

Epitheliocystis disease; results and progress

Task 25.1, D25.4

DIVERSIFY Meeting, Tenerife 2018

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INSTITUTE OF MARINE BIOLOGY, BIOTECHNOLOGY AND AQUACULTURE

HELLENIC CENTRE FOR MARINE RESEARCH

Epitheliocystis

Infectious disease affecting a wide range of wild and cultured fish

Global distribution

First observed in 1920

Described and named in 1969

Caused by intracellular pathogens

Inclusions in gill and skin epithelium of the fish

Despite efforts, no epitheliocystis-related agent has been isolated in culture until today

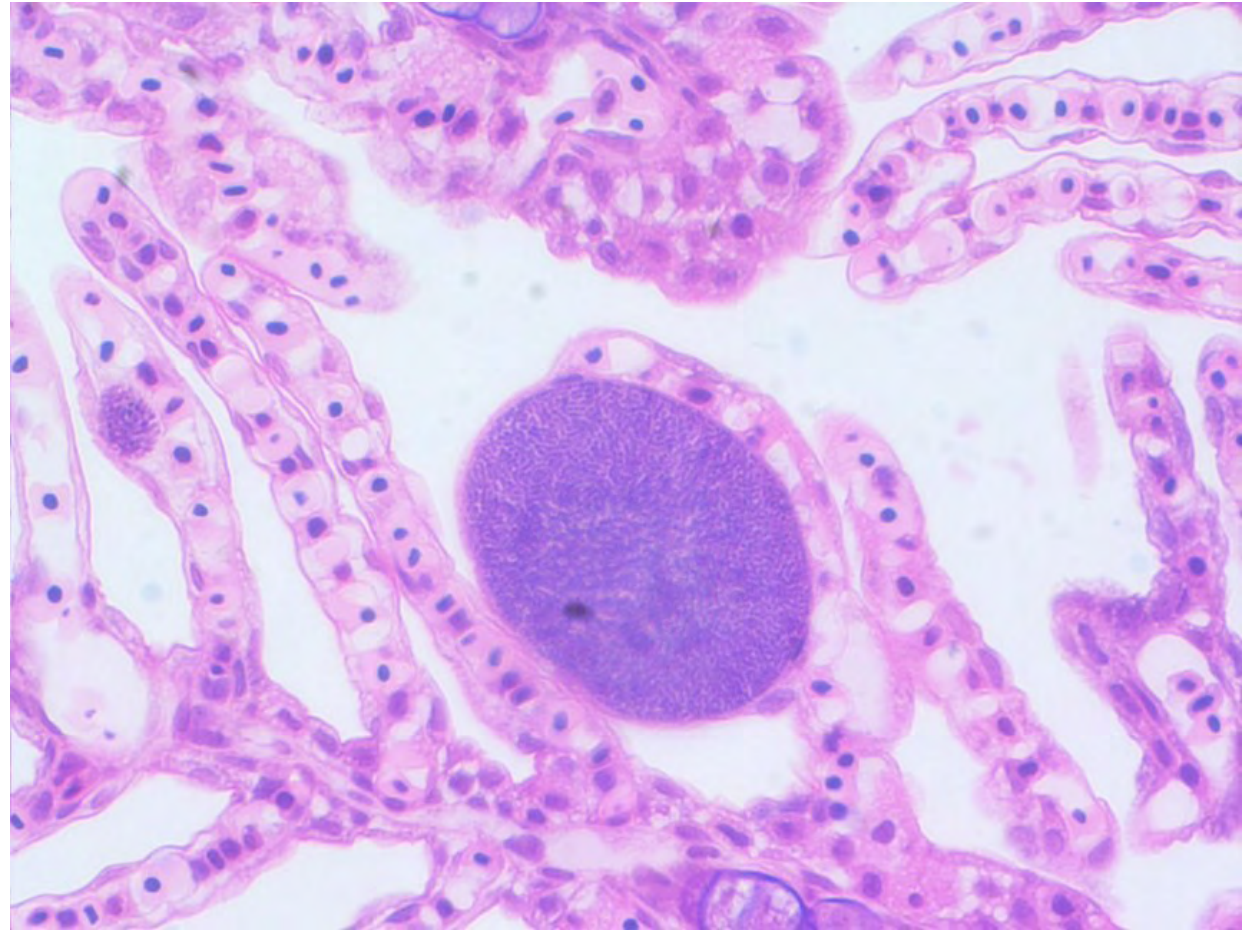
Diagnosis

Easy !

Wet mounts

Histology

PCR



Two stereotypes about epitheliocystis

It is a benign disease

It is caused by Chlamydia

Epitheliocystis and HCMR. A recurrent problem

100% mortality in greater amberjack (*Seriola dumerili*) larvae

80% mortality in common dentex (*Dentex dentex*) larvae

>50% mortality in sharpsnout seabream (*Diplodus puntazzo*)

In some cases 100% mortality overnight

Experimental investigation

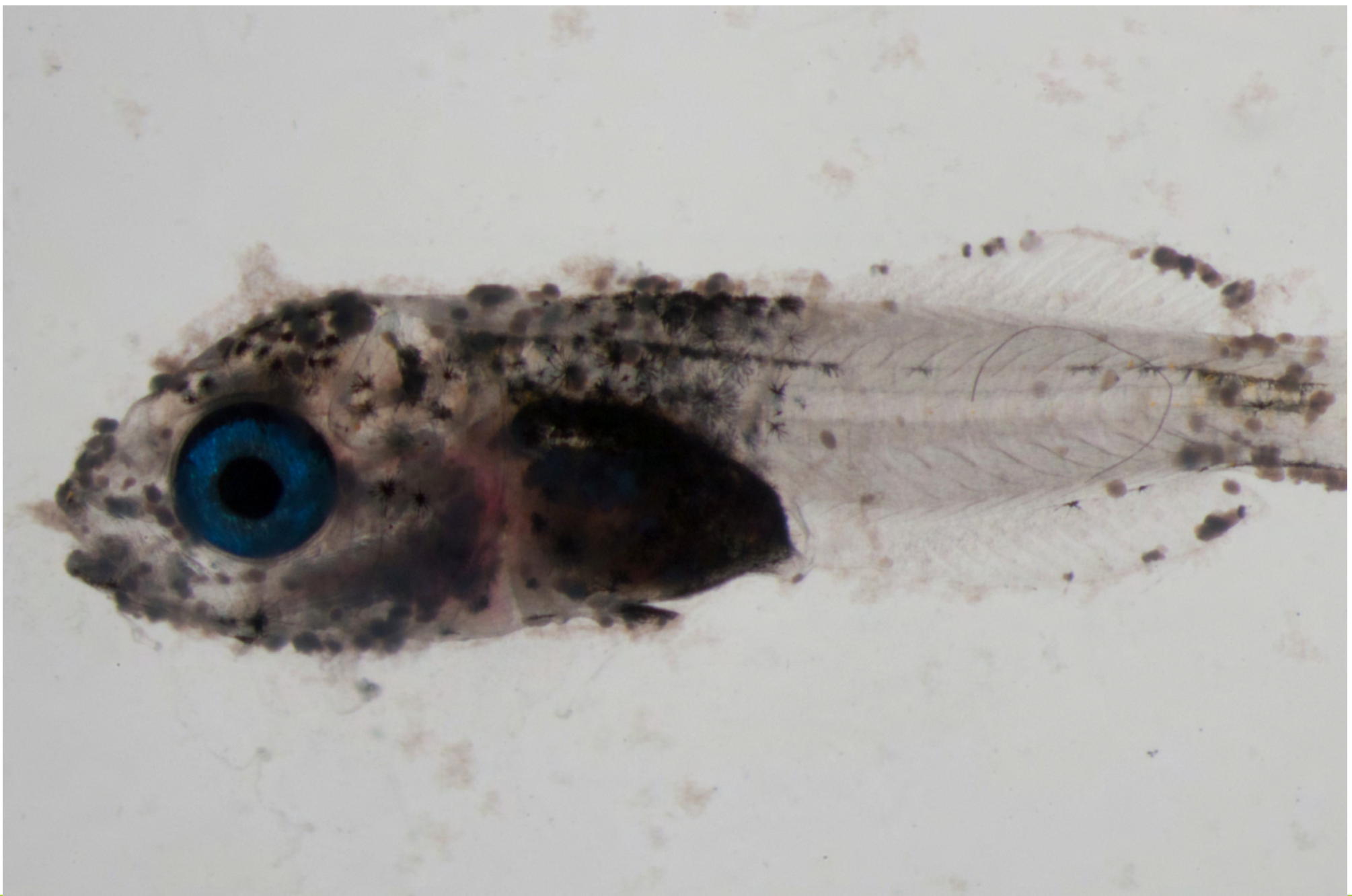
Aquaexcel Project

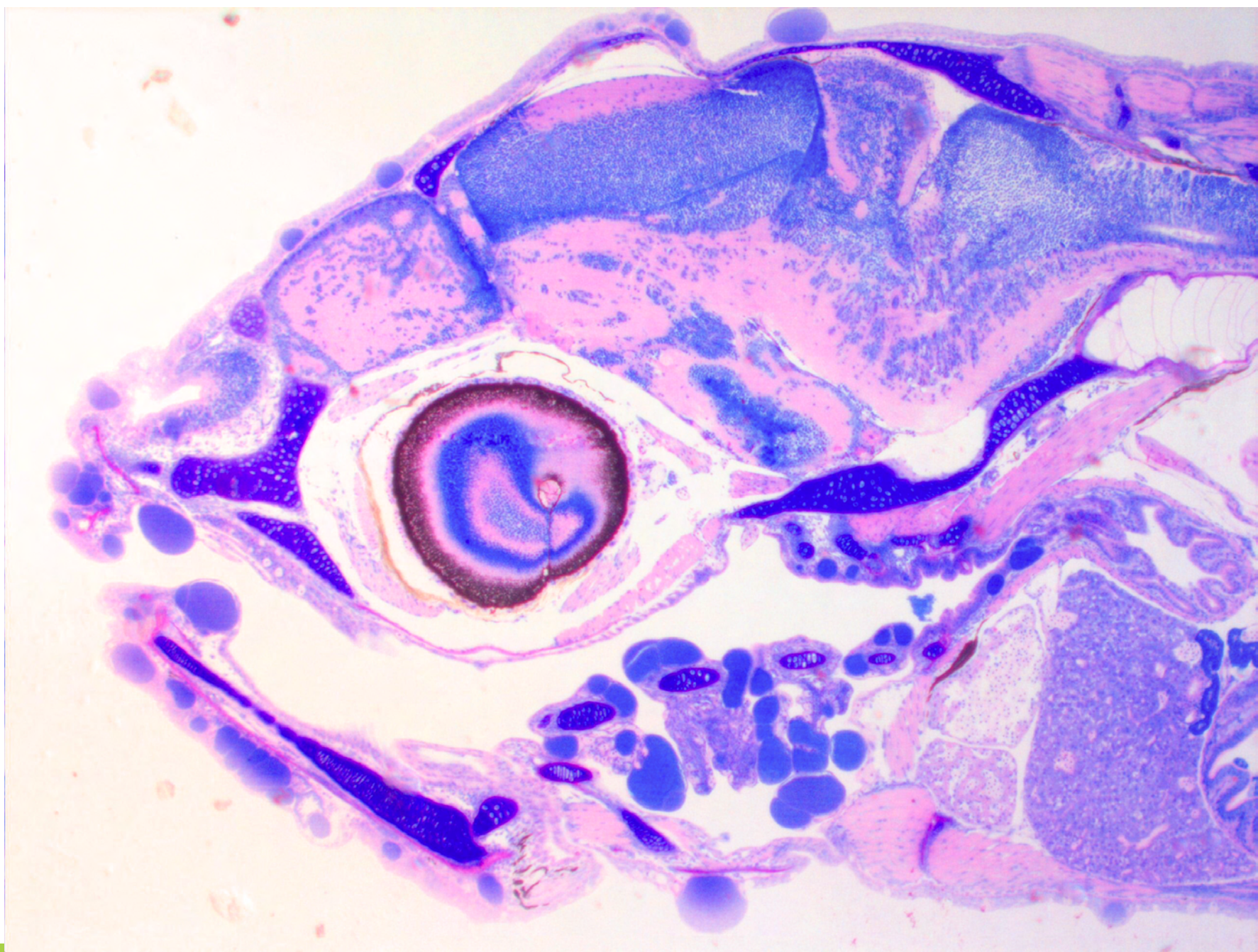
Prof. Lloyd Vaughan, Veterinary School, University of Zurich

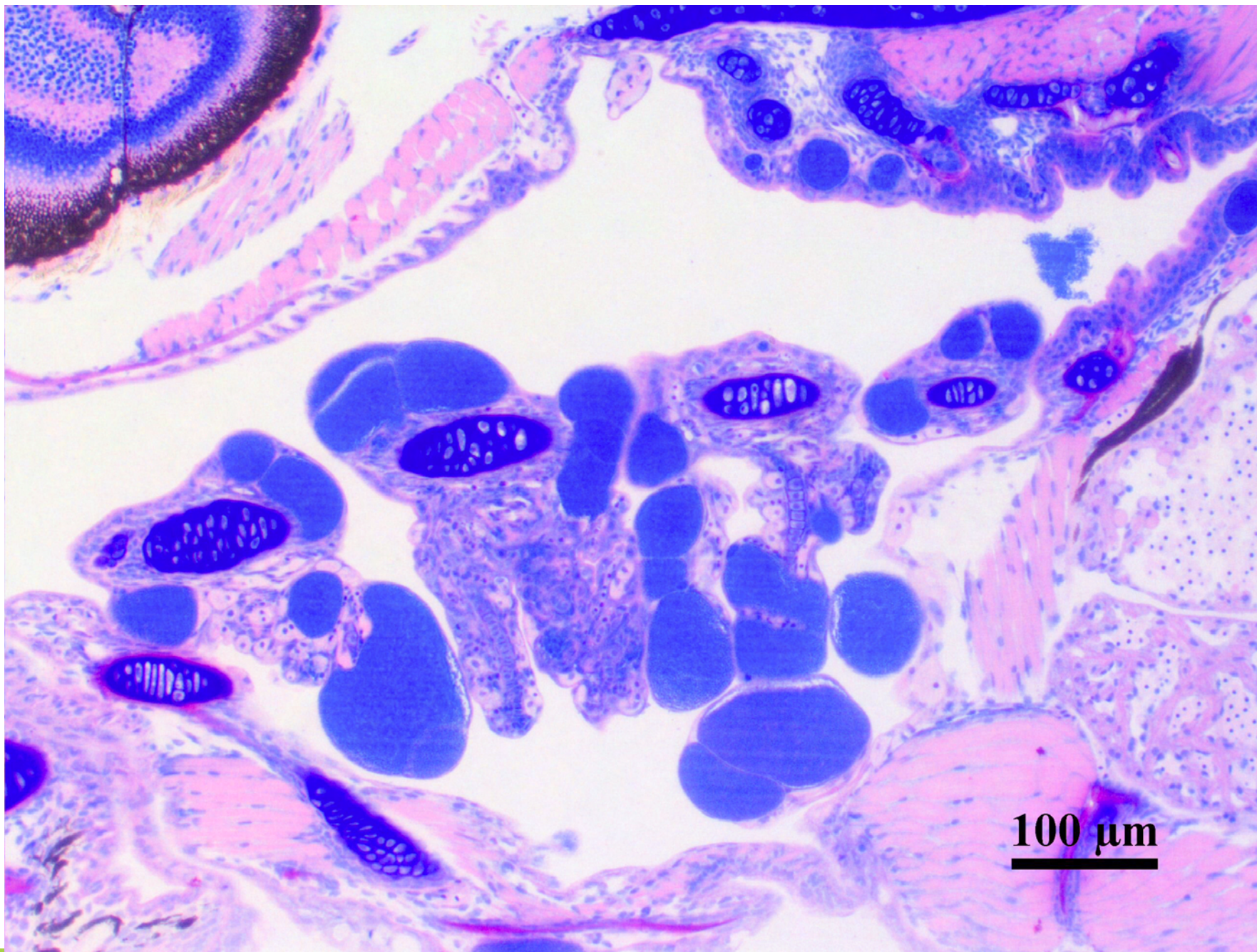
Mesocosm culture of sharpsnout seabream– Natural infection

Plan B: collect samples (gilthead seabream) from Greek commercial fish farms

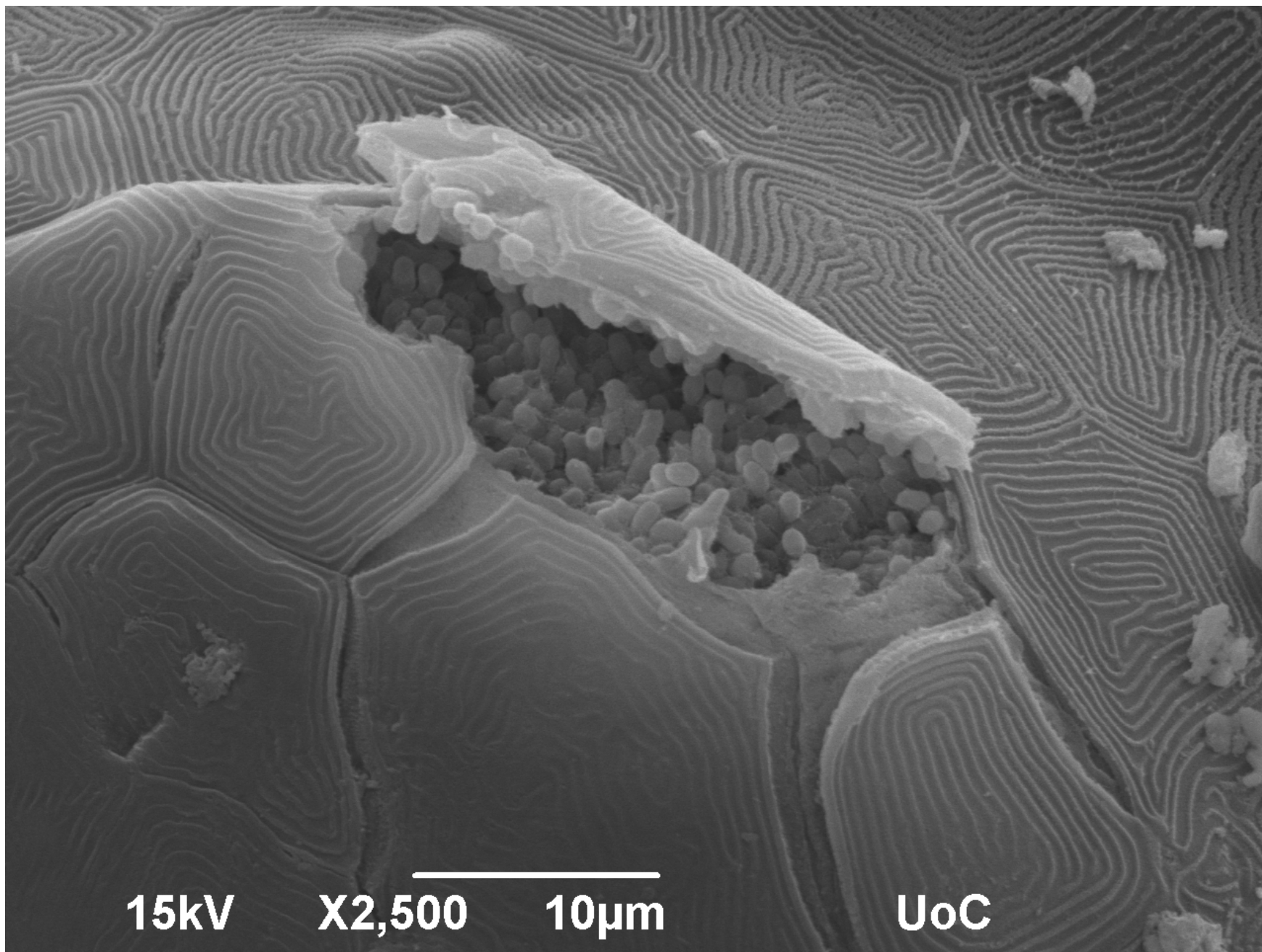
October-November 2012







100 μm

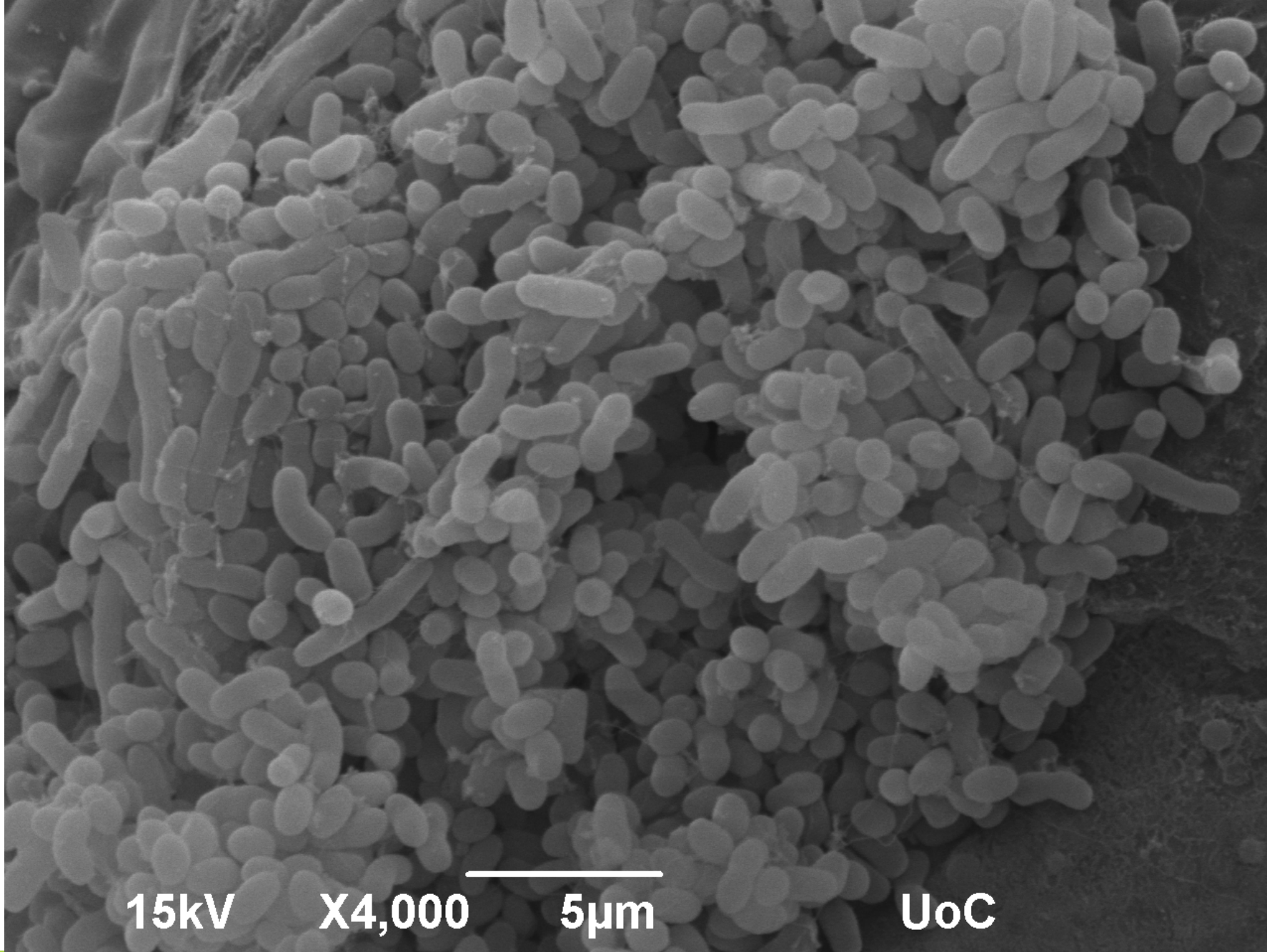


15kV

X2,500

10 μ m

UoC



15kV

X4,000

5µm

UoC

Identification pipeline

PCR for chlamydia positive

Sequencing pointed to known chlamydia pathogens

In situ for chlamydia was negative

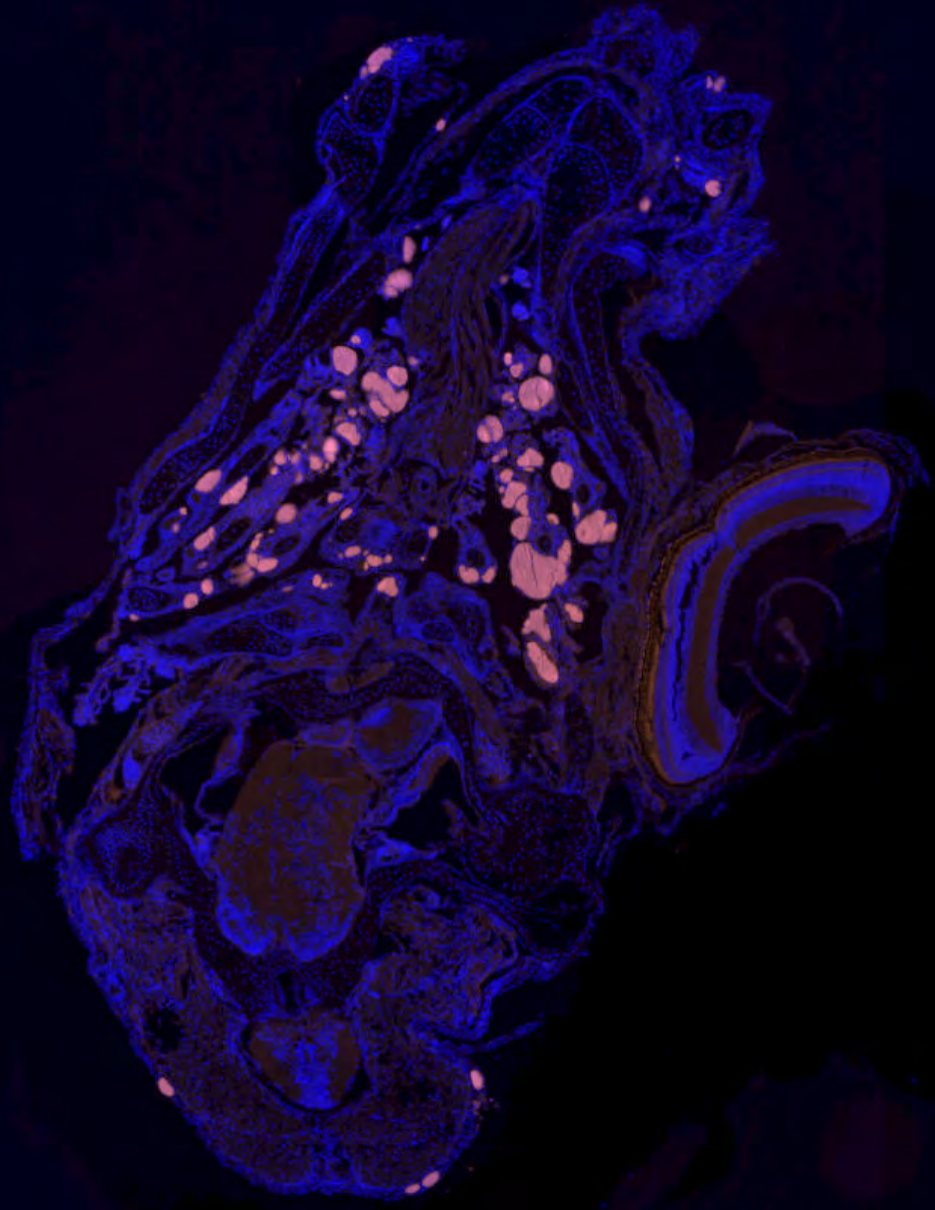
DNA extraction from single cysts

PCR 16s followed by sequencing

Endozoicomonas sp.

Construction of probes for *Endozoicomonas*

New in situ



Novel pathogen

Candidatus Endozoicomonas cretensis n.sp

γ -proteobacteria

Full genome sequencing from single cysts

First genome draft published

SCIENTIFIC REPORTS 

OPEN

Environmental marine pathogen isolation using mesocosm culture of sharpnose seabream: striking genomic and morphological features of novel *Endozoicomonas* sp.

Received: 05 August 2015
Accepted: 02 November 2015
Published: 07 December 2015

Pantelis Katharios^{1,*}, Helena M. B. Seth-Smith^{2,3,*}, Alexander Fehr³, José M. Mateos⁴, Weihong Qi², Denis Richter³, Lisbeth Nufer³, Maja Ruetten³, Maricruz Guevara Soto^{3,5}, Urs Ziegler⁴, Nicholas R Thomson⁶, Ralph Schlapbach² & Lloyd Vaughan³

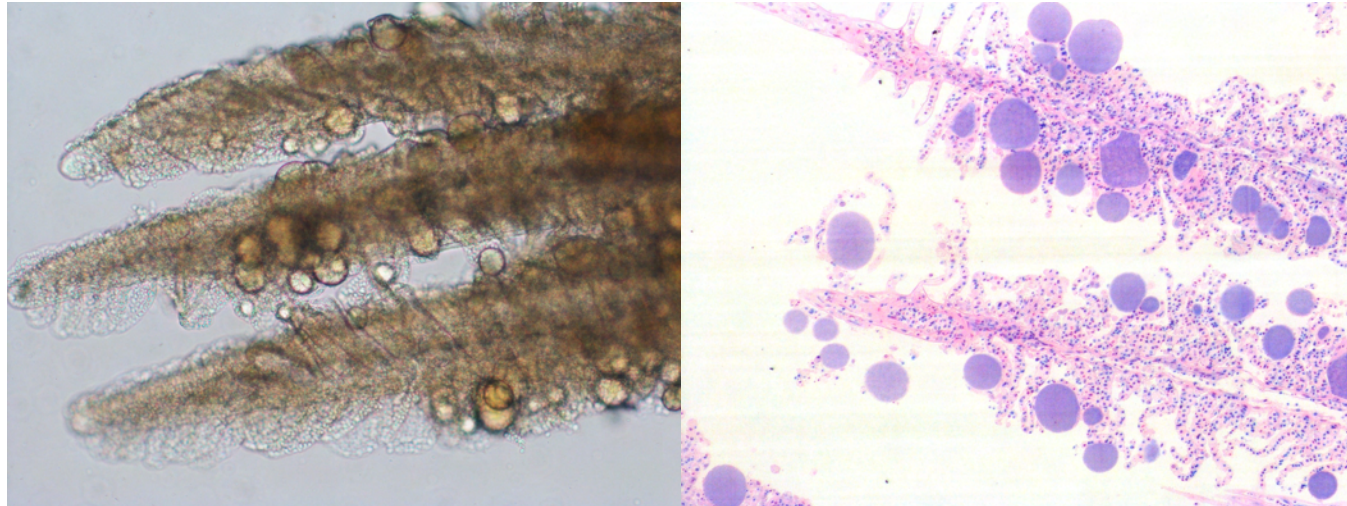
Epitheliocystis in gilthead sea bream

Juvenile breams (2-5g)

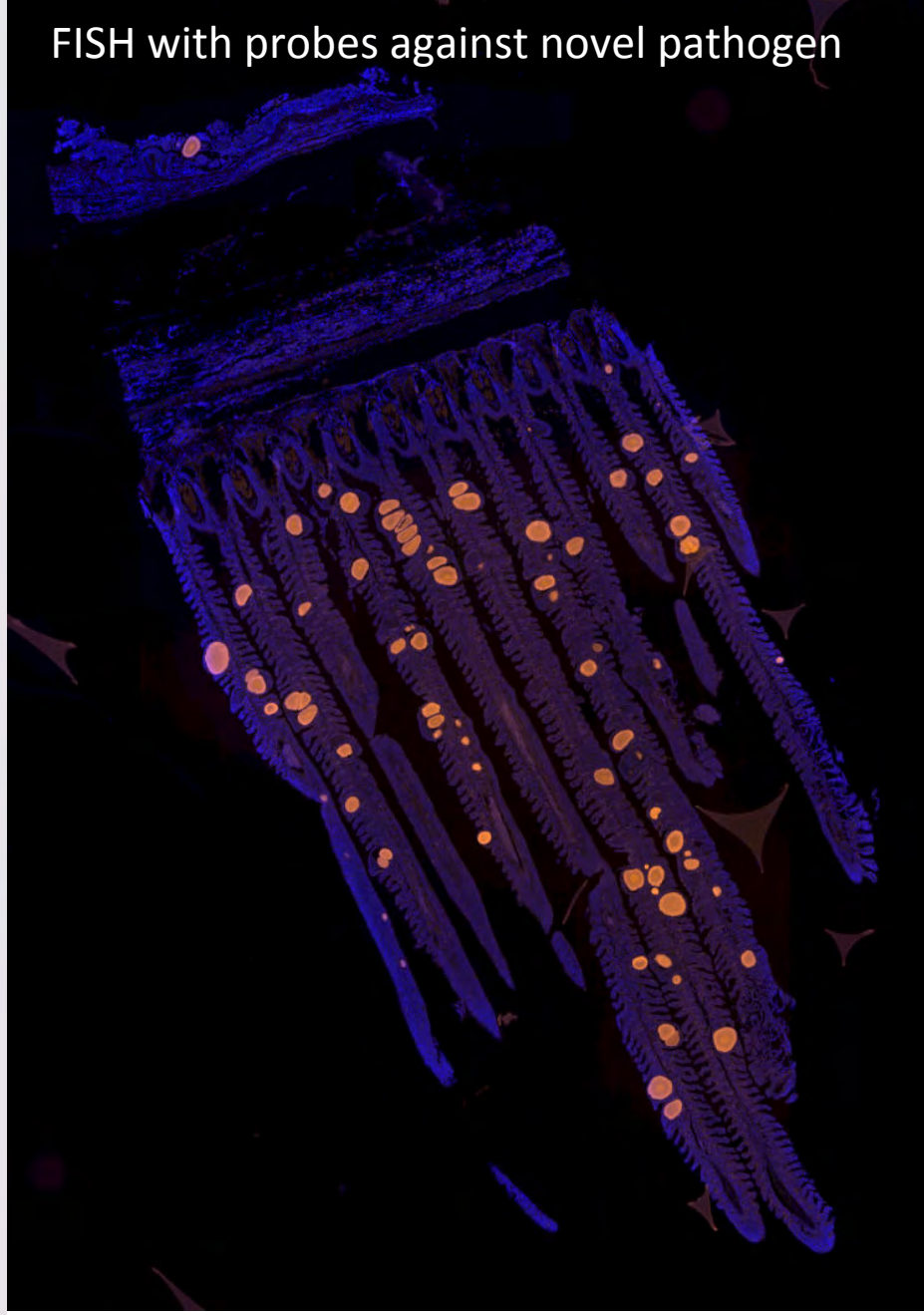
Similar case

PCR for chlamydia positive

In situ negative



FISH with probes against novel pathogen



Novel pathogens

Novel genus

2 new species

Candidatus Ichthyocystis hellenicum n.sp

Candidatus Ichthyocystis sparus n.sp

β -proteobacteria

Full genome sequencing from single cysts

First genome draft published

The ISME Journal (2016), 1–13
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www.nature.com/ismej



ORIGINAL ARTICLE

Emerging pathogens of gilthead seabream: characterisation and genomic analysis of novel intracellular β -proteobacteria

Helena MB Seth-Smith^{1,2}, Nancy Dourala³, Alexander Fehr², Weihong Qi¹, Pantelis Katharios⁴, Maja Ruetten², José M Mateos⁵, Lisbeth Nufer², Roseline Weilenmann², Urs Ziegler⁵, Nicholas R Thomson⁶, Ralph Schlapbach¹ and Lloyd Vaughan²
¹Functional Genomics Center Zürich, University of Zürich, Zürich, Switzerland; ²Institute for Veterinary Pathology, Vetsuisse Faculty, University of Zürich, Zürich, Switzerland; ³Selonda Aquaculture, Athens, Greece; ⁴Institute of Marine Biology, Biotechnology and Aquaculture, Hellenic Center for Marine Research, Heraklion, Crete, Greece; ⁵Center for Microscopy and Image Analysis, University of Zürich, Zürich, Switzerland and ⁶Pathogen Genomics, The Wellcome Trust Sanger Institute, Hinxton, Cambridge, UK

GBE

GENOME BIOLOGY AND EVOLUTION

Host-Associated Genomic Features of the Novel Uncultured Intracellular Pathogen *Ca. Ichthyocystis* Revealed by Direct Sequencing of Epitheliocysts

Weihong Qi¹, Lloyd Vaughan², Pantelis Katharios³, Ralph Schlapbach¹, and Helena M.B. Seth-Smith^{1,2,*}

¹Functional Genomics Center Zurich, University of Zurich, Switzerland

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³Hellenic Center for Marine Research, Institute of Marine Biology, Biotechnology and Aquaculture, Heraklion, Greece

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Accepted: May 5, 2016

Data deposition: This project has been deposited at DDBJ/EMBL/GenBank under the study accession PRJEB7439.

Following the false lead

In both cases Chlamydia were co-infecting the fish

Smaller and fewer cysts were caused by fish pathogenic Chlamydia were also simultaneously present in the same tissue

Through qPCR analysis it was estimated that the chlamydia:ichthyocystis ratio was from 1:5 to 1:3000

PCR targeting chlamydia was always positive and sequencing was pointing to known pathogens

If *in situ* was not used, then the true pathogens would have been misidentified

Going after the chlamydial pathogens

Gilthead seabreams with Ichthyocystis infections

PCR Chlamydiae-specific 16s primers

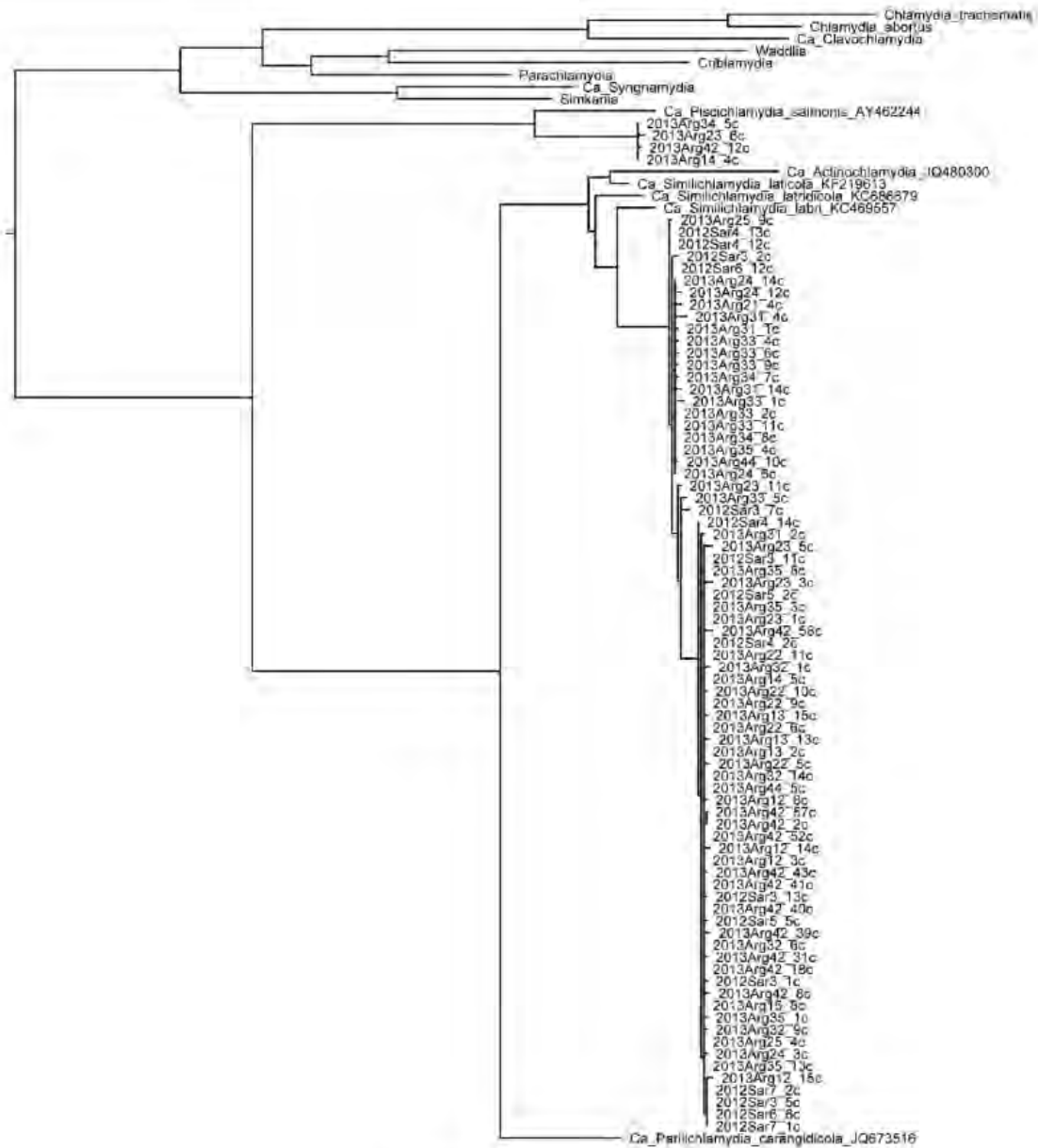
Amplicons cloned into Topo vecto pCR2.1

Blast in Genbank database

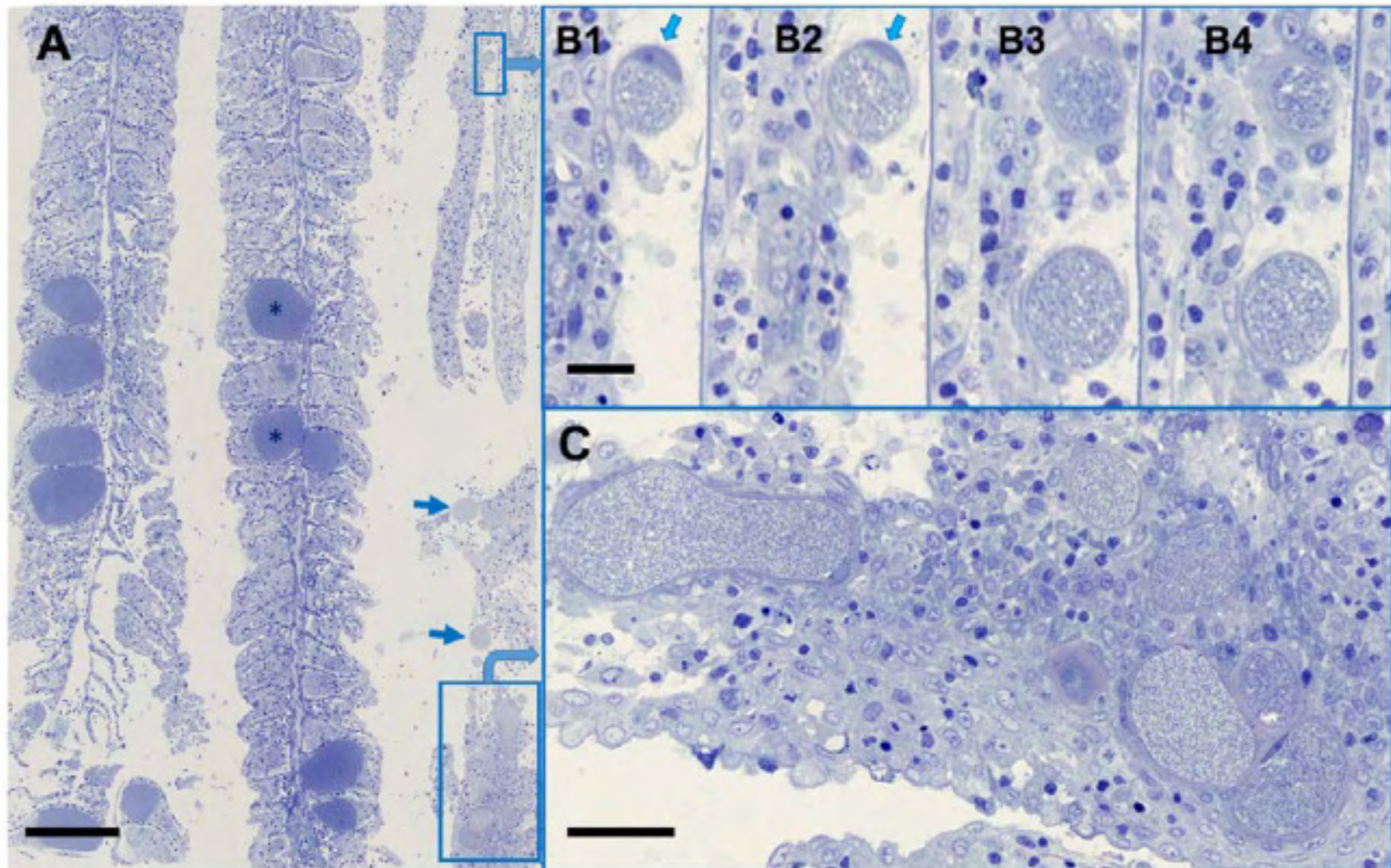
Phylogenetic analysis

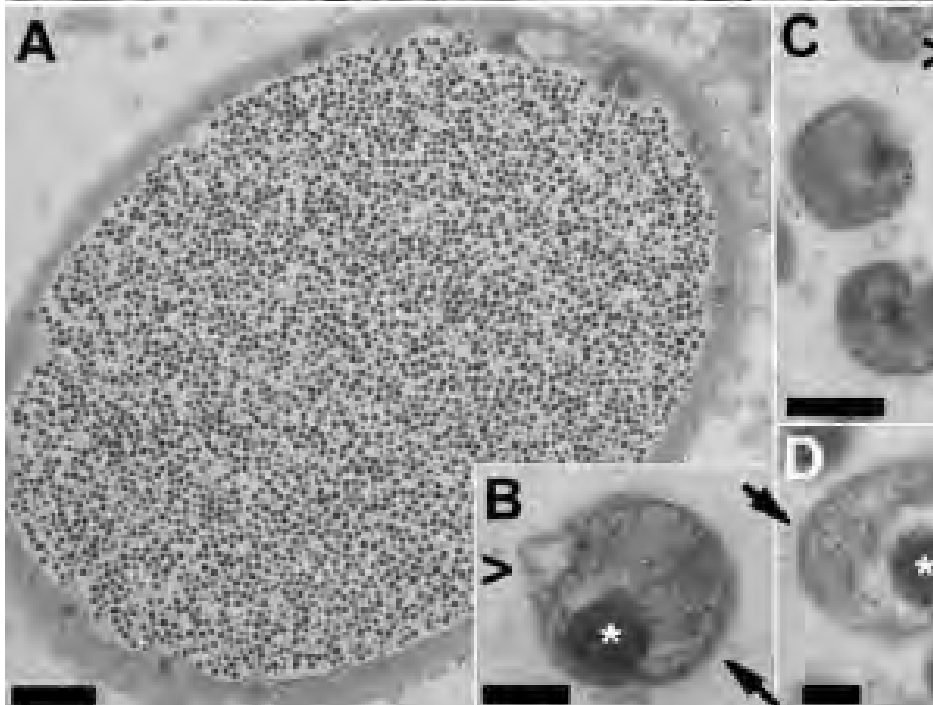
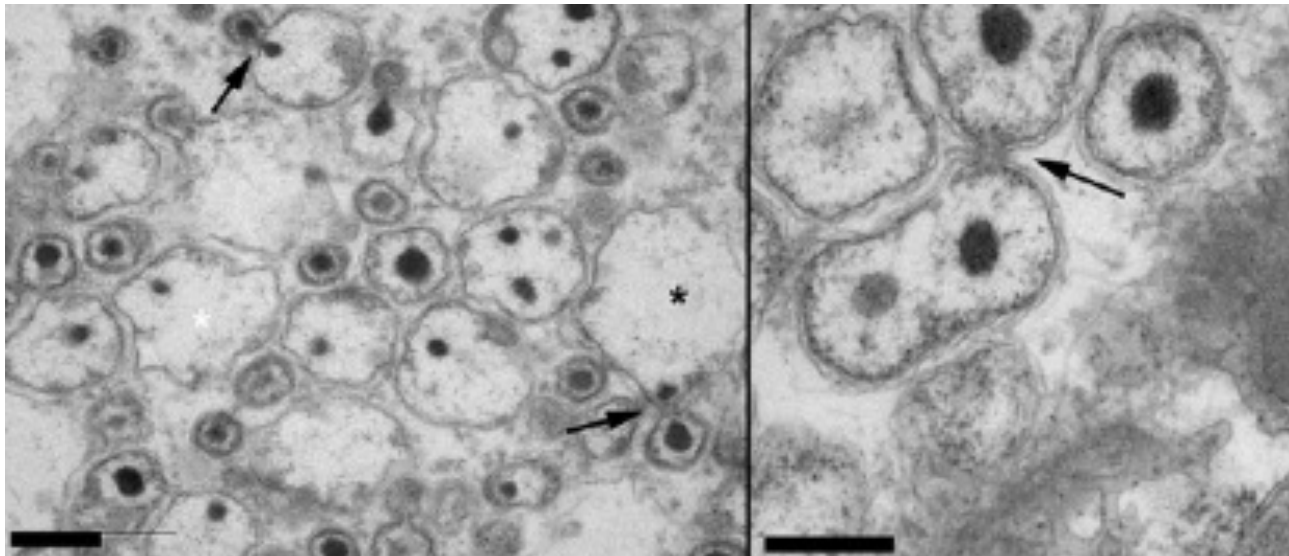
Fluorescent in situ hybridization

Histology and TEM



0.04

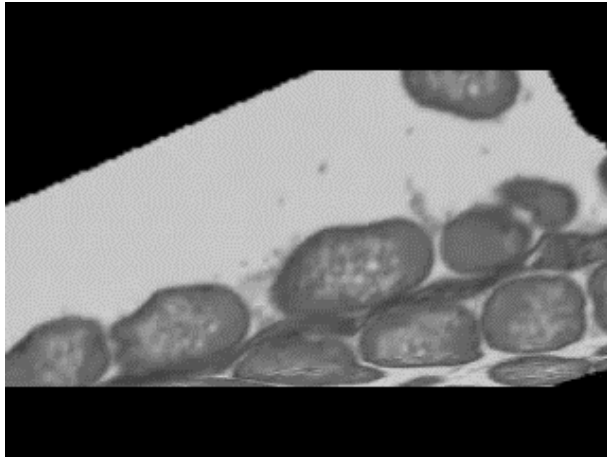




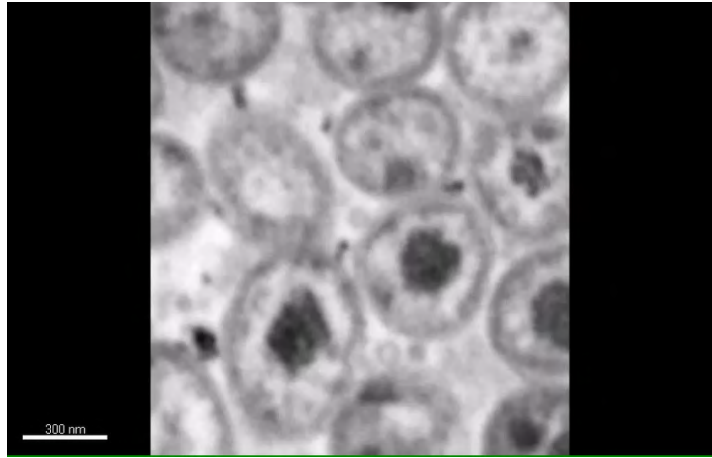
***Ca. Similichlamydia* in Epitheliocystis Co-infection of Gilthead Seabream Gills: Unique Morphological Features of a Deep Branching Chlamydial Family**

Helena M. B. Seth-Smith^{1,2†}, Pantelis Katharios³, Nancy Dourala⁴, José M. Mateos⁵,
Alexander G. J. Fehr¹, Lisbeth Nufer¹, Maja Ruetten^{1,6}, Maricruz Guevara Soto^{1,7} and
Lloyd Vaughan^{1,6*}

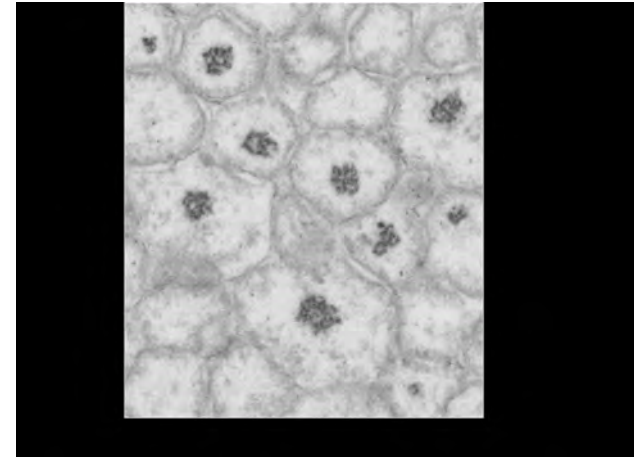
OPEN ACCESS



Ca. Endozoicomonas cretensis
 γ -proteobacteria
0.8 x 2.5 μm



Ca. Ichthyocystis sp
 β -proteobacteria
0.5 x 0.7 μm



Ca. Similichlamydia sp
Chlamydiae

DIVERSIFY project

Study epitheliocystis disease in greater amberjack

Mesocosm experimental setup

Isolate (?) and characterize the pathogen

Test the tools developed for early detection and diagnosis

Survey fish farms in Greece



Mesocosm trials

3 trials

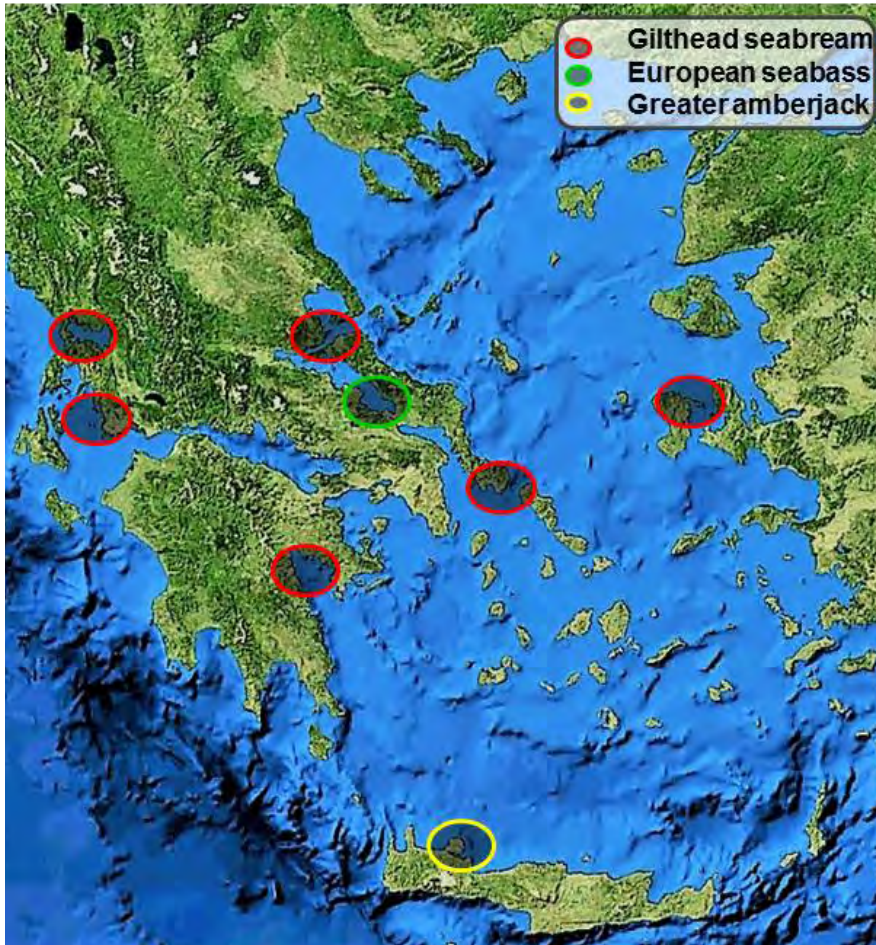
Greater amberjack, seabass, meagre

Fish did not get the infection

Screening in planktonic organisms and water

Environmental chlamydia not related to the disease

Survey in Greece



Gilthead seabream

- Astakos
- South Evoia
- North Evoia
- Chios
- Argolida
- Vonitsa

European seabass

- Larymna

Greater amberjack

- Souda

Identification pipeline

Amplification of 16S rRNA gene fragments with specific primers for

- *Ca. Ichthyocystis* spp.
- *Endozoicomonas* sp.
- Chlamydiae

Visualization of the results

Agarose-gel with Et-Br

Identification of pathogens

- Sequencing of the PCR products (Sanger)
- Comparison of the sequences with other sequences of NCBI GenBank (BLAST)

Phylogenetic analysis

- Alignment of sequences with ClustalW
- Phylogenetic trees were made using Tamura-Nei model with Neighbor joining analysis at 1000 bootstrap.

Results

PCR results

All (33) samples were positive for *Ca. Ichthyocystis* spp.

15 samples were positive for Chlamydiae

2 samples were positive for *Endozoicomonas* spp.

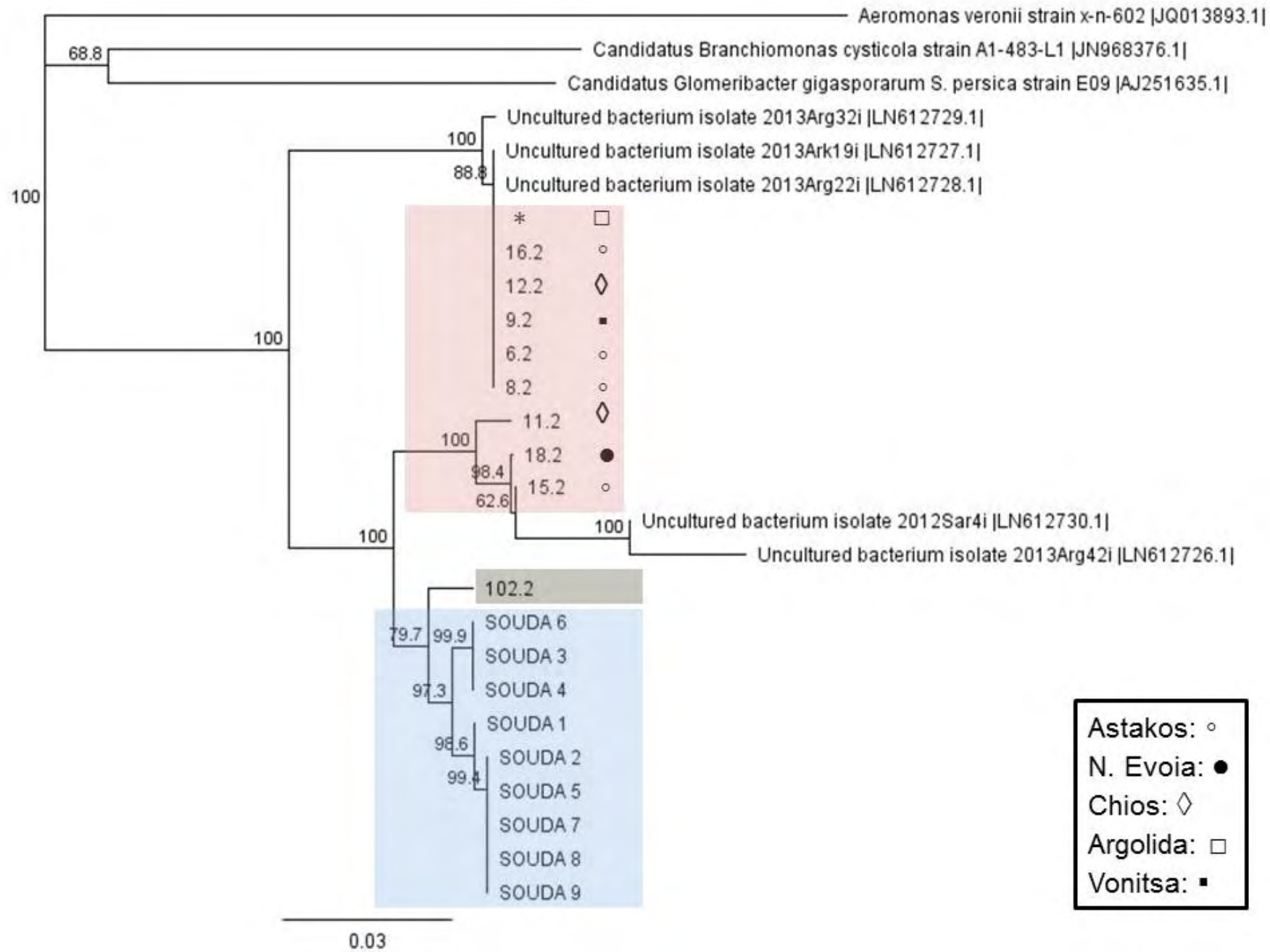
Chlamydiae

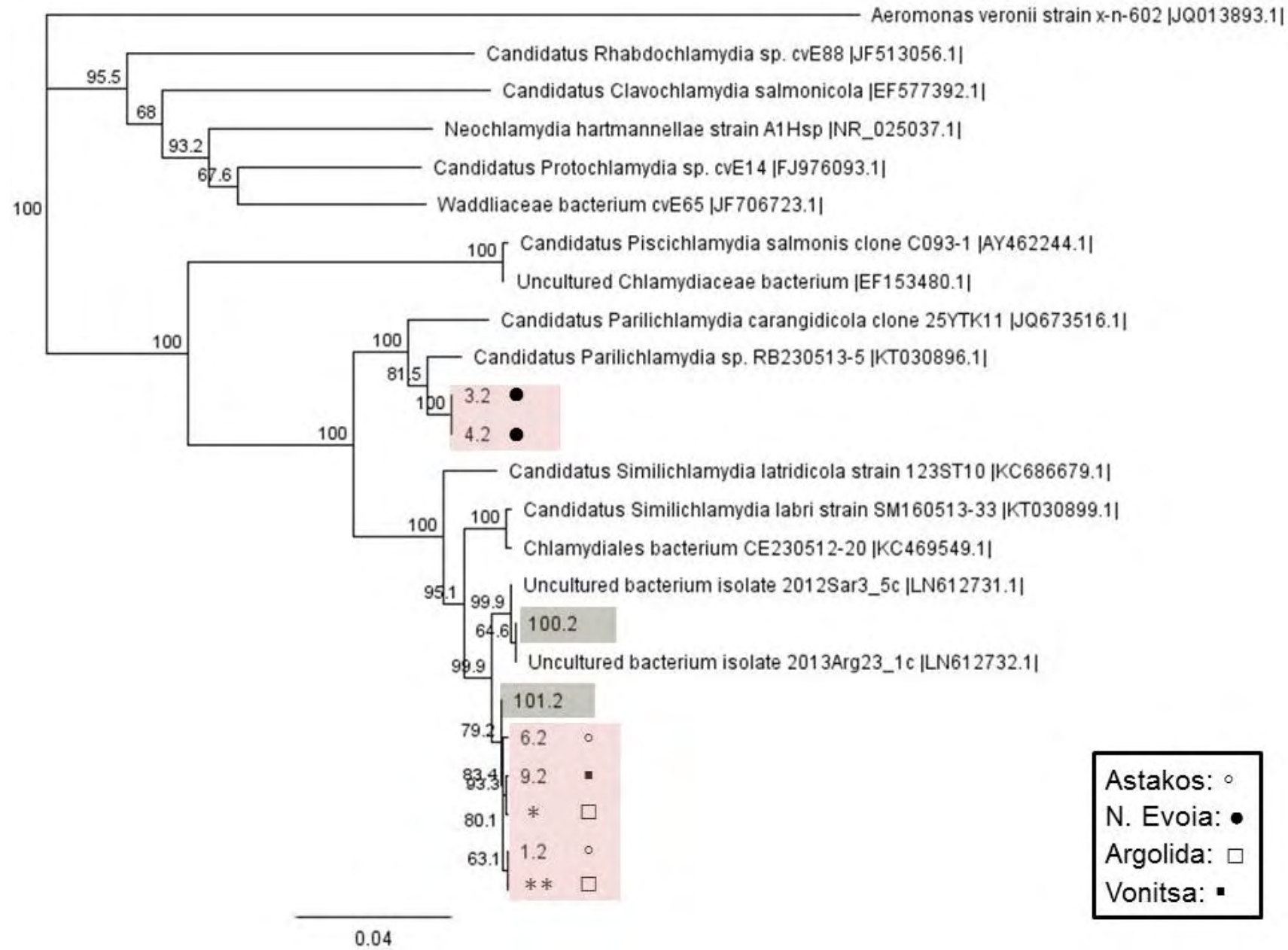
Gilthead seabream: Astakos, S. Evoia, N. Evoia, Argolida, Vonitsa

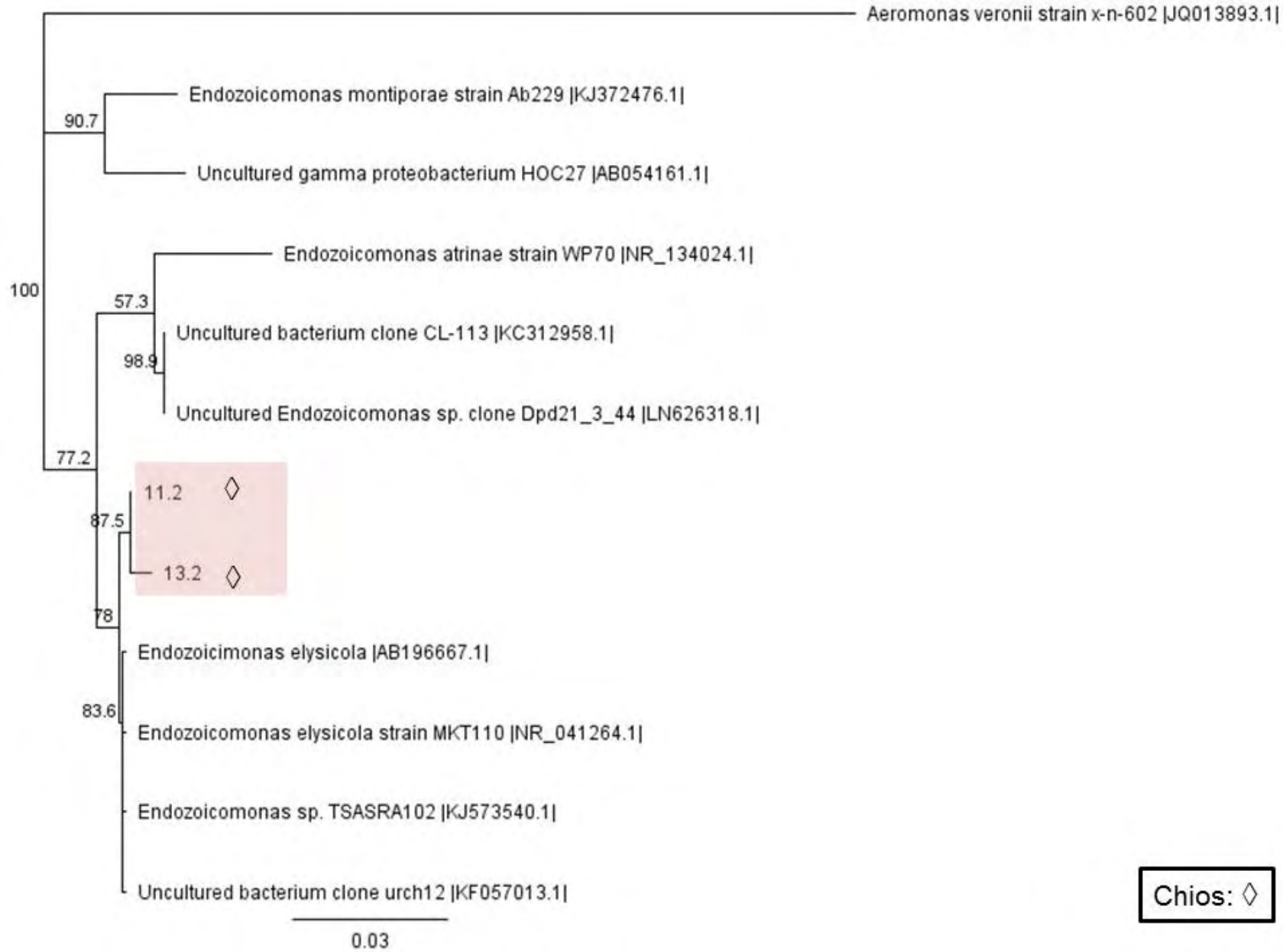
European seabass: Larymna

Endozoicomonas

Gilthead seabream: Chios







Chios: ◇

Epitheliocystis agents in Greece

	Chlamydia	Ichthyocystis	Endozoicomonas
Taxonomic class	Chlamydiae	β -proteobacteria	γ -proteobacteria
# species found	3	2-3	1-2
# fish species infecting	3	4	2
Genome size	<1 Mb	~2.3 Mb	~5.8Mb
Life style	Obligate intracellular	Obligate intracellular	In transition

December outbreak in Souda



Sampling

2 samplings (December and January)

Fish gills (infected and uninfected), spleens

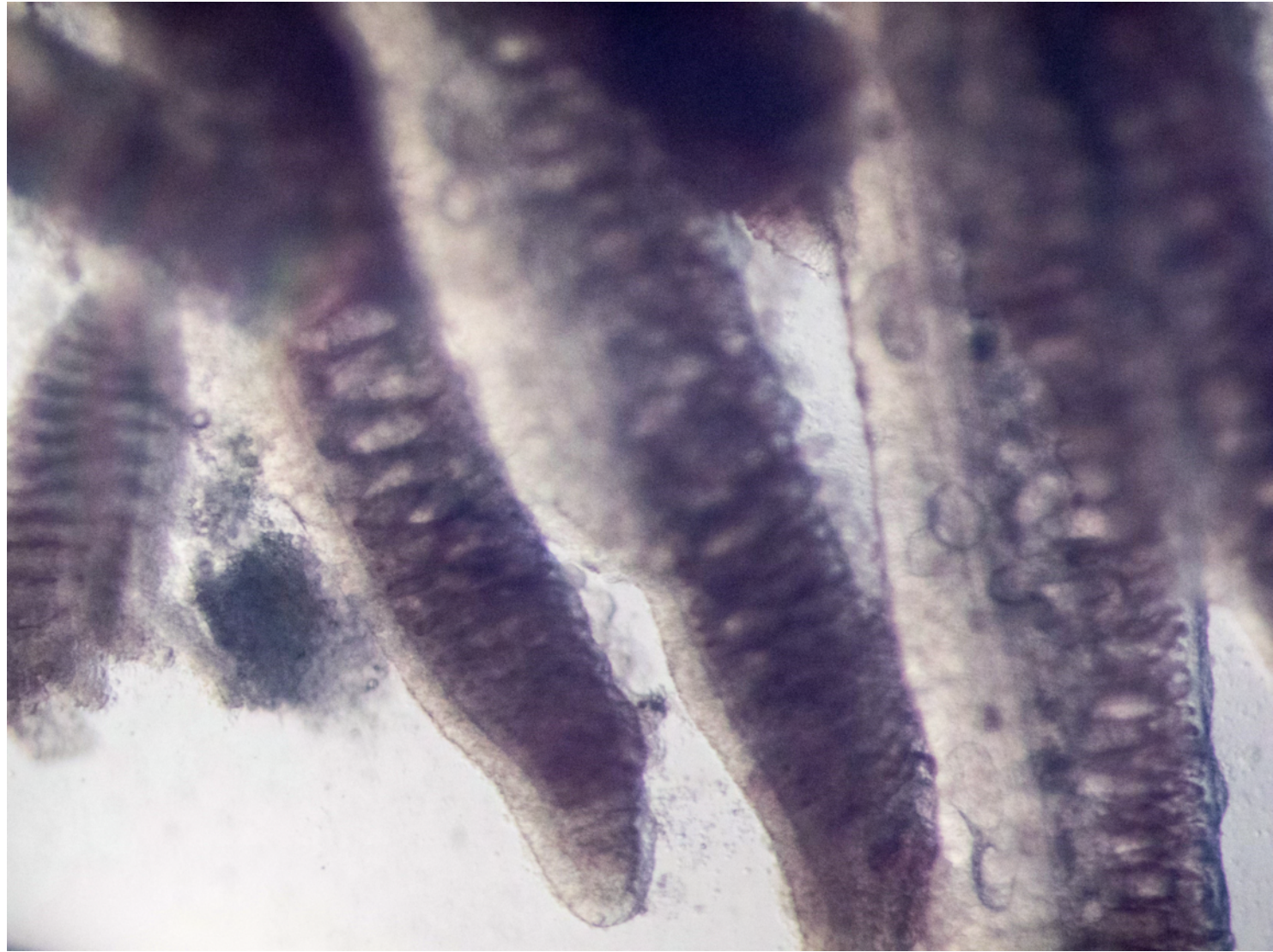
Commensal organisms (clams, oysters, mussels, anemone, ascidia, sponges)

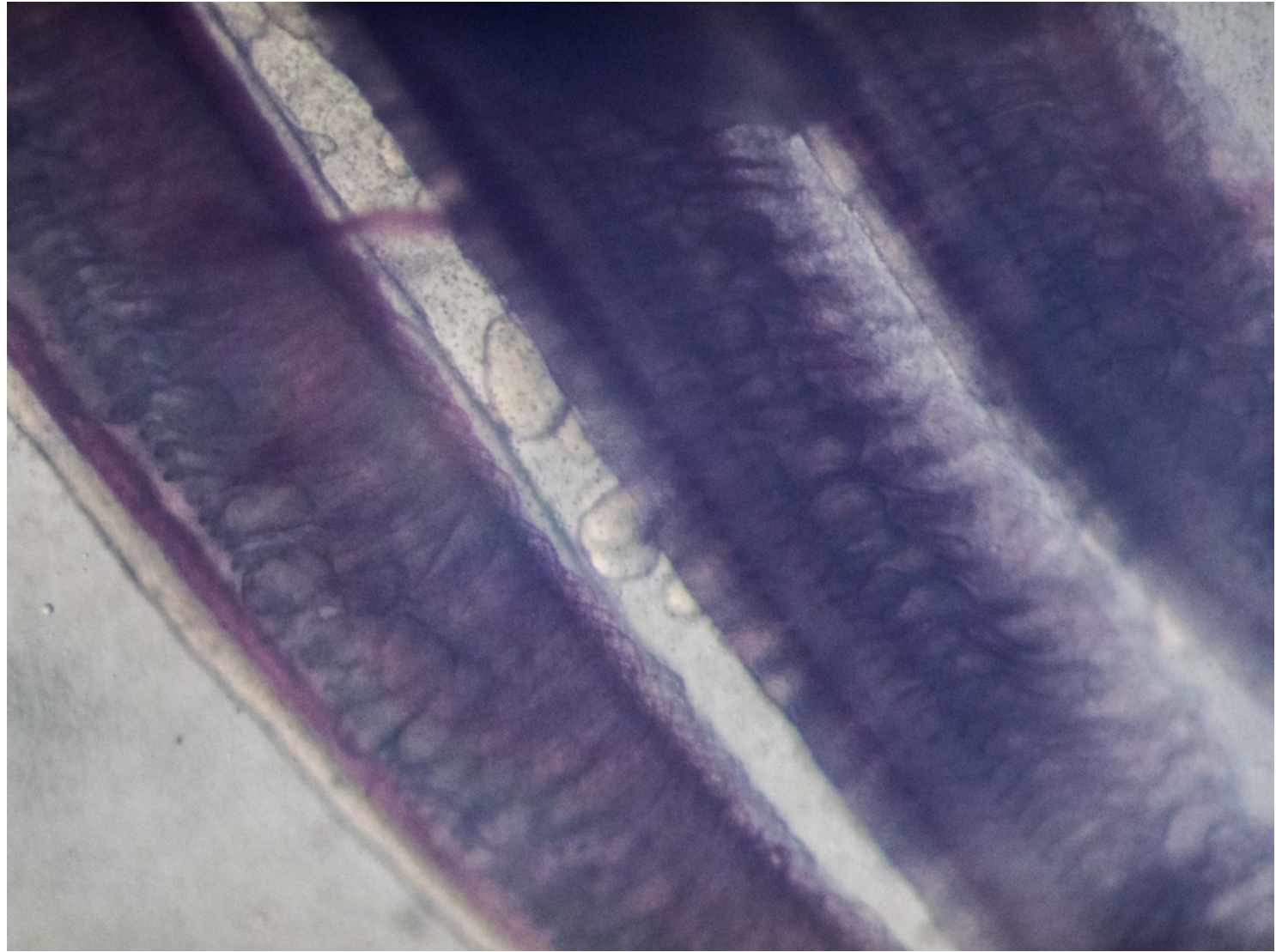
Samples for PCR, histology, TEM, genomic analysis, gene expression

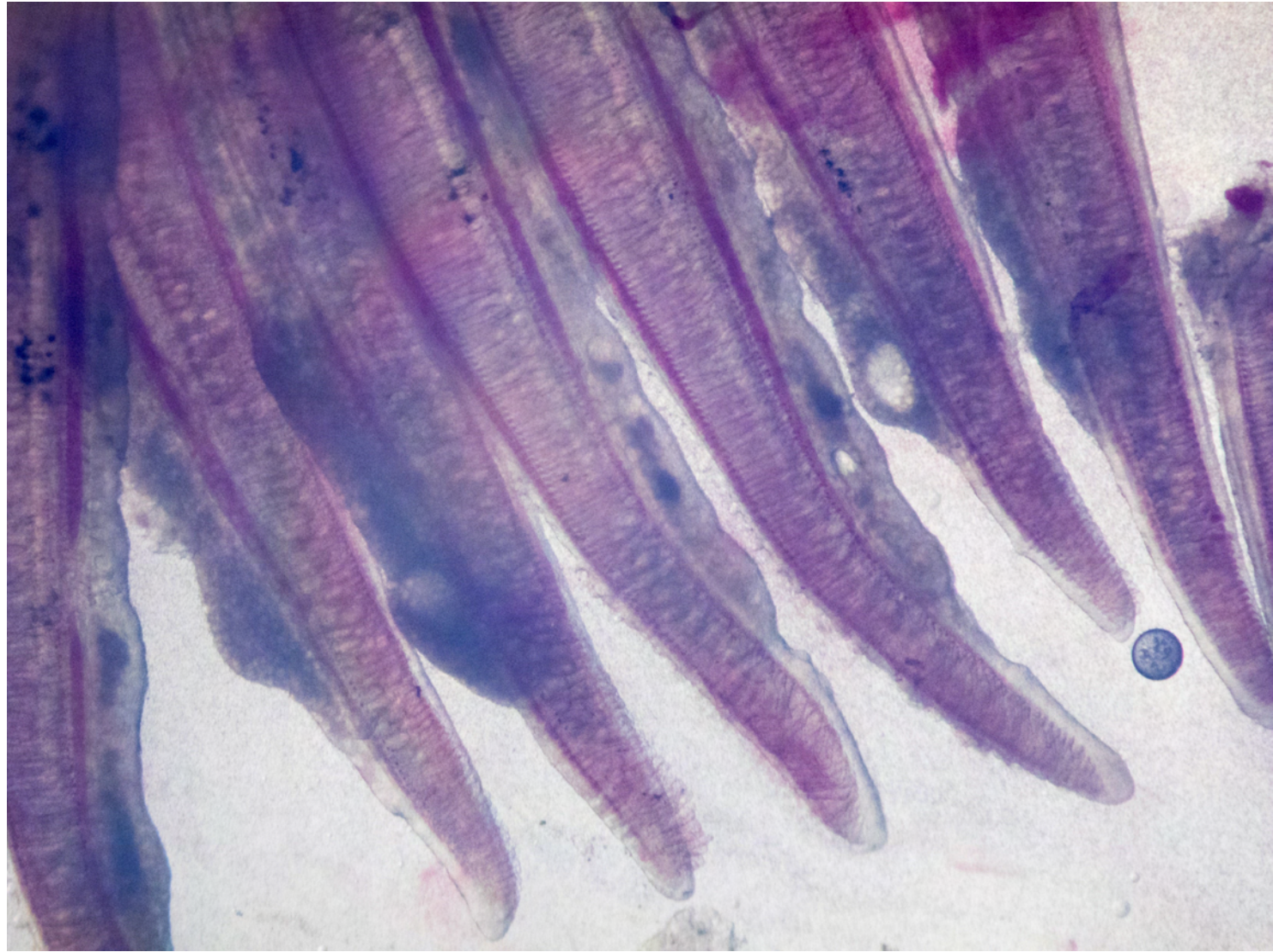
Fresh samples for infection experiments using zebrafish

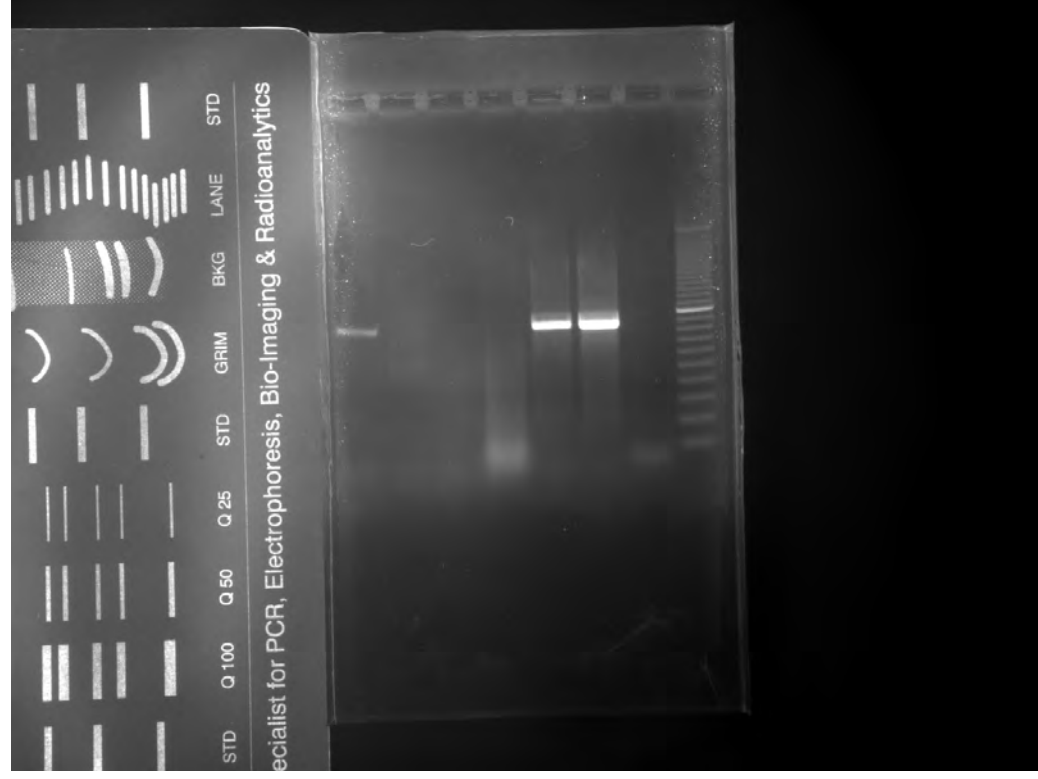
Samples sent to Chris Secombes for immune gene expression analysis

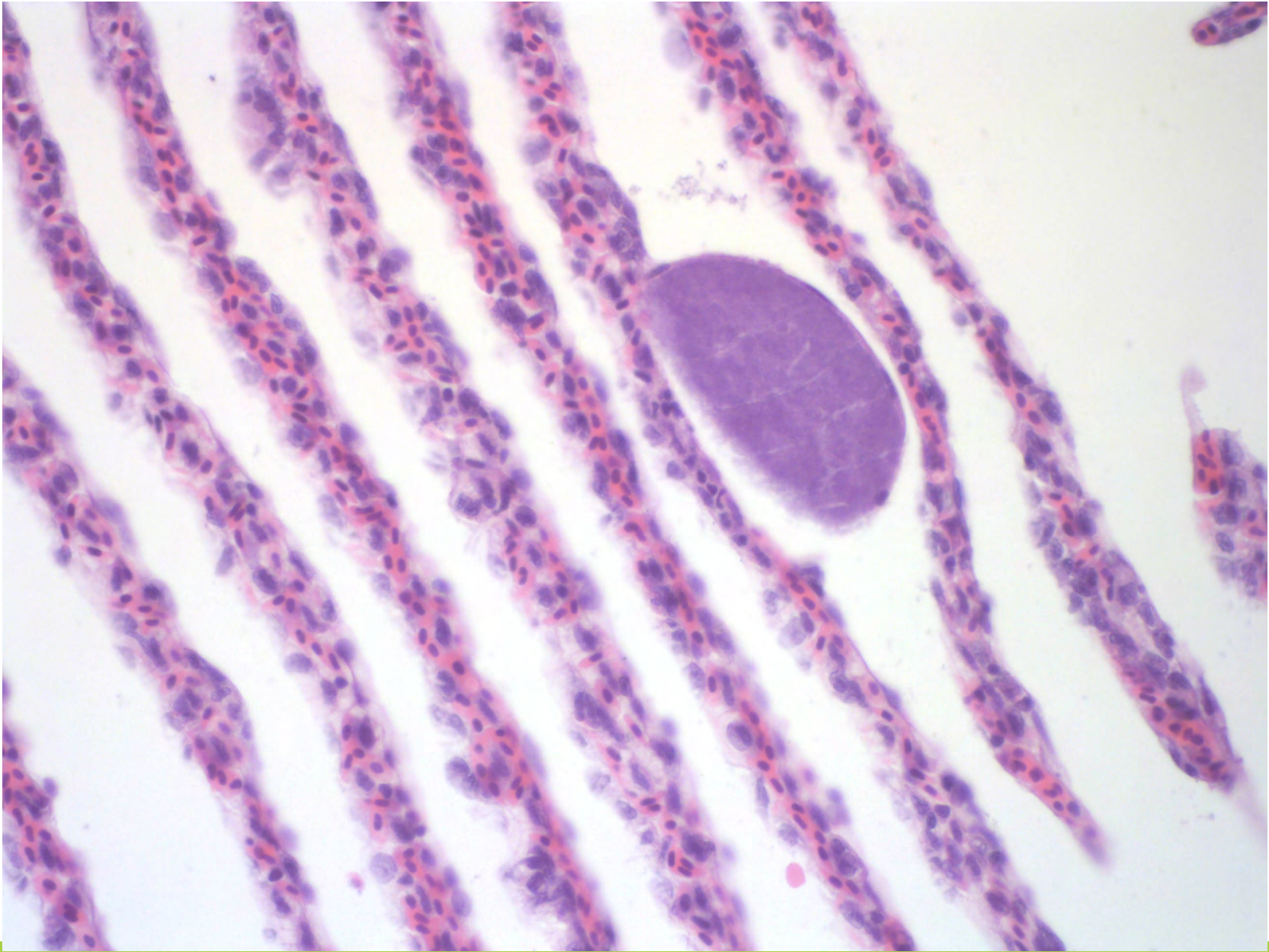
Whole Genome Sequencing

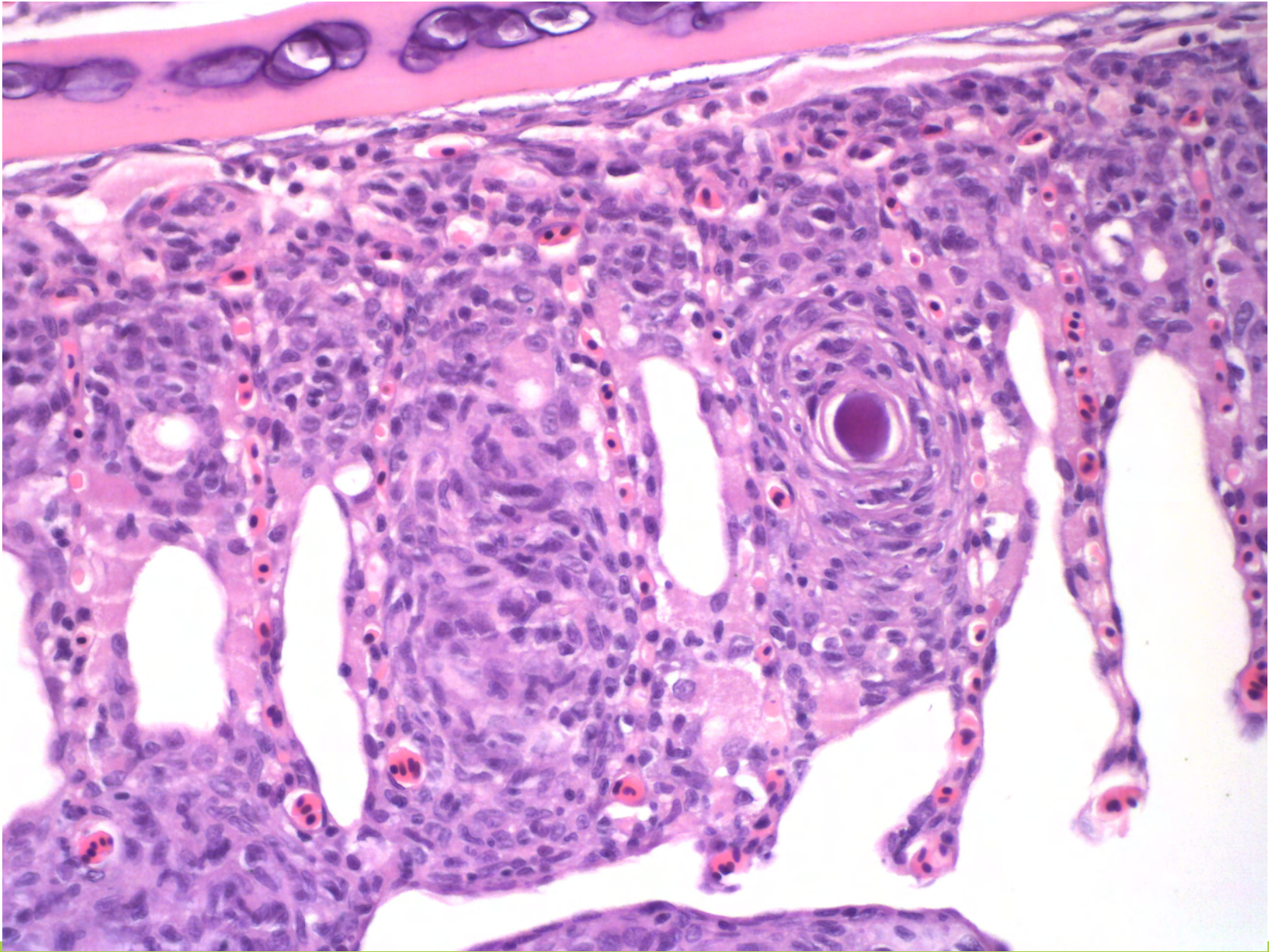


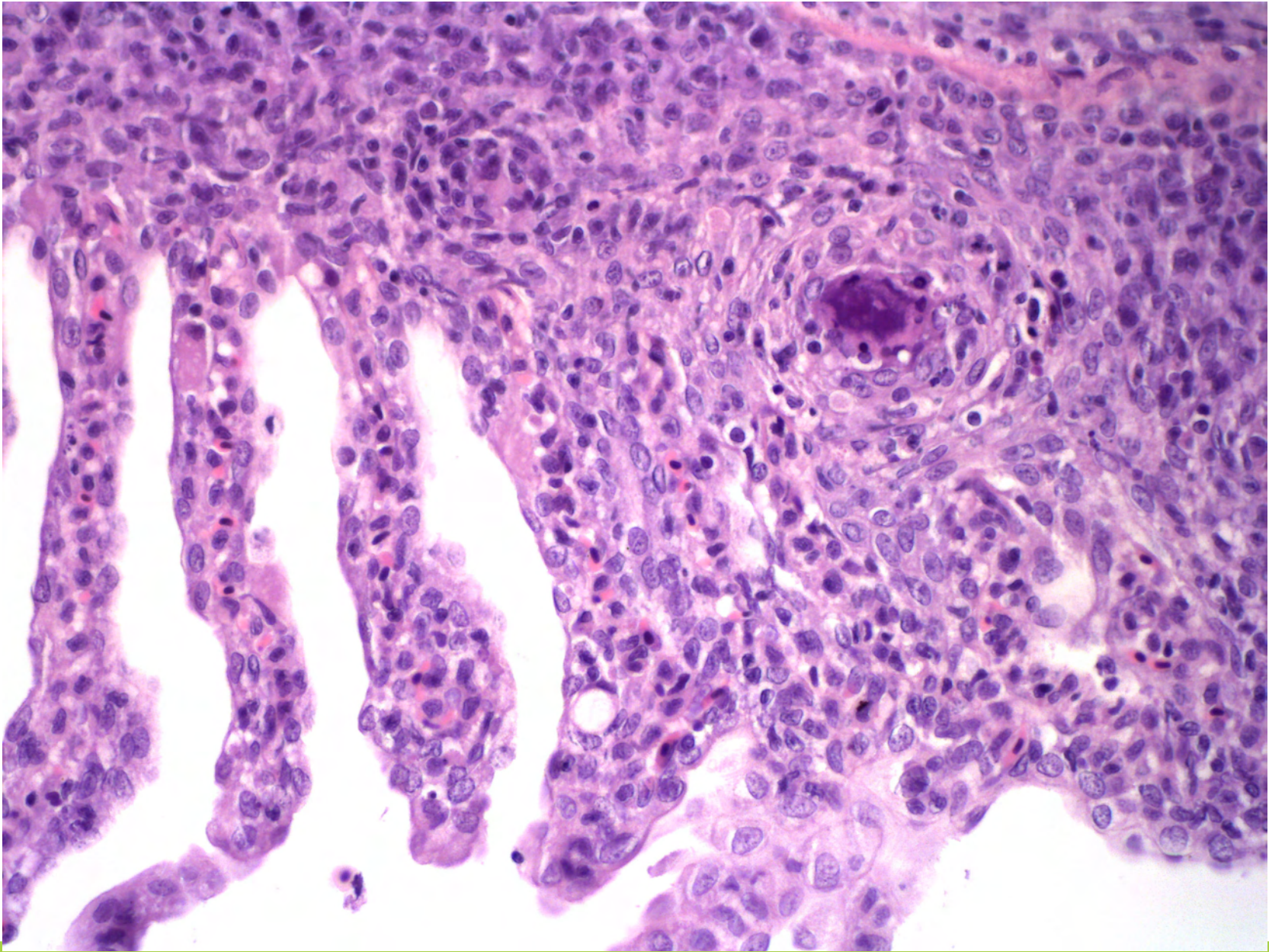


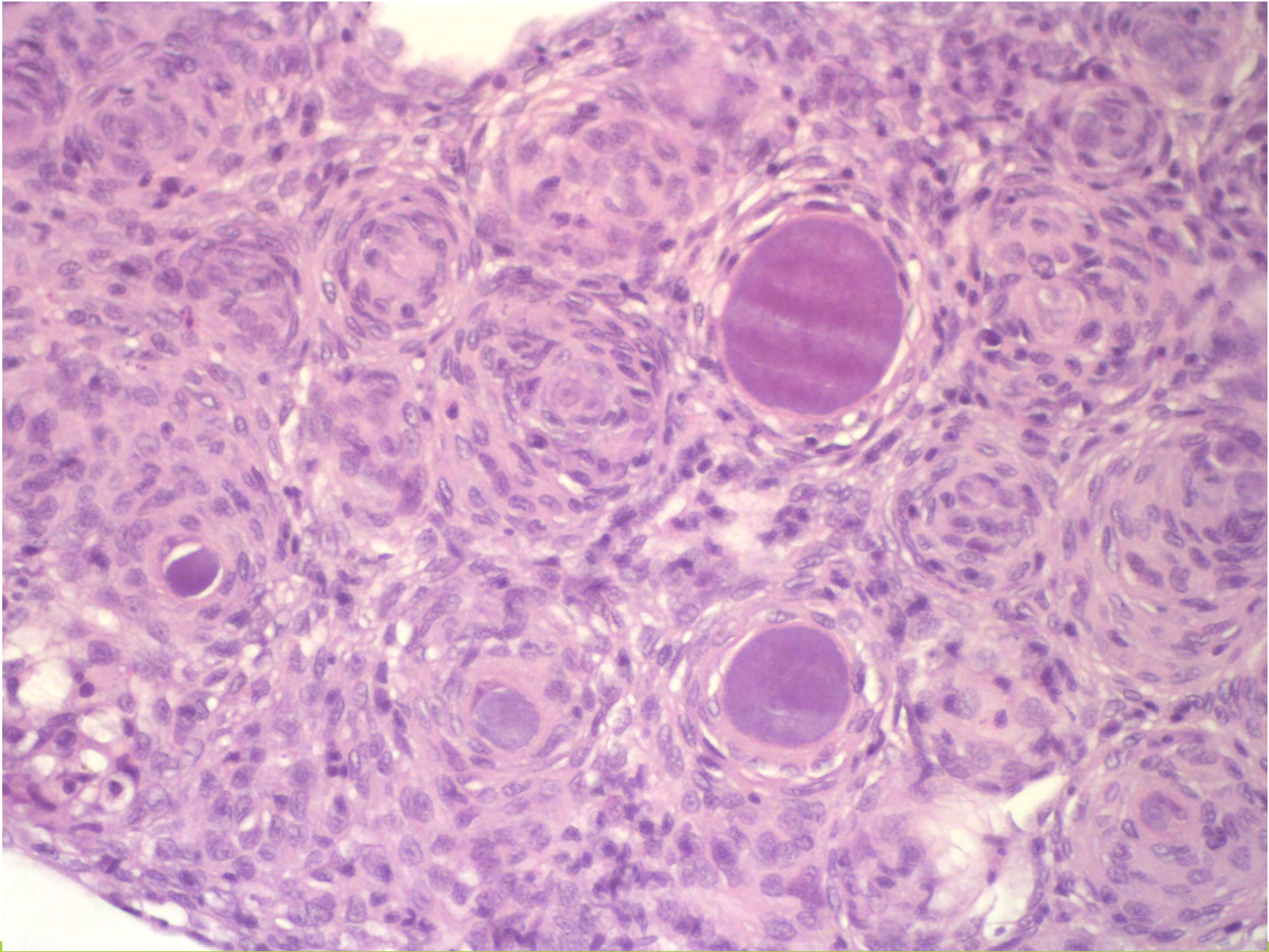












Results so far

Confirmed epitheliocystis outbreak in cultured greater amberjack

Disease caused by *Ca. Ichthyocystis* sp. (possibly novel species)

Zebrafish infection unsuccessful

All commensal organisms examined are negative for the bacteria

...next steps

Whole genome sequencing

Phylogenetic analysis

Gene expression (fish for immune response, bacteria for virulence regulation)

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5. Pathovet AG, Tagelswangen, Switzerland
6. Andromeda SA, Patras, Greece
7. Selonda Aquaculture, Athens, Greece
8. University of Aberdeen

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