

Article

Protozoa as the “Underdogs” for Microbiological Quality Evaluation of Fresh Vegetables

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Abstract: The monitoring of the microbial quality of fresh products in the industrial environment has mainly focused on bacterial indicators. Protozoa, such as *Giardia duodenalis*, *Cryptosporidium* spp., *Toxoplasma gondii*, and *Cyclospora cayetanensis*, are routinely excluded from detection and surveillance systems, despite guidelines and regulations that support the need for tracking and monitoring these pathogens in fresh food products. Previous studies performed by our laboratory, within the scope of the SafeConsume project, clearly indicated that consumption of fresh produce may be a source of *T. gondii*, thus posing a risk for the contraction of toxoplasmosis for susceptible consumers. Therefore, preliminary work was performed in order to assess the microbiological quality of vegetables, highlighting not only bacteria (*Escherichia coli*, *Listeria monocytogenes*, and *Salmonella* spp.), but also the zoonotic protozoa *G. duodenalis* and *Cryptosporidium* spp. Although all samples were found to be acceptable based on bacteriological parameters, cysts of *G. duodenalis* and oocysts of *Cryptosporidium* spp. were observed in vegetables. Moreover, it was possible to genetically characterize *G. duodenalis* positive samples as assemblage A, a genotype that poses risks to human health. Although these are preliminary results, they highlight the need to include protozoa in the microbiological criteria for foodstuffs, as required by EU Law No. 1441/2007, and to improve inactivation and removal procedures of (oo)cysts in fresh produce and water.

Keywords: vegetables; foodborne pathogens; *Giardia duodenalis*; *Cryptosporidium* spp.; *Escherichia coli*; *Listeria monocytogenes*; *Salmonella* spp.



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1. Introduction

Each year, nearly 600 million people become ill with foodborne diseases, leading to over 420,000 deaths [1]. Microbial contamination of food products can be caused by bacteria, viruses, or parasites, and these pathogens represent the most important cause of enteric illness [2]. During the past decade, there has been a remarkable increase in the consumption of raw or partially cooked fresh produce, due to changes in the eating habits of consumers who are more conscious of the direct and indirect health benefits of eating fruit and vegetables [3]. Hence, these types of foods have also become important vehicles for transmission of foodborne pathogens, and the number of outbreaks of foodborne diseases associated with fresh produce has increased [4–6]. Examples of major multi-national foodborne outbreaks include an *Escherichia coli* outbreak due to contaminated fenugreek

seeds and an outbreak of listeriosis due to the consumption of frozen vegetables contaminated with *Listeria monocytogenes* [7,8]. In addition to bacteria, several protozoan parasites can also contaminate fresh produce. *Giardia duodenalis*, *Cryptosporidium* spp., *Toxoplasma gondii*, and *Cyclospora cayetanensis* have been associated with several global outbreaks of foodborne illnesses related to the consumption of contaminated fresh vegetables and fruits, and are the most important emerging parasitic protozoans that commonly infect humans and animals [9–14].

Despite guidelines and regulations that support the need for tracking and monitoring protozoa in fresh food products, their routine exclusion from detection and surveillance systems can be ascribed to the limited number of studies investigating parasite contamination of vegetables, the absence of consensual and accurate methodologies for their identification, and the wide acceptance among the expert authorities that the bacteriological aspects of the evaluation are more relevant for food safety [14]. Consequently, the monitoring of the microbial quality of fresh products in the industrial environment has mainly focused on bacterial indicators, and EU and national laws highly regulate the presence and microbiological limits of, in particular, the bacteria *E. coli*, *L. monocytogenes*, and *Salmonella* spp. [15–18]. In contrast, protozoan pathogens are rarely tested in fresh produce for food safety purposes. They are frequently neglected because they do not cause an immediate deleterious clinical response, but rather a chronic and long one. In addition, the associated diseases are also more common and prevalent in poor countries or communities. However, the harmful effects of these pathogens on humans in developed countries, especially on immunocompromised individuals, children, and pregnant women, and particularly their recent impacts, have increased the focus and attention of authorities [14,19–23].

Giardia duodenalis is the causative agent of giardiasis, resulting in acute watery diarrhea in both humans and animals. Eight major genotypes of *G. duodenalis* (assemblages) have been identified to date (A–H); however, assemblages A and B (and to a lesser extent assemblage E) are considered to have zoonotic interest due to their risk of infecting humans [24,25].

Cryptosporidium spp. are widespread protozoan parasites that infect humans and animals. Human cryptosporidiosis is the second commonest cause of diarrhea in children after rotavirus and can be caused either by the zoonotic *Cryptosporidium parvum* or anthroponotic *Cryptosporidium hominis* [9,26].

Toxoplasma gondii is an intracellular coccidian protozoan, for which domestic and wild felids are the only known definitive hosts responsible for oocyst dissemination in the environment. Toxoplasmosis is usually asymptomatic in immunocompetent individuals, but may cause severe infections in immunocompromised individuals, fetuses during pregnancy, and newborns [27,28].

Cyclospora cayetanensis is a human coccidia parasite that causes cyclosporiasis, a disease that causes acute diarrhea and other gastroenteritis symptoms. Humans are the only known host for *C. cayetanensis* oocysts, and their zoonotic role remains to be determined [29,30].

Two of the main reasons for the biological success of these parasites are their low infectious dose and their excretion in feces into the environment as resistant stages (cysts or oocysts) [31,32]. The (oo)cysts possess the ability to survive for a long period of time (weeks, months, or even years) in the environment, in extreme conditions of temperature and humidity, due to their structure, which protects them in exogenous stages. This allows the parasites to resist the common inactivation methods used for bacteria and viruses, such as heating, irradiation, and chemical disinfection [33–38].

Fresh vegetables may be contaminated at several steps along the food production chain, such as during crop production, harvesting, and processing, or directly by infected food handlers. Potential sources include water (irrigation water or water used for washing produce), wastewater discharge, soil contaminated with fecal waste from warm-blooded animals/livestock, organic fertilizers, and human handling during downstream processing and packaging steps [5,38–41].

In a recent work performed by our team [42], within the scope of the SafeConsume project (<https://safeconsume.eu/>, accessed on 11 July 2022), we designed a laboratory approach for the detection of *T. gondii* oocysts in vegetables and berry fruits based on the knowledge gained with Method 1623.1/EPA for the detection of the *Cryptosporidium* oocyst and *Giardia* cyst [43,44]. Moreover, we also described a new recombinant *T. gondii* oocyst wall protein 1-derived fragment, rTgOWP1-f. Strong microscopic evidence of the ability of rabbit polyclonal antibodies, anti-rTgOWP1-f, to identify environmental *T. gondii* oocysts, suggests rTgOWP1-f may be a potential biomarker for the detection of environmental oocysts [45]. Our studies clearly indicate that consumption of fresh produce may be a source of *T. gondii* infection in humans and a potential risk for consumers. However, technical refinement is still required for routine application at the industrial level or for food testing in laboratories for the detection of *T. gondii* oocysts.

Taking into account the experience and learning obtained from the SafeConsume work package 2 (WP2) laboratory studies, and the “One Health” approach, a parallel preliminary study was undertaken in order to assess the microbiological quality of fresh vegetables, highlighting not only bacteria (*E. coli*, *L. monocytogenes*, and *Salmonella* spp.) covered by the legislation [17], but also the zoonotic protozoa *G. duodenalis* and *Cryptosporidium* spp.

2. Materials and Methods

Five bulk and 11 packaged and ready-to-eat (RTE) vegetables were collected from local producers in Portugal and Spain, or provided by retail sellers, between September 2018 and June 2019 (Table 1). The fresh produce comprised lettuce (*Lactuca sativa*), watercress (*Nasturtium officinale*), coriander (*Coriandrum sativum*), and parsley (*Petroselinum crispum*). Ready-to-eat mixed salads included different varieties of lettuce, arugula (*Eruca vesicaria sativa*), endive (*Cichorium endivia*), chicory (*Cichorium intybus*), carrot (*Daucus carota sativus*), red cabbage (*Brassica oleracea* var. *capitata* f. *rubra*), and lamb’s lettuce (*Valerianella locusta*).

Table 1. Origin, collection date, and product presentation of the vegetable samples.

Sample Number	Product	Origin	Collection Date	Product Presentation
1	Watercress	Portugal	27 September 2018	bulk
2	Coriander	Portugal	27 September 2018	bulk
3	Parsley	Portugal	27 September 2018	bulk
4	Watercress	Portugal	2 October 2018	RTE
5	Mixed salad	Portugal	2 October 2018	RTE
6	Coriander	Portugal	2 October 2018	packaged
7	Parsley	Portugal	2 October 2018	packaged
8	Watercress	Portugal	12 October 2018	RTE
9	Mixed salad	Portugal	12 October 2018	RTE
10	Coriander	Portugal	12 October 2018	packaged
11	Parsley	Portugal	12 October 2018	packaged
12	Mixed salad	Portugal	4 February 2019	RTE
13	Mixed salad	Spain	17 June 2019	RTE
14	Mixed salad	Spain	17 June 2019	RTE
15	Arugula	Spain	17 June 2019	RTE
16	Lettuce	Portugal	18 June 2019	bulk

RTE: ready-to-eat.

The concentration and recovery of *G. duodenalis* and *Cryptosporidium* spp. (oo)cysts from the vegetable samples were performed by Filtration/Immunomagnetic Separation (IMS)/Fluorescence Assay (FA) (Method 1623.1: *Cryptosporidium* and *Giardia* in water; US EPA 816-R-12-001-Jan 2012). Method 1623.1 is a validated and approved method used for surface water analysis [43]. This method was previously fully established in our laboratory, initially to gain knowledge regarding the contamination of water with *Cryptosporidium* spp. and *Giardia* spp. in the northern region of Portugal [44,46]. We also previously designed a long-term program aiming to identify the sources of surface water and environmental

contamination, working with the water-supply industry, and to provide cattle and human fecal screening and molecular characterization [47–49].

Briefly, vegetable samples weighing between 350 and 900 g each were manually washed in large volumes of distilled water (between 10 and 80 L) (Table 1). A 1 µm Filtamax[®] filter (IDEXX, Westbrook, ME, USA) applied to a peristaltic pump at a pressure of three bar was used for washing water filtration. Elution was performed in a Filtamax[®] manual wash station and concentrated into 3 mL of phosphate-buffered solution (PBS) with 0.01% of Tween 20 (Merck KGaA, Darmstadt, Germany), after centrifugation at 1000× *g* for 10 min at room temperature. Magnetic beads conjugated with specific antibodies (Dynabeads[™] GC-Combo; Thermo Fisher Scientific, Waltham, MA, USA) were added to the concentrate, and potential magnetized *Cryptosporidium* oocysts and *Giardia* cysts were recovered from the extraneous material using a magnet. Fluorescein isothiocyanate (FITC) conjugated anti-*Cryptosporidium* spp. and anti-*Giardia* spp. monoclonal antibodies were used according to the instructions of the manufacturer, Aqua-Glo[™]G/C (Waterborne, Inc., New Orleans, LA, USA), on slides containing the potential samples' parasites, and observed by epifluorescence microscopy at 200× magnification. The total number of (oo)cysts per slide/sample was screened by two different microscopists to cross-check the results.

DNA was extracted from slides of positive samples containing two or more *Cryptosporidium* oocysts and *Giardia* cysts. Genomic DNA was extracted by scraping the slides to collect the (oo)cysts using the QIAamp[®] DNA Mini Kit (Qiagen GmbH, Hilden, Germany) according to the manufacturer's instructions and as described previously by Almeida et al. [46]. Two-step nested PCR was performed to amplify a portion of the small subunit rRNA gene of *Cryptosporidium* with primers and PCR conditions described by Xiao et al. [50]. For *G. duodenalis*, β-giardin was the gene chosen for amplification by semi-nested PCR [51]. Positive amplification products were purified using the GRS PCR & Gel Band Purification kit (GRiSP Research Solutions, Porto, Portugal) according to the manufacturer's instructions and sequenced using Sanger sequencing services from GATC Biotech (Eurofins Genomics, Ebersberg, Germany). Sequence comparison was made with already published sequences using the NCBI Basic Local Alignment Search Tool (BLAST).

For bacterial analysis, samples were transported to the laboratory in portable, insulated cold-bags and stored at 4 °C until analysis, normally between 1 and 2 days after collection. Twenty-five grams of each sample was added to 225 mL of sterile buffered peptone water (Biokar Diagnostics, Beauvais, France), and homogenized in a stomacher (Interscience, Saint Nom la Brèche, France) for 2 min. Appropriate decimal dilutions were prepared in Ringer's solution (Biokar Diagnostics) for microbial enumeration: colony counts at 30 °C on Plate Count Agar according to ISO 4833-1:2013 [52]; *Enterobacteriaceae* on RAPID[®] Enterobacteriaceae medium (Bio-Rad, Hercules, CA, USA; ISO 16140 [53], 2016, ISO 21528-2, 2017 [54]); *E. coli* according to ISO 16649-2:2001 [55]; *L. monocytogenes* according to ISO 11290-2:2017 [56]; and yeasts and molds according to ISO 21527-1:2008 [57]. Detection of *L. monocytogenes* (ISO 11290-1:2017 [58]), *Salmonella* spp. (ISO 6579-1:2017 [59]), and *E. coli* was also performed. After appropriate incubation, colonies were counted and/or confirmatory tests performed and the colony forming units (CFU)/g calculated.

3. Results and Discussion

Between one and four *G. duodenalis* fluorescent cysts (Figure 1A) were observed in microscopic slides of four of the 16 samples (25.0%; 95% CI: 9.7 to 49.9%) (Table 2). The same percentage (25.0%) was obtained for the number of *Cryptosporidium* spp. (Figure 1B) oocysts (between one and eight oocysts) (Table 2); however, none of the *Cryptosporidium*-positive slides were confirmed by PCR. On the contrary, three of the four *Giardia*-positive slides were also confirmed by PCR. β-giardin fragment sequencing analysis revealed the presence of *G. duodenalis* assemblage A having more than 98% homology with *G. duodenalis* assemblage A isolates available in the Genebank database (KJ668152.1, KP687765.1, KF963547.1, EU642897.1, EU200934.1). The determination of genotypes is important for

the identification of risk to human health, and potentially, the source of contamination. Assemblage A is one of the genotypes that undeniably causes human infections [24].

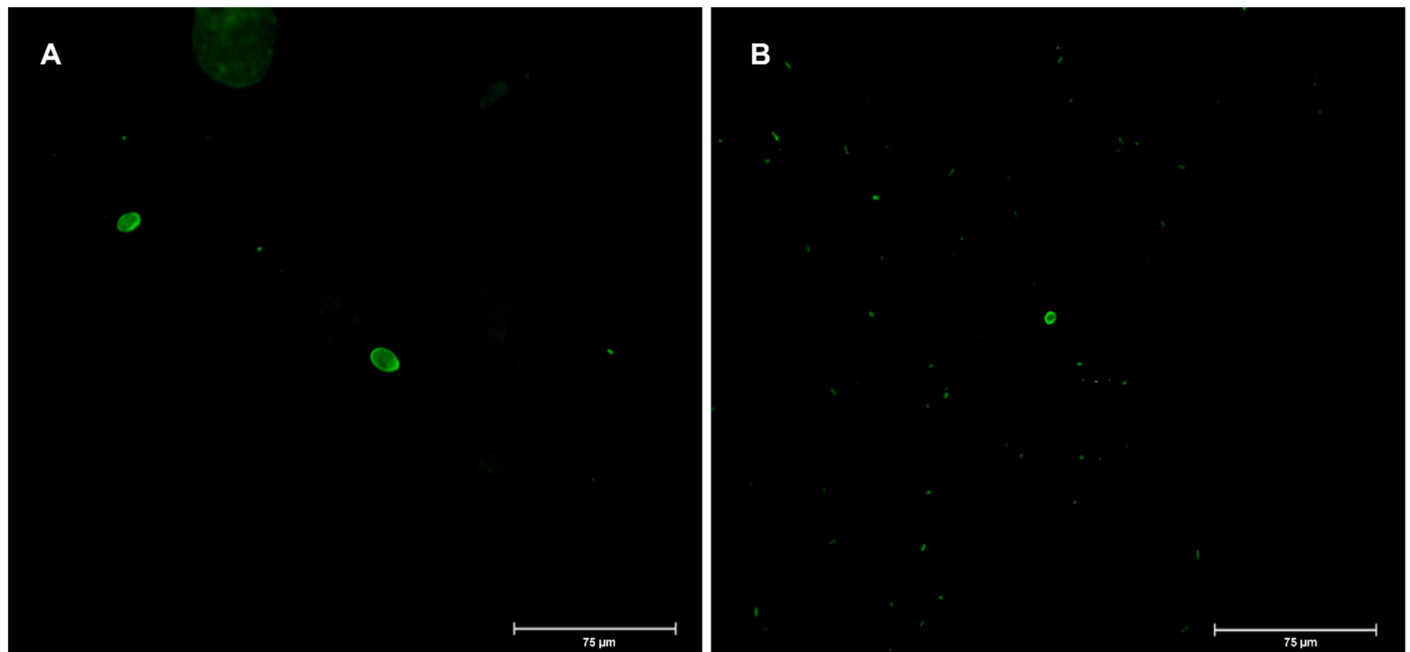


Figure 1. Epifluorescence microscopy images (400× magnification) of *Giardia* cysts (A) and *Cryptosporidium* oocysts (B) found in sample 12.

Table 2. Protozoa and bacteria positive samples.

Sample	Product	Product Presentation	No. Cysts <i>Giardia</i>	PCR <i>Giardia</i>	No. (oo)cysts <i>Cryptosporidium</i>	PCR <i>Cryptosporidium</i>	Genotyping (Assemblage)		
2	Coriander	bulk	2	Negative	0	n.a.	n.a.		
8	Watercress	RTE	1	Positive	0	n.a.	A		
9	Mixed salad	RTE	1	Positive	0	n.a.	A		
12	Mixed salad	RTE	4	Positive	8	Negative	A		
13	Mixed salad	RTE	0	n.a.	1	Negative	n.a.		
15	Arugula	RTE	0	n.a.	1	Negative	n.a.	LMO Present	LMO, detection Positive in 25 g
16	Lettuce	bulk	0	n.a.	2	Negative	n.a.	< 4.0 × 10 ¹	

RTE: ready-to-eat; n.a.: not applicable; LMO: *Listeria monocytogenes*.

Globally, significant variations in the prevalence of these parasites have been observed in vegetables, ranging between 0% and 53% for *G. duodenalis* and between 0% and 63% for *Cryptosporidium* spp. (reviewed by Berrouch et al. [60]). Several factors can easily explain this discrepancy, such as different sampling procedures and detection methods having different recovery efficiencies and detection limits, distinct geographic locations having variable seasonal climatic conditions, and the degree of local development. The type of vegetable, in addition to the quality of water used for irrigation and washing, the proximity of livestock, and the presence of organic fertilizers, are also extremely relevant [60].

Unexpectedly, positive samples were only observed for the RTE type of produce. Thus, it is acceptable to discuss the management of the washing systems used in the food industry, even based on these preliminary data. Indeed, Trevisan et al. [61] noted that the use of treated wastewater for irrigation increases the chance of contamination of RTE/packaged vegetables. Other studies mention that contamination of RTE products by food handlers

during packing is also a transmission route of *Cryptosporidium* and *Giardia* [61–63]. Therefore, it is clear that current RTE production treatment does not guarantee that fresh produce will be free from protozoa, and that alternative water treatments and increased hygienic measures for food handlers are necessary to ensure the safety of RTE vegetables.

Fresh produce has repeatedly been implicated in foodborne outbreaks and recalls, with *Salmonella* spp., *E. coli*, and *L. monocytogenes* the most implicated bacteria [64]. However, with the exception of a lettuce sample (sample no. 16) in which *L. monocytogenes* was detected, but at levels below 4×10 CFU/g (Table 2), all the sampled products were considered safe based on the microbiological criteria set by the European Commission Regulation (EC) No. 2073/2005 for “Pre-cut fruit and vegetables (ready-to-eat)”, i.e., the absence of *Salmonella* spp. and *L. monocytogenes* in 25 g samples and counts of *E. coli* not exceeding 1000 CFU/g. This is in agreement with previous studies, which demonstrated that the prevalence of these pathogens in this type of product is generally low [65–67]. Although the sample that tested positive for *L. monocytogenes* was unwashed lettuce, “Even if salad leaves and herbs are labelled with an instruction to wash before consumption, they are considered ready-to-eat. Although washing can reduce microbial contamination on the plant’s surface, it is not effective to eliminate microorganisms of concern or reduce them to an acceptable level because some pathogens can become internalised within the plant’s tissue.” [68].

The other microbiological parameters are recognized as hygiene and spoilage indicator micro-organisms in RTE foods. Most of the parameter results fell within the Satisfactory-Questionable categories according to the values recommended by the Portuguese National Health Institute [69]. Yeasts and molds were exceptions, with some products being classified as “Not Satisfactory” (yeast and mold counts exceeding, respectively, 10^6 CFU/g and 10^3 CFU/g) (Supplementary Table S1).

The knowledge gained in this field during the course of SafeConsume WP2 highlighted the need to include protozoa in the microbiological criteria for foodstuffs, as required by EU Law No. 1441/2007. This is particularly the case of *T. gondii* and the enteric pathogens *G. duodenalis*, *Cryptosporidium* spp., and *C. cayetanensis*, which are the most common protozoa associated with foodborne illness outbreaks. The absence of fecal contamination in vegetables and the guarantee of food safety for consumers can definitely no longer be defined only by the analysis of detection limits for bacteria, i.e., *E. coli*, *L. monocytogenes*, and *Salmonella* spp. However, microbial risk assessment evaluation and definition of strategies for the management and control of these parasites requires total collaboration between academia and industry, and investments by major companies and governments. These efforts are required to support and foster epidemiological and surveillance studies, and for the development of new technologies and “golden standard procedures” for protozoan removal (or recovery) and detection along the food chain. Monitoring of irrigation and processing water and the efficiency of wastewater treatment plants should be regularly performed. In addition, analysis of soil and manure used as fertilizer should be frequently undertaken, and animal access to crops and personal hygiene regulations applied to food handlers should be controlled. For instance, in addition to the use of sodium hypochlorite as a disinfectant, improved, safer methods, such as UV treatment, treatment with chlorine dioxide or ozone, membrane filtration, and a multi-barrier approach could be used as treatments for (oo)cyst inactivation and removal [33,37,70]. Furthermore, improvement in tools used for genotyping is necessary for the rapid and reliable detection of protozoans of concern for human health and to clarify their transmission patterns. The development of accurate methods for determining the viability and infectivity of (oo)cysts would also be invaluable in surveillance and outbreak studies, and for estimating risk to consumers. Finally, controlling foodborne (and waterborne) diseases should always be accomplished from a collaborative One Health perspective, involving a multidisciplinary team of public health and food experts, epidemiologists, veterinarians, environmental scientists, microbiologists (bacteriologists, virologists, and parasitologists), farmers, industry managers, lawmakers, and media communication specialists.

4. Conclusions

Although preliminary, the findings of the present study highlight the importance of implementing strategies for the detection, inactivation, and removal of parasite (oo)cysts in fresh produce and water, at all stages of food production, including pre- and post-harvest processes. The consumption of raw and/or undercooked vegetables, and particularly salads, has increased due to changes in the eating habits of consumers, who are switching to healthier diets. As a result, there is an urgent need to include protozoa in the microbiological criteria of EU regulations relating to foodstuffs. In addition, it is also important to make consumers aware of hygiene measures at home when washing and cooking raw vegetables.

Supplementary Materials: The following supporting information can be downloaded at: <https://www.mdpi.com/article/10.3390/app12147145/s1>.

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