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## Assessment of gastrointestinal parasites in wild chimpanzees (*Pan troglodytes troglodytes*) in southeast Cameroon

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### Abstract

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**Ethical standards** The research conducted here complies with host country and institutional policies of ethical research on non-human primates.

**Conflict of interest** The authors declare no conflict of interest.

We tested 114 faecal samples from wild simian immunodeficiency virus (SIV)-positive ( $n=43$ ) and SIV-negative ( $n=71$ ) chimpanzees (*Pan troglodytes troglodytes*) in southeast Cameroon for the presence of gastrointestinal parasites by direct smear. We observed cysts from different protozoa (*Entamoeba coli* and *Entamoeba histolytica/Entamoeba dispar*, *Endolimax nana*, *Iodamoeba butschlii*, *Chilomastix mesnili*, *Balantidium coli* and *Blastocystis* cells) and trophozoites from *Troglodytella abrassarti* and *Balantidium coli*. Eggs from different helminths (strongylids, *Ascaris lumbricoides*, *Abbreviata caucasica*, *Trichuris* sp., *Capillaria* sp., *Enterobius anthropopeci*, *Bertiella* sp., *Hymenolepis diminuta* and an undetermined fluke) were also observed. Finally, we observed eggs that could not be properly identified and classified. We did not observe any differences between the SIV+ and SIV- samples except for the unidentified eggs. The studied chimpanzees were highly parasitised by strongylid (85.1 % of prevalence), *Troglodytella* (43.8 %) and *Blastocystis* (2.9 %), and the frequency of the other parasites ranged from 0.9 to 8.8 %. These high levels of parasite infections could represent an additional burden in a population where there is a high rate of the SIV virus in circulation.

## Keywords

Chimpanzees; Parasites; Coprology; SIV; Helminths; Protozoa

## Introduction

Chimpanzees (*Pan troglodytes*) are classified as a highly endangered species (<http://www.iucnredlist.org/details/15936/0>) due to expanding human activities that have direct (through poaching and bush meat hunting) or indirect (through logging, mining and human-induced habitat fragmentation) detrimental effects on their survival. In addition, infectious diseases, such as Ebola haemorrhagic fever, paramyxovirus, anthrax and pneumonic streptococcosis, have additional impacts on the already fragile populations (Formenty et al. 1999; Leendertz et al. 2006; Chi et al. 2007; Kaur et al. 2008). Moreover, chimpanzees are also widely infected with the simian immunodeficiency virus (SIVcpz), the ancestor of HIV-1, the prevalence of which can reach up to 40 % in certain populations. Until recently, it was commonly accepted that SIVs were not pathogenic for their primate hosts (Hirsch 2004; Pandrea et al. 2008; Silvestri et al. 2007; Pandrea and Apetrei 2010). However, several studies have indicated that SIVcpz infection can lead to a severe immunodepression and ultimately to an AIDS-like syndrome, challenging this theory of non-pathogenicity. Indeed, CD4 T-cell depletion was retrospectively observed in necroscopy samples from infected chimpanzees; additionally, one of the tested chimpanzees showed skeletal muscle and hepatocellular atrophy and abdominal lesions due to a massive nematode infestation (Keele et al. 2009; Terio et al. 2011). A decrease in CD4+ T cells, which is associated with various protozoan, helminthic and fungal infections, was also reported for a naturally infected chimpanzee in a sanctuary in Cameroon (Etienne et al. 2011). These studies show that chimpanzees may be much more affected by SIV infections than previously thought. Thus, immunodepressed chimpanzees may be more susceptible to other pathogens, including gastrointestinal parasites. Indeed, several parasites that have been described in chimpanzees, including *Balantidium coli* (Kim et al. 1978; Nakauchi 1999), *Isospora belli* (Rijpstra 1967), *Cryptosporidium* sp. (Mbaya and Udendeye 2011; Locatelli et al. 2012;

Gonzalez-Moreno et al. 2013), *Microsporidia* (Sak et al. 2011; Locatelli et al. 2012) and *Blastocystis hominis* (Petrášová et al. 2010; Mbaya and Udendeye 2011), are known to be pathogenic in immunosuppressed HIV-infected humans (Karp and Auwaerter 2007). Here, we performed a pilot study on a highly SIV-infected chimpanzee population in southeast Cameroon to study the extent to which they are infected with gastrointestinal parasites.

## Material and methods

### Study site and samples collection

A total of 220 faecal samples were collected between 2003 and 2011, as previously described, around the village of Mambele in southeast Cameroon, an area where chimpanzees are known to be highly infected with SIVcpz (Keele et al. 2006; Van Heuverswyn et al. 2007) (Fig. 1). Briefly, stool samples were collected on tracks or in the vicinity of night nests. The GPS position was recorded, and the species was inferred in the field according to shape, size and texture of the faecal samples as well as presence of footprints or nearby nests. For each sample, approximately 20 g of stool was collected in a 50-ml tube containing 20 ml of RNAlater<sup>®</sup> (Ambion, Austin, TX) The samples were kept at ambient temperature at the base camp for a maximum of 3 weeks. These samples were then shipped to a laboratory in Montpellier and stored at  $-80^{\circ}\text{C}$ .

### DNA extraction from faecal samples and species identification

The host species was confirmed by mtDNA analyses, as described previously (van der Kuyl et al. 1995; Keele et al. 2006; Van Heuverswyn et al. 2006). Briefly, a QIAamp stool DNA MiniPrep Kit (Qiagen, Valencia, CA) was used to extract the faecal DNA. Two millilitres of the faecal sample was used to obtain a final elution volume of 100  $\mu\text{l}$  of faecal DNA. A ~450- to 500-bp fragment spanning the hyper variable D-loop region was amplified using the primers L15997 and H16498, and/or a 386-bp fragment spanning the 12S gene was amplified using the primers 12S-L1091 and 12S-H1478. The sequences obtained using a 3130xl Genetic Analyzer (Applied Biosystems, France) were aligned using the SeqMan DNASTar (Lasergene, Madison, USA) software. The species from which the samples were obtained was confirmed by performing neighbour-joining analysis using the CLUSTAL X 2.0 program (Thompson et al. 1997).

### Detection of HIV cross-reactive antibodies in faecal samples

IgGs were recovered after dialyses of the faecal samples using methods previously adopted for antibody detection in faecal samples of gorillas and chimpanzees (Keele et al. 2006; Van Heuverswyn et al. 2006). Briefly, RNAlater<sup>®</sup>-precipitated immunoglobulins were resolubilised by diluting the faecal/RNAlater<sup>®</sup> mixtures (1.5 ml) in PBS-Tween 20 (7.5 ml) followed by inactivation of this solution for 1 h at  $60^{\circ}\text{C}$  and centrifugation (3,500g for 10 min) to eliminate the presence of salt contained in the RNAlater<sup>®</sup> medium. The samples were then dialysed against PBS overnight at  $4^{\circ}\text{C}$ . The reconstituted extracts were tested for HIV cross-reactive antibodies using the INNO-LIA HIV confirmation test (Innogenetics, Ghent, Belgium) according to the manufacturer's instructions. Samples were scored as INNO-LIA positive when they showed cross-reactivity with at least one HIV antigen.

## Coprology

Once the species identification of the host was confirmed, we assessed the parasite content of the samples by using the method described in Drakulovski et al. 2013. For each sample, 1 ml of 50:50 stool/RNALater® mix was thawed at room temperature. Each sample was then vortexed briefly. Then, 100 µl was mixed with 100 µl of physiological serum. Next, 50 µl of the sample–physiological serum mix were smeared on a slide and observed directly under a microscope (Olympus BX41). A second slide was prepared using Para-selles KOP Color II (Fumouze diagnostics, France) dye (10 µl of dye for 50 µl of stool/physiological serum mix). When the slides results were both negative or when a significant discrepancy was found, either in type or in number of parasitic elements, between coloured and not coloured slides, two additional smears were performed. The species, number and the shape of the parasitic elements, as well as their ability to be dyed, were recorded. The parasitic elements were photographed using a Nikon Coolpix 5400 digital camera.

## Results

### Species confirmation and SIV infection rates

Among the 220 faecal samples collected between June 2003 and February 2011 and identified in the field as potentially originating from chimpanzees, mitochondrial DNA and phylogenetic analyses confirmed that 114/220 samples were indeed from the *Pan troglodytes troglodytes* subspecies. The remaining samples were either from other non-human primate species (11 *Cercocebus agilis*, 9 *Cercopithecus cephus*, 4 *Cercopithecus nictitans*, 4 *Lophocebus albigena*) of human origin (3 samples) or could not be identified, most likely because of DNA degradation.

A total of 43 (37.7 %) of the 114 chimpanzee samples cross-reacted with one or more HIV-1 antigens in the INNO-LIA HIV confirmation test.

### Gastrointestinal parasitic content

Precise species identification of some parasite elements, especially strongylid eggs and amoeba cysts, was difficult because of the hyperosmotic nature of the RNAlater® medium and the low temperature (−80 °C) the samples were preserved in (Drakulovski et al. 2013). Therefore, all the strongylid eggs were regrouped under that same label. For *Entamoeba* sp. amoeba, some cysts were identifiable as either *Entamoeba coli* or *Entamoeba histolytica/Entamoeba dispar* (Fig. 2), but in most cases, the internal features that are used for identification were too degraded to determine the species. Therefore, we decided to regroup all such cysts under the label *Entamoeba* sp. The same limiting factor made the detection and identification of smaller amoebas (*Endolimax*, *Iodamoeba*, *Chilomastix*) difficult. As a consequence, their prevalence was most likely underestimated.

We found eggs from different helminth species at the following rates of infection: strongylid eggs of different sizes, possibly belonging to different species, in 87/114 samples (85.1 %); eggs of *Ascaris lumbricoides* (1/114, 0.9 %), *Abbreviata caucasica* (1/114, 0.9 %), *Trichuris* sp. (4/114, 4.5 %), *Capillaria* sp. (2/114, 1.8 %), *Enterobius anthropopeci* (3/114, 2.7 %), *Bertiella* sp. (3/114, 2.7 %), *Hymenolepis diminuta* (1/114, 0.9 %), and an unidentified fluke

(2/114, 1.8 %); and different larvae. Regarding the presence of protozoa, we found cysts of *Entamoeba* sp. (10/114, 8.8 %), *Endolimax nana* (5/114, 4.4 %), *Iodamoeba, butschlii* (2/114, 1.8 %), *Chilomastix mesnili* (6/114, 5.2 %) and *Balantidium coli* (1/114, 0.9 %). We also observed *Troglodytella abrassarti* (50/114, 43.8 %), *Balantidium coli* trophozoites and *Blastocystis hominis* cells (25/114, 21.9 %) (Table 1 and Fig. 2).

In addition, unknown elements were observed in ten samples. They were characterised by an oval shape, 48–50 µm in length by 24–25 µm in width, with a thick wall (Fig. 2) and were stained with KOP Color II dye, as were other parasitological elements. Pollen database (PalDat, [www.paldat.org](http://www.paldat.org)) browsing did not link them to any identified pollen. The modified Ziehl stain method did not stain these samples as it would for coccidian oocysts. Thus, by default, these elements were classified as unknown helminth eggs. Their overall prevalence was 10.5 %, with several samples presenting a high number of “eggs” (2–3/microscope field at ×40).

The rates of parasite infection in SIV-positive versus SIV-negative chimpanzees were presented in Table 1. None of the rates were significantly different between SIV+ and SIV– groups except for the small unidentified eggs ( $|\varepsilon|=2.19>1.96$ ).

## Discussion

### Helminth infections

In this study, we found that the wild chimpanzees in the southeast Cameroon population are infected with 17 different parasite elements: 10 different types of helminths and 7 different types of protozoa. Among the helminth eggs, we found strongylid eggs of different sizes: large eggs (60–80 µm in length), possibly belonging to *Hyostromylus*, *Libyostromylus*, *Trichostromylus*, *Nematodirus*, *Ternidens*, *Oesophagostomum* and/or *Gongylonema* genera; and small eggs (40 to 60 µm of length), possibly belonging to *Strongyloides* and/or *Ankylostoma* genera (Rothman et al. 2006) (see Fig. 2). The rate of global strongylid infection was 85.1 %. This rate was significantly higher than the 10.9 to 41.3 % strongylid prevalence in wild *Pan troglodytes schweinfurthii* populations in Rubondo Island National Park, Tanzania (Petrášová et al. 2010; Petrzelková et al. 2010), 19.53 % prevalence in wild *Pan troglodytes verus* population in Fongoli, Senegal, (Howells et al. 2011) and 13.04 % prevalence found in captive *Pan troglodytes elliotii* from Calabar, Nigeria (Mbaya and Udendeye 2011). However, this rate of infection was similar to what was found in *Pan troglodytes schweinfurthii* populations of Gombe National Park, Tanzania (Gillespie et al. 2010), and Kibale National Park, Uganda (Krief et al. 2005; Muehlenbein 2005), with a strongylid prevalence ranging from 74.32 up to 100 %. The rate of infection found was also relatively higher than in free-ranging baboons and vervet monkeys in Ethiopia (Legesse and Erko 2004); forest baboons in Senegal (Howells et al. 2011); red-tailed, blue and L’Hoest’s monkeys in Uganda (Gillespie et al. 2004); blue monkeys in South Africa (Gillespie et al. 2004) and mountain gorillas in Uganda (Rothman et al. 2008). This high level of strongylid infection could represent a threat to the health of the chimpanzee population, as strongylids and Ankylostomatidae are known to be mildly to highly pathogenic in humans and are responsible for a wide range of symptoms, from mucosal ulcerations to abscesses, anaemia,

weight loss and ultimately death in non-human primates (NHPs) (Gutierrez 2000; Rothman et al. 2006; Krief et al. 2008; Terio et al. 2011).

Concerning the other helminth infections (*Ascaris*, *Trichuris*, *Abbreviata*, *Capillaria*, *Enterobius*, *Bertiella*, Fluke and *Hymenolepis*), the prevalence was low and ranged from 0.9 to 4.5 %, depending on the species, and was similar to or below the average rate of infection found in other chimpanzee communities in Tanzania, Nigeria, Uganda, and Central African Republic (Lilly et al. 2002; Krief et al. 2003; Gillespie et al. 2010; Howells et al. 2011; Petrášová et al. 2010; Petrzelková et al. 2010; Mbaya and Udendeye 2011).

### Protozoan infections

In our samples set, we found up to seven different protozoan elements belonging to the *Entamoeba*, *Endolimax*, *Iodamoeba*, *Chilomastix*, *Balantidium*, *Troglodytella* and *Blastocystis* genera. The amoeba cyst prevalence was low (8.8 % for the *Entamoeba* sp. and ranging from 1.75 to 5.2 % for the smaller amoebas) and similar to what was detected in the chimpanzee communities in Fongoli, Senegal (Howells et al. 2011), Rubondo Island, Tanzania (Petrzelková et al. 2010), and Kibale National Park, Uganda (Muehlenbein 2005), and slightly lower than what was found in Gombe National Park, Tanzania (Gillespie et al. 2010), and Dzanga-Ndoki National Park, Central African Republic (Masi et al. 2012). The smaller amoebas (*Endolimax*, *Iodamoeba*, *Chilomastix*) do not represent a health hazard: They are commensal non-pathogenic parasites. However, even if the prevalence of the *Entamoeba* group (including the highly pathogenic *Entamoeba histolytica*) was low, its presence could have consequences, as amoebic dysentery outbreaks are potentially lethal in chimpanzees (Miller and Bray 1966).

In respect to ciliates detection, we found no *Giardia intestinalis* shedding in our sample set. Instead, we found high prevalence of *Troglodytella abrassarti* (prevalence 43.8 %), consistent with other studies of chimpanzees (Pomajbíková et al. 2010a). We also found one positive sample for *Balantidium coli* cysts and trophozoites. *Balantidium coli* is considered to be mildly pathogenic and is rarely detected in wild chimpanzees (Pomajbíková et al. 2010b). However, clinical balantidiasis has been reported in captive African great apes (Teare and Loomis 1982; Lee et al. 1990; Lankester et al. 2008). The possible clinical significance in a high rate SIV-infected population will be discussed below.

Finally, we also detected *Blastocystis* elements at relatively high prevalence (21.9 %). This prevalence was likely underestimated because *Blastocystis* are small elements easier to observe after concentration methods not applicable to our sample set. Indeed, a pilot study using molecular detection methods on the chimpanzee stool samples collected from the same area showed presence of *Blastocystis* in 87–88 % of the samples (Locatelli et al. 2012). Infection with *Blastocystis* has already been reported in the captive and free-roaming chimpanzee population of the Afi Mountain Conservation area, Calabar, Nigeria (Mbaya and Udendeye 2011), and in Rubondo Island, Tanzania (Petrášová et al. 2010; Petrášová et al. 2011), with prevalence ranging from 13.04 to 60.9 %. They were also detected in other NHP species including vervets and baboons in Ethiopia (Legesse and Erko 2004) as well as orangutans in Borneo, Indonesia (Labes et al. 2010). The infectious role of *Blastocystis* remains unclear so far, as asymptomatic shedding is highly common. However, *Blatocystis*

presence is also associated with gastrointestinal symptoms in humans including diarrhoea and abdominal pain (Qadri et al. 1989). *Blastocystis* could also be responsible for chronic gastrointestinal illnesses, including irritable bowel syndrome (Boorom et al. 2008; Poirier et al. 2012). However, the clinical significance of *Blastocystis* infection for NHPs populations is as unclear as it is for the human population and, in certain cases, may be dependent on the subtype involved (Hussein et al. 2008; Parija and Jeremiah 2013).

### Gastrointestinal parasites in SIV-positive versus SIV-negative chimpanzees

This is the first study that compares the gastrointestinal parasites of wild-living chimpanzees in respect to their SIV serological status. Concerning the helminths, there was no significant difference in the prevalence of strongylid, *Ascaris*, *Abbreviata*, *Trichuris*, *Capillaria*, *Enterobius*, *Bertiella*, fluke or *Hymenolepis* infections between the groups. However, there was a slightly significant difference ( $|z|=2.19>1.96$ ) in the prevalence of individuals shedding the unidentified helminth eggs. Helminths are poor lentivirus infection sentinels, as their rate of infection is not particularly influenced by HIV-induced immunodepression in humans (Wiwanitkit 2006; Karp and Auwaerter 2007). Thus, it is not clear why such a difference exists in this specific case, especially for a still unidentified helminth. We can only speculate that it is most likely more related to the feeding habits of the chimpanzee population and ecological/environmental factors that are more likely to influence a helminth infection (Gillespie et al. 2010) rather than their immunological status.

Regarding protozoan infection, there were no significant differences between SIV+ and SIV – samples sets for any of the detected amoeba (*Entamoeba* sp., *Iodamoeba*, *Chilomastix* or *Endolimax*), ciliates (*Troglodytella* or *Balantidium*) or heterokonts (*Blastocystis*).

Results for gastrointestinal amoebas were not unexpected, as these protozoa are not considered to be opportunistic, and their prevalence is not influenced by HIV serological status in humans (Wiwanitkit 2006; Karp and Auwaerter 2007). The ciliate *Balantidium coli* is not considered to be an opportunistic parasite in humans with AIDS either. However, this protozoon can be responsible for chronic, life-threatening diarrhoea in HIV+ patients (Clyti et al. 1998; Cermeño et al. 2003). As such, its presence in a SIV hotspot could be of concern for the health and survival of the apes' population. The other ciliate detected in our sample set, *Troglodytella abrasarti*, is commonly found in great ape (gorillas, chimpanzees and bonobos) faeces (Pomajbíková et al. 2010a). This ciliate was previously considered to be pathogenic and associated with diarrhoeas (Mortelmans et al. 1971). However, more recent studies have suggested a symbiotic function and participation in nourishment degradation ((Pomajbíková et al. 2010a; Profousová et al. 2011). Indeed, occurrence and shedding intensity of *Troglodytella* trophozoites seem linked to the level of starch in the food intake in chimpanzees (Petrželková et al. 2012). Thus, this entodiniomorphid does not seem to be harmful in healthy apes. However, its health impact on SIV-infected chimpanzees is still to be determined, especially given that ape-to-ape transmission may occur (Modrý et al. 2009). Finally, concerning *Blastocystis* sp., there are contradictory observations concerning the *Blastocystis* sp. prevalence in HIV+ versus HIV– patients. In some studies, increased prevalence of these protozoa has been described in an HIV+ population (Storgaard et al. 1996), but other studies did not find this correlation (Morgan et al. 1996). Whether these

observations could be extrapolated to chimpanzees, where the prevalence of *Blastocystis* is high (18.6 %) in a heavily SIV-infected population, should be a concern, especially in young animals because *Blastocystis* has been associated with chronic diarrhoeas in HIV+ children (Cegielski et al. 1993; Idris et al. 2010).

### Inter-species (apes to humans and humans to apes) transmission risk

We performed our study on faecal samples collected from wild chimpanzees. Because these primates were not habituated to humans, contact with humans and inter-species transmission of pathogens should be minimal. The absence of *Giardia intestinalis* in our samples supports this theory. Indeed, *Giardia intestinalis* is the model example of a gastrointestinal parasite transmitted by faecal contamination of the environment (water and farming). The baseline prevalence of *Giardia intestinalis* contamination of free-living chimpanzees in undisturbed areas was set at 3 % (Gillespie et al. 2009). As we have prevalence of 0 % in our sample set (Table 1), it would be easy to conclude that interactions between humans and great apes in the area are below minimal. However, several facts are challenging this observation. First, three faecal samples collected on chimpanzee trails or near a nest were of human origin. Secondly, one sample was detected positive for *Balantidium coli*. While this protozoa is almost never detected in wild apes (Pomajbíková et al. 2010b), *Balantidium coli* is known to infect a wide range of hosts from pigs (wild or domesticated) to rats but also humans. Terrestrial or semi-arboreal NHPs are likely to enter in contact with contaminated soil or faeces where habitats overlap (Levecke et al. 2007; Schuster and Ramirez-Avila 2008). Gastrointestinal pathogen exchanges between humans and apes have been already documented in African regions (Goldberg et al. 2008; Rwego et al. 2008). Thus, this infected individual could be the sentinel case showing that interaction between humans or human livestock and wild chimpanzee occurs around Mambele, leading to a risk of contamination for the apes. Finally, farmers frequently report chimpanzees raiding crops around the village of Mambele. Indeed, GPS data showed that some stool samples were collected less than 1,500 to 1,000 m from the centre of the village (see Fig. 1). The proximity to human settlements can increase the chances of parasite contamination from chimpanzees to humans, especially for strongylid/Ankylostomid. Cross-species transmission of *Strongyloides fülleborni* has been described in Africa (Hira and Patel 1977; Hira and Patel 1980; Evans et al. 1991) and could be a concern with the vicinity of such a highly strongylid-infected chimpanzee population.

### Conclusion

This is the first study assessing parasitic coprology in wild chimpanzees from southeast Cameroon and the first study that compares the frequency of parasites between SIV+ and SIV- chimpanzees. Because SIVcpz has a negative impact on the health and survival of chimpanzees, it is important to document the gastrointestinal pathogens present in this chimpanzee population. Studying this chimpanzee population showed that it is highly infected with gastrointestinal nematodes of the strongylid type (85.1 % of prevalence), *Troglodytella* (43.8 % of prevalence) and *Blastocystis* (21.9 % of prevalence) as well as several others helminths and protozoa. We did not find any significant difference in the prevalence between SIV+ and SIV- samples; however, this chimpanzee population is



clearly infected with mildly to potentially highly pathogenic helminths and protozoa that could be lethal in HIV-infected humans and with pathogens responsible for deleterious symptoms in HIV+ children and possibly also for SIVcpz-infected chimpanzees.

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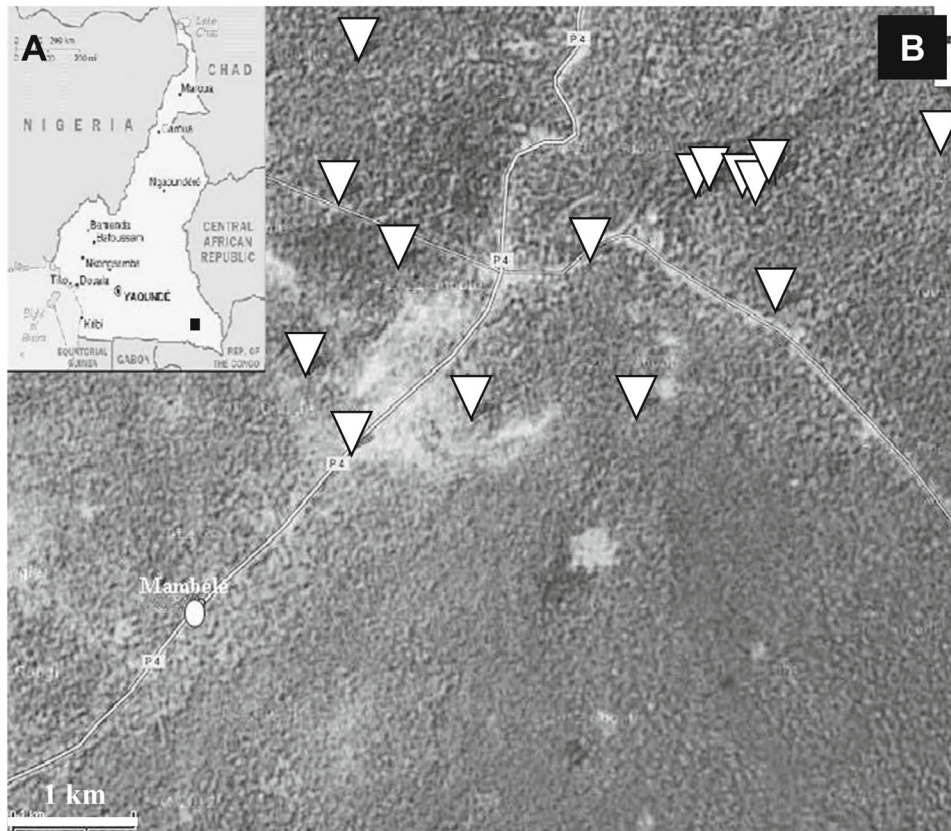
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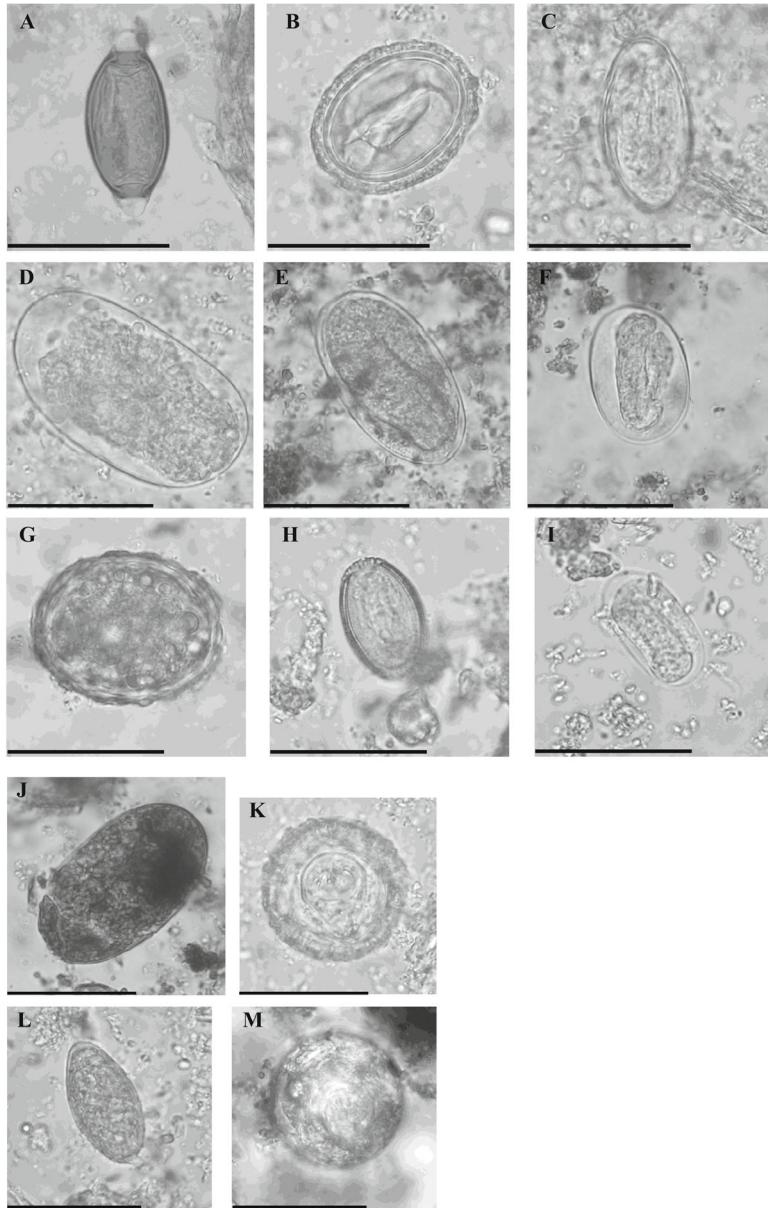
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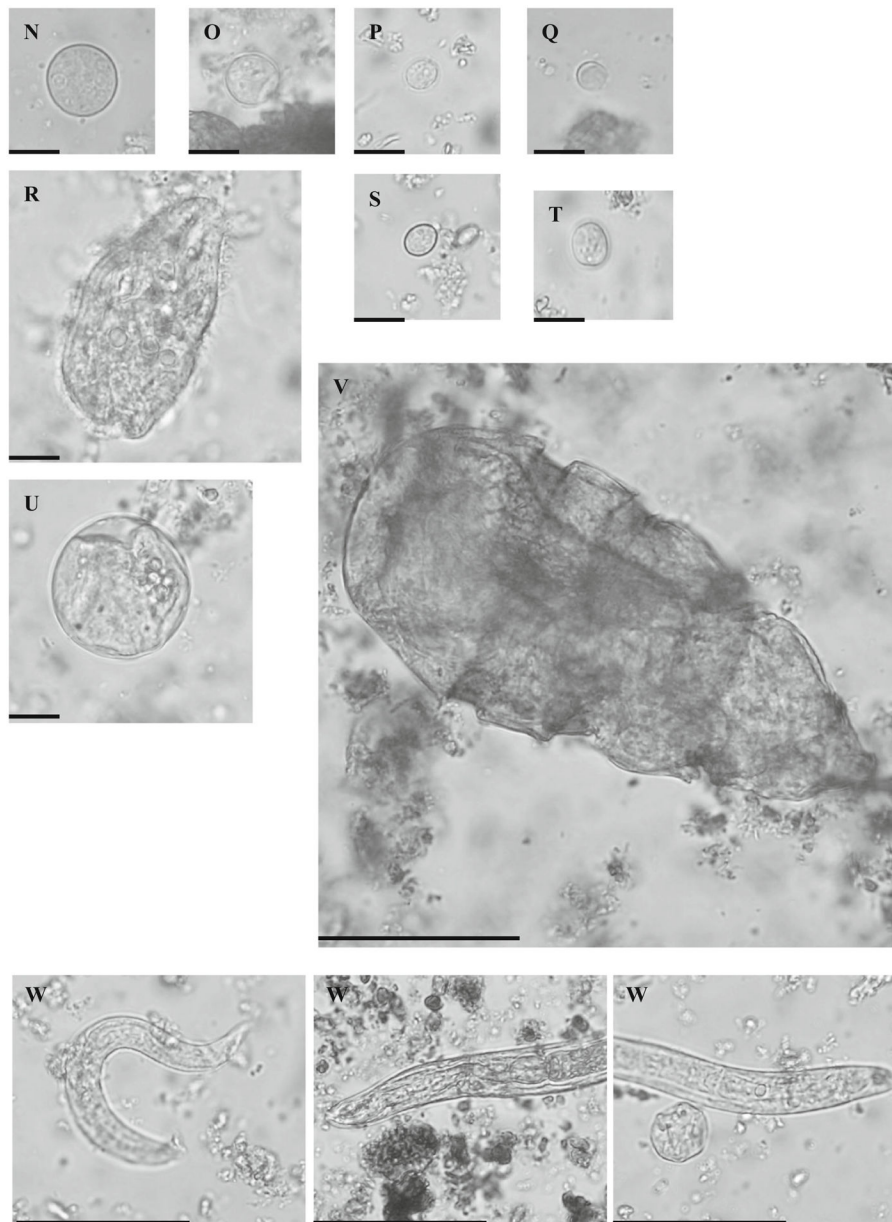
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**Fig. 1.** Sample collection points. **a** Cameroon map. Sample collection area is indicated by *square*. **b** Mambelé area view. GPS collection points are indicated by *inverted triangle*. Mambelé village centre is indicated by *circle*. The distance bar indicates 1 km





**Fig. 2.** Parasite elements observed by direct microscopy in stool samples collected from wild chimpanzees between 2003 and 2011 in the Mambele area of Cameroon. **a** *Trichuris* sp. egg. **b** *Abbreviata caucasica* egg. **c** *Enterobius anthropopeci* egg. **d** Large strongylid egg (>50  $\mu\text{m}$  in length). **e** Large strongylid egg with formed lava (>50  $\mu\text{m}$  in length). **f** Small strongylid egg with formed larva (<50  $\mu\text{m}$  in length). **g** *Ascaris lumbricoides* egg. **h** *Capillaria* sp. egg. **i** Small strongylid egg (<50  $\mu\text{m}$  in length) with formed larva and thin wall, probably Ankylostomid egg. **j** Undetermined fluke egg. **k** *Bertiella* sp. egg. **l** Unidentified egg. **m** *Hymenolepis diminuta* egg. **n** *Entamoeba coli* cyst. **o** *Entamoeba histolytica/Entamoeba dispar* cyst. **p** *Endolimax nana* cyst. **q** *Blastocystis* cell. **r** *Balantidium coli* trophozoite. **s** *Chilomastix mesnili* cyst. **t** *Iodamoeba butschilii* cyst. **u**



*Balantidium coli* cyst. **v** *Troglodytella abrassarti* trophozoite. **w** Various nematode larvae.  
Size bars indicate respectively 50 µm for elements **a** to **m** and **v** to **w** and 10 µm for elements **n** to **u**

Table 1

Prevalence of various parasite elements found in chimpanzee stool sample set collected from 2003 to 2011 in the Mambéle area of Cameroon

Parasite elements (eggs or cysts)	Number of SIV+ samples positive for parasite elements (/43)	Prevalence among SIV+ samples	Number of SIV- positive for parasite elements (71)	Prevalence among SIV- samples	Overall prevalence (SIV++ SIV- samples)
Strongylidae	35	<b>81.4 %</b>	62	<b>87.3 %</b>	<b>85.1 %</b>
<i>Ascaris</i>	1	2.3 %	0	0 %	0.9 %
<i>Abbreviata</i>	0	0 %	1	1.4 %	0.9 %
<i>Trichuris</i>	1	2.3 %	3	4.2 %	4.5 %
<i>Capillaria</i>	2	4.6 %	0	0 %	1.8 %
<i>Enterobius</i>	0	0 %	3	4.2 %	2.7 %
<i>Bertiella</i>	2	4.6 %	1	1.4 %	2.7 %
<i>Hymenolepis</i>	0	0 %	1	1.4 %	0.9 %
Fluke	1	2.3 %	1	1.4 %	1.8 %
Unidentified egg	8	18.6 %*	4	5.6 %*	10.5 %
<i>Entamoeba</i>	3	6.9 %	7	9.9 %	8.8 %
<i>Endolimax</i>	1	2.3 %	4	5.6 %	4.4 %
<i>Iodamoeba</i>	1	2.3 %	1	1.4 %	1.8 %
<i>Chilomastix</i>	2	4.6 %	4	5.6 %	5.2 %
<i>Giardia</i>	0	0 %	0	0 %	0 %
<i>Balanitidium</i>	0	0 %	1	1.4 %	0.9 %
<i>Troglochyrella</i>	19	<b>44.2 %</b>	31	<b>43.6 %</b>	<b>43.8 %</b>
<i>Blastocystis</i>	8	18.6 %	17	<b>23.4 %</b>	<b>21.9 %</b>

Prevalence of the various helminths (A) or protozoa (B) elements was calculated for SIV+, SIV- and total chimpanzee samples. High prevalence (>20 %) is indicated in bold. Significant difference ( $(P > 1.96)$ ) between SIV+ and SIV- groups is indicated by asterisk