

8 | Parasitology | Full-Length Text

The fourth-stage autoinfective larva of *Strongyloides stercoralis*: redescription and diagnostic implications

Huan Zhao,¹ Anson V. Koehler,² Cameron Truarn,³ Damien Bradford,³ David W. New,³ Rick Speare,¹ Robin B. Gasser,² Harsha Sheorey,⁴ Richard S. Bradbury¹

AUTHOR AFFILIATIONS See affiliation list on p. 9.

ABSTRACT Human strongyloidiasis is often underdiagnosed or misdiagnosed, which can relate to a lack of knowledge or recognition of the importance of particular developmental/larval stages of *Strongyloides stercoralis* in making an accurate diagnosis using parasitological methods (a morphological approach or morphological features/characters). Here, we report the identification of *S. stercoralis* autoinfective fourth-stage larvae (L4a) in naturally infected humans, encountered in two clinical cases in Australia. These larvae were identified in sputum (Case 1) and bronchoalveolar lavage (Case 2) specimens by direct wet-mount microscopy. The L4a of *S. stercoralis* can be morphologically differentiated from autoinfective third-stage larvae by its conical and pointed tail and a relatively mature genital primordium with an enlarged genital rudiment and the formation of a vulva within cuticle layers. This study emphasizes the need to consider these morphological features of the L4a stage for an accurate diagnosis of *S. stercoralis* infection. A detailed morphological description of this stage is given to guide laboratory practitioners and researchers in the identification and differentiation of this unique but neglected life-cycle stage of *S. stercoralis*.

KEYWORDS Strongyloides stercoralis, Strongyloidiasis, diagnosis, autoinfection, life cycle stages, larva

A ll nematodes have four larval stages (1). Strongyloides stercoralis (Bavay 1876) is a medically important parasitic nematode with a unique life cycle (2). This nematode has a remarkable ability to perpetuate within its host for decades and, when triggered by immunosuppression, can cause life-threatening systemic disease (3). The clinical diagnosis of *S. stercoralis* infection is challenging, and the microscopic detection and morphological identification of developmental stages of this parasite in feces or other biological specimens (e.g., sputum) remain a major diagnostic modality. Identification and differentiation of *S. stercoralis* from morphologically similar nematodes require the expertise of experienced morphologists. Recognizing and identifying morphological nuances in various stages of *S. stercoralis* are essential for an accurate and timely diagnosis of this neglected parasitic infection.

The life cycle of *S. stercoralis* is unusually complex. Parasitic adults are parthenogenetic females that reside in the intestinal mucosa. The eggs that they produce hatch within the crypts of Lieberkühn into rhabditiform larvae (L1r), which then migrate to the intestinal lumen and are passed in the feces (3, 4). The L1r molt into L2r and then, depending on environmental conditions, these L2r either develop into infective filariform larvae (L3i; homogonic cycle) or undergo four molts to become sexually reproducing free-living adult females and males, whose progeny are all L3i, which must infect a new host or die (heterogonic cycle) (3, 4). Environmental L3i from homogonic and heterogonic routes invade a host percutaneously to complete their life cycle (2–4). **Editor** Bobbi S. Pritt, Mayo Clinic Minnesota, Rochester, Minnesota, USA

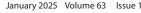
Address correspondence to Richard S. Bradbury, richard.bradbury@jcu.edu.au.

Rick Speare is deceased but was involved in the taxonomic investigations of the first case. Permission to include his name as a co-author was granted by his widow.

The authors declare no conflict of interest.

Received 8 July 2024 Accepted 21 October 2024 Published 5 December 2024

Copyright © 2024 Zhao et al. This is an open-access article distributed under the terms of the Creative Commons Attribution 4.0 International license.





Strongyloides stercoralis also undergoes an autoinfective cycle, wherein some L1r within the intestine transform into autoinfective filariform larvae (L3a). These larvae penetrate the host's lower gut or perianal skin to re-establish infection (3–5). Autoinfection is believed to be the mechanism responsible for the chronicity of infection in hosts and for extensive multiplication during hyperinfection (3–5). The *in vivo* migratory routes of L3i and L3a involve both the cardio-pulmonary-esophageal pathway and random navigation to reach the small intestine, where they develop into a new generation of parasitic females (3–5).

Fourth-stage filariform larvae (L4) are known to occur in many human-parasitic skin-penetrating nematodes, including the hookworms (6), and have been observed in other *Strongyloides* species, such as *Strongyloides* ratti (7), *Strongyloides* venezuelensis (8), and *Strongyloides* felis (9). Remarkably, while four larval stages are recognized in the free-living life cycle of *S. stercoralis*, a fourth stage is not usually recognized in descriptions of the external infective and the autoinfective life cycles of this important human and animal parasite (4). Early experimental studies using canine (10) and non-human primate (11) models support the existence of at least one pre-adult, or "juvenile," parasitic stage of *S. stercoralis*. These juvenile females were recognized as morphological intermediates between filariform L3 and parasitic adults and have been detected in the respiratory tract (10, 11). This information indicated that larval maturation can occur during pulmonary migration, supported by the recovery of adult worms from this site in experimental dogs (11–14) and in naturally infected humans (15, 16). However, until now, this pre-adult form has only been demonstrated experimentally, and its description was made based on an incorrect assumption at the time that parasitic males existed (10).

Diagnosis of "uncomplicated" strongyloidiasis typically relies on the identification of *S. stercoralis* L1r in the stool (17). Less commonly, filariform larvae may be found in respiratory specimens, such as bronchoalveolar lavage (BAL) fluid and sputum (17, 18). In clinical cases of hyperinfection, pulmonary involvement is more pronounced, and autoinfective filariform larvae may be abundant in these specimens (17). Thus, the detection and identification of autoinfective *S. stercoralis* larvae from clinical specimens are crucial for the early recognition of complicated strongyloidiasis.

Thus far, two filariform stages of *S. stercoralis*, i.e., L3i and L3a, have been morphologically characterized. They are distinguished from other major life stages of *S. stercoralis* by several body dimensions, most distinctly the esophagus and the tail (4) (Table 1). Morphological differences between L3i and L3a are subtle, yet discernible, with the former appearing longer and more slender (4) (Table 1).

This study presents an unusual autoinfective stage of *S. stercoralis* identified in respiratory tract specimens of two patients in Australia. Our findings indicate the need to expand diagnostic parameters to include this clinically important yet neglected filariform stage of *S. stercoralis*.

MATERIALS AND METHODS

Cases

This is a retrospective study of diagnostic data for two clinical cases of disseminated strongyloidiasis occurring in Australia within the past 10 years. The patients were from South East Asia and East Africa, respectively, and had resided in Australia for a number of decades. They reported no recent travel history to endemic countries prior to admission.

As part of the routine investigation for respiratory infections, a sputum specimen was collected from Patient 1, and bronchoscopy was performed to obtain BAL fluid from Patient 2. Upon collection, respiratory specimens were sent to a hospital microbiology laboratory within 4 hours for microscopic analysis.

Laboratory processing and microscopy

Specimens were processed immediately (within 1 hour) upon arrival. The BAL specimen was centrifuged at $500 \times g$ for 20 minutes, and the sputum specimen was centrifuged

Life stage	Mean length	Mean width ^a	Width ^a /	Esophagus/	Esophagus	Reproductive system	Tail
	(range) (µm)	(range) (µm)	length (%)	length (%)			
Parasitic female	2,420 (2,100–	37 (30–40)	1.5	23.8	Elongated	Straight reflected ovary; uteri short, usually	Narrowly tapered
	2,700)				cylindrical	containing no more than six eggs; vulva	to a cone-shaped
					(filariform)	about two-thirds body length from anterior	tail
					esophagus	end, with pair of prominent muscles	
						surrounding transverse opening; seminal receptacles absent	
Free-living female	1,140 (920–1,700)	62 (52–85)	5.4	12.6	Rhabditoid	Didelphic with opposed equal uteri and	Gradually tapered
					esophagus	reflected ovaries; vulva near the middle of the	e to a finely pointed
						body; seminal receptacles present	tail
Free-living male	900 (810–1,000)	43 (40–50)	4.8	13.1	Rhabditoid	Straight tubule structure; two small	Gradually tapered
					esophagus	sickle-shaped spicules and a single	to a finely pointed
						gubernaculum	tail; often curved ventrally.
Rhabditiform (L1r)	210 (180–240)	14.5 (14–15)	6.9	30	Rhabditoid	Lateral rhomboid genital primordium halfway	Gradually tapered
larvae					esophagus	down the larval body	to a finely pointed
							tail
Infective filariform	563 (490–630)	15.8 (15–16)	2.8	43	Elongated	A small genital rudiment visible at about the	Truncated notched
(L3i) larvae					cylindrical	midpoint of the intestine	tail
					(filariform)		
					esophagus		
Autoinfective	269 (234–317)	11 (10–13)	4.1	42–52	Elongated	A small genital rudiment visible at about the	Truncated notched
third-stage					cylindrical	midpoint of the intestine	tail
filariform (L3a)					(filariform)		
larvae					esophagus		
Autoinfective	nd ^c	nd	3.7–5.2	37–46	Elongated	Elongated genital rudiment midway of the	Narrowly tapered
fourth-stage					cylindrical	intestine; vulva formed as a transverse slit	to a point or to a
filariform (L4a)					(filariform)	at the midpoint of the intestine but has an	cone-shaped tail
larvae ^b					esophagus	overlying layer of cuticle	

TABLE 1 Morphological characteristics of the major life stages of Strongyloides stercoralis (8, 9, 18–21)

^a At the widest point.

^bData derived from the present study.

^cnd, no data.

at $500 \times g$ for 10 minutes. Supernatants were discarded, and two wet mount slides were prepared from each specimen and examined by microscopy at 100- and 400-times magnification. Measurements were not taken in this routine pathology setting, but the identified larvae were photographed, and the percentage ratios of body length, maximum width, and esophagus length were determined by examination of those photographs.

Molecular and phylogenetic analyses

The remaining sputum (n = 1) and BAL (n = 1) specimens underwent DNA extraction by first digesting with proteinase K (10 µg/µL) (Promega, USA) and lysis buffer for 2 hours at 56°C and then using the Promega DNA Clean Up Kit, following the manufacturer's protocol. PCR of the mitochondrial cytochrome c oxidase subunit I (cox1) gene was performed as previously described (22, 23), and the amplicons were subjected to Sanger sequencing. Sequence data were BLASTN searched against the GenBank database containing all available nematode sequences. These sequences were placed in a maximum likelihood tree generated in MEGA11 (24) along with other sequences of *Strongyloides* and *Necator americanus* as an outgroup.

RESULTS

Microscopic findings

Multiple larvae were recovered from the respiratory specimens (Fig. 1 and 2). These larvae closely resembled the L3a stage of *S. stercoralis* in several aspects. Specifically, they appeared slender, with a maximum width-to-length ratio of 3.7%–5.2% (Table 2). The buccal cavity was shallow, with a small pore-like mouth structure. The esophagus was cylindrical, filariform, extending 37%–46% of the body length (Table 2). The anus was evident sub-terminally, with small lip-like swellings along the posterior edge of the transverse opening.

However, the reproductive systems of these larvae appeared to be more mature than those of the L3a stage, but not as developed as the parasitic female. Specifically, an enlarged genital rudiment was visible around the midpoint of the intestine. In one larva, the vulva formed as a transverse slit but with an overlying layer of the cuticle (Fig. 2A). This larva had a notably lower width-to-length ratio (3.7%), and its esophagus was comparatively shorter relative to the body length (37%) than L3a. Ovary or uteri structures were inconspicuous. Moreover, the tails of these larvae were not notched but narrowly tapered to a cone shape or a point, resembling those observed in the parasitic adult stage (Fig. 1 and 2).

These larvae appear to represent a transitional form between the filariform L3 stage and the parasitic adult stage of *S. stercoralis* (Fig. 3). Specifically, these recovered larvae are morphologically consistent with the *S. stercoralis* autoinfective fourth-stage filariform larvae described by Faust in 1933 (10).

Several L3a were identified from the BAL specimen (Fig. S1). No other parasite eggs, larvae, cysts, or trophozoites were identified in either of the two specimens by microscopy.

Molecular findings

Sequencing confirmed that the larvae belonged to *S. stercoralis* (GenBank accession numbers PP946346 and PP946347), both having a 99.62% (964 bp) and 99.68% (393 bp) identity, respectively, to known *S. stercoralis* sequences (GenBank accession numbers MN509458 and ON954823). A maximum likelihood phylogenetic tree placed these sequences within the *S. stercoralis* clade (Fig. 4).

DISCUSSION

The L4a stage has until now been neglected in the scientific and clinical literature, and its detection in these two clinical cases caused confusion even among the experienced morphologists/parasitologists authoring this paper when it was first encountered. The presence of thin and tapered nematode larvae in sputum and BAL fluid

TABLE 2	Morphological	features of Strona	/loides stercoralis	autoinfective for	ourth-stage larvae

Case	Width ^a / length (%) ^b	Esophagus/ length (%) ^b	Esophagus	Excretory system	Reproductive system	Cuticle	Tail
Case 1	4.9–5.2	46	Filariform esophagus	Anus subterminal, with small lip-like swelling along the posterior edge of transverse opening	Elongated genital rudiment midway of the intestine	Finely striated cuticle	Narrowly tapered to a point
Case 2	3.7–4.6	37–46	Filariform esophagus	Anus subterminal, with small lip-like swelling along the posterior edge of transverse opening	Elongated genital rudiment midway of the intestine; vulv formed as a transverse slit at the midpoint of the intestine but has an overlying layer of cuticle	2	Narrowly tapered to a cone-shaped tail

^a At the widest point.

^bAverage of five measurements.

Full-Length Text

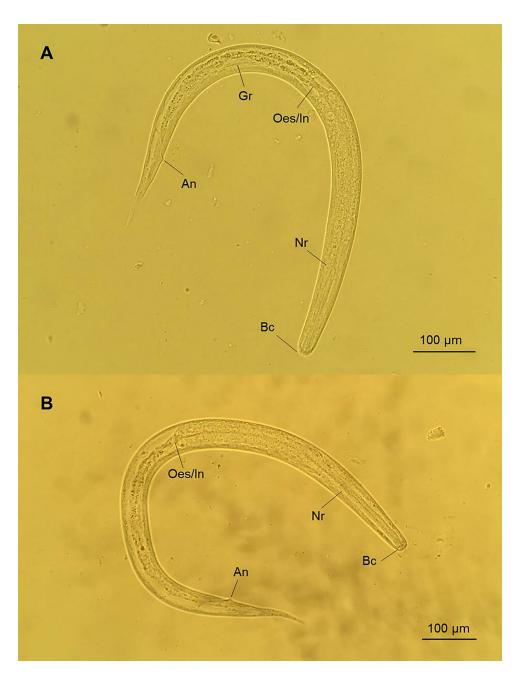


FIG 1 *Strongyloides stercoralis* autoinfective fourth-stage larvae (A and B) recovered from the sputum of Patient 1. An, anus; Bc, buccal cavity; Gr, genital rudiment; Nr, nerve ring; Oes/In, esophageal-intestinal junction.

indicated *S. stercoralis*, but the unfamiliar morphology meant that this diagnosis could not be confirmed in the first instance. DNA sequencing of well-known genetic markers confirmed the species identity, and further investigation of the literature identified these as the neglected juvenile or pre-adult stage of *S. stercoralis*. Here, we identify these as juvenile or pre-adult forms as the missing fourth-stage autoinfective larva (L4a) of *S. stercoralis*.

Pre-adult parasitic females of *S. stercoralis* have been only documented twice in the literature (10, 11). Faust (10) in 1933 described two premature parasitic stages of *S. stercoralis*, referred to as "the preadolescent female" and "the adolescent female," recovered from the lung of experimentally infected dogs. Marked developments in these two stages included the enlargement of genital primordia and the formation of a vulva

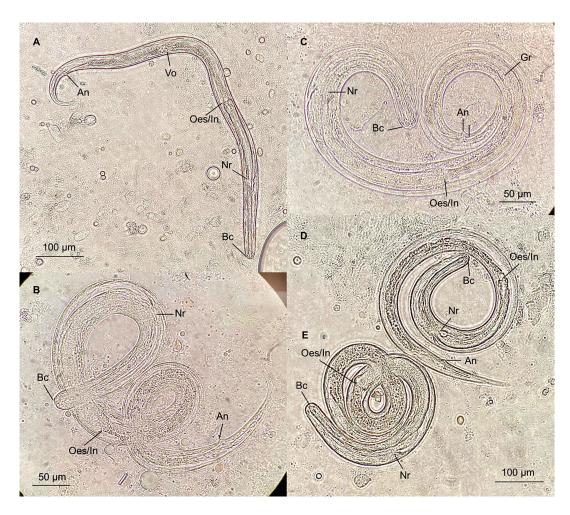


FIG 2 Strongyloides stercoralis autoinfective fourth-stage larvae (A through E) recovered from the BAL fluid of Patient 2. An, anus; Bc, buccal cavity; Gr, genital rudiment; Nr, nerve ring; Oes/In, esophageal-intestinal junction; Vo, vaginal opening.

opening in some cases. These findings accord with our observations of *S. stercoralis* L4a in the present case series. However, Faust (10) noted that the notch of the tail persisted in juvenile females, contrasting our finding of the conical or pointed tail morphology. Additionally, despite an increase in body length, the esophagus of pre-adult females was shorter compared with the L3i stage, while this difference was inconspicuous in our study. Faust's work was conducted in an era when parasitic *S. stercoralis* males were believed to exist. It is possible that his description reflected a combination of parasitic and free-living stages. The only other study reporting *S. stercoralis* parasitic pre-adults was that of Mati et al. in 2014 (11), who used the marmoset experimental model. Faust's criteria (10) were followed for larval identification. It was noted that the esophagus-to-body length ratio of the identified worms was intermediate between the ratios seen in L3i and adult females, consistent with our findings for some L4a. No further morphological characterization was made in the study by Mati et al. (11).

L4 filariform larvae have been previously reported in the life cycle of *S. felis*, a larviparous *Strongyloides* species infecting cats. According to Speare (9), this larval stage was characterized by a non-notched tail morphology and conspicuous reproductive maturation, with the vulva forming as a transverse slit within layers of cuticle. These findings closely mirror our findings for *S. stercoralis* L4a. Phylogenetically, *S. felis* is closely related to *S. stercoralis* (25), and, therefore, it is very plausible that the two species share morphological similarities at different life stages.

Journal of Clinical Microbiology

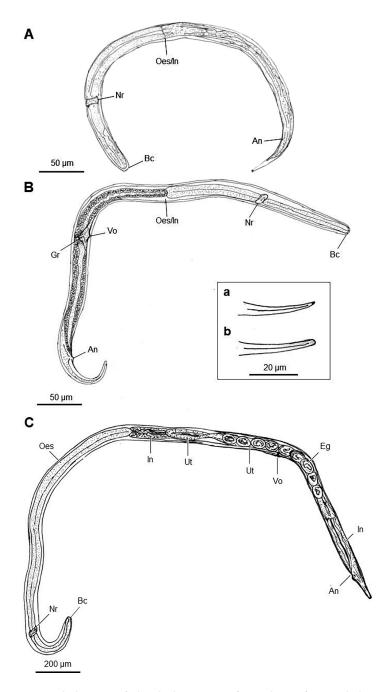
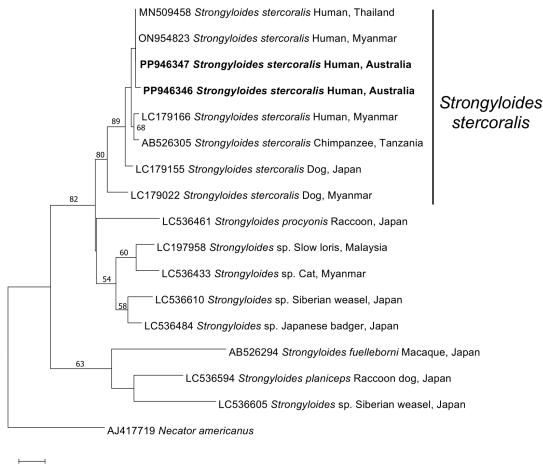


FIG 3 Anatomical drawings of the third-stage autoinfective larva of *Strongyloides stercoralis* (A), fourth-stage autoinfective larva (B), and parasitic adult (C) (4). The tail of L4a is either pointed (a) or cone-shaped (b). An, anus; Bc, buccal cavity; Eg, eggs; Gr, genital rudiment; In, intestine; Nr, nerve ring; Oes, esophagus; Oes/In, esophageal-intestinal junction; Ut, uterus; Vo, vaginal opening.

One limitation of this work is the lack of morphometric data for the *S. stercoralis* larvae identified. Consequently, we were unable to experimentally assess the developmental progression of parasitic stages based on size changes. Despite this issue, our morphological characterization of the L4a stage has substantial diagnostic value. Established criteria for identifying autoinfective larvae rely heavily on the tail morphology. While the characteristic notched tail of the L3a stage differentiates it from other nematodes infecting humans, it should not be the only consideration for the diagnosis of hyperinfective strongyloidiasis. The present study indicates the need for including such diagnostic



0.050

FIG 4 Maximum likelihood tree based on cox1 sequences generated in this study (in bold) and those published in the literature.

criteria for this neglected L4a stage of *S. stercoralis*. Future work is needed for more detailed characterization of this larval stage.

This study was based on the observation of two clinical cases. Given this limited data set, these descriptions of the L4a stage of *S. stercoralis* should be considered preliminary and would benefit from further evidence in broader clinical contexts. The clinical data available for these cases were sparse. While larvae were detected during the initial diagnosis of strongyloidiasis, the impact of anthelmintic therapies on the occurrence of this stage is unclear. Future research is needed to fully understand the implications of the L4a stage in respiratory specimens on the severity, progression, and treatment outcomes of strongyloidiasis.

It is important for the clinical parasitology community to recognize and identify rare diagnostic stages of parasites. In these cases, identification of the infecting agents as *S. stercoralis* was delayed due to the very unusual morphology observed. Early involvement of infectious disease specialists is essential in any suspected case of *Strongyloides* hyperinfection or systemic disease. Given the risk of disseminated strongyloidiasis, particularly in immunocompromised patients, empiric treatment may be recommended before diagnostic confirmation when clinical suspicion is high. This approach can be crucial in preventing severe complications and should be guided by the overall clinical presentation, risk factors, and symptom severity.

The complexities in diagnosing *S. stercoralis* infection highlight the ongoing need to maintain laboratory and morphological skills, especially in light of the global progressive loss of expertise in morphology-based parasitic diagnostics (26). Training

programs should emphasize the differentiation of all *S. stercoralis* life stages from other human-infecting nematodes, such as hookworms. To address diagnostic challenges, the integration of targeted molecular testing, such as *Strongyloides* PCR and sequencing (27), should complement traditional methods in clinical diagnostic practice. Furthermore, proper specimen processing, such as centrifuging BAL specimens to concentrate any parasitic or other diagnostic elements present, is essential to prevent false-negative results.

Conclusion

We report and redescribe the "forgotten" juvenile form of the parasitic female of *S. stercoralis*, observed in two clinical cases in Australia, and identify this form as the fourth-stage autoinfective larva. This developmental stage is a morphological intermediate between the L3a and the parasitic adult stage of *S. stercoralis*. It is important that clinical and veterinary microbiologists, as well as parasitologists, be aware of the morphological features of this stage in order to avoid diagnostic confusion and delayed diagnosis and treatment when this stage is encountered in extra-intestinal specimens from human or animal patients suffering from *Strongyloides* hyperinfection or systemic strongyloidiasis.

ACKNOWLEDGMENTS

R.B.G.'s research work is supported by the Australian Research Council. H.Z. receives an Australian Government Research Training Program Scholarship from James Cook University.

AUTHOR AFFILIATIONS

 ¹School of Public Health and Tropical Medicine, College of Public Health, Medical and Veterinary Sciences, James Cook University, Townsville, Queensland, Australia
²Department of Veterinary Biosciences, Melbourne Veterinary School, Faculty of Science, The University of Melbourne, Parkville, Victoria, Australia
³PathWest Laboratory Medicine, Nedlands, Western Australia, Australia
⁴Microbiology Department, St. Vincents Hospital, Fitzroy, Victoria, Australia

AUTHOR ORCIDs

Huan Zhao (b) http://orcid.org/0000-0002-9753-939X Anson V. Koehler (b) http://orcid.org/0000-0001-8330-6416 Harsha Sheorey (b) http://orcid.org/0000-0003-4891-2925 Richard S. Bradbury (b) http://orcid.org/0000-0002-5524-506X

AUTHOR CONTRIBUTIONS

Huan Zhao, Formal analysis, Investigation, Methodology, Writing – original draft, Visualization | Anson V. Koehler, Formal analysis, Visualization, Writing – review and editing, Investigation, Methodology | Cameron Truarn, Writing – review and editing, Investigation, Methodology | Dawien Bradford, Writing – review and editing, Investigation, Supervision, Methodology | David W. New, Writing – review and editing, Investigation, Supervision | Rick Speare, Formal analysis, Investigation | Robin B. Gasser, Resources, Supervision, Writing – review and editing, Investigation, Methodology | Harsha Sheorey, Writing – review and editing, Investigation, Methodology | Richard S. Bradbury, Conceptualization, Formal analysis, Project administration, Resources, Supervision, Writing – review and editing, Data curation, Investigation, Methodology, Visualization

ETHICS APPROVAL

This study was reviewed by the James Cook University Human Research Ethics Committee and granted exemption to review under the National Statement Section 5.1.17(a).

ADDITIONAL FILES

The following material is available online.

Supplemental Material

Figure S1 (JCM01021-24-S0001.docx). *Strongyloides stercoralis* autoinfective third-stage larva recovered from the bronchoalveolar lavage (BAL) fluid of patient 2.

REFERENCES

- Vlaar LE, Bertran A, Rahimi M, Dong L, Kammenga JE, Helder J, Goverse A, Bouwmeester HJ. 2021. On the role of dauer in the adaptation of nematodes to a parasitic lifestyle. Parasites Vectors 14:554. https://doi. org/10.1186/s13071-021-04953-6
- Page W, Judd JA, Bradbury RS. 2018. The unique life cycle of *Strongy-loides stercoralis* and implications for public health action. Trop Med Infect Dis 3:53. https://doi.org/10.3390/tropicalmed3020053
- Buonfrate D, Hunt VL, Odermatt P, Streit A. 2024. Strongyloides: omics to worm-free populations. Philos Trans R Soc B 379:20220448. https://doi. org/10.1098/rstb.2022.0448
- Buonfrate D, Bradbury RS, Watts MR, Bisoffi Z. 2023. Human strongyloidiasis: complexities and pathways forward. Clin Microbiol Rev 36:e0003323. https://doi.org/10.1128/cmr.00033-23
- Al-Jawabreh R, Anderson R, Atkinson LE, Bickford-Smith J, Bradbury RS, Breloer M, Bryant AS, Buonfrate D, Cadd LC, Crooks B, et al. 2024. *Strongyloides* questions-a research agenda for the future. Philos Trans R Soc Lond B Biol Sci 379:20230004. https://doi.org/10.1098/rstb.2023. 0004
- Loukas A, Hotez PJ, Diemert D, Yazdanbakhsh M, McCarthy JS, Correa-Oliveira R, Croese J, Bethony JM. 2016. Hookworm infection. Nat Rev Dis Primers 2:16088. https://doi.org/10.1038/nrdp.2016.88
- Harvey SC, Gemmill AW, Read AF, Viney ME. 2000. The control of morph development in the parasitic nematode *Strongyloides ratti*. Proc R Soc Lond B 267:2057–2063. https://doi.org/10.1098/rspb.2000.1249
- Little MD. 1966. Comparative morphology of six species of *Strongyloides* (*Nematoda*) and redefinition of the genus. J Parasitol 52:69–84. https:// doi.org/10.2307/3276396
- 9. Speare R. 1986. Studies on the taxonomy of *Strongyloides* (Nematoda; Strongyloididae), James Cook University of North Queensland
- Faust EC. 1933. Experimental studies on human and primate species of Strongyloides. Am J Epidemiol 18:114–132. https://doi.org/10.1093/ oxfordjournals.aje.a117940
- Mati VLT, Raso P, de Melo AL. 2014. Strongyloides stercoralis infection in marmosets: replication of complicated and uncomplicated human disease and parasite biology. Parasit Vectors 7:579. https://doi.org/10. 1186/s13071-014-0579-2
- Fülleborn F. 1914. Untersuchungen über den infektionsweg bei Strongyloides und Ankylostomum und die biologie dieser parasiten, p 26–80. In Archiv für Schiffs- und Tropen-Hygiene. Vol. 18.
- Genta RM, Schad GA, Hellman ME. 1986. Strongyloides stercoralis: parasitological, immunological and pathological observations in immunosuppressed dogs. Trans R Soc Trop Med Hyg 80:34–41. https:// doi.org/10.1016/0035-9203(86)90190-2
- Schad GA, Hellman ME, Muncey DW. 1984. Strongyloides stercoralis: hyperinfection in immunosuppressed dogs. Exp Parasitol 57:287–296. https://doi.org/10.1016/0014-4894(84)90103-6

- Cahill KM, Shevchuk M. 1996. Fulminant, systemic strongyloidiasis in AIDS. Ann Trop Med Parasitol 90:313–318. https://doi.org/10.1080/ 00034983.1996.11813056
- McLarnon M, Ma P. 1981. Brain stem glioma complicated by Strongyloides stercoralis. Ann Clin Lab Sci 11:546–549.
- Buonfrate D, Tamarozzi F, Paradies P, Watts MR, Bradbury RS, Bisoffi Z. 2022. The diagnosis of human and companion animal *Strongyloides stercoralis* infection: challenges and solutions. A scoping review. Adv Parasitol 118:1–84. https://doi.org/10.1016/bs.apar.2022.07.001
- Kim J, Joo HS, Ko HM, Na MS, Hwang SH, Im JC. 2005. A case of fatal hyperinfective strongyloidiasis with discovery of autoinfective filariform larvae in sputum. Korean J Parasitol 43:51–55. https://doi.org/10.3347/ kjp.2005.43.2.51
- 19. Little MD. 1966. Seven new species of *Strongyloides (Nematoda*) from Louisiana. J Parasitol 52:85–97.
- Schad GA, Smith G, Megyeri Z, Bhopale VM, Niamatali S, Maze R. 1993. Strongyloides stercoralis: an initial autoinfective burst amplifies primary infection. Am J Trop Med Hyg 48:716–725. https://doi.org/10.4269/ ajtmh.1993.48.716
- 21. Grove Dl. 1996. Human strongyloidiasis. Adv Parasitol 38:251–309. https: //doi.org/10.1016/s0065-308x(08)60036-6
- Derycke S, Vanaverbeke J, Rigaux A, Backeljau T, Moens T. 2010. Exploring the use of cytochrome oxidase c subunit 1 (COI) for DNA barcoding of free-living marine nematodes. PLoS One 5:e13716. https:// doi.org/10.1371/journal.pone.0013716
- Hu M, Chilton NB, Gasser RB. 2003. The mitochondrial genome of *Strongyloides stercoralis* (Nematoda) - idiosyncratic gene order and evolutionary implications. Int J Parasitol 33:1393–1408. https://doi.org/ 10.1016/s0020-7519(03)00130-9
- Tamura K, Stecher G, Kumar S. 2021. MEGA11: molecular evolutionary genetics analysis version 11. Mol Biol Evol 38:3022–3027. https://doi.org/ 10.1093/molbev/msab120
- 25. Jitsamai W. 2019. Prevalence of enteric helminths and protozoa and identification of hookworm, threadworm and *Giardia* spp. in cats in Bangkok and vicinity, ThailandChulalongkorn University
- Bradbury RS, Sapp SGH, Potters I, Mathison BA, Frean J, Mewara A, Sheorey H, Tamarozzi F, Couturier MR, Chiodini P, Pritt B. 2022. Where have all the diagnostic morphological parasitologists gone? J Clin Microbiol 60:e0098622. https://doi.org/10.1128/jcm.00986-22
- Barratt JLN, Lane M, Talundzic E, Richins T, Robertson G, Formenti F, Pritt B, Verocai G, Nascimento de Souza J, Mato Soares N, Traub R, Buonfrate D, Bradbury RS. 2019. A global genotyping survey of *Strongyloides stercoralis* and *Strongyloides fuelleborni* using deep amplicon sequencing. PLoS Negl Trop Dis 13:e0007609. https://doi.org/10.1371/journal. pntd.0007609