Molecular taxonomy of the Spirochonidae (Chonotrichia, Ciliophora) from the Baïkal Lake. First results.

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ABSTRACT. The Baïkal Lake appeared some twenty-five million years ago, and is actually the biggest freshwater lake in the World.
It harbours more than 600 plant and 1500 animal species, including more than 250 freshwater amphipod species. These crustaceans host a number of epibionts, among which the ciliates are represented by various kinds of Peritrichia, Suctoria and Chonotrichia.
According to the Jankowski book, («Chonotricha», 1973), at least 10 species (in 3 genera) of the Spirochonidae family live on diverse appendages of Baïkal amphipods.
In order to complete the cytological datas, we planed to obtain molecular data.

In july 2018, during a 9-days scientific cruise on the Baïkal Lake, we have collected amphipods from sediments dredgings.

More than 100 specimen were examined and dissected under binocular, 15 of them carrying spirochonids.

The gills were micro-photographed and stored in ethanol.

In 2019, DNA was extracted, SSU-amplified and sequenced. The first molecular datas are presented here.

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In memory of Denis Lynn (1947-2018)



GENEVA – IRKUTSK - BAIK



THE BAÏKAL SPIROCHONIDAE PROJECT The "Baïkal Spirochonidae Project" was initiated at the ICOP_2017_Prague, by Lynn – Fahrni – (Sherbakoy). THE JOB: to get a phylogenomic tree corresponding to the huge morphological diversity described by JANKOWSKI (1973).



GVA : Geneva IRK : Irkutsk

Geneva - Irkutsk: ca. 7'000 km 6 time zones : GVA + 6 h Flight : Geneva - Moscow / Moscow – Irkutsk

IRKUTSK-BAÏKAL LAKE



Baïkal Lake is a paradise for amphipods (250 species) and for the chonotrich Family Spirochonidae . According to Jankowski, the family counts 2-3 genera (*Cavichona, Spirochona, Serpentichona*?), with about 25 different species, leaving on some 30 different host species.

Goals: **Exploring the Jankowski's work on the Spirochonidae Family** The main goals of the study are:

Obtaining a clear phylogenic tree of the family;
Clarification of some synonymies (*Serpentichona / Spirochona - anthus*),
(*Cavichona / Spirochona – elegans*), (*Cavichona / Spirochona - elegantula*);
Clarification of the "genus" status of some (several) "similar" species;
Possible new genera or species;
Relationships between hosts and epibionts;
Possible co-evolution between hosts and epibionts (Sherbakov).

Tasks: The tasks were roughly distributed as follow:

Fahrni: - collection of Spirochonidae living on amphipods from Baïkal Lake (July 2018);
 Lynn: - DNA extraction, sequencing, phyllogenomy of the collected Spirochonidae;
 Sherbakov: - organisation of the sampling cruise on the Baïkal Lake;
 - phylogenomic of the collected amphipods hosts.

Funding: Lynn, Fahrni (?), Sherbakov.

Actual state

- I have participated to the 2018 sampling cruise on Baïkal Lake (12-21.07.2018).
 I have recolted gills bearing chonotrichs (photos) from some 18 amphipods hosts; gills are preserved in EtOH (for DNA extraction), in osmium (for protargol staining), and in glutaraldehyde (for SEM examination). Cytological studies in progress.
- I keep the EtOH preserved gills samples.
 Dr. Sherbakov keeps the 18 amphipods specimen (in EtOH) for molecular identification and phylogenomic study.

The tragedy : Denis Lynn died accidentally on 26.06.2018.

29.10.2018 : J. Fahrni restart the molecular work, hosted in the laboratory of Ass. Prof. J. Pawlowski * *Dpt. Of Genetic and Evolution, University of Geneva, Switzerland

Cavichona elegans.



The Spirochonidae (Jankowski, 1973)

3 genera : Cavichona, Spirochona, Serpentichona;

13 (14). Large forms with notable pseudostyle.

14 (13). Small forms without distinct pseudostyle

- 25 species : - C. abyssalis ; acanthogammari ; brevistyla ; carinuri ; cincinnata ; crassa ; elegans ; elegantula ; exilis ; laticollis ; multifida ; oligocineta ; protecta ; rhabdomorpha ; tenue ; virgata .
 - Spirochona brevis ; gemmipara ; globulus ; gnatopodialis ; halophila ; marina ; simplex ; tuba .
 - Serpentichona anthus .

JANKOWSKI's DETERMINATION TABLES				
JANKOWSKI's DETERMINATION TABLES Identification of the genus Serpentichona (p. 276) Jankowski_1973 Identification of the genus Serpentichona (p. 276) Jankowski_1973 Control Serpentichona is monotypical, hence a table is absent. (vortex = funnel ???) Jankowski's diagnosis: • Very large forms, with length up to 300 µm. • The body is sharply elongait, rather narrow, non flatened, broadened in upper part. • Pf's large, wide, rather hourd the underdevelopment of fumal (usually 2½ md less, rarely up to 3¼ turns) is characteristic. • The ventrial edge of PF uneven, with central cut (as in <i>Corkehona</i>) and series of additional cut to the right and left of central cut, visible already in early ontogenesis. • Cornice is no tunited as in spirochones and not double as in corkehone, under several (up to 8) self-depended cornice.		-		
- Neck is good developed. - Predictive very long, measures up to ½, rarely up to 2.3 length of body. - Peldicula with longitudinal nibbed folds. The genus is differ: - from Sprochona by presence of ventral cuts, folds of PF and by elongation of pseudostyle; - from Carkchona by polymerization of ventral cuts, by presence of folds, and trend to underdevelopment of funnel. Keys for identification of genera from family Spirochonidae (p. 238) 1 (2). Ventral edge of peristomal funnel (PF) is regular, without folds and cuts; pseudostyle distinct, and elongated in the most of species. 3 (4). There is single cut. Carkchona (p. 254).	and a			
4 (3). There are numerous cuts. .3. Genus Serpentichona (p. 276). Keys for identification of species of genus Spirochona (p. 239) Jankowski_1973: page#; fig# (fig-page) 2). Freshwater species with moderate PF. (PF = pre-oral funnel = peristomal funnel) (2). The body is dorso-ventrally flattened. 4. S. stmplex. 248; 110 (250) (2). The body is short flattened, with circular section. 3. S. globulus. 246; 108 (249) 5 (4). The body is longate, L/D > 2:1. 3. S. globulus. 246; 108 (249)				
6 (7). Very big side-pockets ; series of additional dorsal folds of PF present. .6. S. tuba. 251; 1/2 (252) 7 (6). The side-pockets are of the usual mode; additional dorsal folds of PF absent. .8. S. brevis. 246, 104 (245) 9 (8). Large forms (about 70 µm); PF with 2½ revolutions. .2. S. brevis. 246, 104 (245) 9 (8). Large forms (about 90 µm); PF with 2½ revolutions. .1. S. genmipara. 244; 103 (244) 1 (10). The neck is very wide, up to ½ of body width; the lower end of body is distinctly elongate. .5. S. gnathopodtalis 249; 111 (251)	đ		Рас. 117. <i>d</i> - <i>Г</i> = 1 по: Сварченскай, 1	Салісіона сіоран с Ілістрополого (Б.– Г.– Салісіона сіоран с Ілісіонаратага согосова (Б.– Г.– 1996), Л.– В. — С. сіпсіквара с Електоратага vagi; З.– грабна у вояка С. abyaala.
	the second se			

.....8. S. marina. 253; 115 (254)7. S. halophila. 253; 114 (253)



Puc. 122. Carichons oligocinets c Eulins

Cavichona oligocineta.

Eulimnogammarus sp.

Serpentichona_morph-2.docx

Genus Cavichona morphology

Cavichona cincinnata.

Eucarinogammarus wagi



Genus Cavichona morphology

ssa.	Cavichona carinurii.	Cavichona elegantula	Cavichona protecta
	Fre 18. Carbon arriad a Carbon	The first descent of the second descent of the second descent of the second descent des	Image: wide wide wide wide wide wide wide wide

Odontogam. calcaratus

(JANKOWSKI, 1973)

Brandtia lata



Pat. 124. Carichena multi/ida e Eulennepanmarus spanase

Cavichona multifida.

Eulimnogammarus cyaneus

(JANKOWSKI, 1973)

Preliminary molecular work

BAÏKAL 2018 – Amphipods samplings

TUBE #	DATE	AMPHIPODS	TOTAL	Spd	Freq.
	TH 12.07.18	1 ???			
		3 Pallasea			
		3 Eulimnog.			
		4 Acanth	11	1	9,1 %
	FRI 13.07.18	1 Pallasea			
		2 Macroectopus			
		1 Gmelinoides	4	1	25 %
	SA 14.07.18	1 Eulimnog.			
		1 Acanthog.			
		1 Pallasea			
		3 Macroectopus	6	0	0
	SU 15.07.18	2 Pallasea			
01		10 Acanthog.			
02	140.45.07.40	6 Pallasea	18	2	11,1 %
11-13	MO 16.07.18	3 fff	5	U	0
14 - 19	1017.07.18	1 Brandtia 7 Gradianidae			Freq. 9,1 % 25 % 0 11,1 % 0 41,7 % 15 % 6.3 % 6.3 %
		7 Gmelinoides			
		2 Eulimnog.			
		1 Odontos	12	5	417%
20-22	WE18.07.18	5 222			42,7 78
20 22		4 gammarids ?			
		1 Acanthog.			
		1 Eulimnog, cvaneus			
		3 Eulimnog. verrucosus			
		6 Brandtia	20	3	15 %
26, 27	TH 19.07.18	4 Eulimnog. vittatus?			
		15 Eulimnog. ?			
		5 Acanthog.			
		1 ???			
		1 Odontog. white			
		3 Pallasea			
		3 Brandtia	32	2	6.3 %
28 - 30	FR 20.07.18	3 Eulimnog. sp.			
		1 Eulimnog. cyaneus			
		2 Eulimnog. violaceus			
		1 Acanthog.			
		3 Brandtia			
		2 ??? white			
		2 Pallasea	14	3	2,4 %
21 tubes			106	17	16 %
	CA 24 07 40	Park to Justicely STATENED			

BAÏKAL – MOLECULAR 1

BLAST IDENTIFICATION OF THE CILIATES AND THEIR HOSTS (BAÏKAL_July 2018)

Cavichona cincinnata.

Eulimnogam. verrucosus | Eucarinogammarus wagi

A	mplif_date	Sample	PCR/no	Minigel	Seq.	BLAST	PCR/no	Minigel	BLAST	HOST	
			CIL				CoxV				
1	8.03.2019	BK 19		+	+	Spg_Lynn 99%		+	FJ756323	Eul-cyaneus 87%	
		BK 21		+	+	Spg_Lynn 99%		+	FJ756323	Eul-cyaneus 87%	
		BK 26		+	+	Spg_Lynn 98%		+	FJ756344	Pall_kessleri 95%	
		BK 27		+	+	Spg_Lynn 99%		+	AY926654	Brandtia lata 94%	
		BK 30		(-)	(-)	Spg_Lynn 99%		+	AY926654	Brandtia lata 96%	
			CIL				CoxV				
1	3.06.2019	BK 15.1	63	-	-	-	83	(+)		Eul-vittatus/verrucosus	
		BK 19.1	64	-	-	-	84	(+)			
		BK 20.1	65	(+)	(+)	Suct. Acineta	85	+		Pallasea/Babr/Nyphargus	
		BK 20.2	66	-	-	-	86	+		Pallasea/Babr/Nyphargus	
		BK 28.1	67	-	-	Suct. Acineta	87	+		Eul-virridis/verrucosus	
		BK 28.2	68	-	-	Suct. Acineta	88	+		Eul-verruc/Garajewia/G-balc.	
		BK 29.1	69	+	+	Suct. Acineta	89	++			
		BK 29.2	70	+	+	Suct. Acineta	90	++		Acanthogam-victorii	
		BK 29.3	71	+	+	Peritr. Vorticella	91	++		Acanthogam-victorii	
		BK 32.1	72	+	-	-	92	+		Brandtia_lata/latissima	
		BK 33.1	73	(+)	-	-	93	(+)		Brandtia_lata/latissima	
			CIL								
2	4.06.2019	BK 15.1	102	-		-					
		BK 19.1	103	-		-					
		BK 20.2	104	-		-					
		BK 29.1	105	+		Acineta flava					
		BK 29.2	106	++		Acineta flava					
		BK 32.1	107	(+)		Spg_Lynn					



Cavichona morph-2a.docx

in black: Available Chonotrichia sequences (with Dysteria as root);
 in green: Baïkal sequences (this study);
 in blue: S. gemmipara _Lynn_2016.



BAIKAL – MOLECULAR 3

RESULTS

- 1) The number of Baikal spirochonids sequences obtained is low : (5 sequences / 106 amphipods analysed) !
- 2) With 18S (CIL), the 5 sequences are 99% identical between them
- 3) With 18S (CIL), the 5 sequences are 99% identical to the S.gemmipara_Lynn_2016 seq.;
- 4) With 18S (CIL), the 5 sequences are 99% identical to the S.gemmipara_Fahrni_2019 seq. (unpubl.);
- 5) The 18S (CIL) sequences of several other ciliates (Suctoria, Peritrichia) were obtained ;
- The 18S (CIL) gene fragment is not appropriate to discriminate different species (if any ?);
 6) Other gene fragments are tested (V9 ; ITS; 28s) ; works in progress ;
- 7) Five D1-D2 (28S) sequences were obtained; they contain very SLIGHT differences between them ! (see " BAIKAL MOLECULAR 2 ")
- 8) All gene fragments tested gave results with positive control species (Paramecium, Vorticella, ...);
- 9) The amphipods molecular identication with the CoxV fragment is satisfying .

CONCLUSIONS

- 1) General contamination with *S.gemmipara*_Fahrni_2019 material ? unlikely (see tree_7) ?
- 2) The D1-D2 sequences must be checked; then sequenced again;
- 3) The D1-D2 amplifications and sequencing must be done again;
- New DNA extractions are needed;
- 5) Getting new amphipods (and spirochonids !) from an other Baïkal_2019 cruize ?



cruize-sampling_3.doc

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-in blue: total number (by day) of amphpipods collected and examined.
 -in red: total number (by day) of amphipods bearing Spirochonidae.
 -in black: frequencies (in percent) of amphipods bearing Spirochonidae.

C. Ko

baïkal – molecular_2.docx

LITERATURE

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- LYNN, DH. 2016. The small subunit rRNA gene sequence of the chonotrich Chilodochona carcini Jank., 1973, confirms chonotrichs as a dysteriid-derived clade (Phyllopharyngea, Ciliophora). Intern. J. of Syst. and Evol. Microbiol., 66, 1–6.

2) D1-D2/28S RIBOSOMALTREE (tree_9) Baikal only !

- in black : Baïkal sequences (this study) ;

CONCL.: The D1-D2 fragments contain very SLIGHT differences !

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(Dr. D. Sherbakov's help needed.)

6) Looking for others gene fragments needed

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