Identification and characterization of the causative agents of fish diseases

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Thai aquaculture production







Fish, 71.22%

Shrimp, 13.17% Mollusks, 7.85%





Production 2.56 M ton Value 145,414 M Baht (DOF 2014)

Fish aquaculture in Thailand

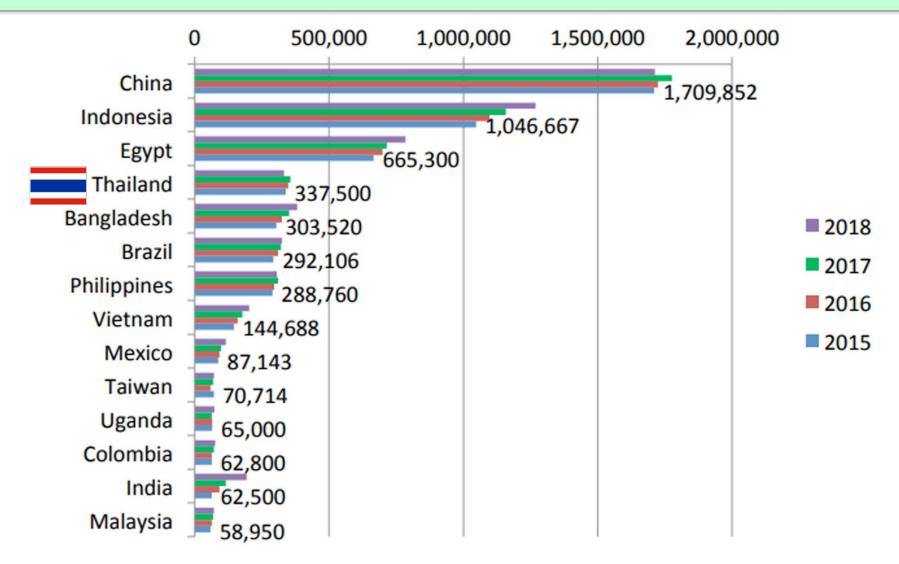
Freshwater species	% Production
Nile tilapia Catfish	45.8 29.8
Silver barb	11.2
Snakeskin gourami	5.7
Striped catfish	4.6
Snakehead murrel	1.8
Giant gourami	1.0
Mrigal carp	0.2

% Production
80.7
19.3

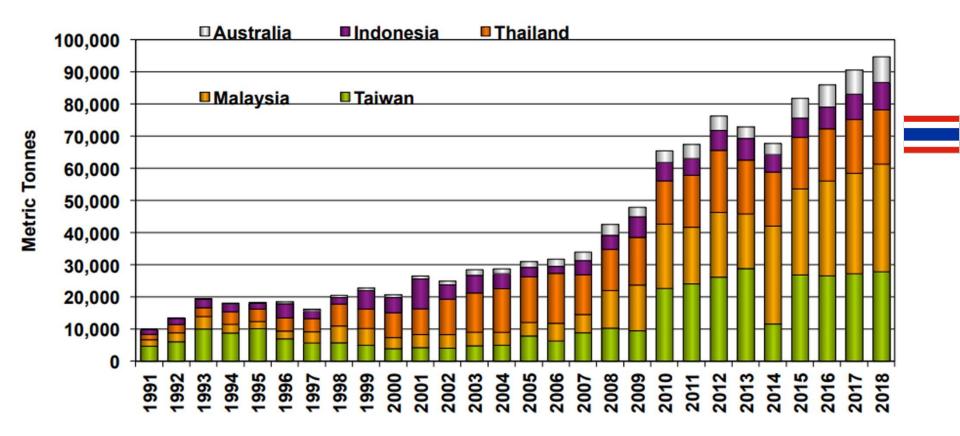
Total production \sim 20,000 tonnes

Total production \sim 400,000 tonnes

Tilapia production by major producing countries

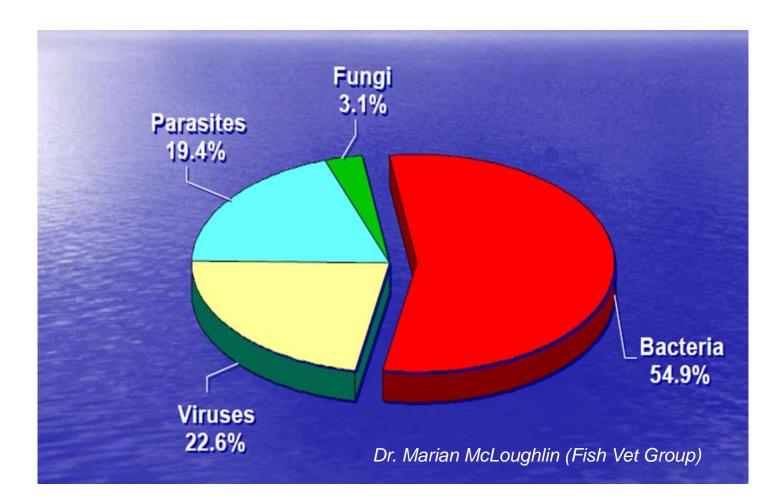


Asian seabass production by major producing countries

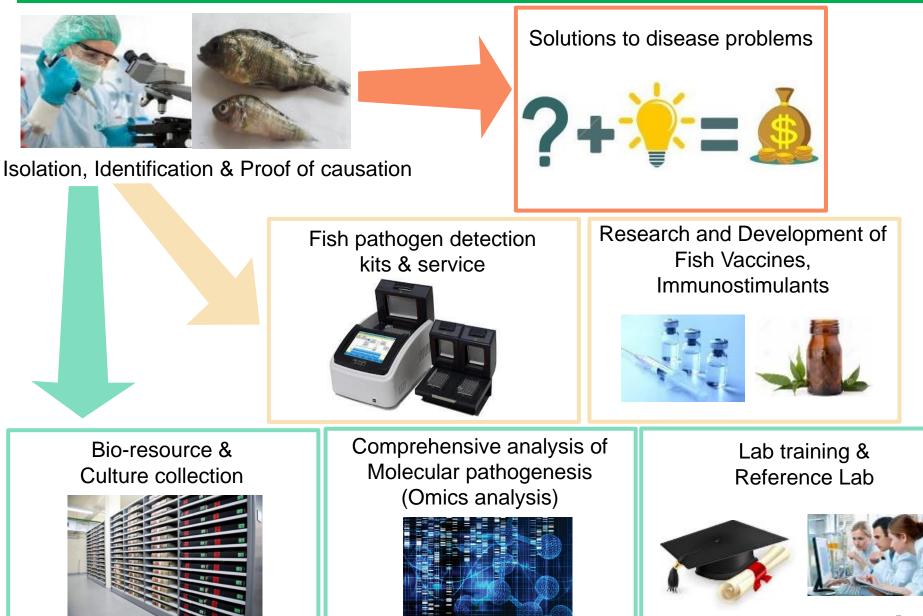


Dr. Ragnar Tveteras, University of Stavanger, Norway

Disease is considered as one of the major problems to aquaculture production globally

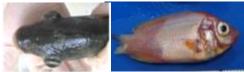


From Research to Solutions



Research on fish diseases





Nile tilapia (Oreochromis niloticus)

Hybrid red tilapia (O. mossambicus x O. niloticus)

- Flavobacterium columnare
- Francisella noatunensis
- Streptococcus agalactiae
- Aeromonas veronii
- Hahella chejuensis
- Tilapia lake virus (TiLV)
- Iridovirus



Barramundi / Asian Seabass (Lates calcarifer)

- Vibrio harveyi
- Scale drop disease virus (SDDV)



Striped catfish (Pangasianodon hypophthalmus)

- Flavobacterium columnare
- Edwardsiella ictaluri



Siamese fighting fish (Betta splendens)

- Skin nodule syndrome
- Big belly syndrome

Research on fish diseases



Nile tilapia (*Oreochromis niloticus*) Hybrid red tilapia (*O. mossambicus x O. niloticus*)

- Hahella chejuensis
- Tilapia lake virus (TiLV)



Barramundi / Asian Seabass (Lates calcarifer)

• Vibrio harveyi

• Scale drop disease virus (SDDV)

Novel disease in tilapia hatcheries

Red eggs

- Since 2000
- Eggs turn to red and fail to hatch
- Cumulative loss of 10% and up to 50% in cold season
- The cause is unknown, but appears to be infectious



Egg collection

Artificial incubation

Red eggs

Novel disease in tilapia hatcheries

Infectious disease of red eggs caused by Hahella chejuensis



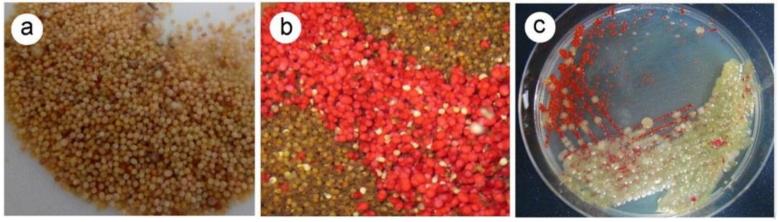
Normal eggs

Red egg disease (10-50% loss) (We named it "Hahellosis")

Red bacteria on TSA (*Hahella chejunensis*)

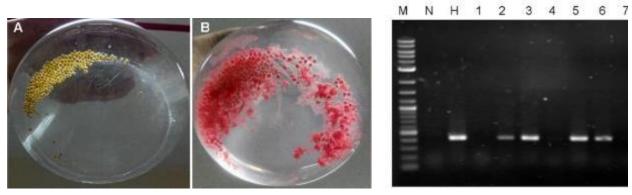
Novel disease in tilapia hatcheries

Infectious disease of red eggs caused by Hahella chejuensis



Normal eggs

Red egg disease (10-50% loss) (We named it "Hahellosis") Red bacteria on TSA (*Hahella chejunensis*)



PCR detection

In situ hybridization

Bioassay

Red egg disease "Hahellosis"

Knowing the enemy leading to the solutions

- Hahella chejuensis is a marine bacterium
- It is a halophilic bacteria
- Red egg problem occurs at temperature below 24 °C

- Reduce salinity from 7 ppt to 4 ppt - Content of the filter system to sunlight

Problem reduced to < 1%





Red egg disease "Hahellosis"

Economic impact

รายการ	จำนวน
(1) จำนวนไข่ปลาที่ผลิต	2,000,000,000 ใบ
การผลิตของฟาร์มที่ประสบปัญหา 2 ฟาร์ม (เฉพาะฟาร์มที่ส่งตัวอย่างในงานวิจัย)	
จำนวนไข่ปลา 1,000,000,000 ใบ/ฟาร์ม/ปี X 2 ฟาร์ม	
(2) จำนวนไข่ปลาในช่วงเวลาที่มีปัญหาไข่ปลานิลไม่ฟักตัวและเปลี่ยนเป็นสีแดง (ช่วง	500,000,000 ใบ
อากาศเย็น) 3 เดือน	
2,000,000,000 ใบ / 12 เดือน X 3 เดือน	
(3) จำนวนการลดการสูญเสียไข่ปลาที่ติดเชื้อไม่ฟักตัว	150,000,000 ใบ
(ก่อนมีโครงการสูญเสีย 30-50% หลังมีโครงการสูญเสียน้อยกว่า 1%	
=500,000,000 ×30%)	
(4) มูลค่าการลดจำนวนไข่ปลาที่ติดเชื้อไม่ฟักตัว	<u>18,750,000 บาท</u>
(จำนวนการลดการสูญเสียไข่ปลาที่ติดเชื้อไม่ฟักตัว X มูลค่าลูกปลา 0.125 บาท/ตัว	•)



550,000 USD per year

Tilapia lake virus (TiLV)

Novel Orthomyxo-like virus

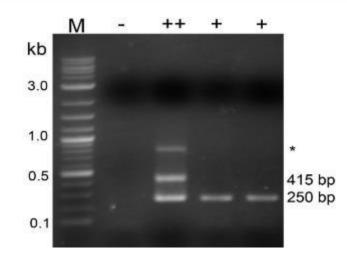


- Syncytial Hepatitis of Tilapia (SHT) in Ecuador (Ferguson et al. 2014)
- TiLV in Israel (Eyngor et al. 2014)
- Complete genome of TiLV (Bacharach et al. 2016)
 - TiLV reported in Colombia, Egypt,
 Thailand, Taiwan, Malaysia, India,
 Philippines, Uganda, Tanzania
 (Kembou Tsofack et al. 2017; Fathi et al. 2017;
 Nicholson et al. 2017; Surachetpong et al. 2017;
 Dong et al. 2017; OIE 2017; Amal et al. 2017;
 Behera et al. 2017, Mugimba et al. 2018)

Tilapia lake virus (TiLV)

Occurrence in Thailand and alternative RT-PCR protocol





- Primers from Eyngor et al. 2014
- Semi-Nested PCR (Dong et al 2017)

Free positive control plasmid for non-commercial use in response to 36 requests from 20 countries

Tilapia lake virus (TiLV)

Publications & Training



Contents lists available at ScienceDirect

Aquaculture

journal homepage: www.elsevier.com/locate/aquaculture

Emergence of tilapia lake virus in Thailand and an alternative semi-nested RT-PCR for detection

H.T. Dong^{a,b,*}, S. Siriroob^b, W. Meemetta^b, W. Santimanawong^b, W. Gangnonngiw^{b,c}, N. Pirarat^d, P. Khunrae^a, T. Rattanarojpong^a, R. Vanichviriyakit^{b,c}, S. Senapin^{b,c,*}



Contents lists available at ScienceDirect

journal homepage: www.elsevier.com/locate/aquaculture

Short communication

Evidence of TiLV infection in tilapia hatcheries from 2012 to 2017 reveals probable global spread of the disease

H.T. Dong^{a,b,*}, G.A. Ataguba^c, P. Khunrae^a, T. Rattanarojpong^a, S. Senapin^{b,d,**}

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Short communication

Inapparent infection cases of tilapia lake virus (TiLV) in farmed tilapia

Saengchan Senapin $^{a,b},$ K.U. Shyama, Watcharachai Meemettaa, Triwit Rattanarojponge, Ha Thanh Donge,



Training on TiLV diagnosis (PCR and histology)

Research on fish diseases



Nile tilapia (*Oreochromis niloticus*) Hybrid red tilapia (*O. mossambicus x O. niloticus*)

- Hahella chejuensis
- Tilapia lake virus (TiLV)



Barramundi / Asian Seabass (Lates calcarifer)

• Vibrio harveyi

• Scale drop disease virus (SDDV)

Caused by Vibrio harveyi







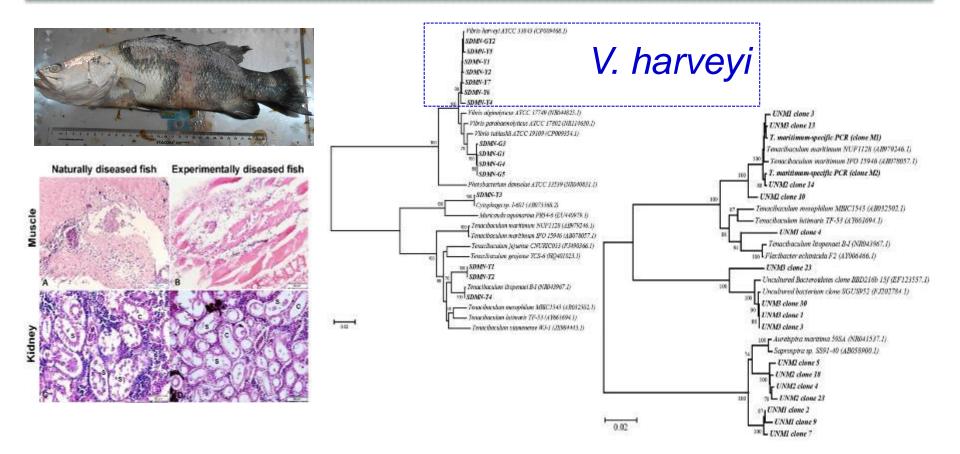


Up to 40% cumulative mortality

Caused by Vibrio harveyi



Caused by Vibrio harveyi

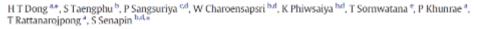


Caused by Vibrio harveyi

CrossMark



Recovery of Vibrio harveyi from scale drop and muscle necrosis disease in farmed barramundi, Lates calcarifer in Vietnam







ORIGINAL ARTICLE

Genome characterization of piscine 'Scale drop and Muscle Necrosis syndrome'-associated strain of *Vibrio harveyi* focusing on bacterial virulence determinants

P. Kayansamruaj^{1,2}, H.T. Dong³, I. Hirono⁴, H. Kondo⁴, S. Senapin^{5,6} and C. Rodkhum^{1,7}

- 2 Department of Aquaculture, Faculty of Fisheries, Kasetsart University, Bangkok, Thaland
- 3 Aquacuture Vaccine Mathem, Department of Microbiology, Faculty of Science, King Mongkut's University of Technology Thorburi, Bangkok, Thaland
- 4 Laboratory of Genome Science, Tokyo University of Marine Science and Technology, Tokyo, Japan
- 5 Center of Excelence for Shrimp Molecular Biology and Biotechnology (Centex Swimp), Faculty of Science, Mahidol University, Bangkok, Thaland
- 6 National Center for Genetic Engineering and Biotechnology (BIOTEC), National Science and Technology Development Agency, Pathumthani, Theiland
- 7 Fish Infectious Diseases (FDs) Special Task Force for Activating Research (STAR), Faculty of Veterinary Science, Chulalongkom University, Bangbok, Thuliand

Highlights

- A pathogenic V. harveyi strain was identified as the main causative agent.
- Coinfections of culturable and unculturable bacteria were uncovered from diseased fish.
- The role of unculturable bacteria needs further investigation.

Highlights

- Genome of V. harveyi Y6 was incorporated by a bacteriophage.
- 17 potential virulence genes were present exclusively in the strain Y6.

¹ Department of Veterinary Microbiology, Faculty of Veterinary Science, Chulalongkom University, Bangkok, Thailand

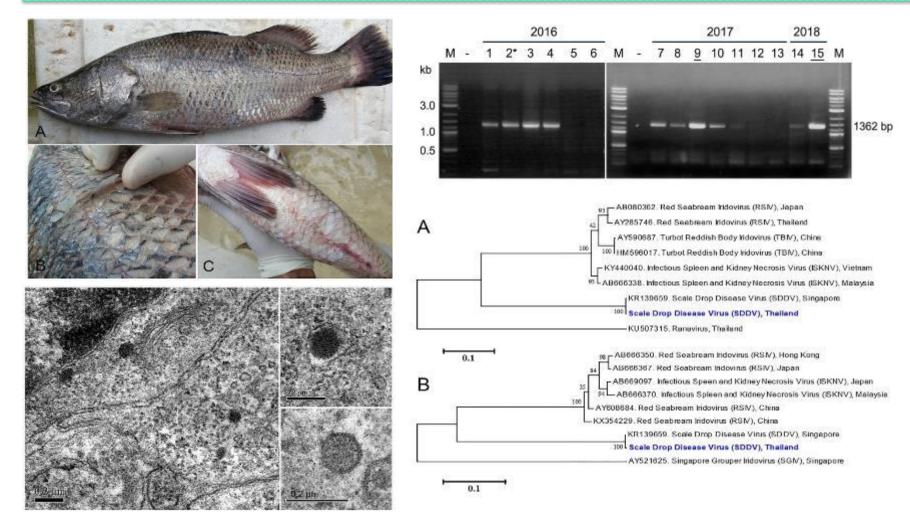
Scale drop disease

Caused by Scale drop disease virus (SDDV)



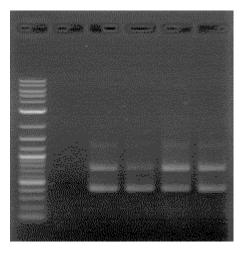
Scale drop disease

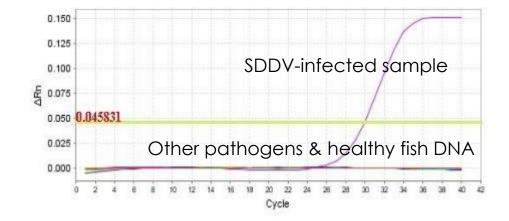
Caused by Scale drop disease virus (SDDV)



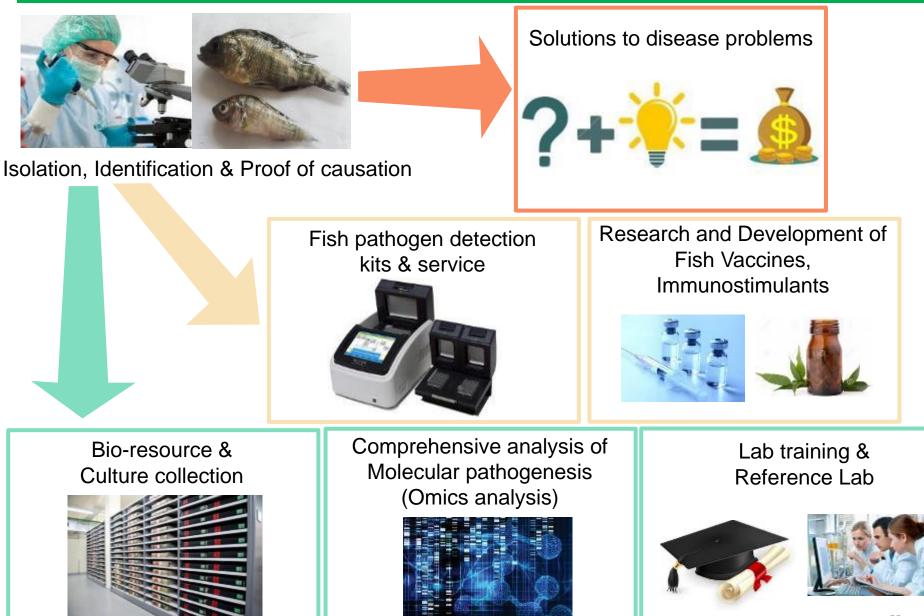
Scale drop disease virus (SDDV)

Occurrence in Thailand and PCR detection protocols





From Research to Solutions



Research Team



Dr. Krishna R Salin

Asian Institute of Technology Private companies & Non-profit organizations

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Information and media from the internetsGranting agencies

