Autochthonous Human Fascioliasis, Belgium

Sandrine Milas,1 Camelia Rossi, Ivan Philippart, Pierre Dorny, Emmanuel Bottieau

Author affiliations: University Hospital Center Tivoli, La Louvière, Belgium (S. Milas); University Hospital Center Ambroise Paré, Mons, Belgium (C. Rossi); Regional Hospital Mons-Hainaut, Mons (I. Philippart); Institute of Tropical Medicine, Antwerp, Belgium (P. Dorny, E. Bottieau)

DOI: https://doi.org/10.3201/eid2601.190190

We report 2 cases of human fascioliasis (HF) in Belgium, likely caused by consumption of vegetables from a garden that was flooded by pasture runoff. Because autochthonous HF is rare and the route of transmission was unusual, HF was not diagnosed until 6 months after symptom onset in both cases.

Human fascioliasis (HF) is a plantborne and waterborne infection caused by the trematodes Fasciola hepatica in temperate areas and F. gigantica in tropical areas (1,2). Fasciola spp. trematodes infect herbivorous mammals and humans. The Fasciola life cycle requires 2 hosts; ruminants carry adult worms and excrete eggs into the environment in feces; lymnaeid snails are invaded and release cercariae, which encyst as metacercariae on aquatic vegetation. Humans become infected by ingesting raw aquatic vegetables or consuming plants or water containing metacercariae (3). Symptoms of fascioliasis are stage-specific and related to hepatic migration by larva or obstruction of the biliary ducts by adult worms (1,2).

In Belgium, only 6 cases of HF have been published since 1960 (4–6). We describe 2 autochthonous cases of HF. The cases were seen in different hospitals and initially were not linked epidemiologically.

Case 1 was in a 72-year-old man with no underlying medical conditions and no history of travel outside the country who was referred to the Regional Hospital of Mons in November 2008. He had fever, abdominal pain, rash, and hypereosinophilia that had lasted for 8 weeks (Table). Fecal egg detection was negative. Several serologic tests targeting parasitic infections were performed (Table); results for Trichinella spiralis were positive, but this diagnosis was discarded in the absence of myalgia and elevated creatine kinase. Eventually, a diagnosis of idiopathic hypereosinophilic syndrome was made. The patient received high doses of corticosteroids, but his condition did not improve. He was reevaluated in March 2009, and HF was considered on the basis of combined clinical, laboratory, and radiologic findings (Table). An indirect hemagglutination test for Fasciola spp. was performed by using ELIHA Distoma (ELITech Group, https://www.elitechgroup.com), and results were positive. The patient received triclabendazole (10 mg/kg/d) for 2 consecutive days. His symptoms abated, and his eosinophil count was nearly normal 1 month later (Table).

Case 2 was in the index case’s neighbor, who experienced similar symptoms that lasted for 3 months before she was seen at the University Hospital Center Ambroise Paré in Mons in December 2008. She also was misdiagnosed initially (Table). In February 2009, 2 stool examinations were negative for parasite eggs. In March, an indirect hemagglutination test for Fasciola was performed and was highly positive. The patient received a single dose of triclabendazole (15 mg/kg) and recovered fully within 5 months (Table).

A detailed anamnesis revealed that both patients consumed unwashed raw vegetables from case-patient 2’s garden, which was flooded with runoff from a neighboring cattle pasture in August 2008. We hypothesize the vegetables were contaminated by metacercariae, either by Fasciola-infested amphibious snails washed into the garden or directly by runoff. We did not perform sampling of the garden. Cases related to garden vegetables contaminated by flooding have been reported previously, such as in Corsica (3).

HF is not a notifiable disease in Belgium. Among 6 published cases, 3 occurred in a cluster related to consumption of homegrown watercress, and 3 non-clustered cases had a questionable autochthonous nature (4–6). Consumption of watercress and dandelions is uncommon in Belgium, but common in France, where ≥300 HF cases occur annually (7). However, Fasciola infection in cattle is common in Belgium; herd prevalence is 37.3% in Flemish dairy cattle (8). In addition, a 2008 survey of snails showed 1.31% of Galba truncatula and 0.16% of Radix spp. were infected by Fasciola trematodes (9).

Both cases in this study experienced an acute invasive stage and a considerable delay in HF diagnosis. Clinicians should be aware of key elements of HF, including potential diet exposure, clinical signs and symptoms, and imaging and laboratory findings. Contrast-enhanced computed tomography scans of the liver sometimes show tortuous subcapsular tracts associated with hypodense nodules and hepatomegaly during the acute phase (2). In industrialized countries, human cases occur singly or in small clusters,

1Current affiliation: University Center of Charleroi, Charleroi, Belgium.
and diagnosis usually is made during the invasive phase by combined clinical, laboratory, and imaging findings. Serologic tests can detect antibodies within 2 weeks after infection but might have low specificity. Fasciola eggs can appear in stool 3–4 months postinfection, depending on the infection burden and the ability of the flukes to produce eggs. Intermittent shedding can occur (1,2). Co-proantigens are detectable 8 weeks after infection and have a high sensitivity, but 1 negative result despite high egg shedding has been reported (2).

Triclabendazole, licensed in Europe only by Novartis (https://www.novartis.com), at 10–15 mg/kg/day in 1 dose or on 2 consecutive days, is the preferred treatment, and patients usually recover rapidly. Resistance increasingly is described in ruminants and treatment failures have been reported in humans (2,10).

Although overlooked in countries in northern Europe, HF should be considered in cases of unexplained eosinophilia associated with liver symptoms, even in the absence of ingestion of freshwater plants. This zoonotic condition highlights the need for good epidemiologic communication between human and animal health workers.

Acknowledgment
We thank Quentin Groom from the Botanic Garden Meise, Meise, Belgium, for his contribution to this article.

About the Author
At the time of this study, Dr. Milas was an infectious disease specialist at University Hospital Center Tivoli, La Louvière, Belgium. She is currently an infectious disease specialist at University Hospital Center of Charleroi, Charleroi, Belgium. Her research interests include general infectious diseases, travel medicine, and infection control.

References
1. Mas-Coma S, Bargues MD, Valero MA. Fascioliasis and other plant-borne trematode zoonoses. Int J Parasitol. 2005;35:1255–78. https://doi.org/10.1016/j.ijpara.2005.07.010
2. Mas-Coma S, Bargues MD, Valero MA. Diagnosis of human fascioliasis by stool and blood techniques: update for the present global scenario. Parasitology. 2014;141:1918–46. https://doi.org/10.1017/S0031182014000869
We report multiple nontypeable genotype II noroviruses circulating in South America; nucleotides differed by >25% from those of other genotypes. These viruses have been circulating in the Americas for ≈20 years and show recombination with other genotypes. Clues to norovirus natural history can guide development of treatment and prevention plans.

Norovirus is a leading cause of acute gastroenteritis (1). The norovirus RNA genome is organized into 3 open reading frames (ORFs). ORF1 encodes for 6 nonstructural proteins, including the RNA-dependent RNA polymerase (RdRp). ORF2 and ORF3 encode for the major capsid protein (VP1) and minor capsid protein (VP2). Norovirus classification was recently updated, and these viruses are now classified into 10 genogroups (GI–GX) and ≈40 genotypes (2). This classification is based on the genetic diversity presented by VP1 and RdRp (3). Human noroviruses are mostly represented by the GI and GII strains.

Advances in genome sequencing approaches enabled us to detect a novel (nontypeable) GII norovirus circulating in Peru in 2008 (4). As part of a larger study to sequence the genomes from noroviruses circulating on different continents and over different decades, we found 7 additional nontypeable GII norovirus strains: 2 in fecal samples (PNV024019 and PNV027026) from children in Peru with diarrhea (5) and 5 in fecal samples collected in Buenos Aires, Argentina (6) (Appendix 1 Table, https://wwwnc.cdc.gov/EID/article/26/1/19-0626-App1.pdf). These strains were detected by routine PCR screening and were either incorrectly assigned as GII.22 or could not be assigned to any genotypes in the