















"Innovation, Translation, and Impact in Tropical Medicine"

Environmental DNA: a different approach for food/water borne helminths studies

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Scenario of NTDs in Endemic areas
Indigenous cases acquired in their own
community are observed among local people

- Related to low sanitary standarts
- Related to low education/information
- Related with the environmental conditions



Foodborne and waterborne diseases, as opisthorchiasis and schistosomiasis, are major threats to human health because their infective forms can be present in the daily life of everyone in endemic areas.

Usually the pathogen detection techniques requires sampling from hosts

- Fecal samples
- Serology is useful for diagnosis by detection of antibodies or antigens
- Other material other than serum samples can be used for detection of Ab or Ag
- Molecular techniques are applicable for diagnosis using any source of DNA





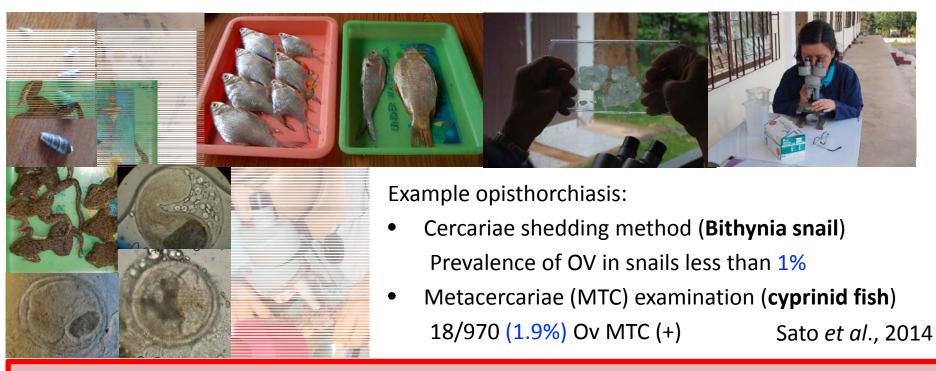
the most part of the time we are focusing in humans...

But... What is the whole situation?

What is the connection of "disease" in an ecosystem

To study these connections: Ecological Survey

Study the species related with the disease in a determined area: hosts, reservoirs, vectors, etc. understand their role in the **ecologic network of the pathogen** to describe the epidemiological scenario of the disease in the area.



Limitations of the classical methods

- Require high skilled techniques, manpower, time to perform, sometimes dangerous
- Difficult to track the distribution changes/apply in wide-ranging studies
- Bias in sampling can occur



To overcome the limitations of classical methods as a new/complementary alternative in ecological surveys

Environmental DNA

Technique to study the biota from environmental samples (water or soil) by detecting species/genus specific DNA (Ficetola *et al.* 2008; Minamoto *et al.* 2012)

DNA found in the environment

With several origins:

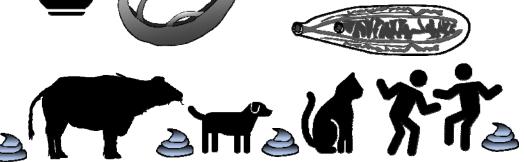
Excretion, secretion, exfoliation, reproduction, decomposition

and states:

intra/extra membranes, particulate and free

Now we are applying this technique for studies in parasitology





Methodology



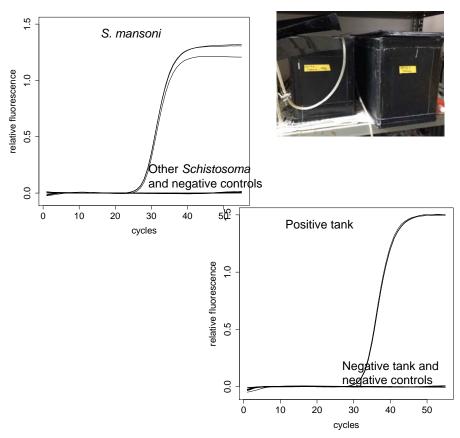
qPCR

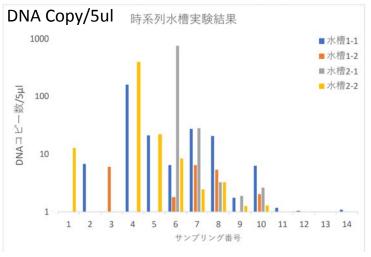
Specific primer/probe design

- ✓ Short Length products: 100 to 200 bp desirable
- ✓ DNA target: *CO1*, ITS (or other)
- ✓ Specificity check in silico: database

Determination of specificity/ sensitivity

- ✓ Compare with conventional PCR using known samples (from specimens and experimental infection water)
- √ Test the system using non-target DNA
- ✓ Confirm PCR positive samples by sequencing





Methodology

In the field

Samples collection (500ml each)

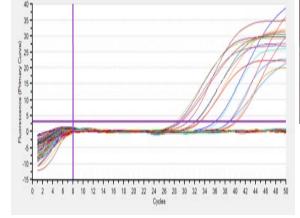
• Filtration of water (0.7 μm pore glass fiber filter)

• Fixation in Ethanol



In the lab

- DNA extraction and purification
- PCR and qPCR
- Sequencing, NGS





eDNA and opisthorchiasis in Lao-PDR

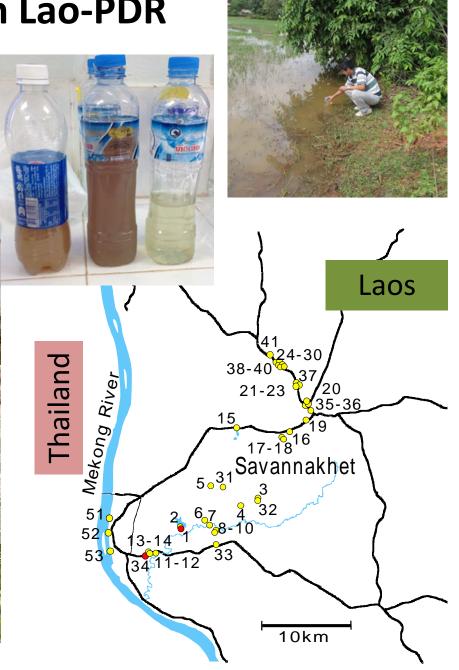
Savhannakhet, Laos

Total 44 points 62 samples

Ponds, rivers, rice field etc.

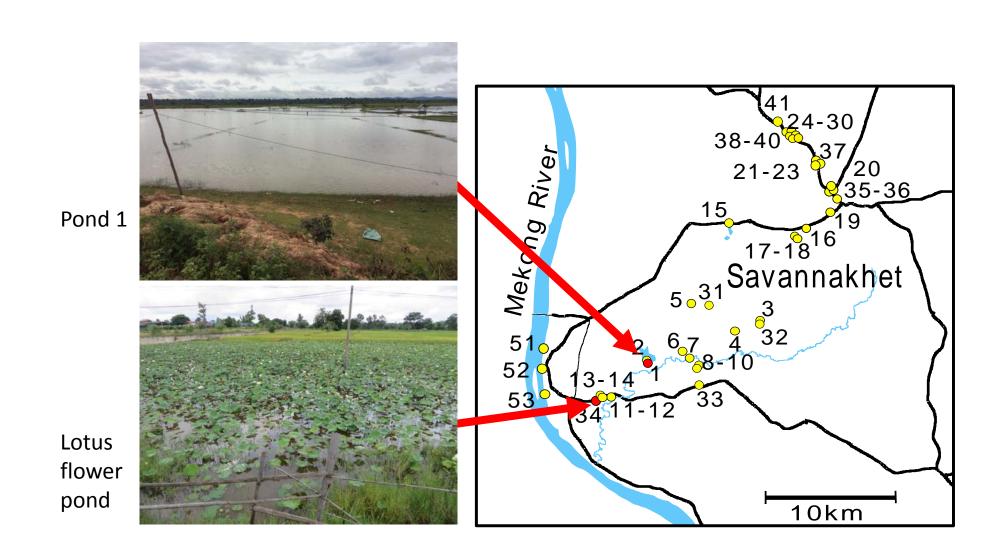


Hashizume et al., Acta Trop 169 (2017)



Results

Ov DNA was detected from 2 sites (pointed red)



Detection of 1st intermediate hosts

- Detection of Bithynid Snail by PCR
- eDNA samples of environmental water
- Difficult primer design (database info)

Detection of 2nd intermediate hosts

- Same DNA samples from environmental water
- PCR assay using the universal primers for fishes
- Next-generation sequencing
- Total 35 fish species were identified
- Including 9 species as host of Ov

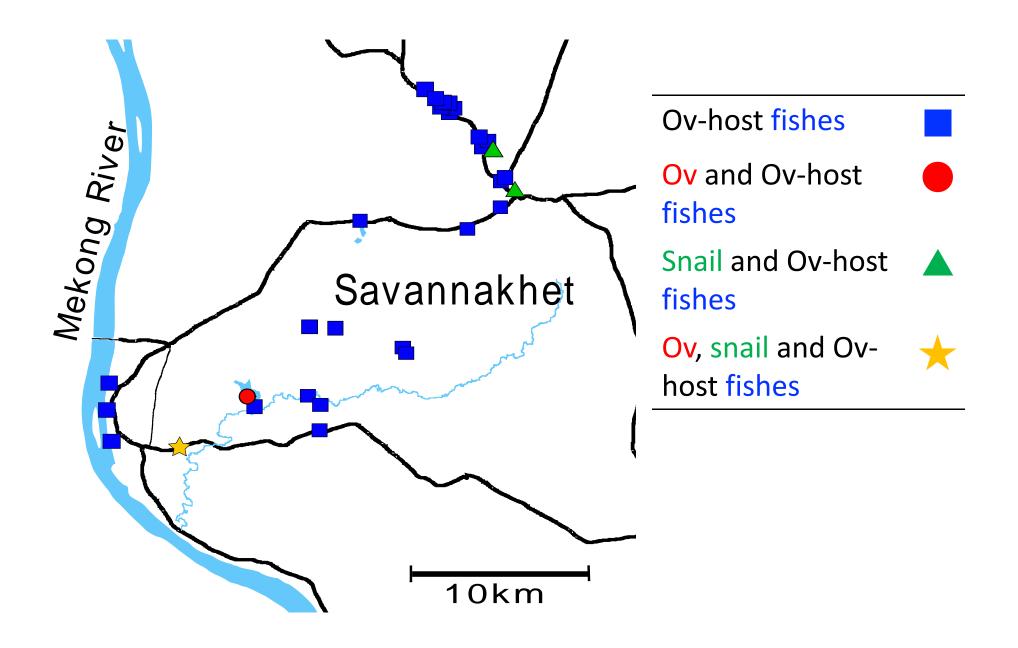
35 fish species detected from environmental water

Scientific name	Family	Scientific name	Family
Barbonymus gonionotus	Cyprinidae	Gambusia affinis	Poeciliidae
Catla catla	Cyprinidae	Clarias fuscus	Clariidae
Cirrhinus cirrhosus	Cyprinidae	Clarias gariepinus	Clariidae
Cyclocheilichthys repasson	Cyprinidae	Ailia sp	Schilbeidae
Cyprinus carpio	Cyprinidae	Pangasius larnaudii	Pangasius
Esomus metallicus	Cyprinidae	Sperata seenghala	Bagridae
Hampala dispar	Cyprinidae	Anabas testudineus	Anabantidae
Henicorhynchus lineatus	Cyprinidae	Nandus nandus	Nandidae
Hypophthalmichthys nobilis	Cyprinidae	Channa marulius	Channidae
Labiobarbus leptocheilus	Cyprinidae	Giuris margaritacea	Eleotridae
Mystacoleucus marginatus	Cyprinidae	Oxyeleotris marmorata	Eleotridae
Osteochilus hasseltii	Cyprinidae	Oreochromis aureus	Cichlidae
Osteochilus salsburyi	Cyprinidae	Oreochromis niloticus	Cichlidae
Probarbus jullieni	Cyprinidae	▲ Oreochromis sp. 'red tilapia'	Cichlidae
▲ Puntius aurotaeniatus	Cyprinidae	Trichopodus trichopterus	Osphronemidae
Raiamas guttatus	Cyprinidae	Betta splendens	Osphronemidae
Acantopsis sp	Cobitidae	Macrognathus pancalus	Mastacembelidae
Clupeichthys aesarnensis	Clupeidae		

Ov-host fishes are written by red-letter,

— were the species detected in No.1 Pond, ▲ were detected in No.34 lotus pond

Epidemiological scenario for opisthorchiasis in Savannakhet



eDNA and Schistosomiasis in Madagascar

Local of study:

Maevatanana area

mae = bad

vatana = beautifull

tanana = village

Local of bichuba = bigbelly

Schistosoma mansoni and S. haematobium

Clinical cases







Sato et al., Int J Inf Dis (2018)



Water sampling

- 7 spots (21 samples of 500ml)
- 3 in *S. haematobium* cases area
- 4 in *S. mansoni* cases area







S. haematobium endemic area



Water used for agriculture, washing clothes, bath and drink. pH8.5-8.8, 25-27°C. no snails found

S. mansoni endemic area



Water used for washing clothes, bath and for drink. pH8.5-8.8, 19-20°C. snails were found

All eDNA samples were subjected to a blind test.

From the tested material **one spot was positive for** *S. mansoni*, **the sample 5**

- The sample 5: Taken in small river where the people use to take bath drink, washing clothes, etc in a *S. mansoni* cases area.
- That was the place where we could confirm *Biomphalaria pfeifferi* snails
- The real time PCR result was confirmed by conventional PCR and direct sequencing.
- We could test successfully our primer-probe set in the field with satisfactory specificity.

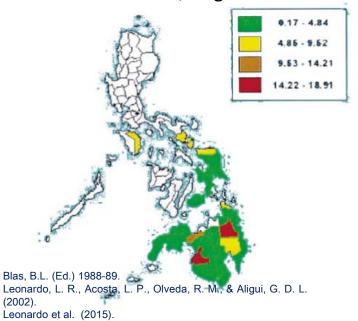


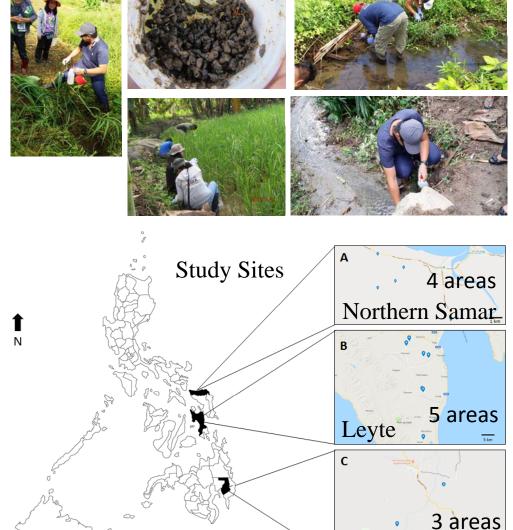
eDNA can be an useful tool for eco-epidemiology studies on schistosomiasis

Development of an eDNA detection method for Philippine isolates of *Schistosoma japonicum* and *Oncomelania hupensis quadrasi* from field water samples

Schistosomiasis japonica in the Philippines

- Endemic in 28 provinces, 14 cities, 189 municipalities, and 2,221 barangays
- 12 million Filipinos are living in endemic areas and 2.5 million are directly exposed to infection
- New endemic foci discovered in 2002 and 2005 respectively in Gonzaga, Cagayan and 2005 in Calatrava, Negros Occidental





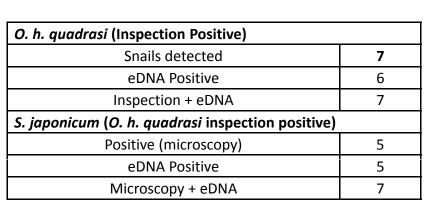
Fornillos et al. (submitted)

Compostela Valley

Results: qPCR of Field Water Samples

Snails observed sites

ı Cooil	Infection Rate	qPCR Results		
+ Snail I Site		<i>S</i> .	O. h.	
		japonicum	quadrasi	
PINOPos	2.08	+	+	
GDNSPos	0.43	+	+	
CBRNPos	1.41	+	+	
SOCSPos	0.24	-	+	
CSLNPos	0.2	_	(-)	
DITAPos	ND	+	+	
TIG	ND	+	+	
Total 7	5	5	6	



Sj: Malacological surveys can be complemented with eDNA detection of both the snail and the parasite.



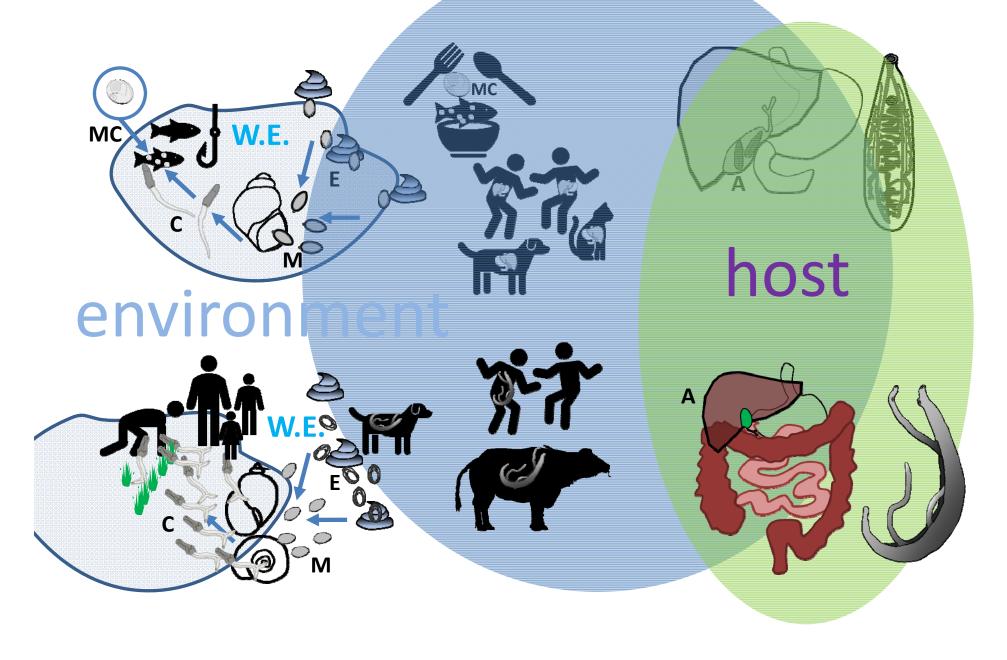


No snails sites (inspection)

	qPCR Results		
Snail Site	<i>S.</i>	O. h.	
	japonicum	quadrasi	
PINONeg	(+)	-	
CBRNNeg	-	-	
SOCSNeg	-	-	
CSLNNeg	-	(+)	
DITANeg	-	-	
WAS	-	(+)	
MGS	(+)	-	
LIBW	-	-	
LIBBH	-	-	
OLR	-	(+)	
MAP	+	-	
NEP	-	-	
Total 12	3	3	

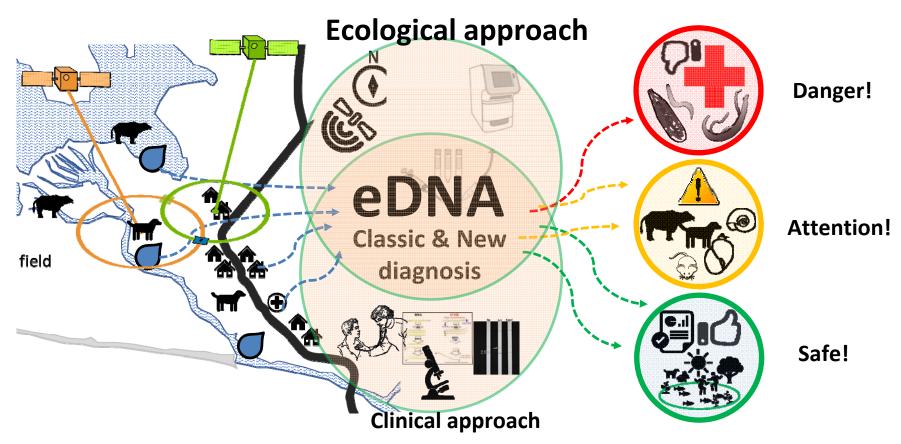
O. h. quadrasi (Inspection Negative)		
No snail	12	
eDNA Positive	3	
S. japonicum (O.h. Negative)		
eDNA Positive	3	

life cycles in a complex network



One-health

Water and food Safety



We hope to contribute with a more accurate surveillance technology to be applied in endemic areas, driving policies for the control of NTDs.

