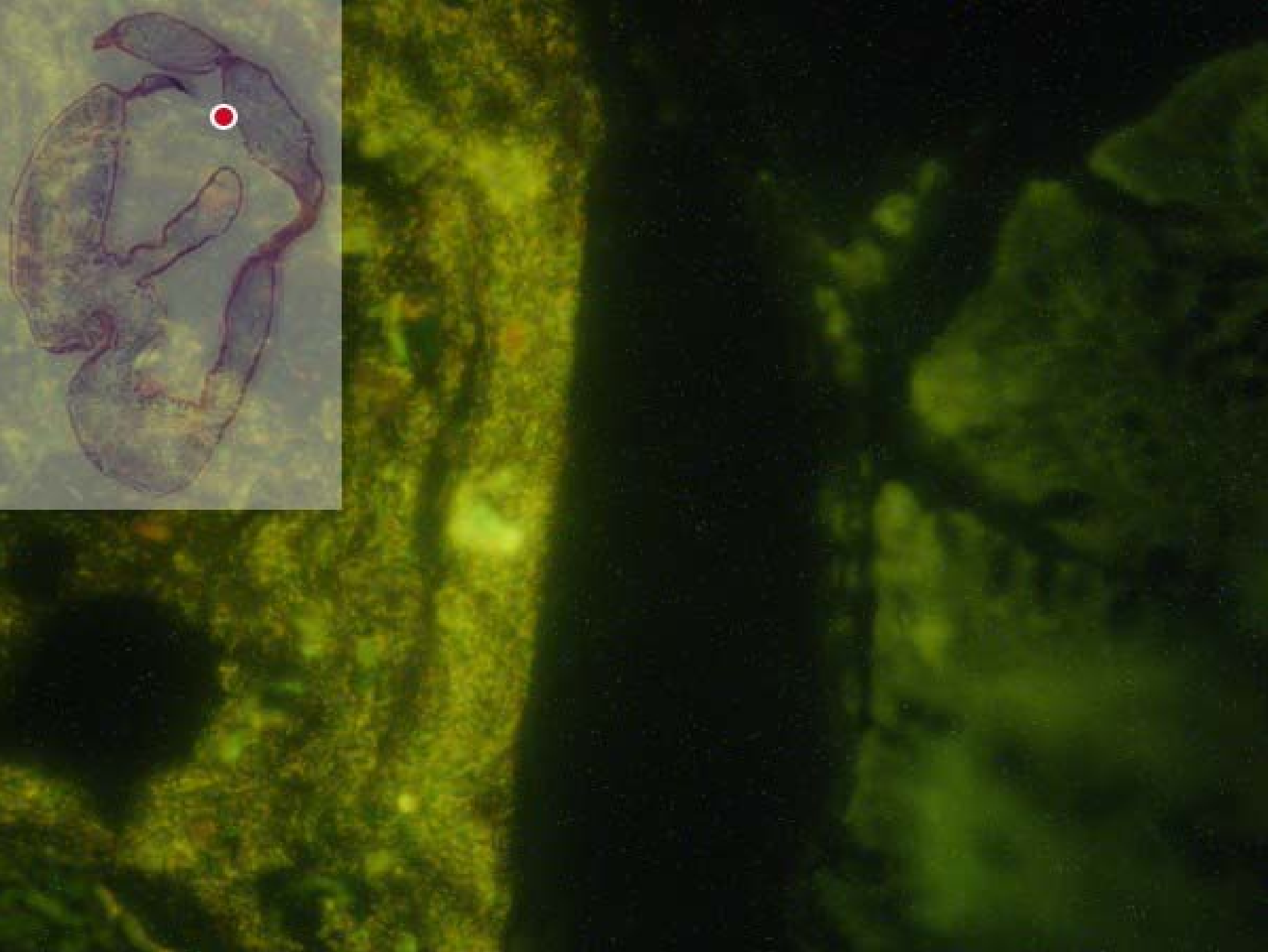


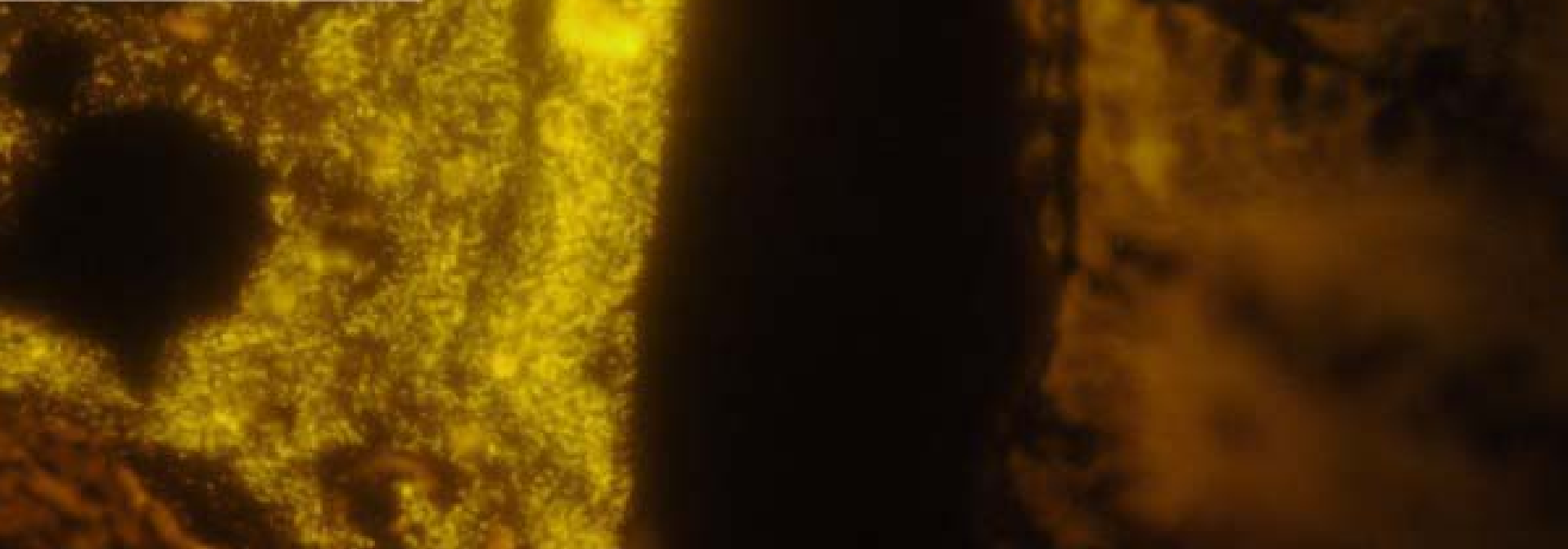
white arrows show location of mucus layer, that can be documented with Alcian blue stain

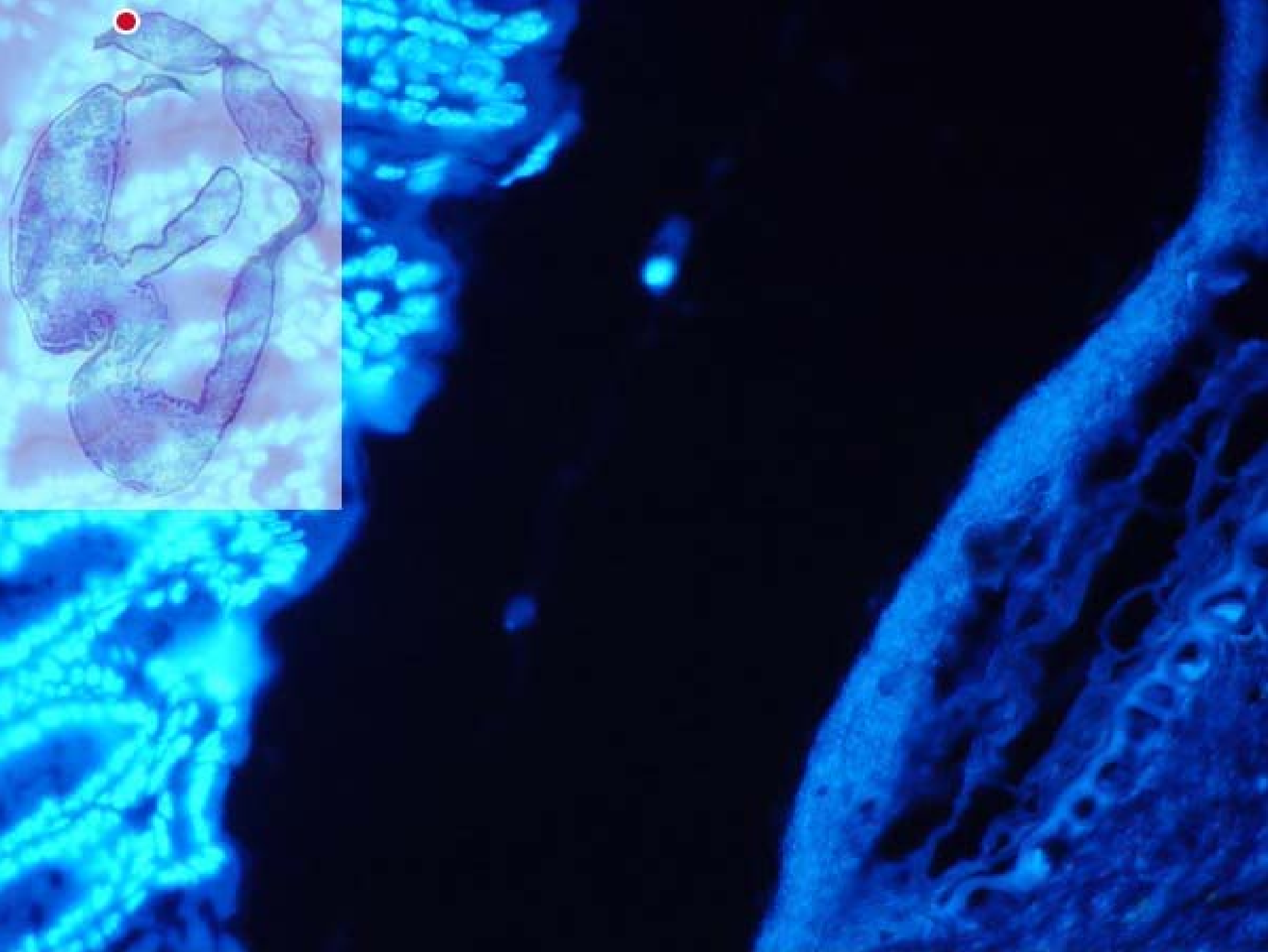
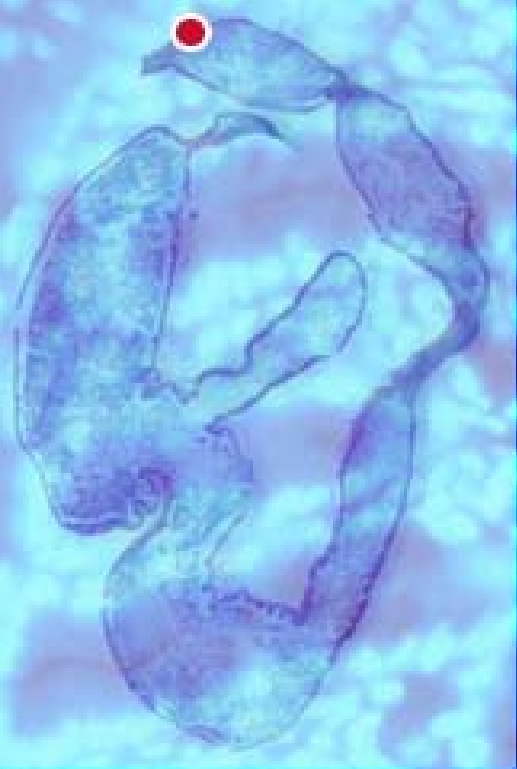


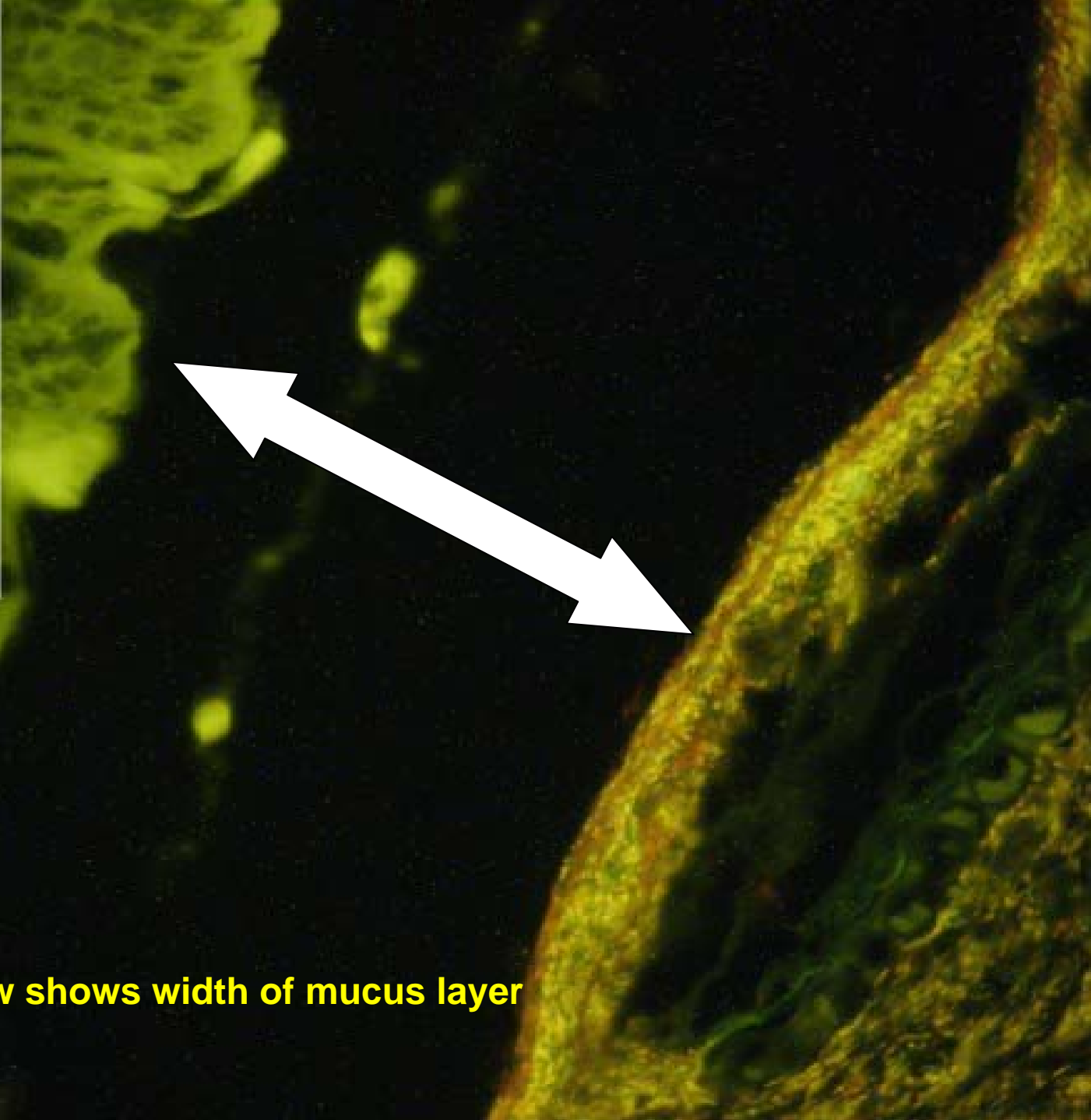






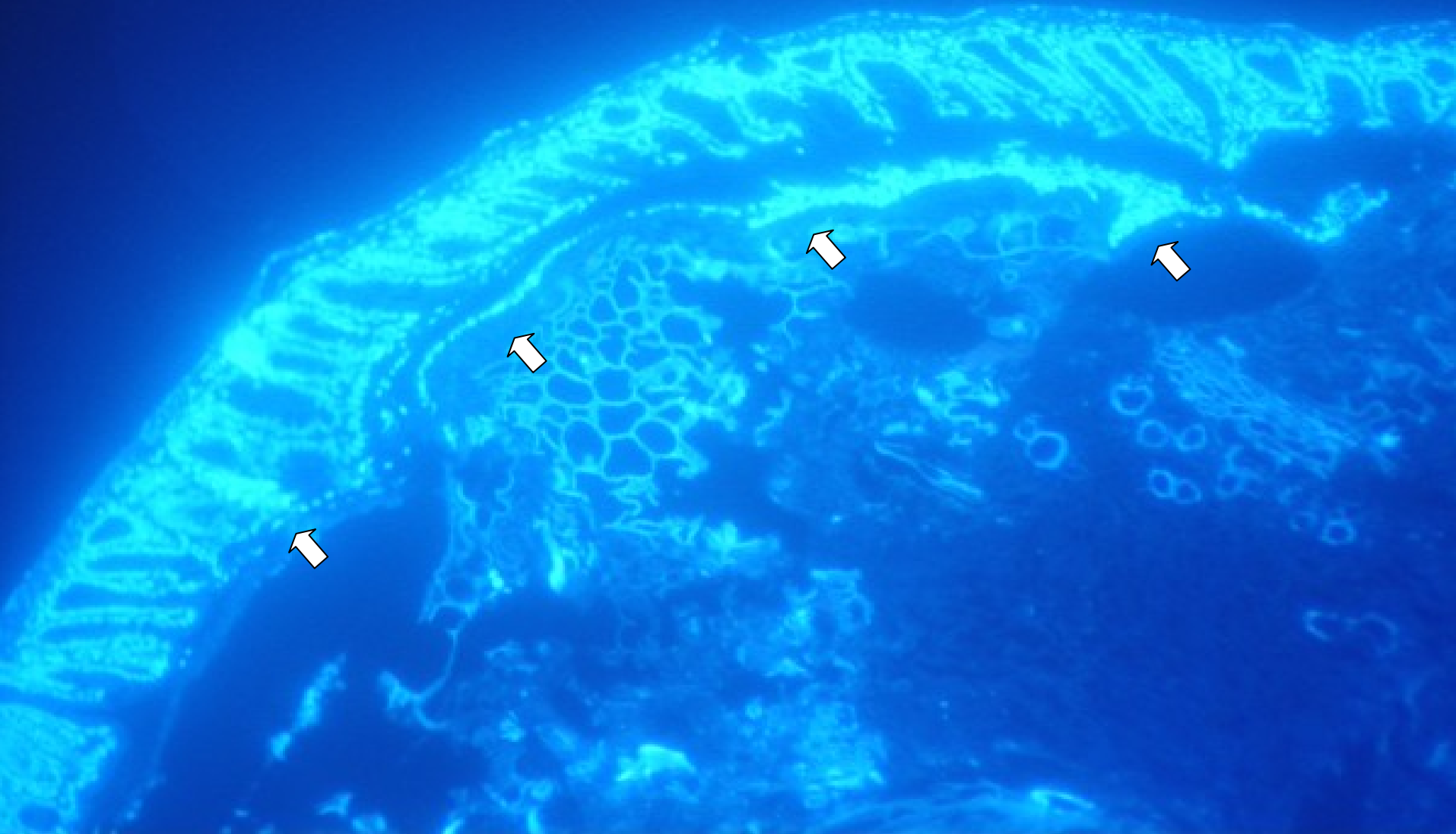


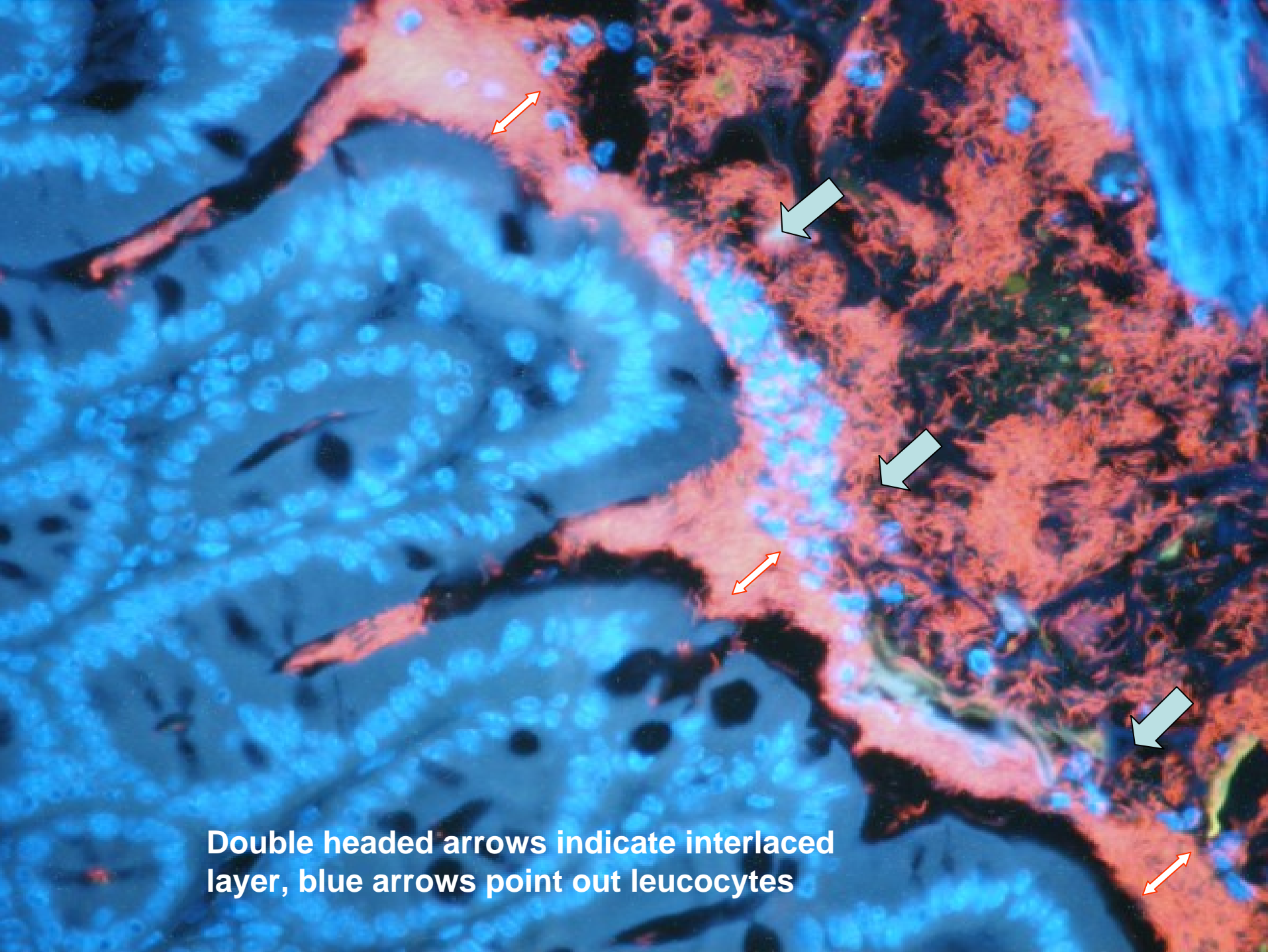




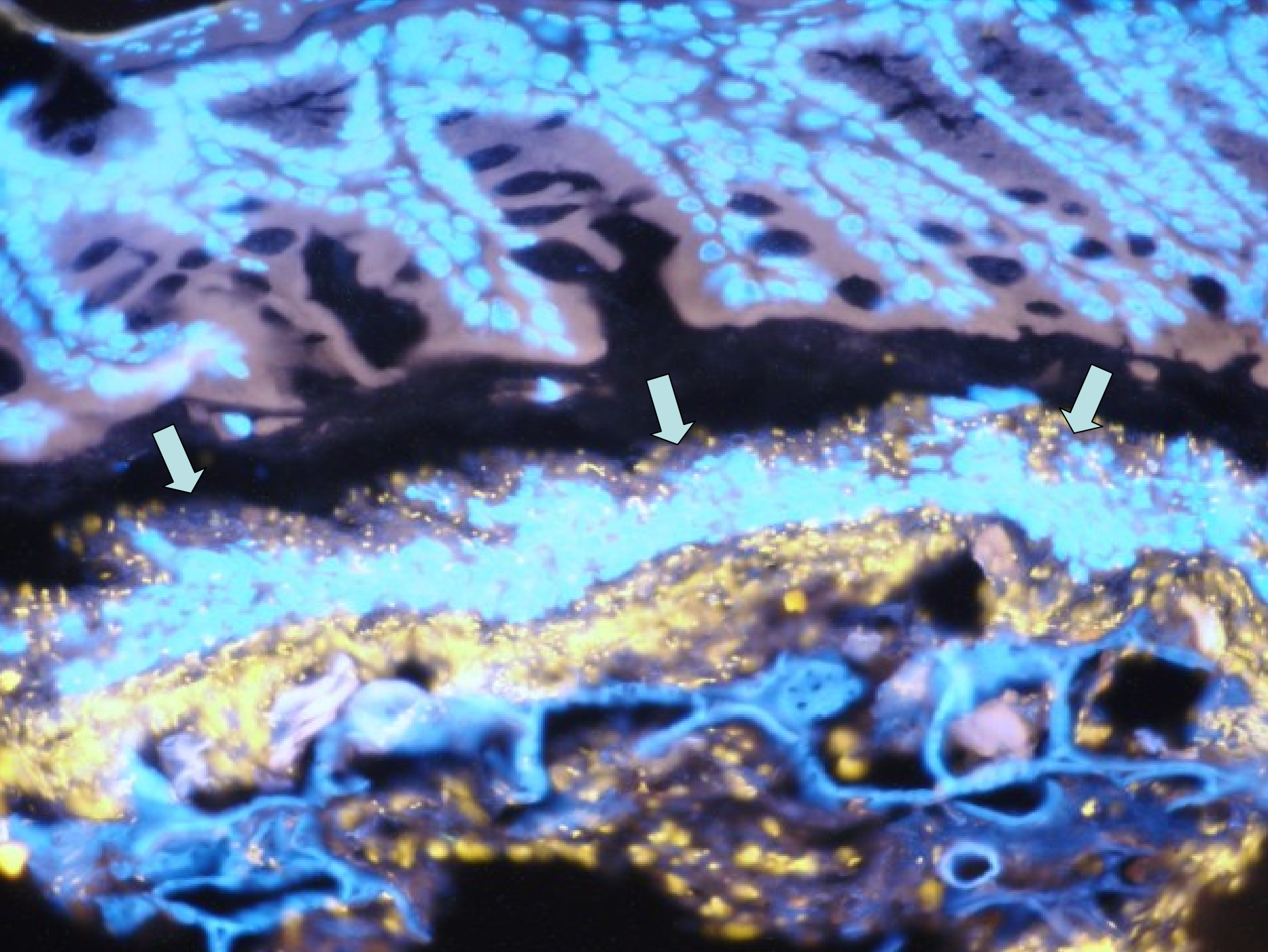
white arrow shows width of mucus layer

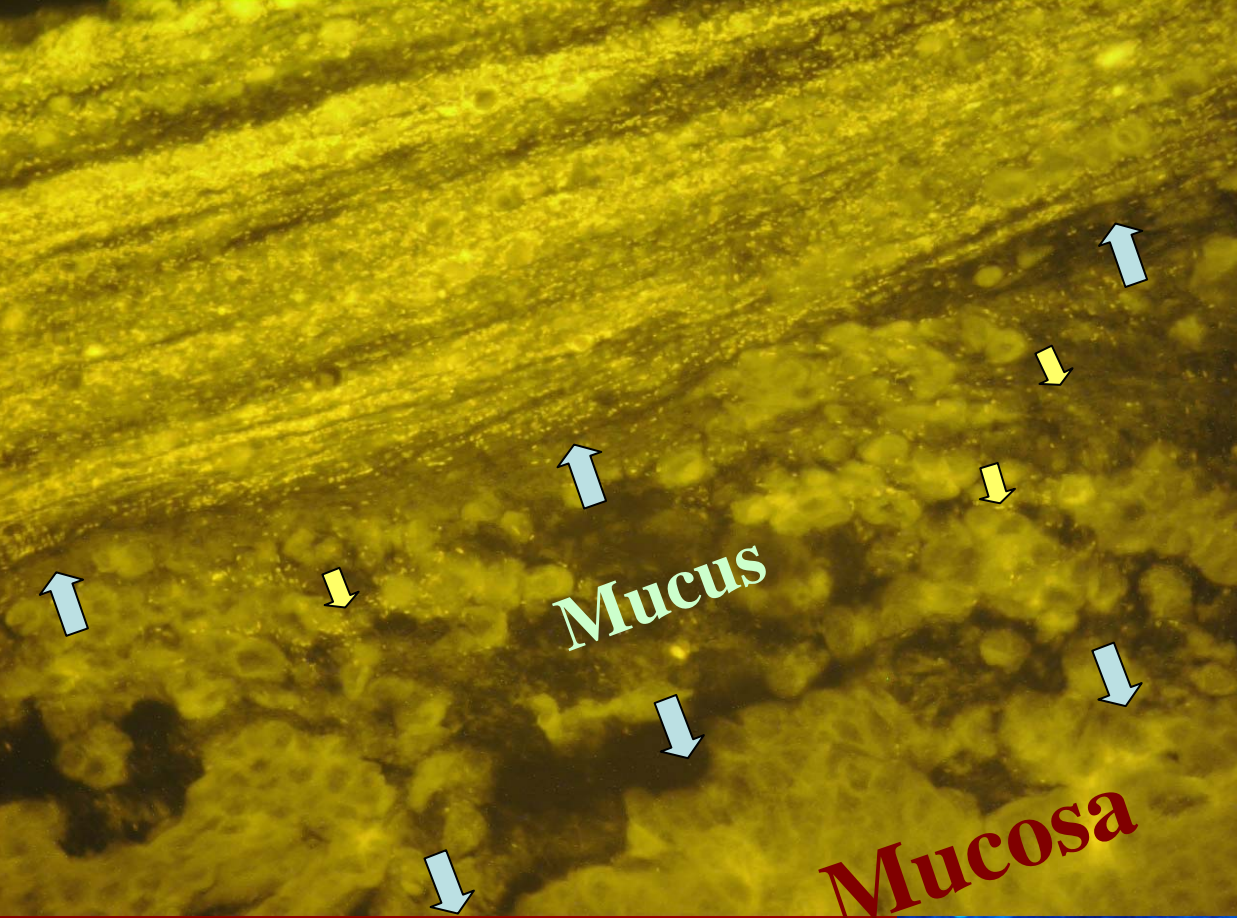
Leukocytes migrate into the lumen of the large intestine (small magnification)



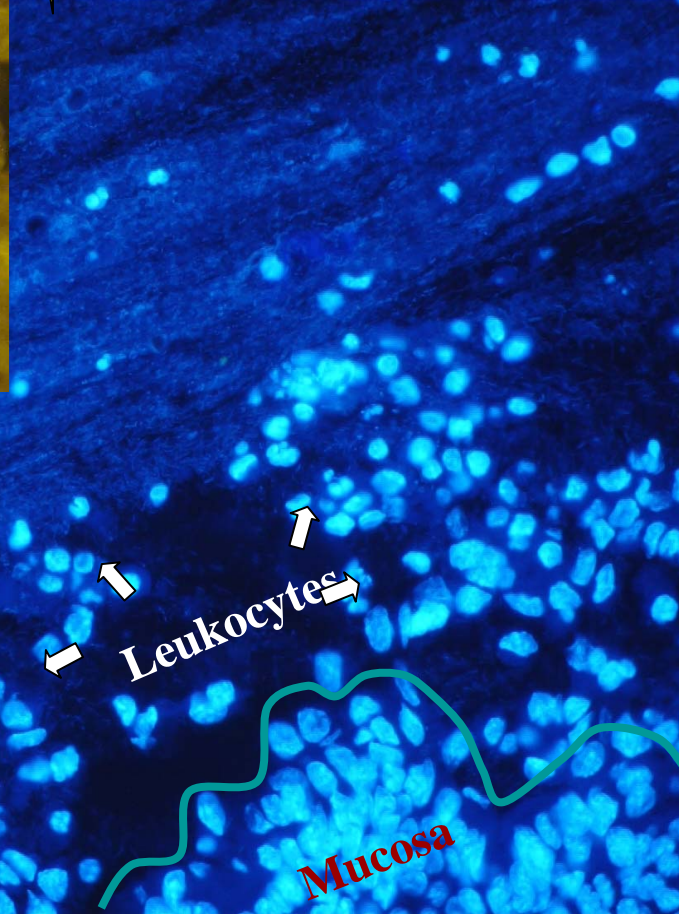


Double headed arrows indicate interlaced layer, blue arrows point out leucocytes

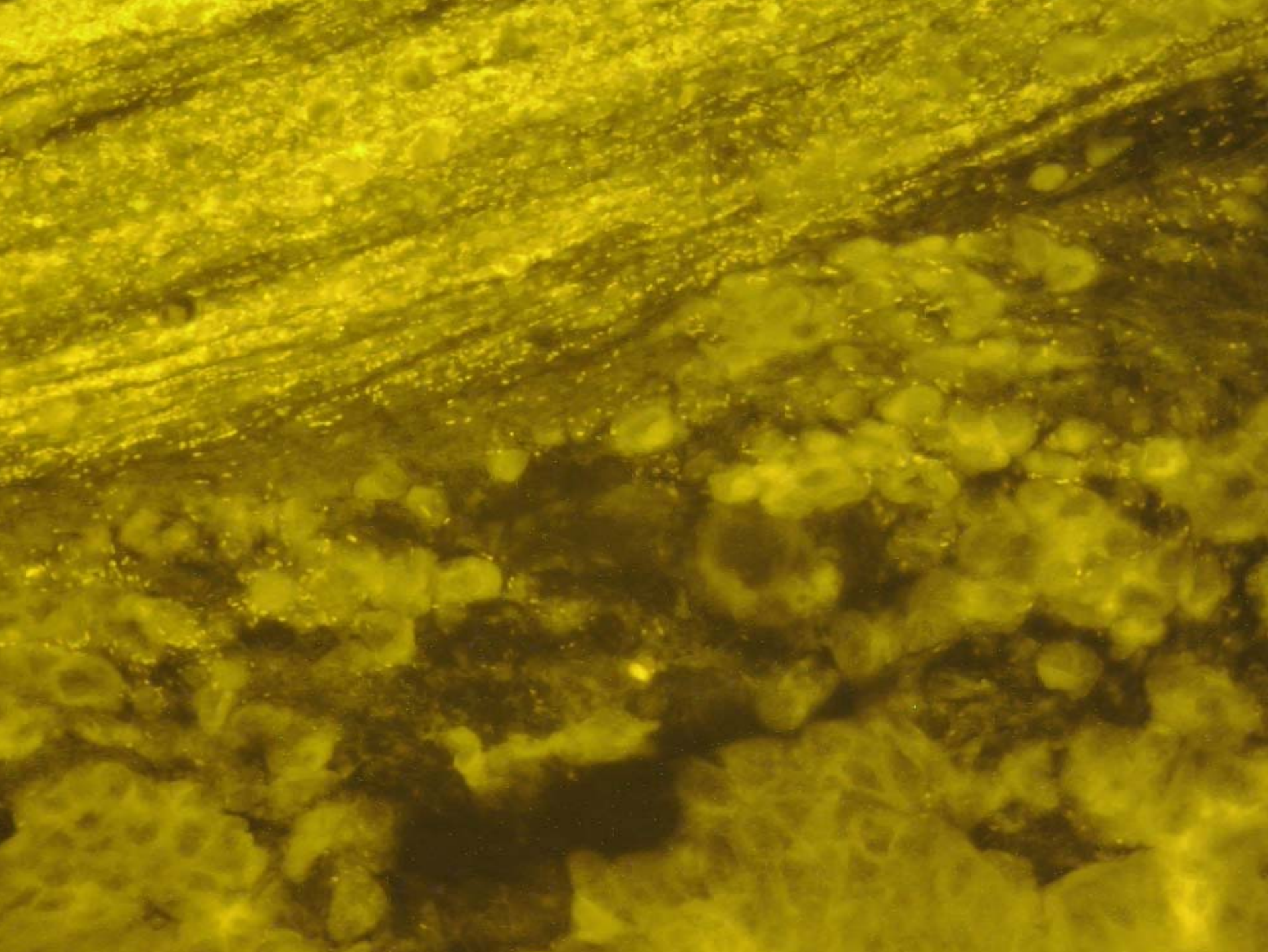




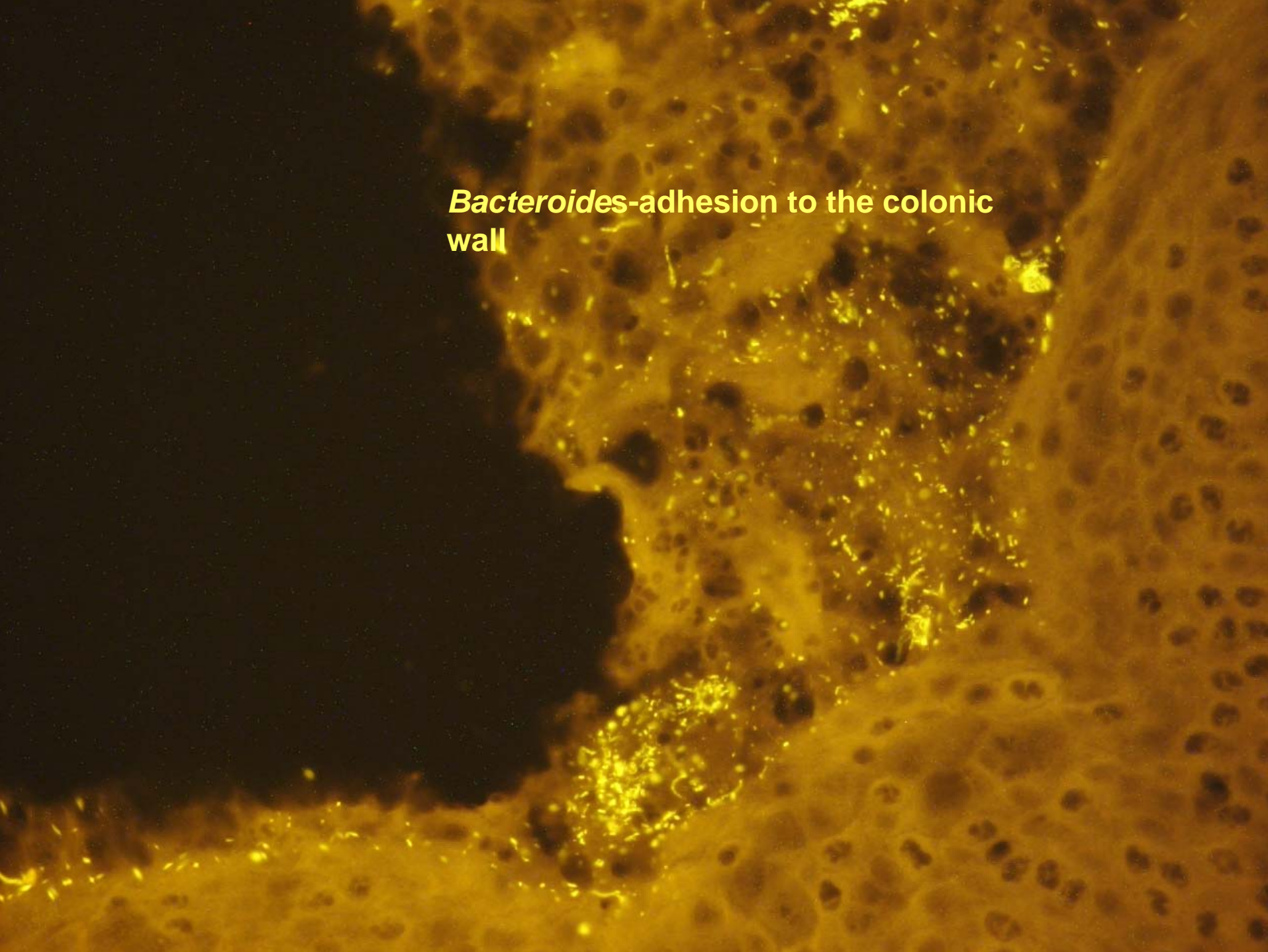
Bacteroides
crosses
mucus



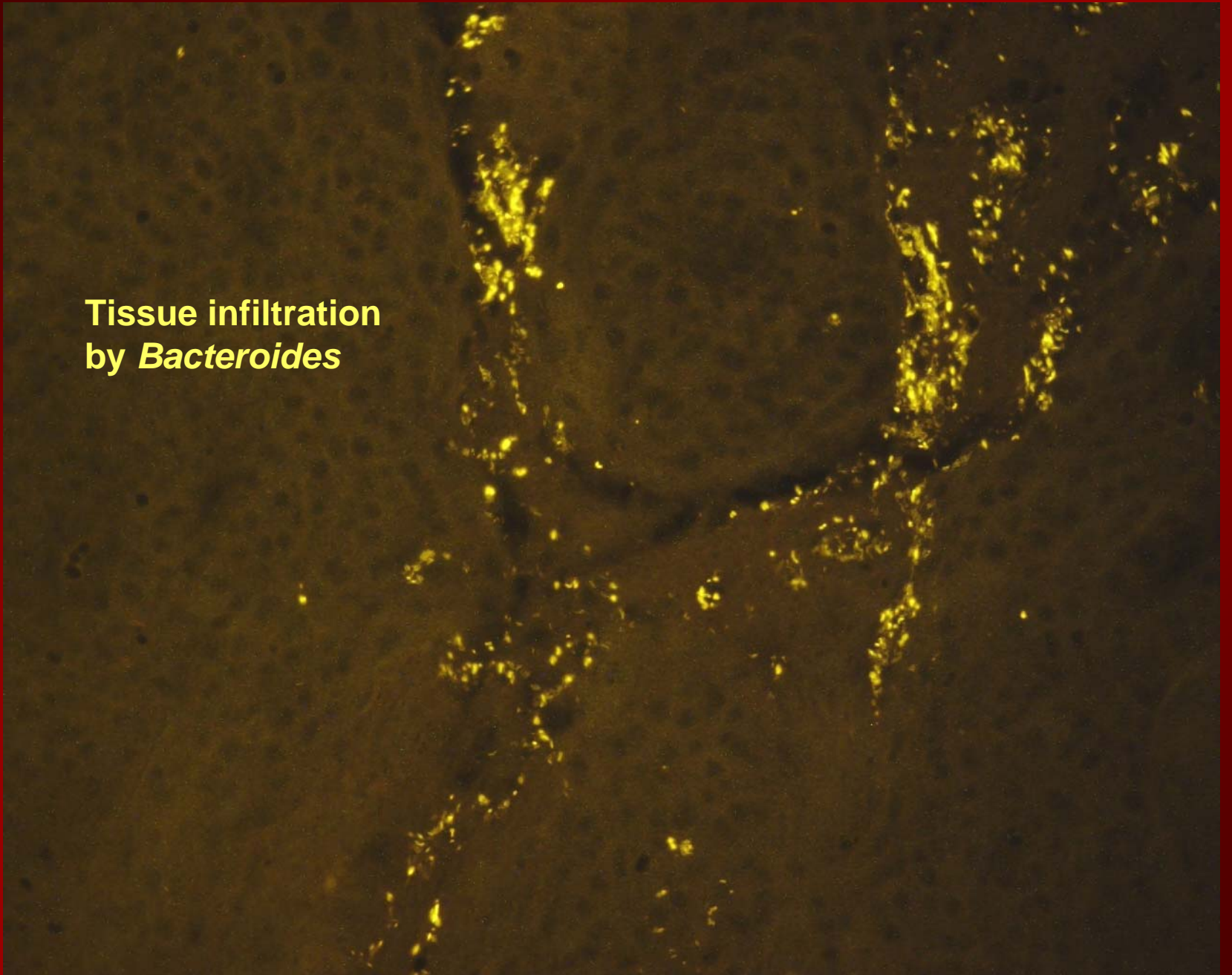
The same microscopic field in **DAPI** shows leukocytes (large blue nuclei) migrating in mucus and hindering *Bacteroides* movement towards mucosa, normally only single leukocytes are present in mucus



***Bacteroides*-adhesion to the colonic wall**



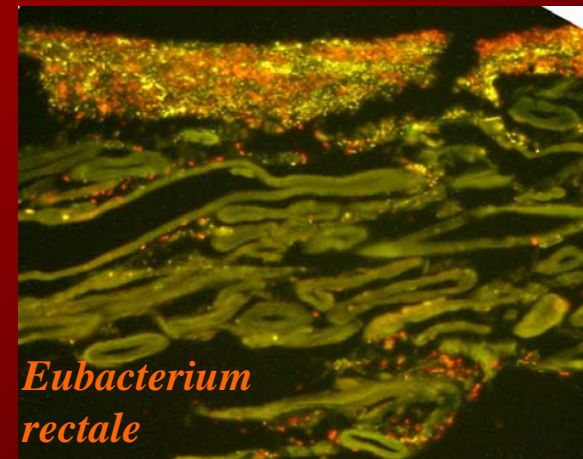
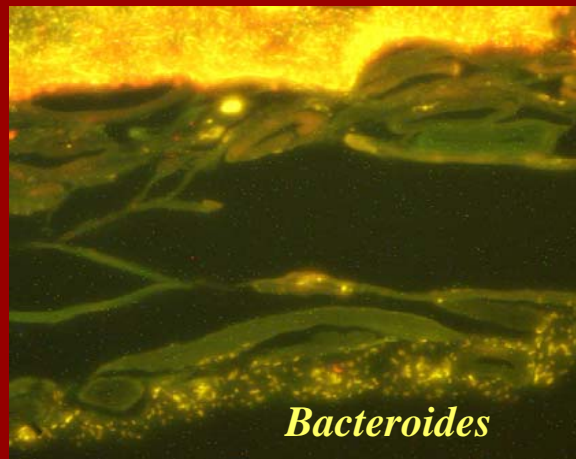
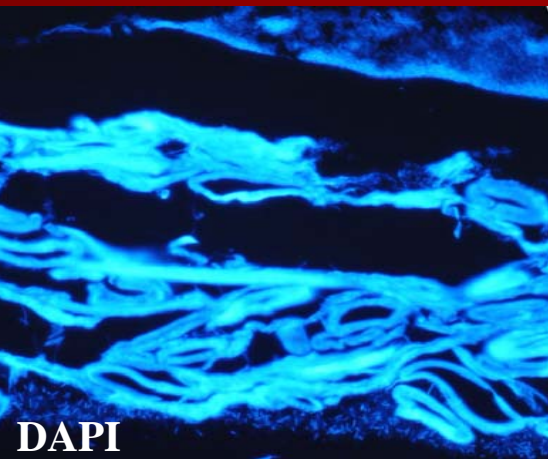
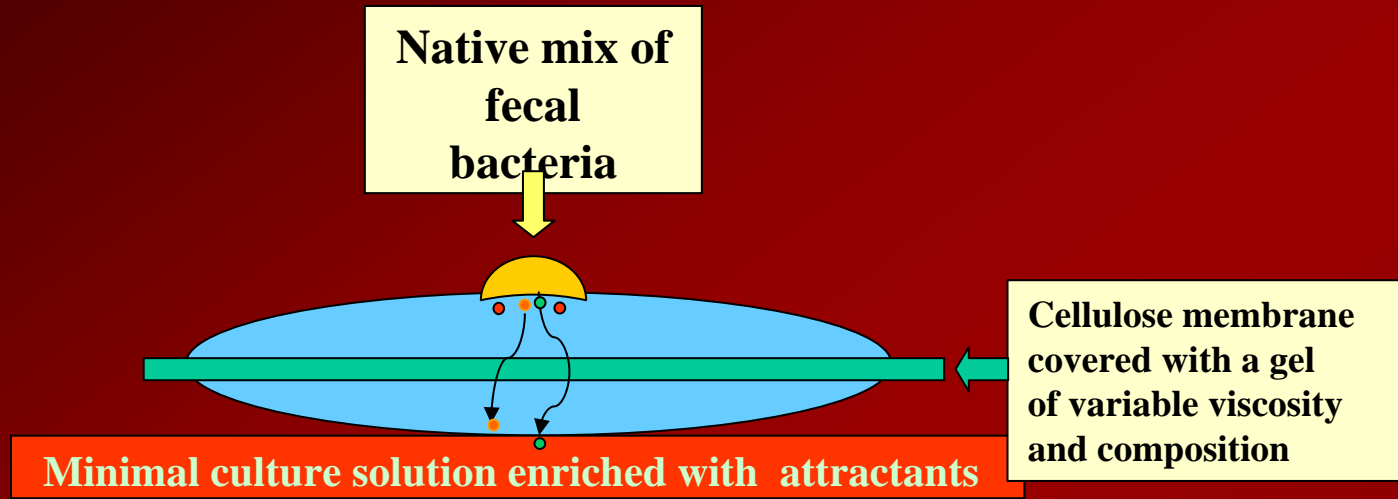
Tissue infiltration
by *Bacteroides*



**How can we explain differences
in distribution of bacteria along
the murine colonic wall?**

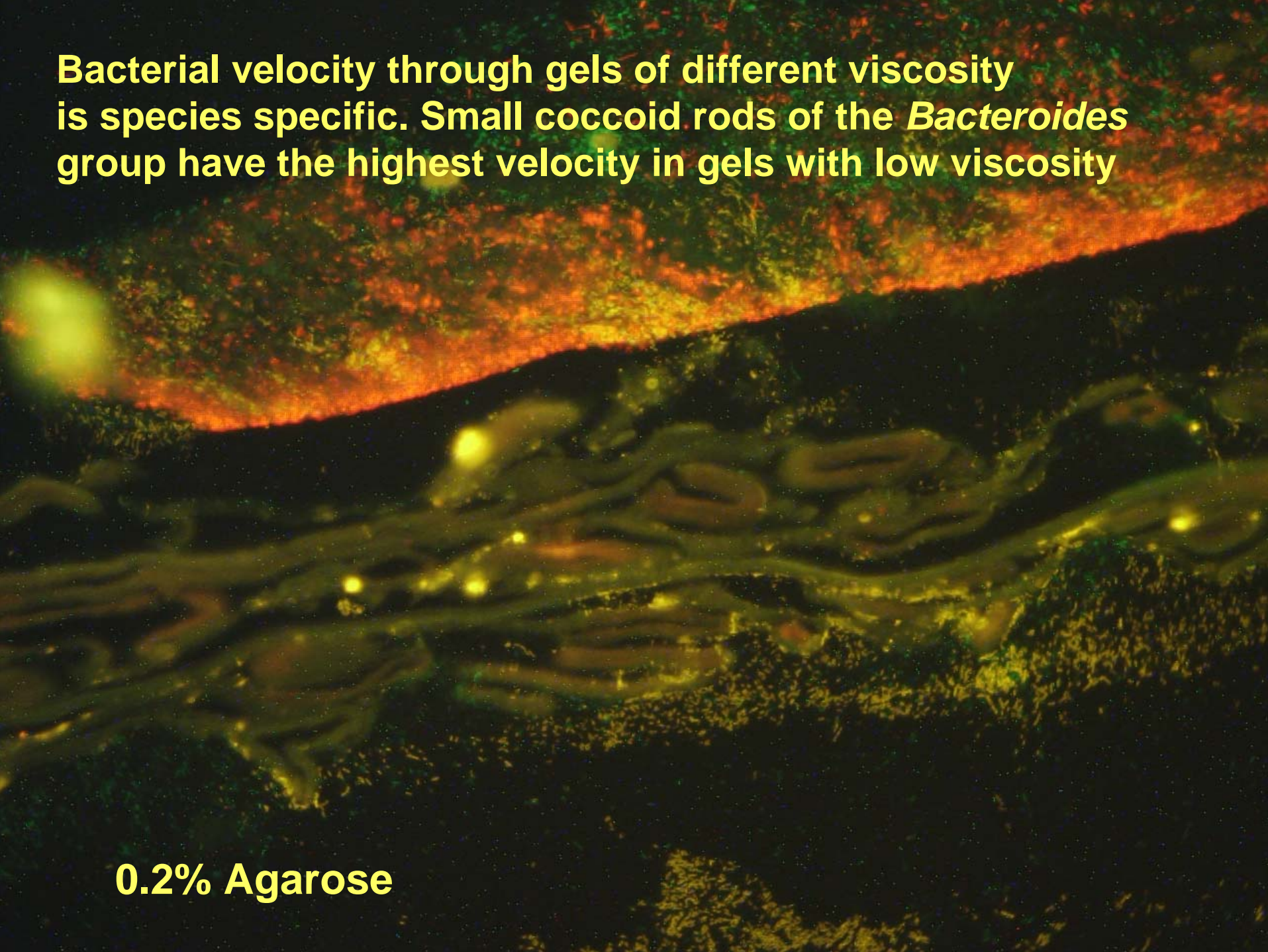
**A full set of figures is available at
www.charite.de/arbmk1**

Mucus simulation in vitro



Examples of mobility

Bacterial velocity through gels of different viscosity is species specific. Small coccoid rods of the *Bacteroides* group have the highest velocity in gels with low viscosity



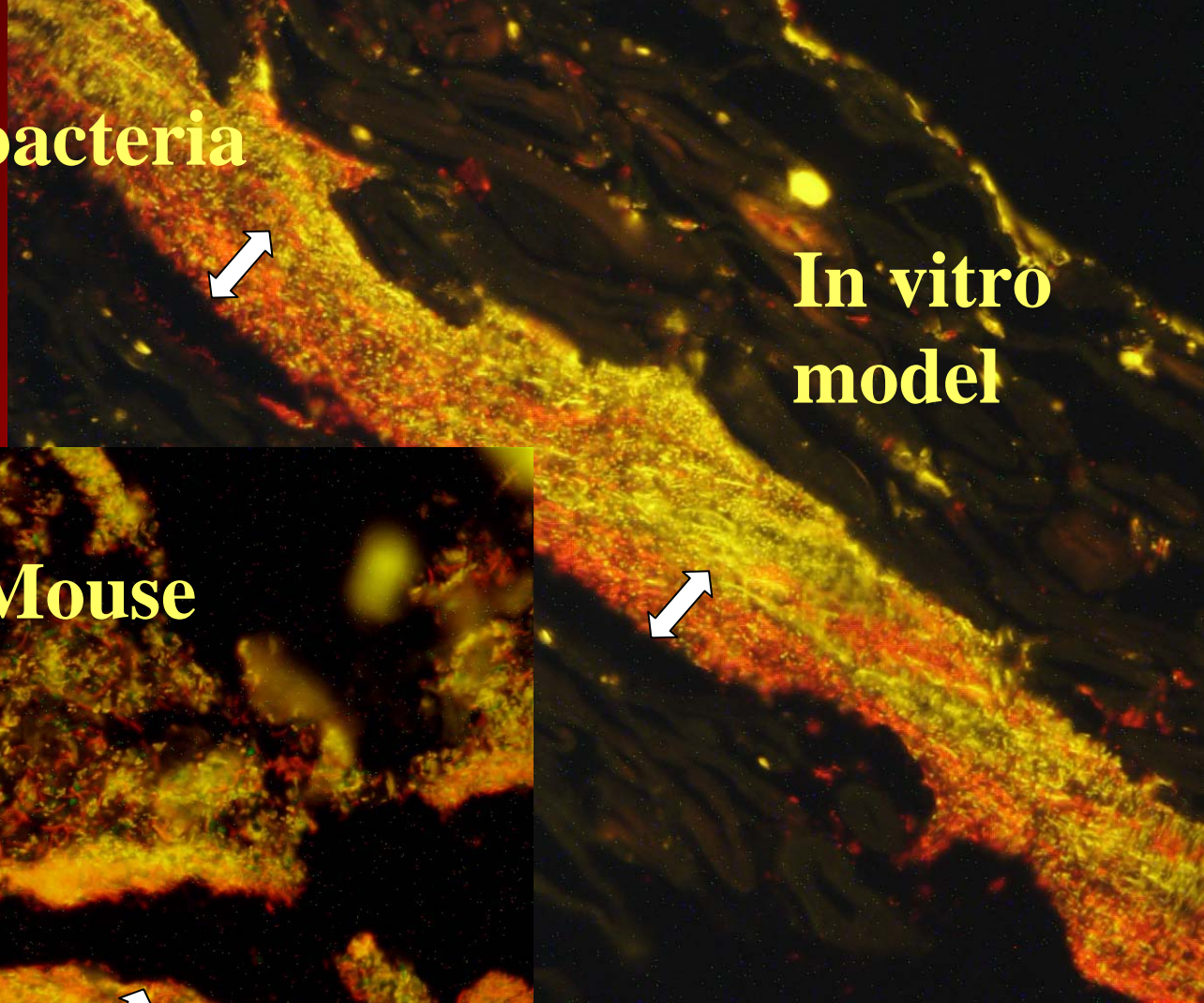
0.2% Agarose

**Long rod of Eubacterium
rectale group (EREC, red)
have the highest velocity
in gels with high viscosity**

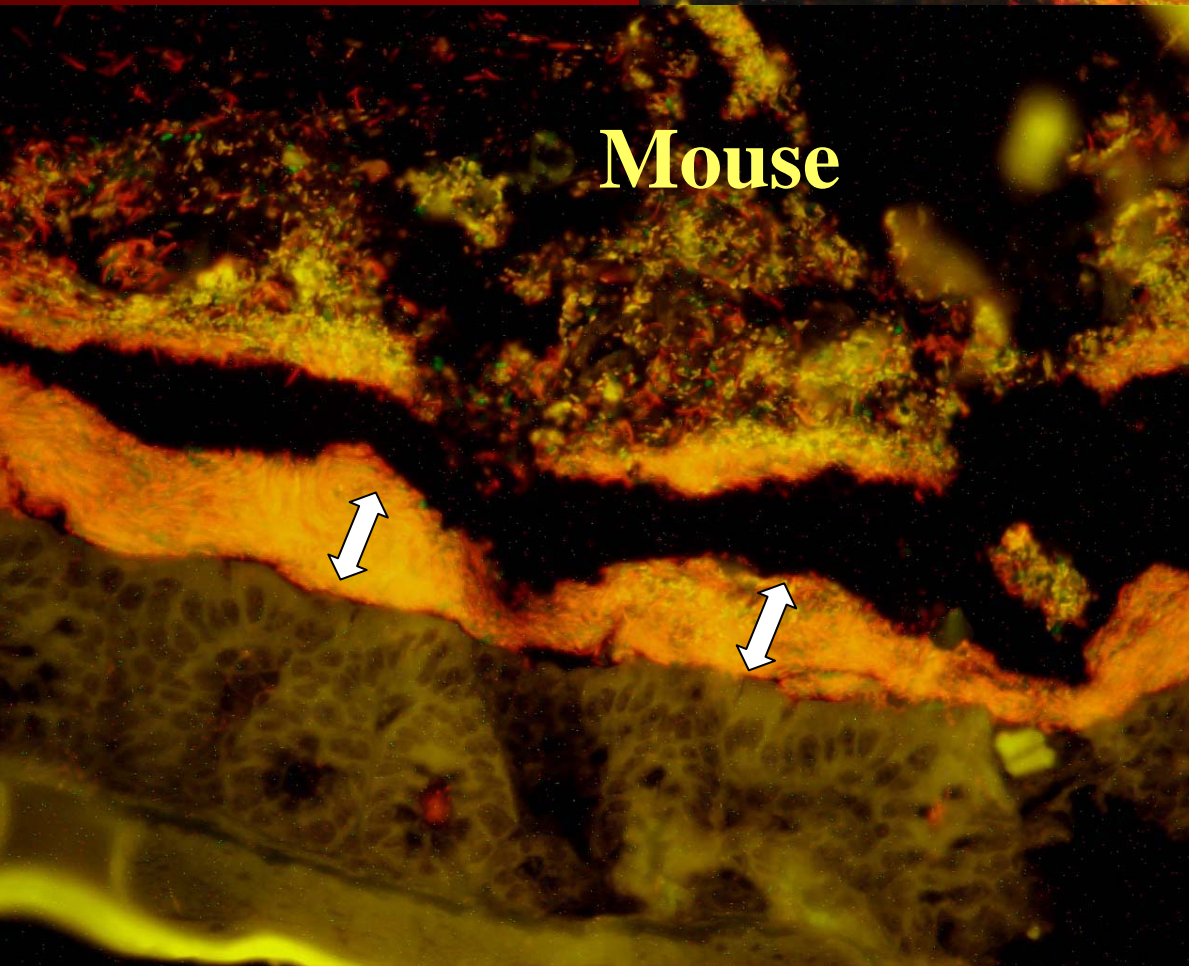
0.5% Agarose

A fluorescence microscopy image showing a bacterial population in a 0.5% agarose gel. The image displays a dense network of long, rod-shaped bacteria, primarily colored red, which are the Eubacterium rectale group (EREC). These rods are oriented in various directions, creating a complex, interconnected pattern. Interspersed among the red rods are smaller, more numerous green and yellow spots, representing other bacterial species. The overall appearance is that of a highly viscous environment where the long rods are the most prominent and likely the fastest-moving component.

**Separation of bacteria
in gels of 0.4%**

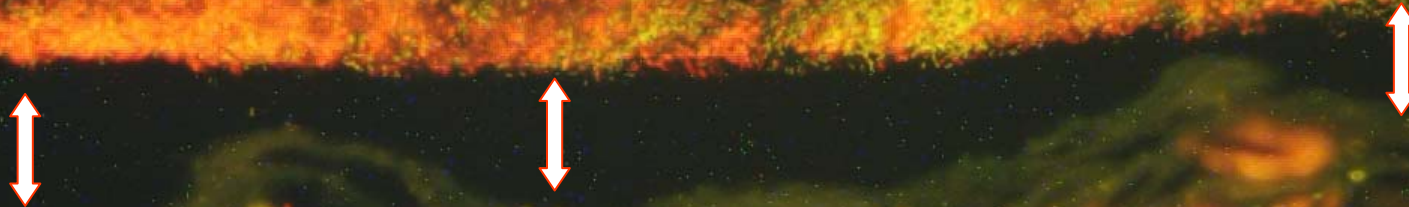


**In vitro
model**



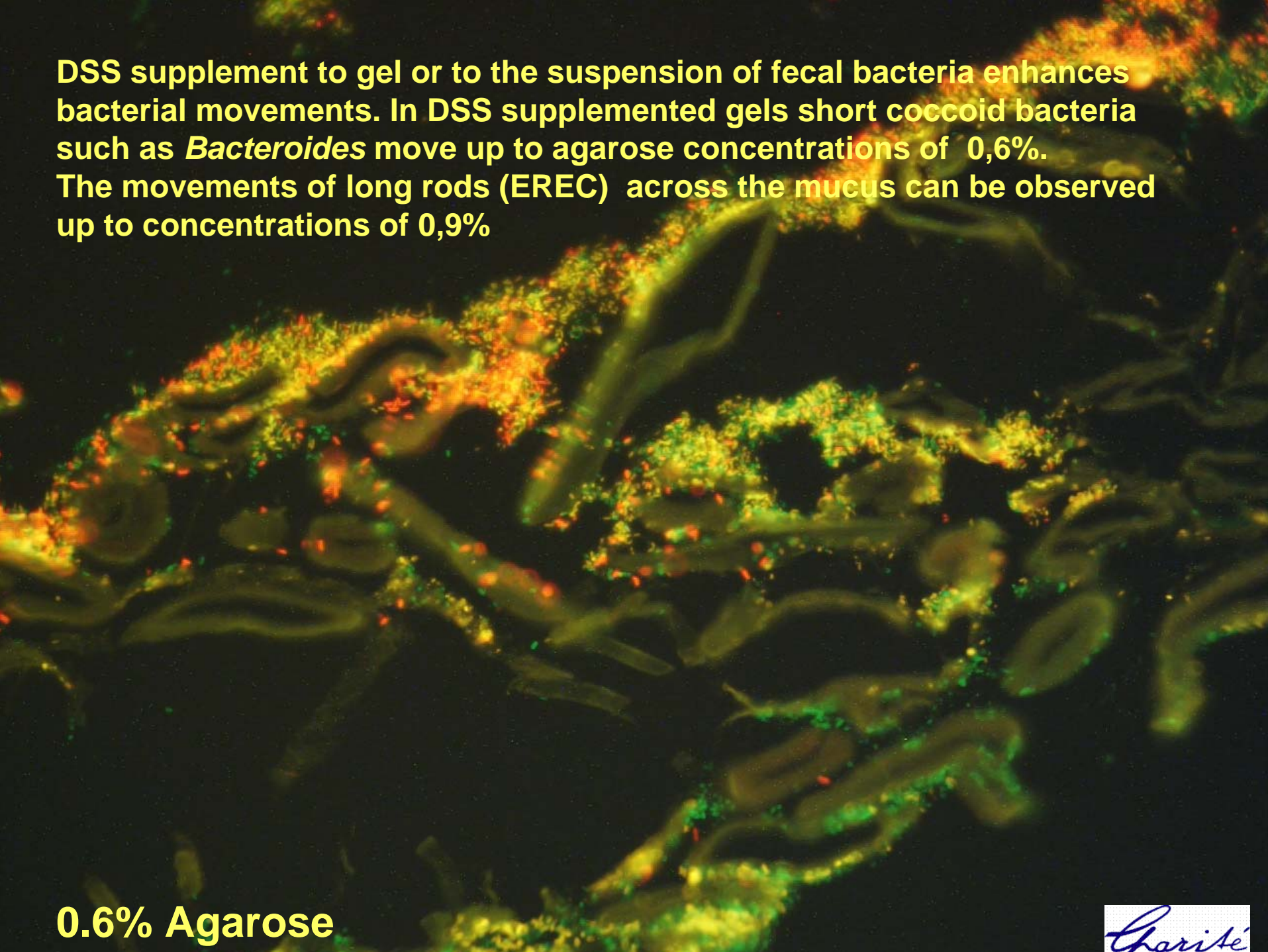
Mouse

0.7% agarose (arrows)



note absence of bacteria below membrane and a gap between bacteria and membrane indicating a lack of bacterial movement across gel layer (double headed arrows)

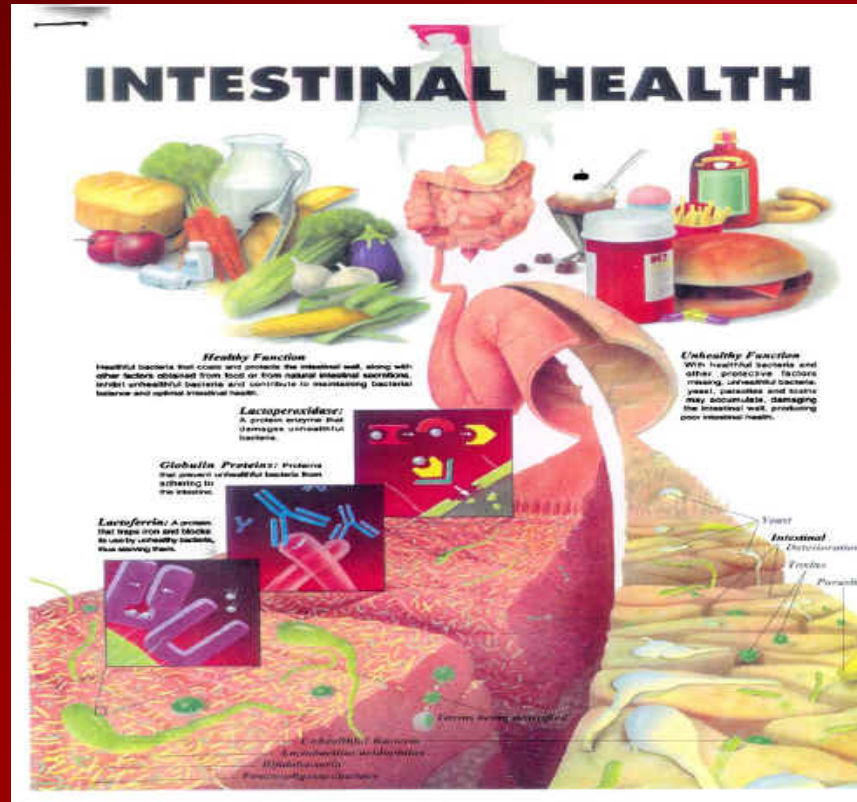
DSS supplement to gel or to the suspension of fecal bacteria enhances bacterial movements. In DSS supplemented gels short coccoid bacteria such as *Bacteroides* move up to agarose concentrations of 0,6%. The movements of long rods (EREC) across the mucus can be observed up to concentrations of 0,9%



0.6% Agarose

Tolerance

normal
Flora



Inflammatory

Response

Enteral
Pathogens



E. coli

Bacteroides

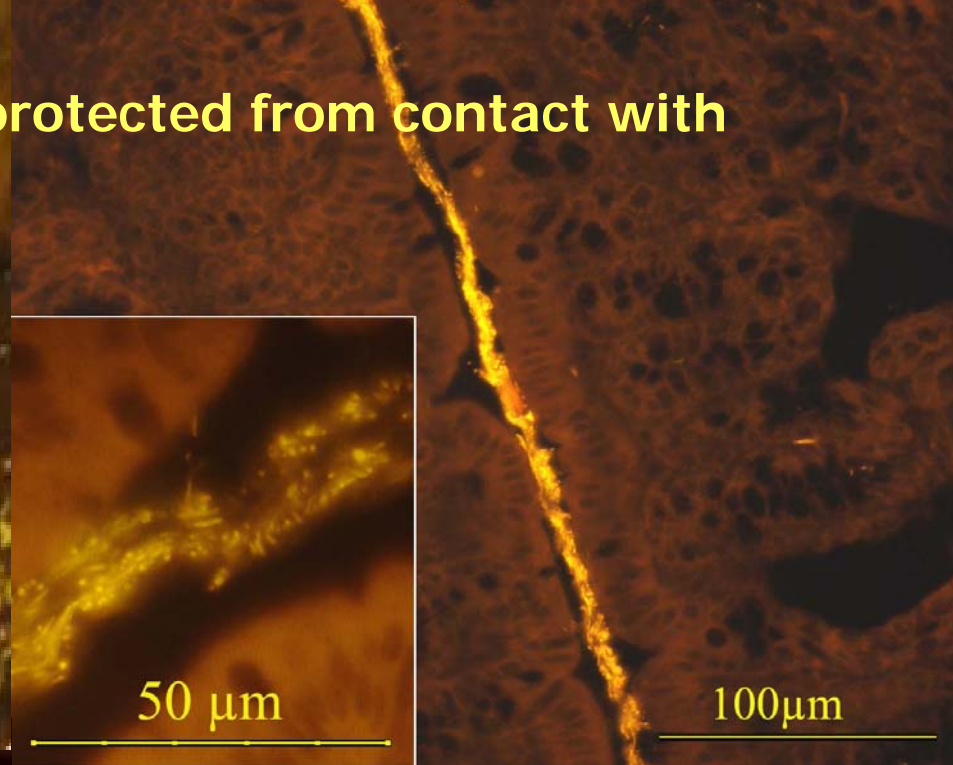
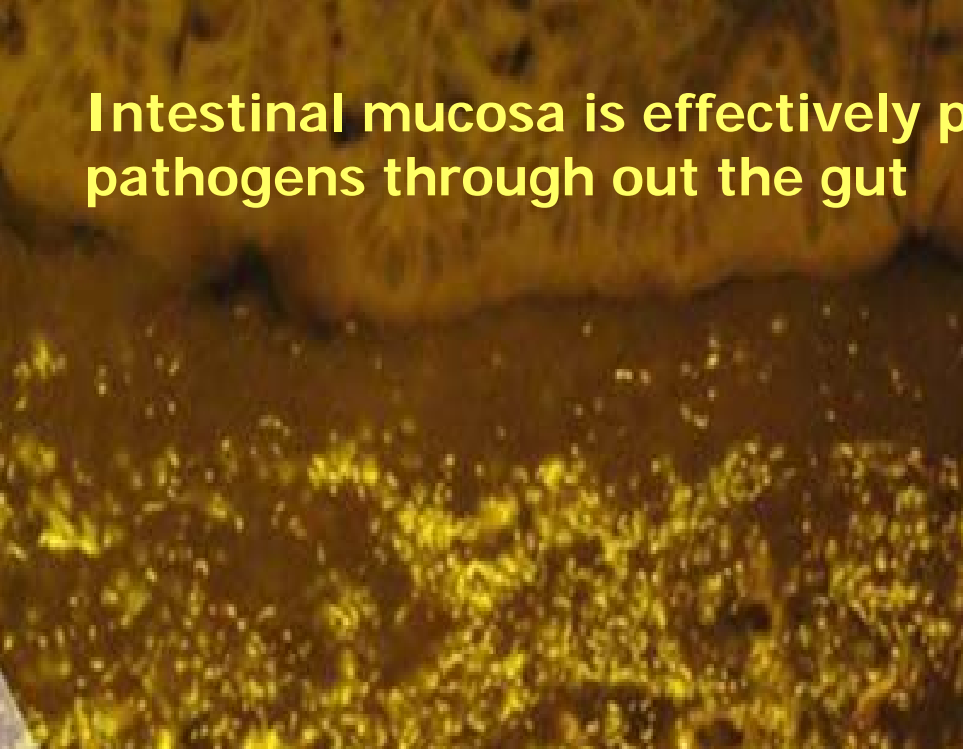
Clostridium difficile

Enterococci

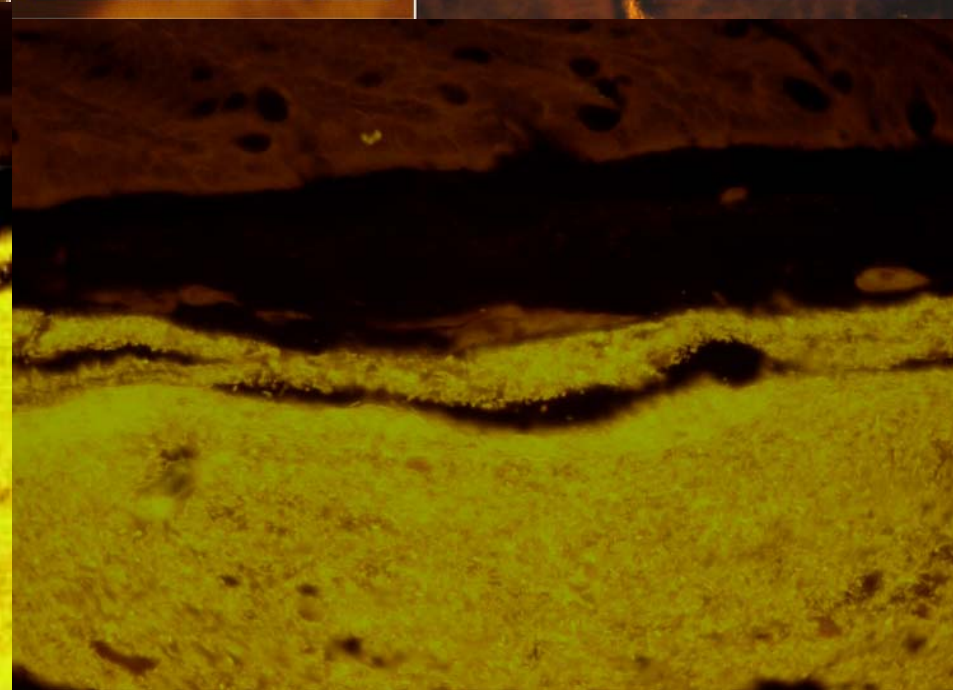
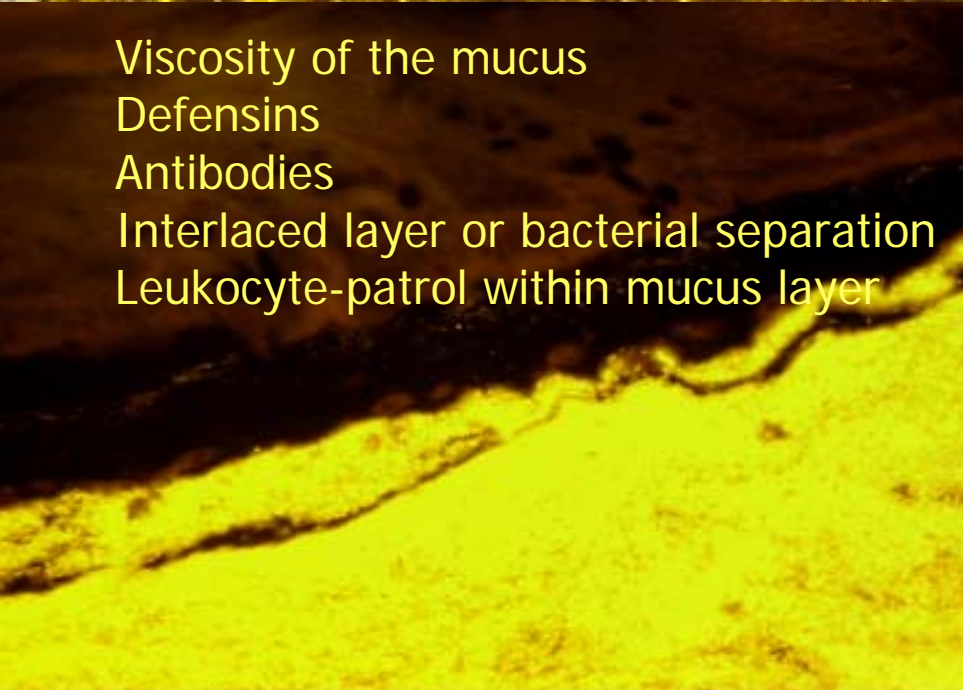
Salmonella

Shigella

Intestinal mucosa is effectively protected from contact with pathogens through out the gut



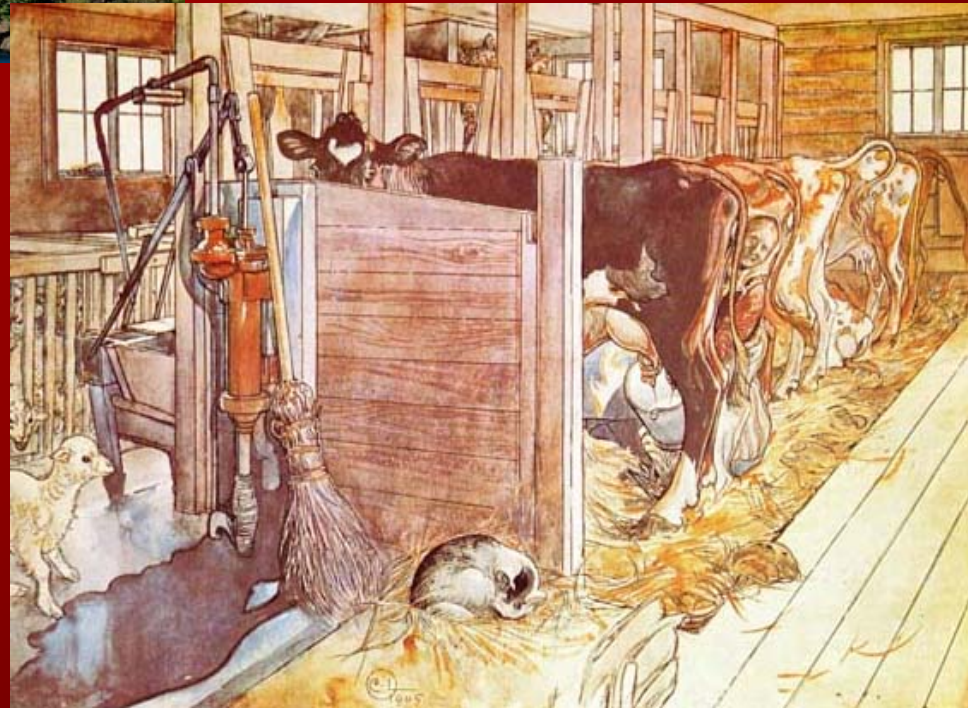
Viscosity of the mucus
Defensins
Antibodies
Interlaced layer or bacterial separation
Leukocyte-patrol within mucus layer

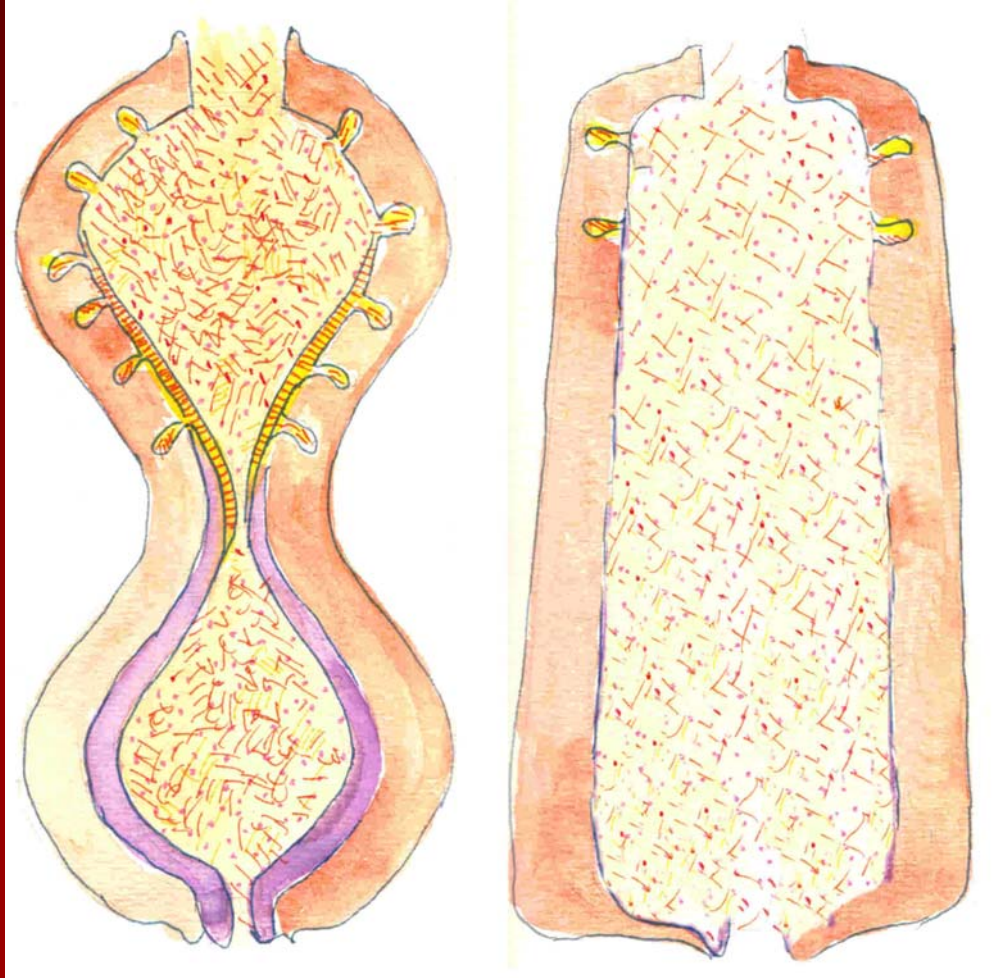




Hygiene hypothesis

GO
BACK!





Soaps and emulsifying substances make our environment clean. They may however have the same effect on the mucus of man as DSS on the mucus of mouse.

Factors affecting mucus barrier

Exogenic:

Detergents:

Bacterial virulence:

Glutens as natural emulsifiers need bacteria to be pathogenetic

Smoking

Endogenic:

Bile acids are normally fully resorbed in ileum but lead to diarrhea if arrive in large intestine

Defensins, Antibodies draining

Probiotics, Prebiotics,

Oligonucleotids Nucleinacidsderivates

Inflammatory response

Genetic

NOD 2 Mutation

[E425](#), Konjak

[E432 bis E436](#), Polysorbat

- E432, Polyoxyethylen-sorbitan-monolaurat (Polysorbat 20)
- E433, Polyoxyethylen-sorbitan-monooleat (Polysorbat 80)
- E434, Polyoxyethylen-sorbitan-monopalmitat (Polysorbat 40)
- E435, Polyoxyethylen-sorbitan-monostearat (Polysorbat 60)
- E436, Polyoxyethylen-sorbitan-tristearat (Polysorbat 65)

[E440](#), Pektine, Amidiertes Pektin

[E442](#), Ammoniumsalze von Phosphatidsäuren

[E444](#), Saccharose-acetat-isobutyrat

[E445](#), Glycerinester aus Wurzelharz/Kolophonester

[E450 bis E452](#), Phosphate

[E459](#), Beta-Cyclodextrin

[E460 bis E469](#) Cellulose und Celluloseverbindungen

- E460, Cellulose, Mikrokristalline Cellulose, Cellulosepulver
- E461, Methylcellulose
- E463, Hydroxypropylcellulose
- E464, Hydroxypropylmethylcellulose
- E465, Ethylmethylcellulose
- E466, Carboxymethylcellulose, Natriumcarboxymethylcellulose
- E468, Vernetzte Natrium-Carboxymethylcellulose
- E469, Enzymatisch hydrolysierte-Carboxymethylcellulose
- [E470a und E470b](#), Salze von Speisefettsäuren
- E470a, Natrium-, Kalium- und Calciumsalze von Speisefettsäuren
- E470b, Magnesiumsalze von Speisefettsäuren
- [E471 bis E472f](#), Mono- und Diglyceride von Speisefettsäuren
- E471, Mono- und Diglyceride von Speisefettsäuren, Monoglycerid
- E472a, Essigsäureester von Mono- und Diglyceriden von Speisefettsäuren
- E472b, Milchsäureester von Mono- und Diglyceriden von Speisefettsäuren
- E472c, Citronensäureester von Mono- und Diglyceriden von Speisefettsäuren
- E472d, Weinsäureester von Mono- und Diglyceriden von Speisefettsäuren
- E472e, Mono- und Diacetylweinsäureester von Mono- und Diglyceriden von Speisefettsäuren
- E472f, Gemischte Essig- und Weinsäureester von Mono- und Diglyceriden von Speisefettsäuren

[E473](#), Zuckerester von Speisefettsäuren

[E474](#), Zuckerglyceride

[E475](#), Polyglycerinester von Speisefettsäuren, Polyglycerinester

[E476](#), Polyglycerin-Polyricinoleat

[E477](#), Propylenglycolster von Speisefetten

[E479](#), Thermooxidiertes Sojaöl mit Mono- und Diglyceriden von Speisefettsäuren

[E481 bis E483](#), Natriumstearoyl-2-lactylat, Calciumstearoyl-2-lactylat, Stearyltrarat

[E491 bis E495](#), Stearin- und Palmitatverbindungen

[E491](#), Saccharosemonoacetat

Conclusions:

The intestinal wall is protected from contact with potentially harmful bacterial groups such as *Bacteroides*, *Enterobacteriaceae*, *Enterococci*, and *Clostridium difficile*, despite extremely high bacterial concentrations in colon.

A mucus barrier and not the epithelial cell layer is the first line of defense against a variety of enteral pathogens.

Inflammatory bowel disease is a polymicrobial infection that is characterized by a sustained broken mucus barrier with subsequent bacterial migration toward mucosa and proliferation of complex bacterial biofilms on the epithelial surface.

As long as the mucus barrier function is impaired, the inflammatory process cannot successfully clear bacteria from the mucosal surface and is harmful.

The rising incidence of IBD over the last century may result from changes in the types and numbers of bacteria within the intestine, growing bacterial burden, and disturbed mucus barrier function.

Further study of how viscosity, defensins, antibodies, antibiotics, probiotic bacteria, leukocytes, and other factors affect mucus barrier function will allow to identify new ways to prevent, treat ulcerative colitis and Crohn's disease.

Ulcerative colitis and Crohn's disease are curable