



Molecular detection of Sarcocystis spp. in meat samples

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Sarcocystis

- Phylum Apicomplexa
- More than 150 species
- Ubiquitous
- Two hosts to complete the life cycle:
 - An intermediate host (prey)
 - A definitive host (predator)

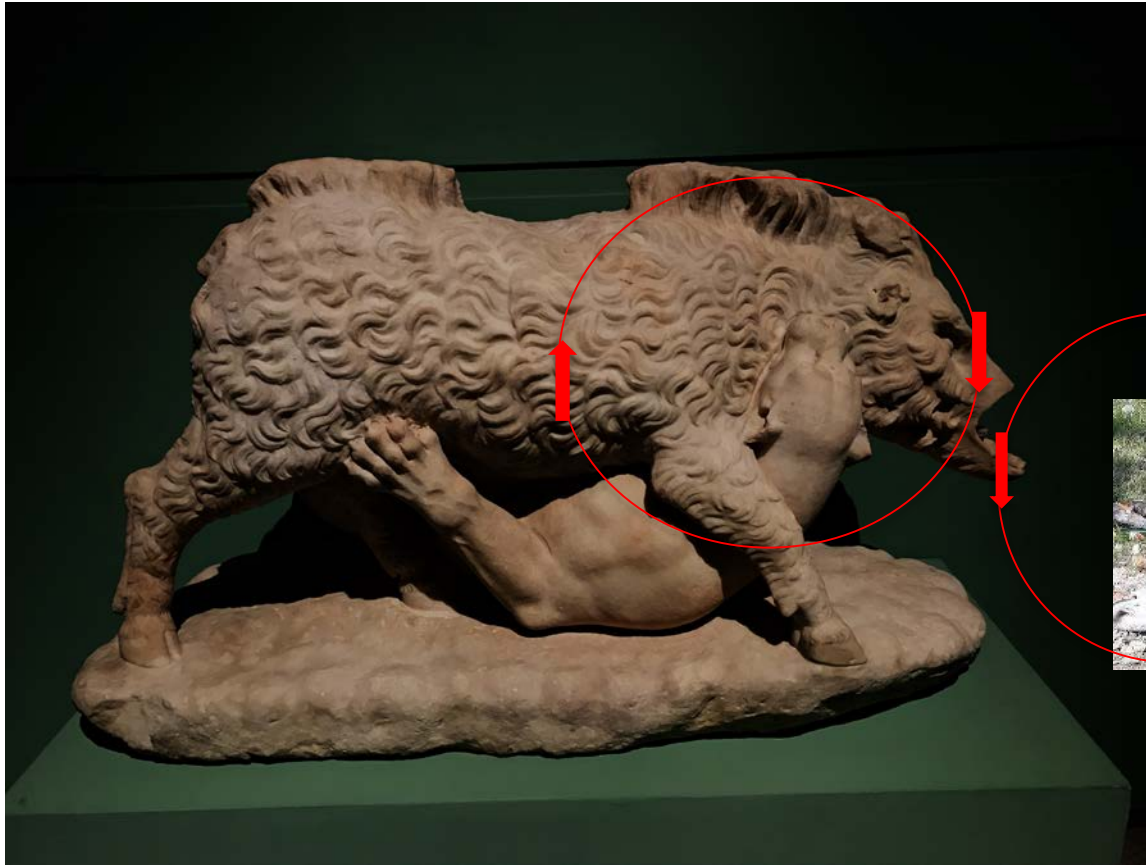
Sarcocystis

Intermediate hosts

- Many herbivorous mammals
- Birds
- Reptiles
- Primates
- Humans

Definitive hosts

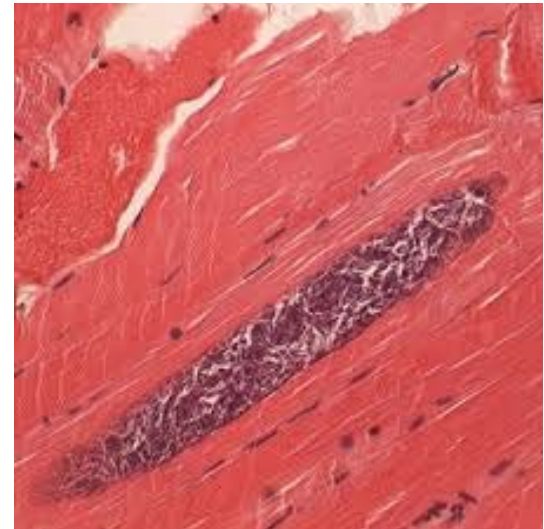
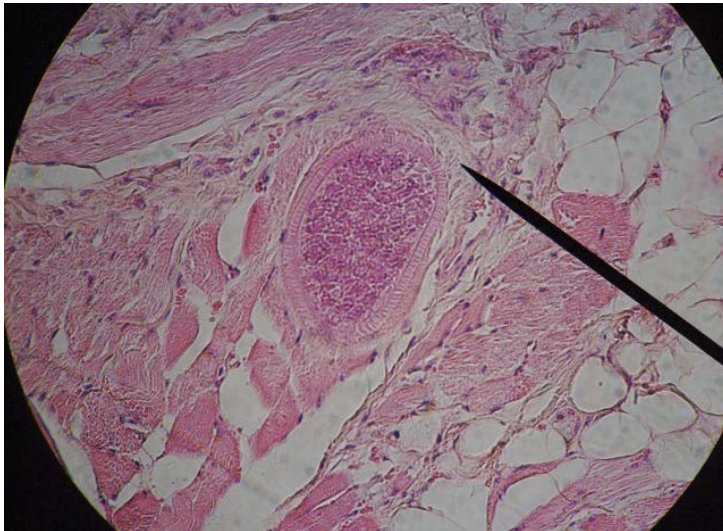
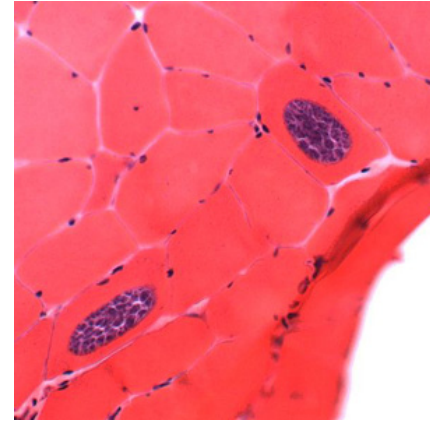
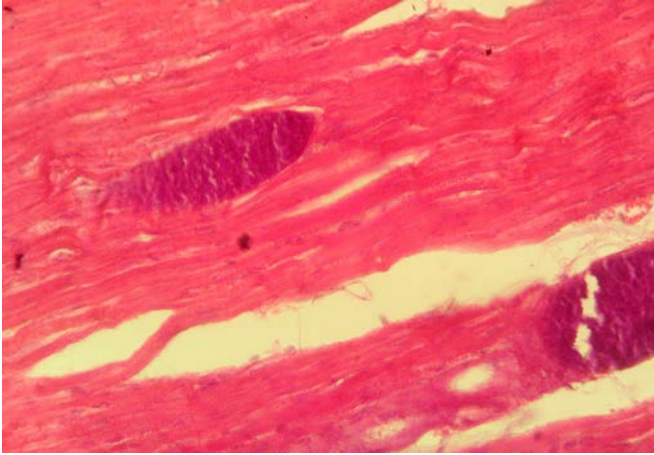
- Carnivorous and omnivorous mammals
- Raptorial birds
- Reptiles
- Humans



Predator-Prey Cycle

Omnivorous animals can play both roles: intermediated or definitive host.

Intermediated hosts develop Intramuscular cysts

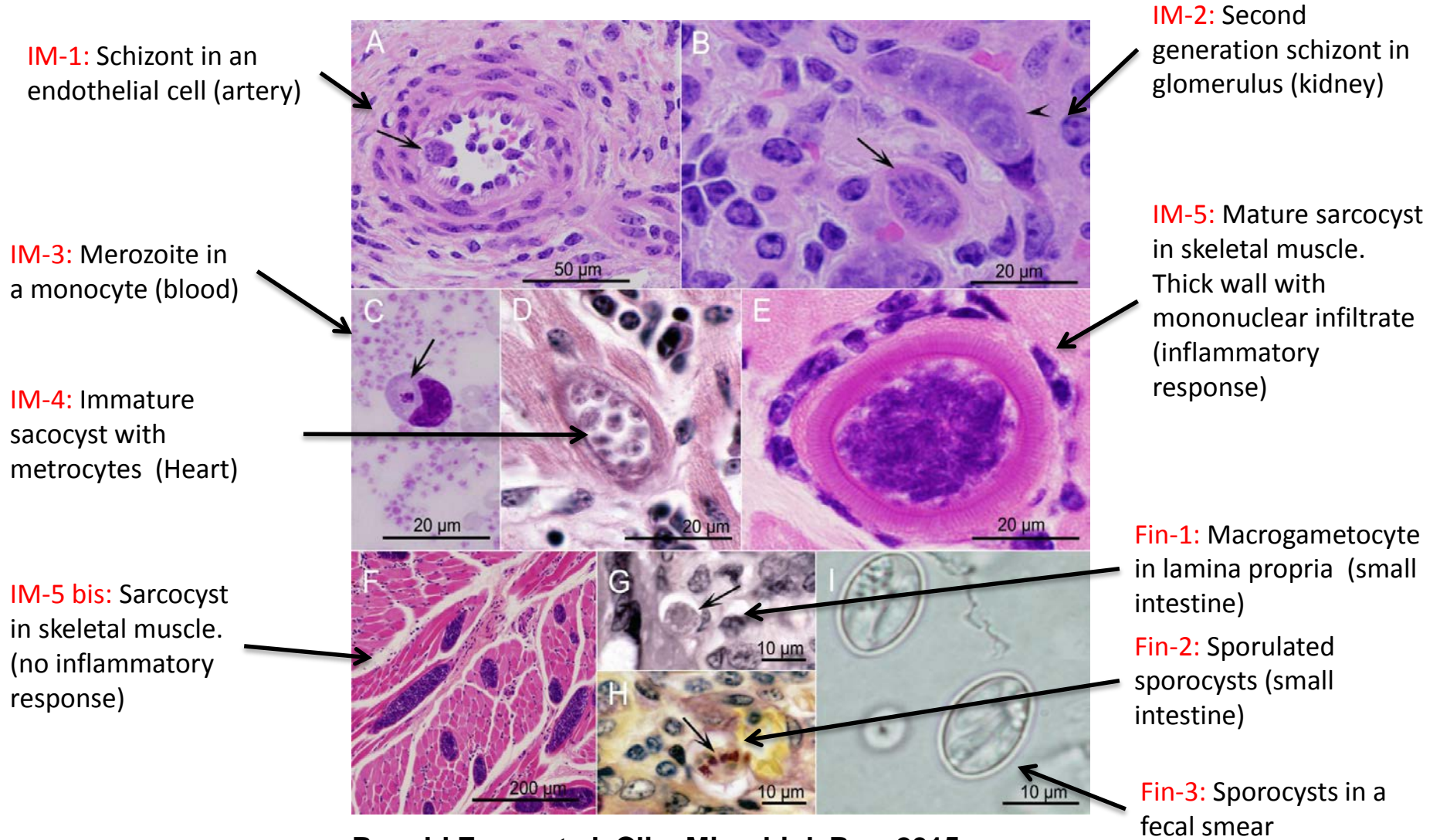


Definitive hosts: sexual cycle in the small intestine up to the excretion of new sporocysts and oocysts in stool

Sporocystis



Sarcocystis stages in tissues of intermediate hosts (A to F) and definitive hosts (G to I).

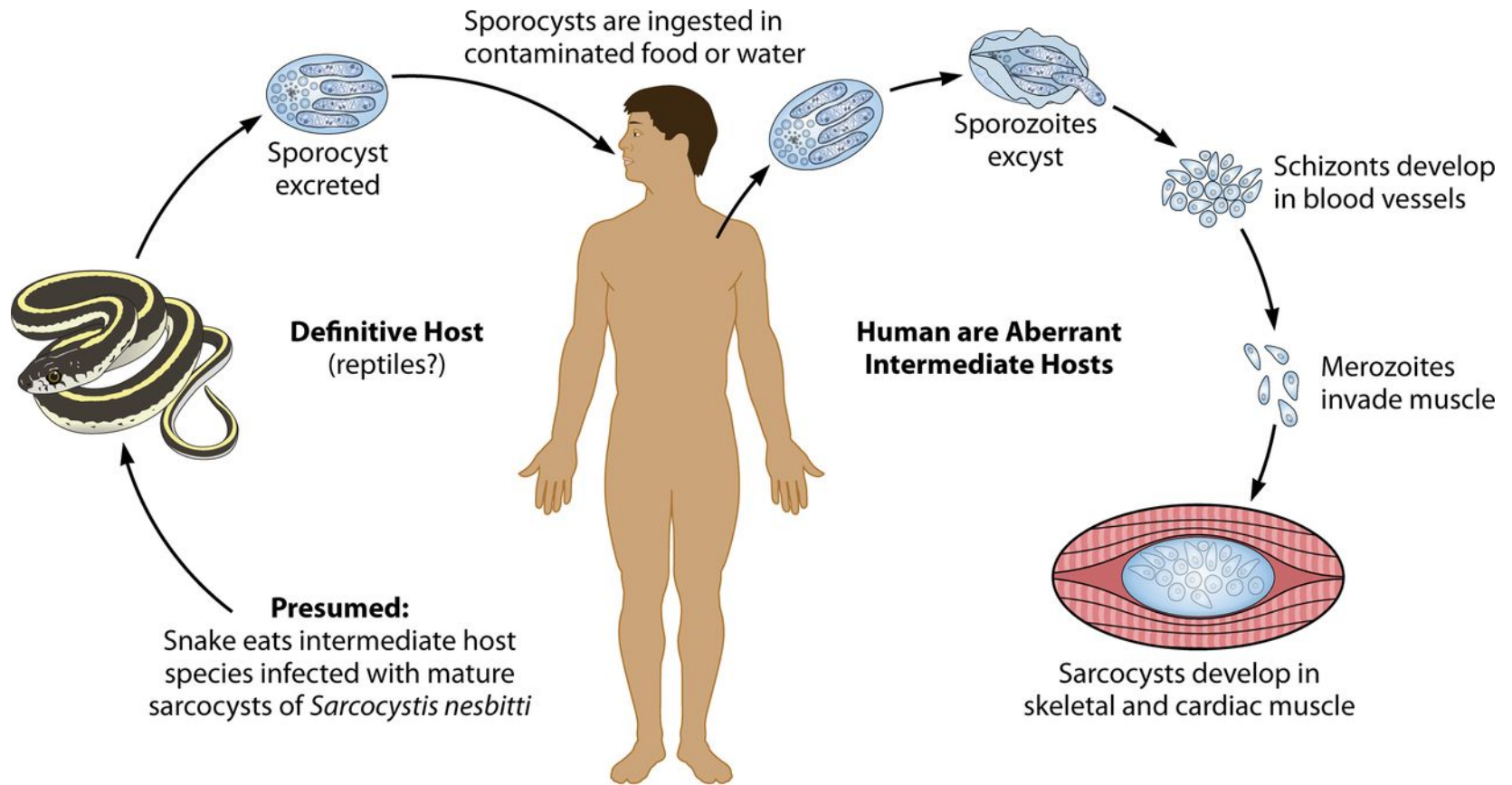


Ronald Fayer et al. Clin. Microbiol. Rev. 2015;
doi:10.1128/CMR.00113-14

Sarcocystis spp. in Humans

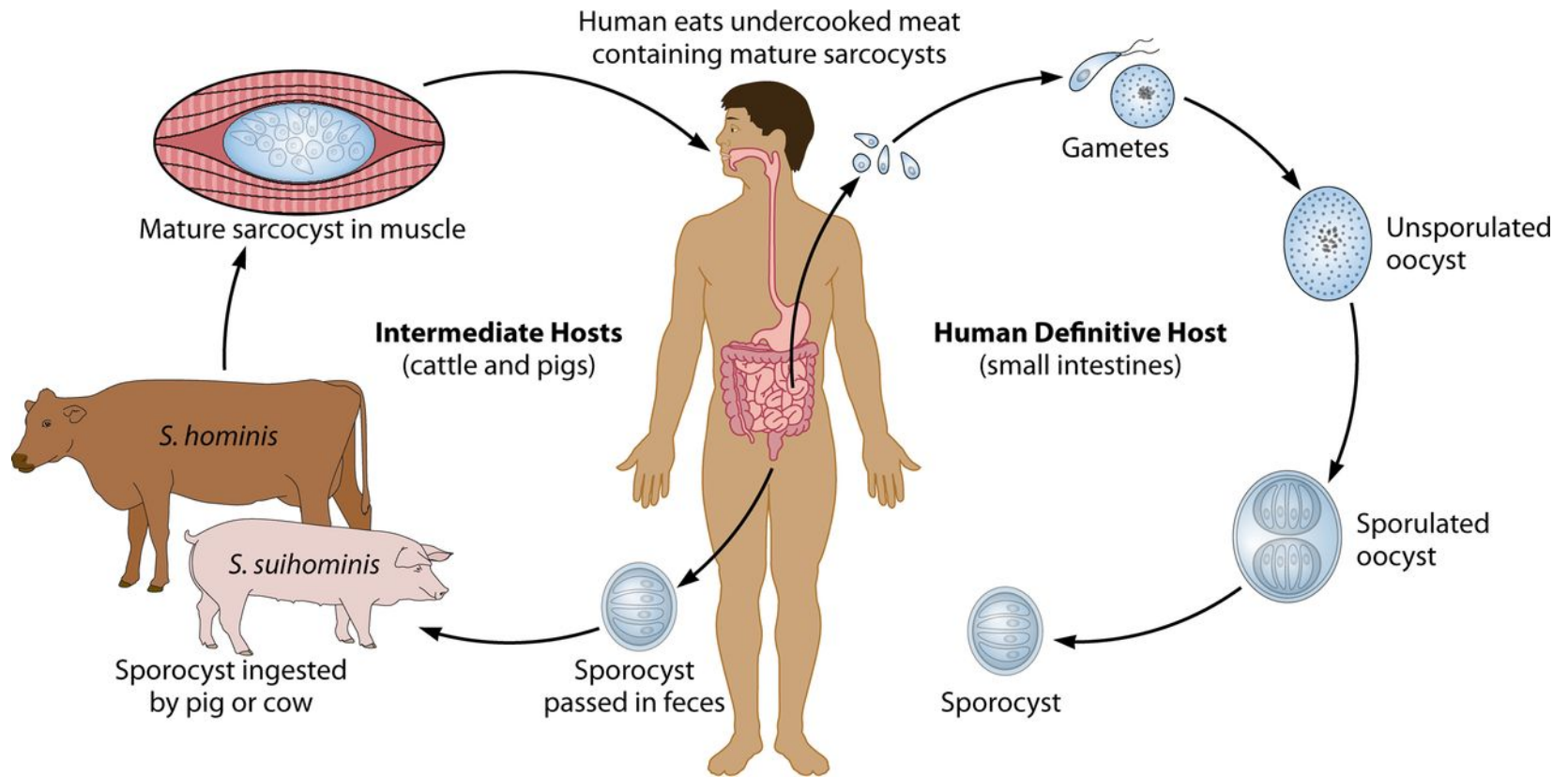
Man as intermediated and definitive
host

Humans as aberrant intermediate hosts for *Sarcocystis* species.



Ronald Fayer et al. Clin. Microbiol. Rev. 2015;
doi:10.1128/CMR.00113-14

Humans as definitive (final) hosts for Sarcocystis species.



Ronald Fayer et al. Clin. Microbiol. Rev. 2015;
doi:10.1128/CMR.00113-14

Sarcocystis: a problematic Taxonomy

Based on intermediated-definitive hosts

- *S. capreolicanis* = roe deer (intermediated)-dog (definitive)
- *S. suishominis* = pig (intermediated)-human(definitive)

Limits:

- Often one of the host is not known
- The same species can infect different intermediated as well definitive hosts

Based on morphological parameters

- Aspect of sporocysts and/oocysts (definitive host)
- Shape and size of tissue cysts (intermediated host)

Limits:

- Morphology of the sporocysts/oocysts is very similar
- Morphology of the cysts is influenced by various factors: (host, tissue, age, etc.)

Molecular Methods for Detection and Identification of Sarcocystis species

- Establishing an unequivocal genetic identity
- Discrimination of species with similar morphology
- Determination of undefined cycles: intermediate and definitive hosts of a given species
- Molecular epidemiology of sarcocystosis among animals and humans
- Unveiling of possible hidden hosts

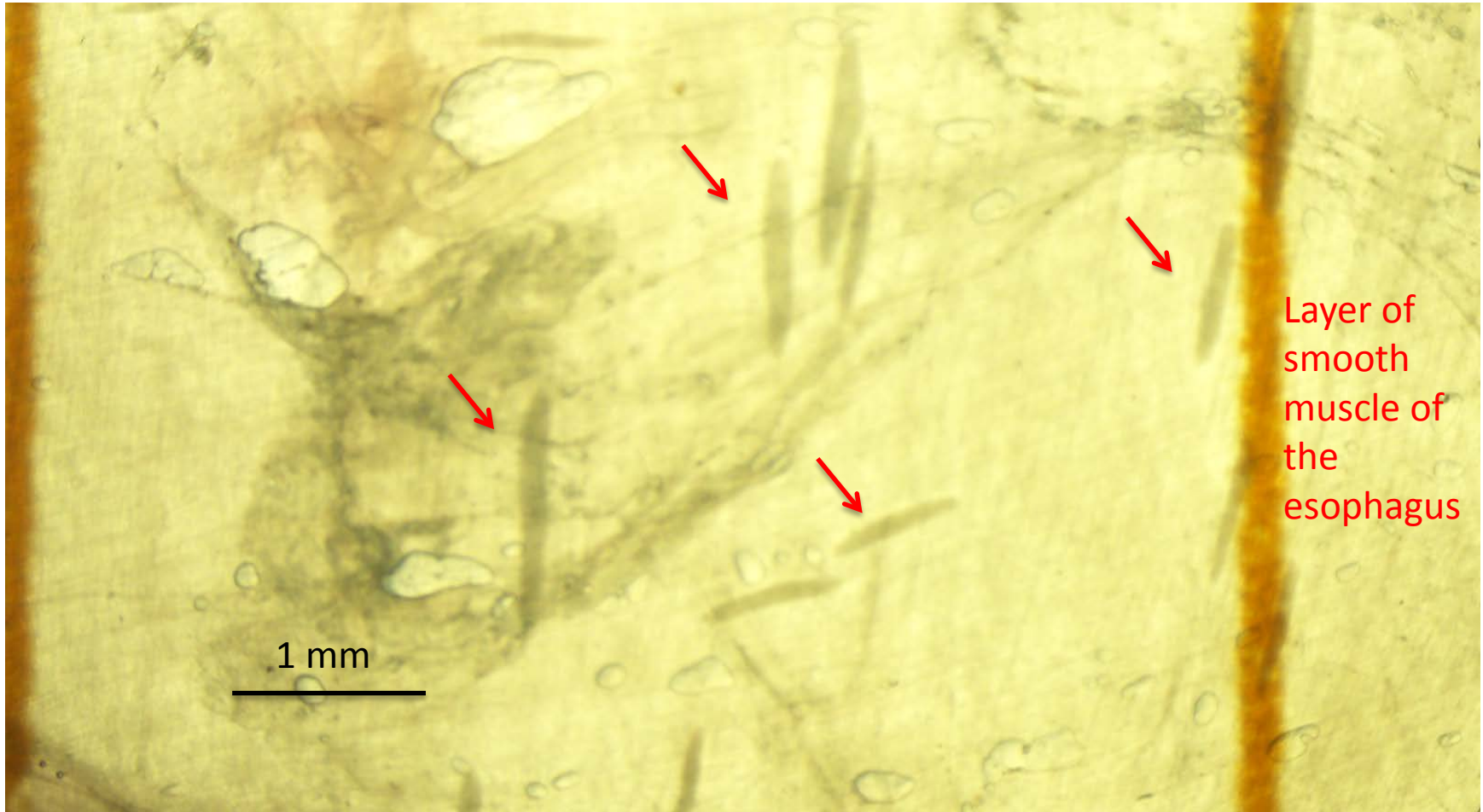
AIMS

- Development of a standard procedure for the molecular identification of Sarcocystis species in meat for human consumption
- Detection of zoonotic species by PCR and sequencing

To Develop a Simple Procedure Based on PCR and Sequencing

- Collection of samples of tissue with sarcocysts
- Collection of DNA samples from identified species
- DNA extraction from cysts or infected tissue
- Selection of reference DNAs to use as controls for PCR assays
- Development of a specific PCR assay
- Sequencing of amplification products
- Identification of species by homology with referenced DNA sequences

Sarcocystis from animal muscles



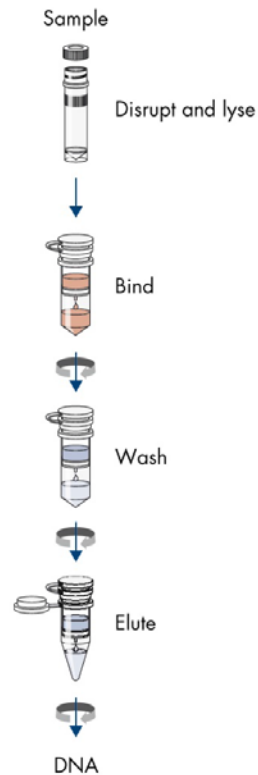
Isolated Sarcocyst in esophagus



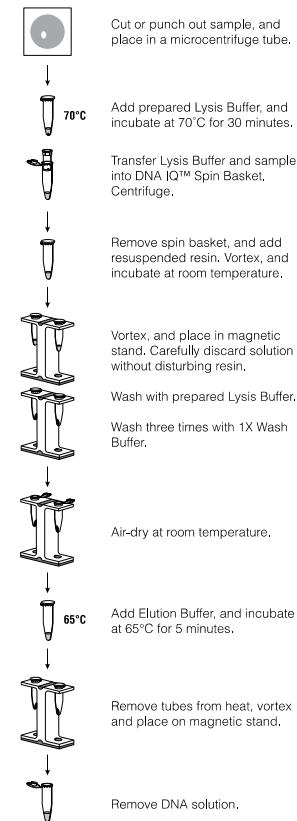
Two different methods for the DNA extraction

QIAamp technology based on silica gel affinity (more suitable for direct extraction from meat)

QIAamp Fast DNA Tissue Procedure



Promega IQ system based on magnetic beads (more suitable for extraction from isolated cysts)



Whole Genome Amplification of positive DNAs for reference samples in PCR assays

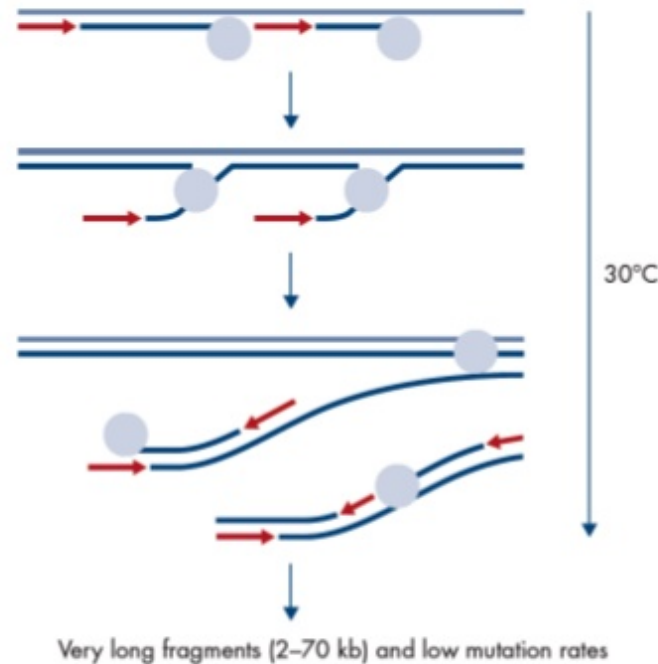


Multiple displacement amplification (MDA) by QIAGEN

MDA technique allows the amplification of an entire genome starting from a small amount of initial DNA.

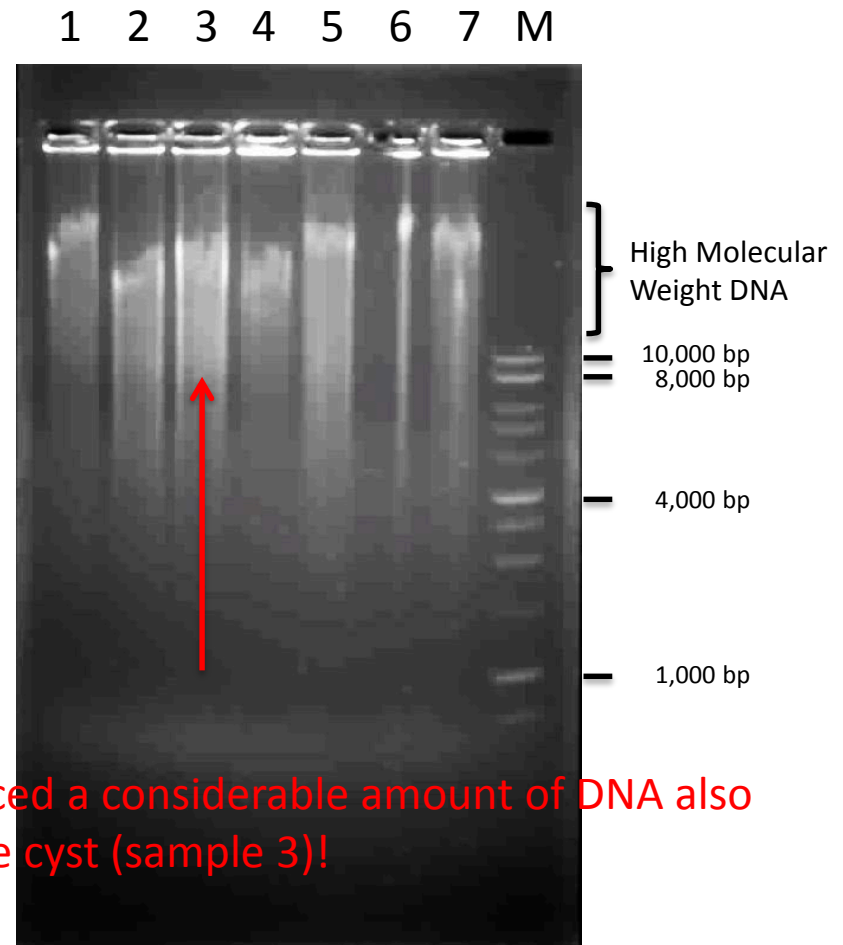
QIAGEN's REPLI-g technology

- Primers (arrows) anneal to the template
- Primers are extended at 30°C as the polymerase moves along the gDNA or cDNA strand displacing the complementary strand while becoming a template itself for replication
- In contrast to PCR amplification, MDA:
 - Does not require different temperatures
 - Ends in very long fragments with low mutation rates

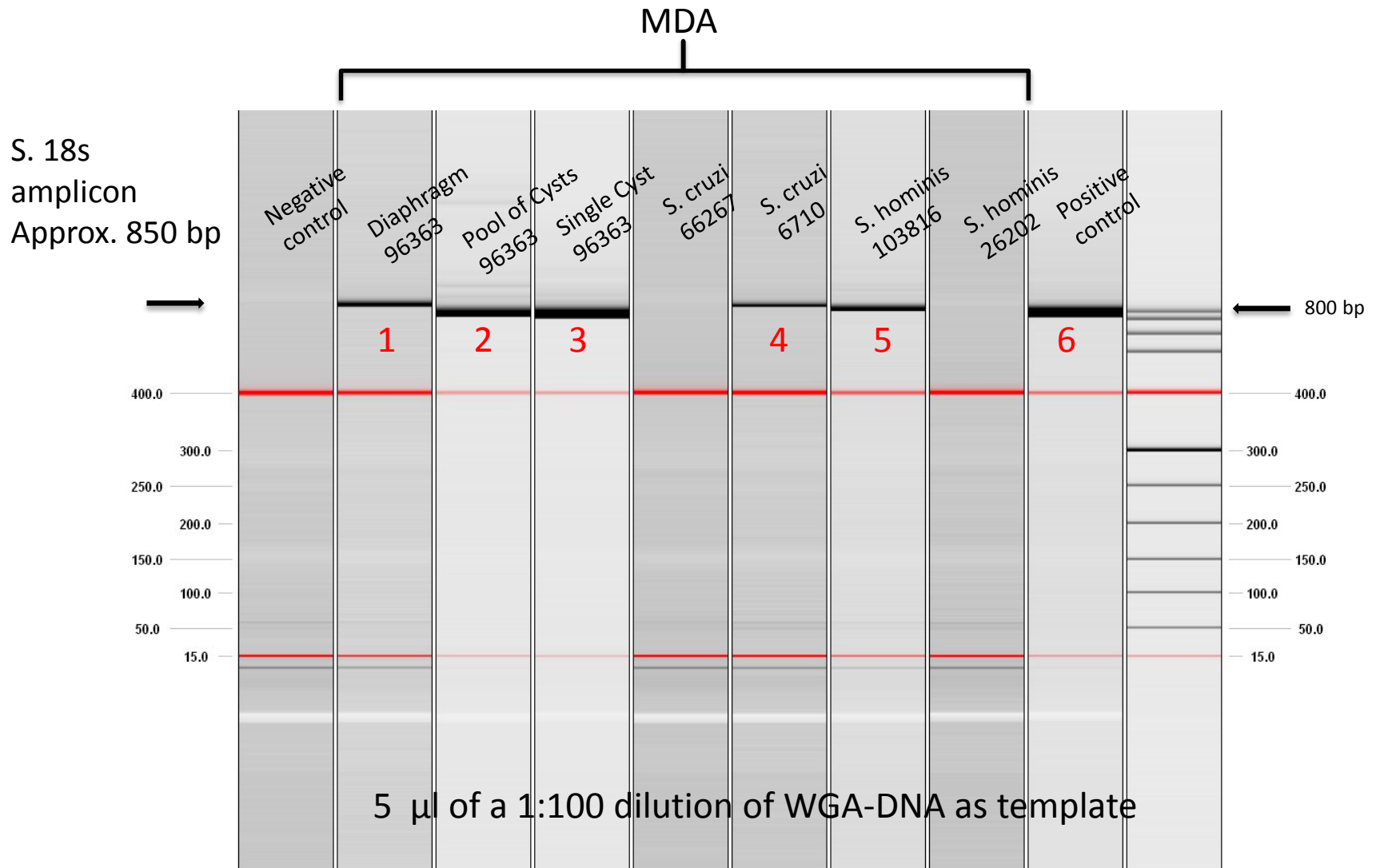


MDA from DNA extracted from isolate cysts and DNA samples of defined species

- 1) DNA extracted from positive diaphragm (muscle tissue) of roe deer (96363)
- 2) Isolate Cysts from diaphragm of roe deer (96363)
- 3) Single Cyst from esophagus of roe deer (96363)
- 4) *S. cruzi* 66267
- 5) *S. cruzi* 6710
- 6) *S. hominis* 103816
- 7) *S. hominis* 26202



PCR amplification of 18S RNA Gene (Moré et al. 2016) and Capillary Electrophoresis of the Amplicons



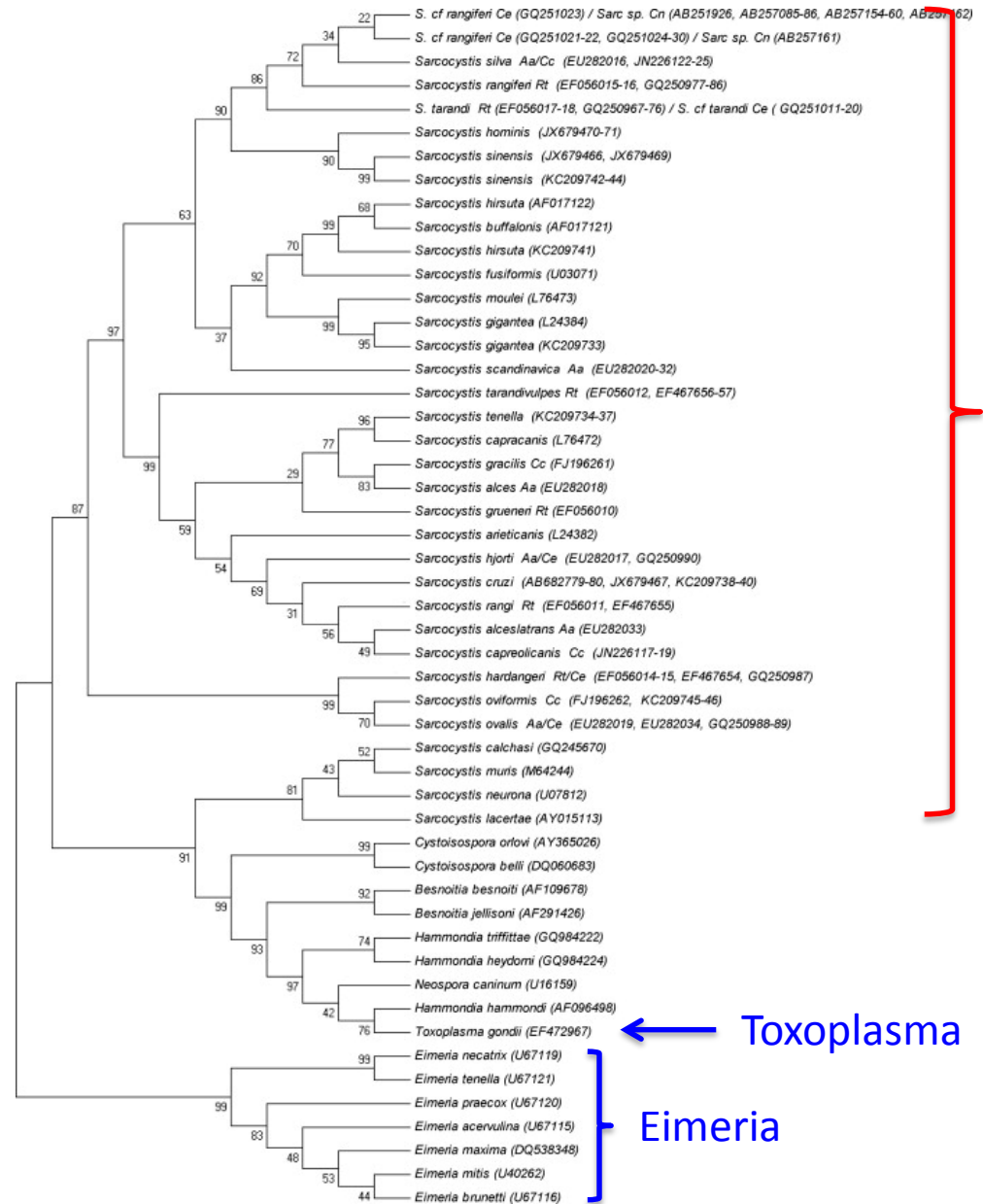
Taxonomy of Sarcocystis spp.

Genetic identification based on the DNA sequences of one or more genetic markers would be the better way to identify the specific species in a given biological sample.

Molecular identification of Sarcocystis species is mainly based on the following genetic loci:

- Ribosomal RNA genes (18s, ITS1)
- Genes of the surface antigen family (SAG 1-4)
- Mitochondrial gene like cytochrome oxidase subunit 1 (cox1)

Recent phylogenetic trees are based on sequences of these loci, however the molecular phylogenesis of Sarcocystis is more complicated than other related parasites like Toxoplasma or Eimeria.



Phylogenetic tree of the Sarcocystidae and related coccidia based on ssrRNA genes. Gjerde et al. 2013

Design of primers to amplify the cytochrome c oxidase subunit I (Cox1)



Phylogenetic relationships among *Sarcocystis* species in cervids, cattle and sheep inferred from the mitochondrial cytochrome c oxidase subunit I gene [☆]

Bjørn Gjerde*

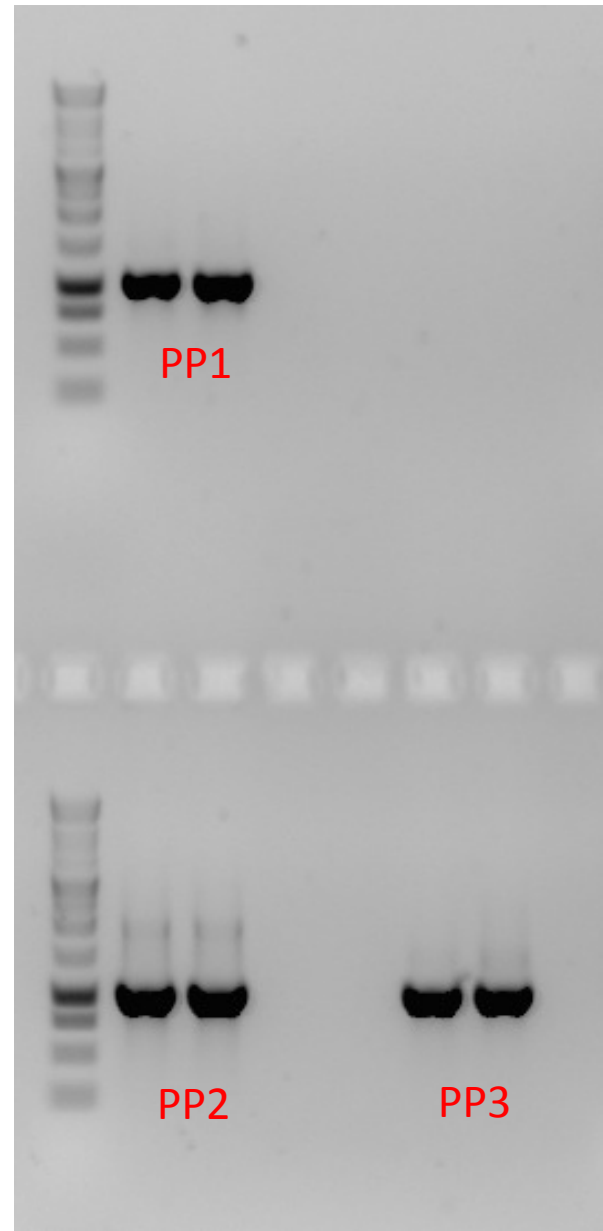
Department of Food Safety and Infection Biology, Norwegian School of Veterinary Science, P.O. Box 8146 Dep., 0033 Oslo, Norway

Cox1 gene has been proposed as the most suitable marker for the identification of the strictly related species of *Sarcocystis*

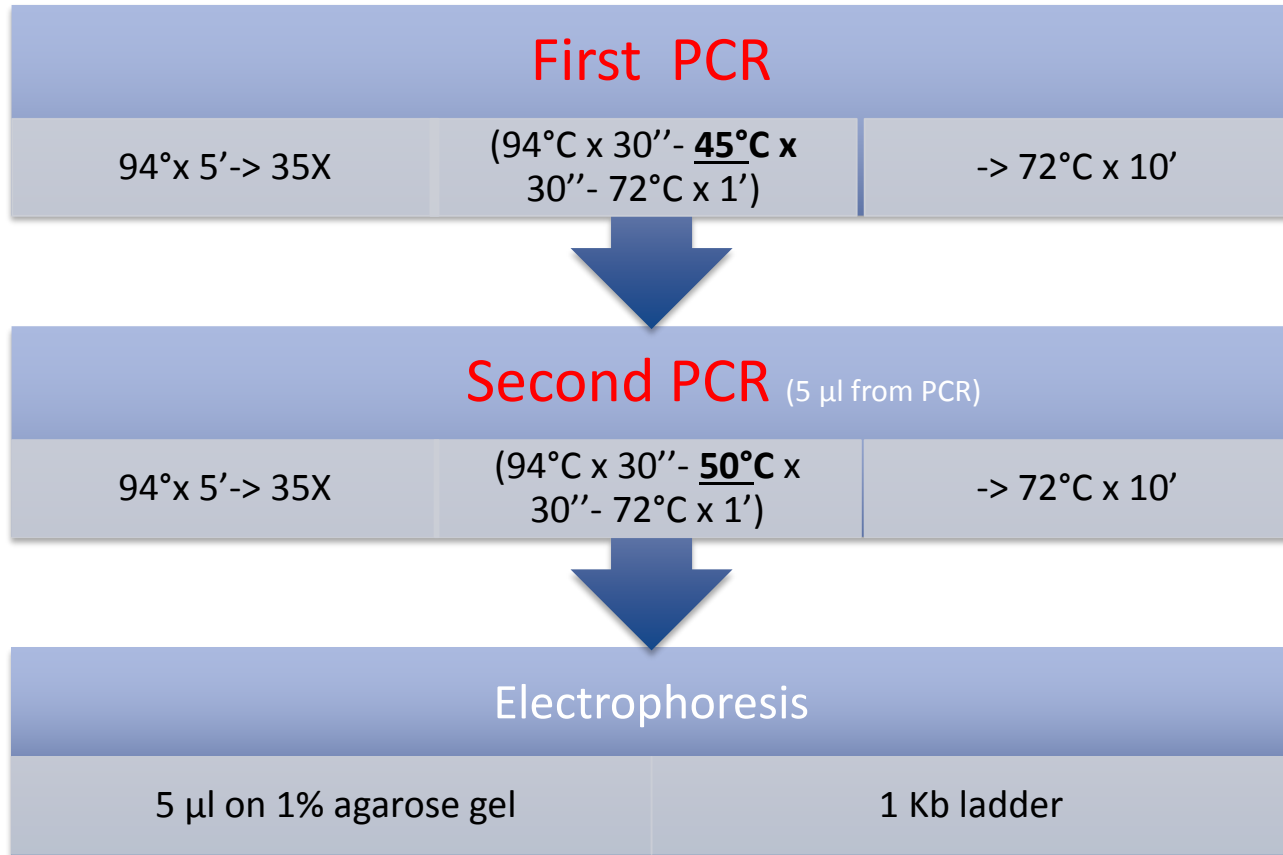
- The actual reconstructions of this gene are obtained assembling different sequences
- To overcome the problem of multiple PCR and sequencing, we tried to find one or more couple of primers able to amplify different species, particularly *S. hominis* and *S. suishominis*
- Design of primers was based on the of *S. cruzi* Cox1 and tested “in silico” to maximise the annealing with other *Sarcocystis* species.

Setting a PCR assay for Sarcocystis Cox1

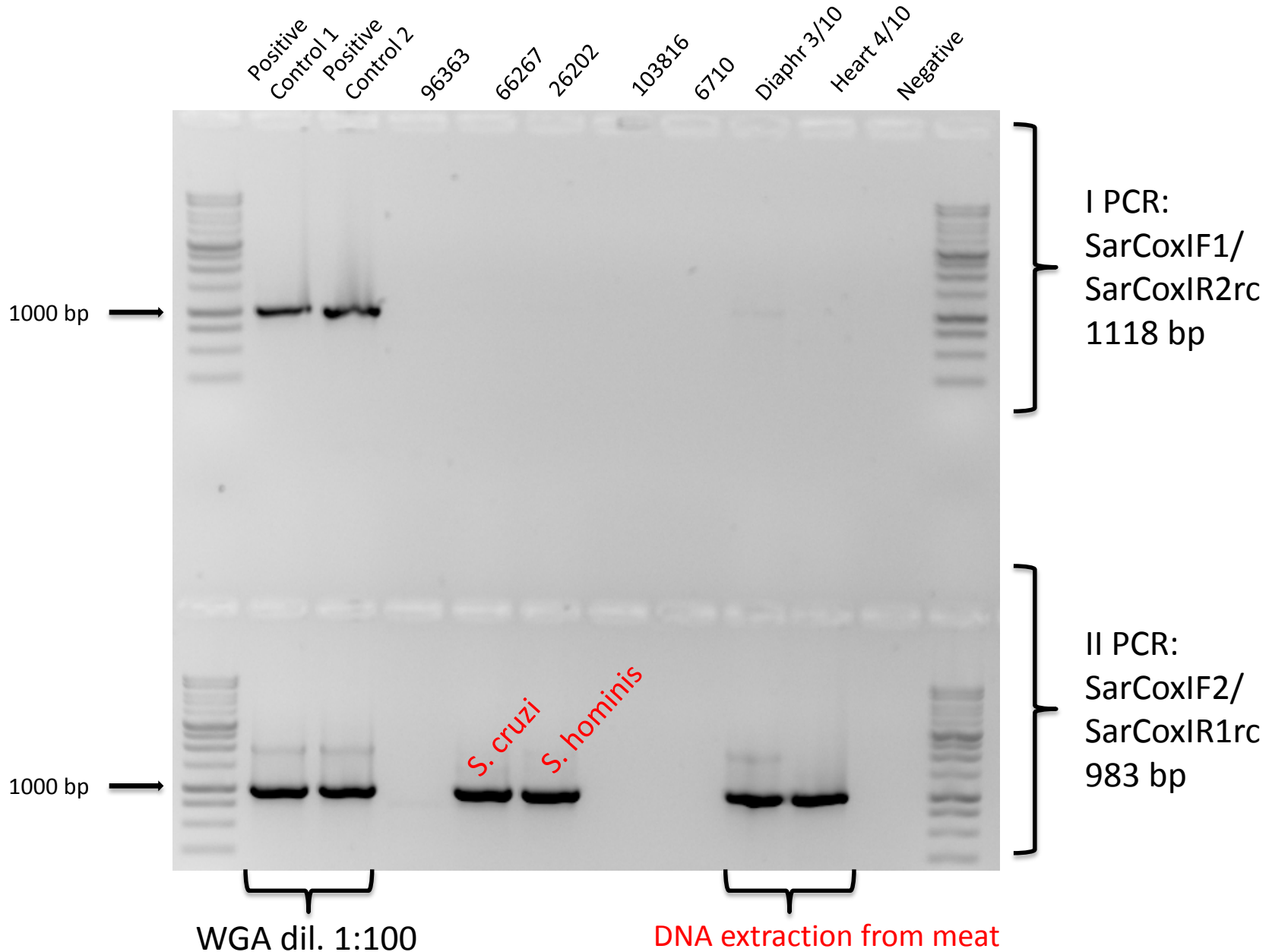
- Identification of primer pairs by Primer-BLAST
<https://www.ncbi.nlm.nih.gov/tools/primer-blast/index.cgi>
- Four primers were selected on the basis of similarity with other Cox1 sequences.
- They were tested in four possible combinations and with different Annealing temperatures
- Three primer pairs (PP) gave optimal results:
PP1 = SarCoxIF1 / SarCoxIR1
PP2 = SarCoxIF1 / SarCoxIR2
PP3 = SarCoxIF2 / SarCoxIR1
- PP1 and PP3 are suitable for nested PCR



Nested PCR For Sarcocystis Cox1 gene

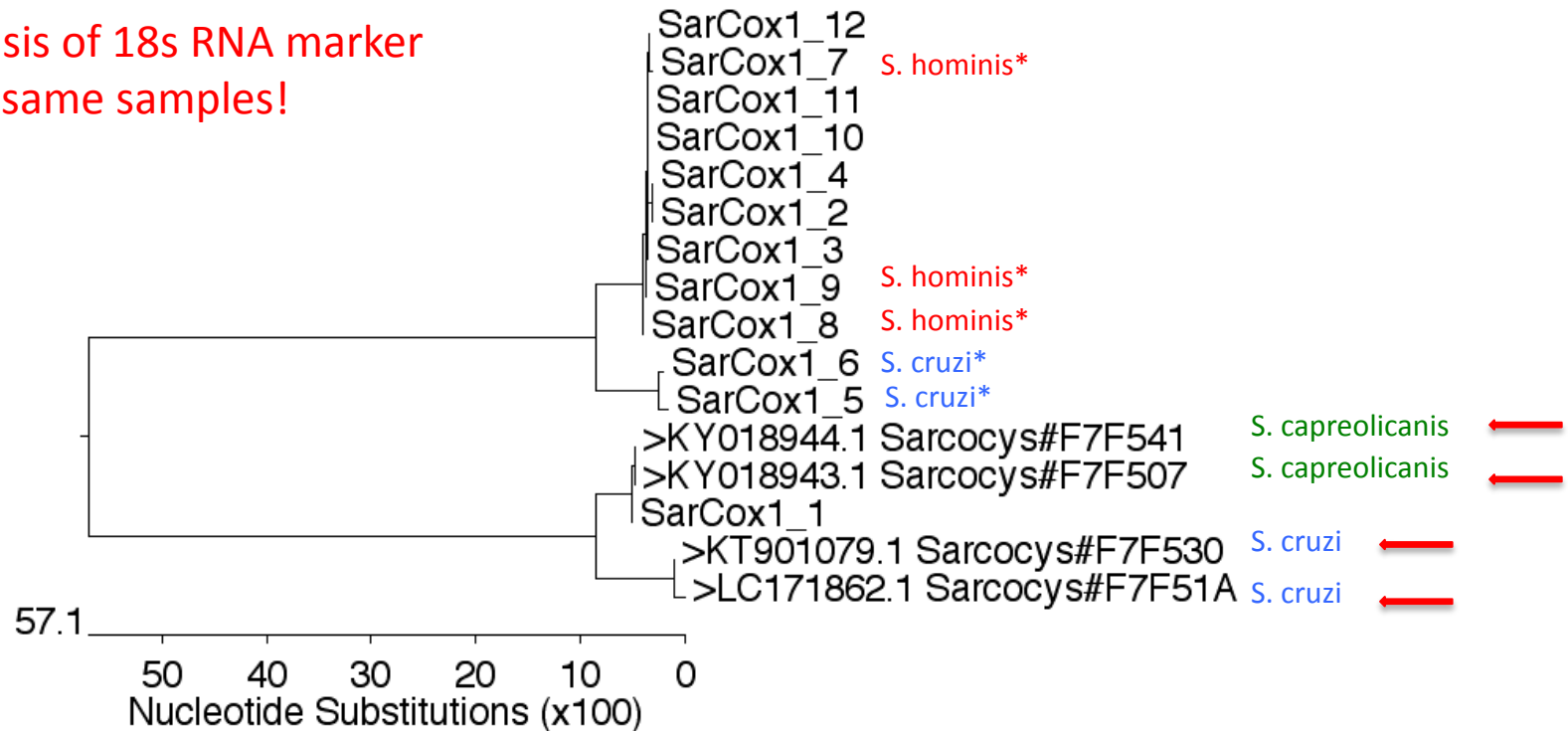


Nested PCR Sarcocystis Cox1 as genetic marker



Phylogenetic Tree based on Cox1 genetic marker

S. gracilis was identified on the basis of 18s RNA marker in the same samples!



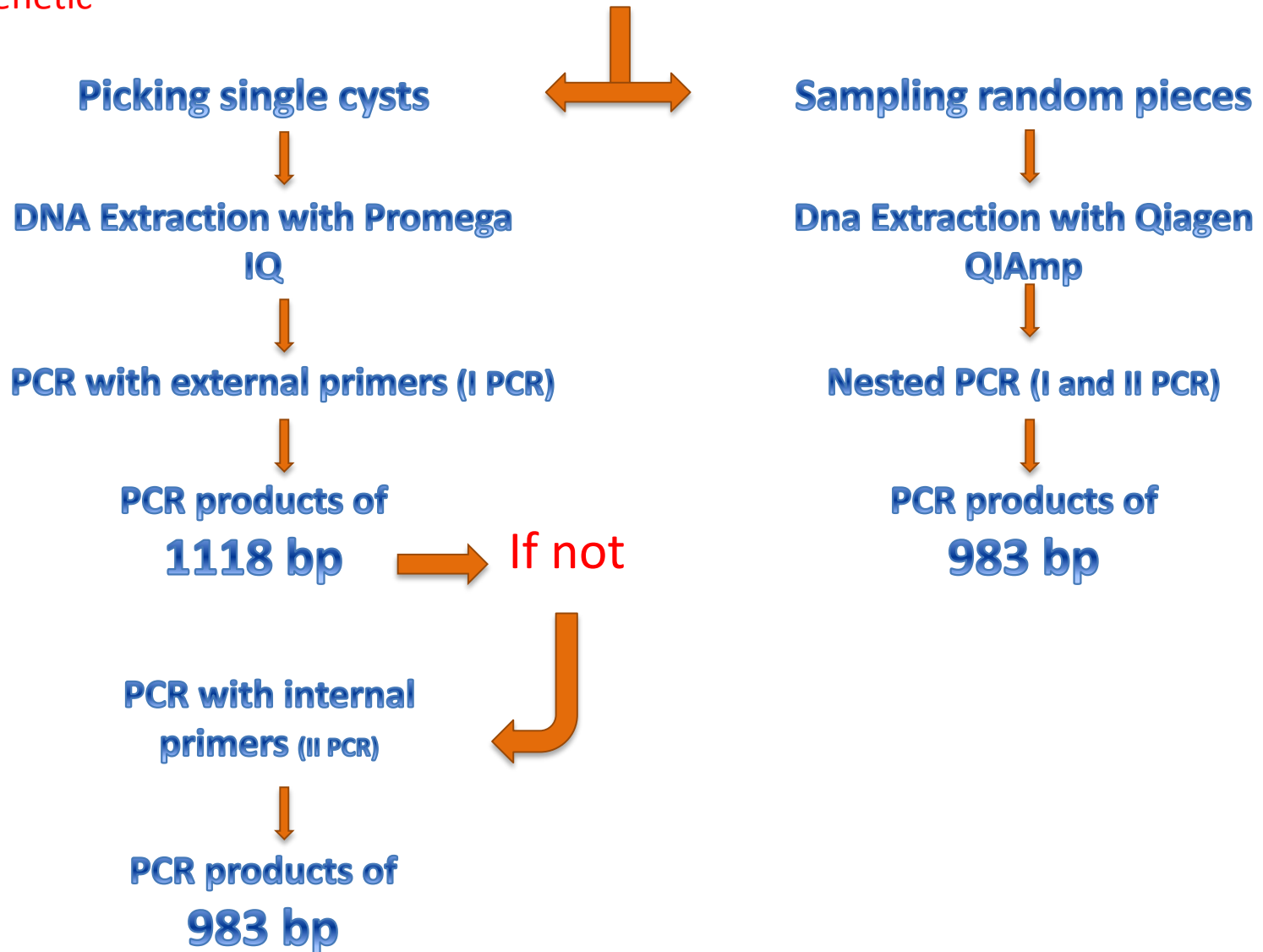
**identification by IZSTO of Turin

Conclusions

- A procedure for DNA extraction from isolated sarcocysts
- A procedure for DNA extraction from meat with sarcocysts
- Application of a PCR assay for 18s RNA genes
- Development of a new Nested-PCR assay targeting Cytochrome-oxidase subunit 1(Cox1)

Flowchart for the identification of Sarcocystis species by Cox1 genetic marker

Meat with Sarcocysts



Aknowledgements

- Irene Bertoletti and Alessandro Bianchi
IZSLER “Bruno Ubertini” Sondrio – Italy
- Silvia Trisorio e Simone Peletto
IZSTO “SS. Genetica e Immunobiochimica”
Torino – Italy
- Edoardo Pozio ISS Roma - Italy

Muscular sarcocystis in humans (Reports from 1981)

- All the recent cases of muscular sarcocystis come from tropical or subtropical countries
- Prevalently from the south-east Asia regions and particularly from Malaysia

TABLE 2 Summary of reports of muscular sarcocystosis in humans not included in the review by Beaver et al. and presented in chronological order of publication

Country or region	Description of human subject(s) or specimen(s)	Sarcocyst location	Biopsy or autopsy ^c	Reference, yr of publication
Malaysia	58-yr-old M ^a	Tongue	B	9, 1981
India	2 persons	Leg Gluteus	B B	10, 1983
Denmark?	4 of 112 specimens ^b	Muscle	A?	11, 1985
Malaysia	45-yr-old M ^a 53-yr-old M ^a	Nasopharynx Nasopharynx	B B	12, 1987
Malaysia	1 person	Skeletal muscle	B	13, 1988
Australia via Thailand	31-yr-old M	Skeletal muscle	B	14, 1990
Egypt	1 person	Skeletal muscle	B	15, 1990
Malaysia	45-yr-old M ^a 67-yr-old F ^a 32-yr-old F ^a	Tongue Tongue Pectoral muscle	B B B	16, 1992
Malaysia	21 persons aged 16 to 57 yr	Tongue	A	17, 1992
Belgium via Brazil, Kenya, and Tanzania	31-yr-old M	Thigh	B	18, 1995
India	40-yr-old M 14-yr-old F 52-yr-old F 23-yr-old F	Thigh Arm Thigh Arm	B B B B	19, 1996
Malaysia	44-yr-old F ^a 19-yr-old F	Thigh Calf	B B	20, 1998
Malaysia	7 adults, M	1 muscle	B	21, 1999
Thailand	66-yr-old M ^a	Larynx	B	22, 2011
India	20-yr-old M	Arm	B	23, 2012
India	50-yr-old M	Neck	B	
India	47-yr-old M	Leg	B	24, 2013
Pangkor Island, Malaysia	89 people	Leg	B	25, 2013 26, 2014 27, 2014
Tioman Island, Malaysia	39 F and 29 M subjects, aged 4 to 72 yr	Skeletal muscle from 15 patients; 1 PCR positive	B	28, 2014 29, 2014
Tioman Island, Malaysia	3 F and 3 M subjects	Febrile myositis syndrome	NTE	30, 2014
Tioman Island, Malaysia	26 cases in Germany reported previously	Symptomatic changes; skeletal muscle	B	31, 2014

^a Cancer patients.

^b Tissues examined by trichinoscopy.

^c NTE, no tissue examined.

Malaysian monkeys are intermediate hosts for *S. nesbitti*

