

Communication

Preliminary Evaluation of Enteric Viruses in Bottled Mineral Water Commercialized in Brazil

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Abstract: In Brazil, mineral water is widely consumed and as of yet there have been no studies done in the country that aim to detect enteric viruses in this water source. The aim of this study was to analyze the presence of the human adenovirus (HAdV), the human rotavirus genogroup A (GARV) and the human enterovirus (hEV) in mineral water samples from six different brands that are commercialized in southern Brazil, using molecular techniques and comparing the results with bacterial indicators. Samples of 1.5 L and 500 mL were analyzed for viruses through PCR and total and thermotolerant coliforms. Additionally, heterotrophic bacteria were assayed using a commercial kit. The most prevalent virus was adenovirus (32.5%) followed by rotavirus (25%) and enterovirus (17.5%). Total and thermotolerant coliforms were absent in all samples and only three samples out of the 60 analyzed presented heterotrophic bacteria contamination. We conclude that, following the example taken into consideration regarding the public supply of drinking water, stricter measures for microbiological control should also be applied to mineral water so that this actually becomes a safer alternative.

Keywords: enteric viruses; HAdV; hEV; GARV; mineral water

1. Introduction

Water quality is essential for human health and should be safely available for the population. Brazil is the sixth largest consumer of mineral bottled water worldwide, with a per capita consumption of 13 L/year (ABINAM—Brazilian Association of Mineral Water Industries) [1]. This may be due to the nation's increasing standards of living and the population's general perception that bottled mineral water is the healthier and safer option for consumption when compared with other sources. Queiroz *et al.* (2013) [2] suggested that bottled water might have the same or worse bacteriological and chemical quality as tap water, concluding that bottled water is not better than tap water when considering health benefits. Santana *et al.* (2003) [3] have demonstrated that up to 25% of commercialized mineral water studied in Brazil did not meet standard regulation. The presence of microbial contaminants depends on source characteristics, bottling methods, asepsis of the packaging material and product storage. The problem is exacerbated by the fact that 64% of mineral water in Brazil is distributed to the final consumer in reusable containers with a 20 L capacity or more (ABINAM) [1]. In Brazil, ordinance No. 274/2005 [4] from the Ministry of Health regulates the microbial quality of natural mineral water and requires that only fecal coliforms and thermotolerant bacteria be tested for. Among the enteric viruses, three of the ones that are most studied as environmental contaminants are the adenoviruses (HAdV, Adenoviridae family, *Mastadenovirus* genus, double-stranded DNA), enteroviruses (hEV, Picornaviridae family, *Enterovirus* genus, single-stranded RNA) and rotaviruses (GARV, Reoviridae family, *Rotavirus* genus) (Fong and Lipp 2005 [5]). These agents are often transmitted by the fecal-oral route, which is associated with gastroenteritis, especially in human and animals (Ahmad *et al.* 2009 [6]). Despite the health impacts caused by outbreaks of these viruses, monitoring and reporting their presence in drinking water, sewage effluents, and recreational waters is not mandatory in many countries, including Brazil. Studies in Brazil aimed at detecting enteric viruses in mineral water have not been carried out yet, even though the presence of these microorganisms in this product has been already reported (Gassilloud *et al.* 2003 [7]; Leclerc and Moreau 2002 [8]). In this context, the goal of the current study was to examine mineral water for the presence of the enteric viruses HAdV, hEV and GARV in mineral water and the possible sources of transmission.

2. Material and Methods

2.1. Mineral Waters

Six Brazilian brands of still water, bottled in polyethylene bottles, were selected for this study. Forty (40) samples of 1.5 L and 500 mL bottles (20 from each volume) of each brand were bought at a local grocery store. An additional twenty (20) mineral water samples were collected aseptically in 500 mL sterile glass bottles from 20 L reusable containers from 5 different departments located at the Feevale University campus during the months of March, June, September and November of 2011. The water containers were available for students and staff and were all from the same supplier. All sampled waters were still within expiration date. According to Brazilian legislation—RDC No. 274/2005 [4]—one sample from every five samples from the same lot (according to labels) was chosen for analysis. Samples were stored in darkness at 4 °C for no more than two days.

2.2. Sample Concentration

Mineral water samples (500 mL) were concentrated using an adsorption-elution method with negatively charged membranes (HA, Millipore, USA), based on the method from Katayama *et al.* (2002) [9] with modifications (Vecchia *et al.*, 2012 [10]). The resulting concentrates were stored at $-80\text{ }^{\circ}\text{C}$ until further processing was done.

2.3. Viral Nucleic Acid Extraction

Viral nucleic acids were extracted from 400 μL of the concentrated sample using the RTP[®] DNA/RNA Virus Mini Kit (Invitex, Berlin, Germany), according to the manufacturer's instructions. The nucleic acids obtained were kept at $-80\text{ }^{\circ}\text{C}$ until further analysis was done.

2.4. Polymerase Chain Reaction Assays for Detecting Virus Genomes

In order to amplify the hEV and GARV genomes, a prior cDNA synthesis step was carried out using the High Capacity cDNA Reverse Transcription[™] commercial kit (Applied Biosystems[™], Carlsbad, CA, USA), with random primers and RNase Inhibitor (Applied Biosciences[™], USA), following the manufacturer's instructions. All PCR reactions were performed as previously described (de Oliveira *et al.* 2012 [11]; Dalla Vecchia *et al.* 2015 [12]) using primers designed to anneal in the highly conserved regions of each viral genome. After the reactions, PCR products were stained with a nontoxic fluorescent dye, SYBR[®] SAFE DNA Gel Stain (Invitrogen[™], Waltham, MA, USA), analyzed by electrophoresis on 2% (w/v) agarose gel and visualized under ultraviolet (UV) light. To determine the analytical sensitivity of the assays, 10-fold serial dilutions of each DNA/RNA standard were employed. EV-PCR has shown to detect a minimum of 0.316 tissue culture infective doses (TCID₅₀) of experimentally contaminated water. For the AdV values were 6.2×10^1 GC (qPCR) and 0.562 TCID₅₀ (PCR). For GARV, the amount of DNA used for amplification was measured by comparison with Low Mass DNA ladder (Invitrogen[™], USA) and the detection limit determined as 200 ng per sample.

Wolf *et al.* (2010) [13] have described the primers used for HAdV quantification by qPCR and Dalla Vecchia *et al.* (2015) [12] have standardized and described the reaction. The qPCR was carried out in iQ5 Real-Time PCR Detection System thermocycler (Bio-Rad Laboratories, Hercules, CA, USA) with iQ[™]5 optical system software, version 2.1. For generating standard curves, 10-fold serial dilutions of standard controls from 10^{-1} to 10^{-5} were prepared, starting at 6.01×10^7 genome copy (gc) per reaction (HAdV-5) and with all standard controls and samples being run in duplicates. There were no template controls (NTC) or negative controls (virus-free water) used in the runs so as to ensure that the assay was not contaminated.

2.5. Bacterial Analysis

For counting the thermotolerant coliforms and total heterotrophic bacteria from the samples, we used the commercial kit AQUACULT M (Laborclin[®], Brazil), following manufacturer's instructions, as per Resolution RDC No. 274/2005 [4] of the Brazilian Ministry of Health (MS).

3. Results

Analyses were performed on samples of mineral waters from 0.5 L and 1.5 L bottles, and 20 L containers. Out of the 40 samples of bottled mineral waters analyzed by conventional PCR, at least one virus was positively detected per sample (Table 1), of which 38.2% were positive for HAdV, followed by 25% for GARV and 17.5% for hEV. In comparison with conventional PCR, the analysis of HAdV by qPCR demonstrated a higher number of positive samples (38.2% *versus* 92.3%), with an average quantification ranging from 7.39×10^3 to 5.78×10^4 gc/L (Figure 1).

Table 1. Detection of GARV, hEV and HAdV in bottled mineral water (0.5 and 1.5 L), by conventional PCR and qPCR.

Mineral Water Brands	GARV PCR	hEV PCR	HAdV PCR	HAdV qPCR
	Positive %	Positive %	Positive %	Positive %
Brand A	80 (8/10)	0 (0/10)	10 (1/10)	90 (9/10)
Brand B	10 (1/10)	0 (0/10)	40 (2/5)	80 (8/10)
Brand C	0 (0/5)	20 (1/5)	40 (2/5)	100 (5/5)
Brand D	20 (1/5)	40 (2/5)	80 (4/5)	100 (5/5)
Brand E	0 (0/5)	80 (4/5)	20 (1/5)	100 (5/5)
Brand F	0 (0/5)	0 (0/5)	75 (3/4)	100 (4/4)
Total	25 (10/40)	17.5 (7/40)	38.2 (13/34)	92.3 (36/39)

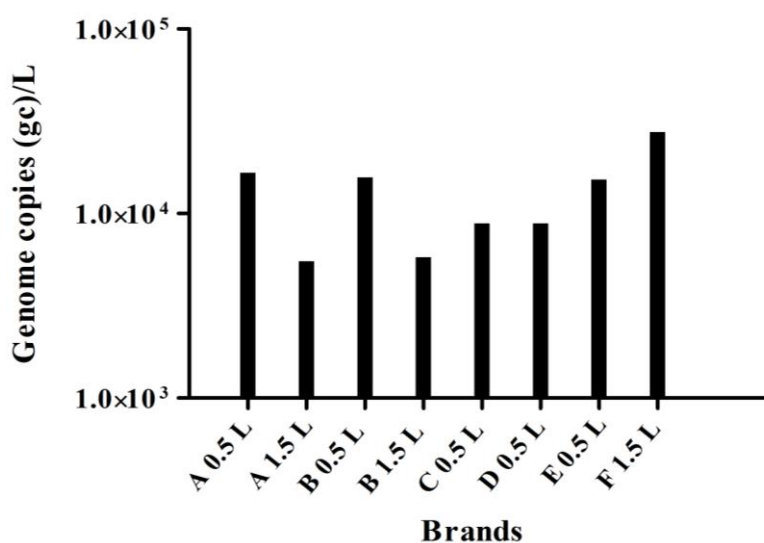


Figure 1. The average HAdV gc/L detected in mineral water samples (in 0.5 L and 1.5 L bottles from the same brand). Values represent the average of four samples of each brand (A–F) and volume.

The analyses of samples collected from 20 L containers ($n = 20$) showed that 100% were negative for hEV and GARV genomes by conventional PCR and 100% were positive for HAdV by qPCR, reaching values from 1.34×10^3 to 9.94×10^4 gc/L (Table 2). HAdV by conventional PCR was not tested in these samples.

Table 2. Number of HAdV gc/L in mineral water collected from 20 L containers over a period of four months in 2011.

Locations	Month/Year				Geometric Average
	March/11	June/11	September/11	November/11	
1 (n = 4)	1.18×10^4	8.71×10^3	6.08×10^4	2.07×10^4	2.55×10^4
2 (n = 4)	3.72×10^4	1.34×10^3	5.71×10^4	3.36×10^4	3.23×10^4
3 (n = 4)	2.89×10^3	6.54×10^4	3.13×10^4	2.71×10^4	3.10×10^4
4 (n = 4)	9.05×10^3	1.20×10^3	6.27×10^3	9.97×10^4	2.88×10^4
5 (n = 4)	7.60×10^3	5.74×10^3	6.00×10^3	8.07×10^4	4.30×10^4

4. Discussion

According to Brazilian regulation of microbiological analysis, all water samples must present absence of *Escherichia coli*, and thermotolerant coliforms in 100 mL. However, there is some tolerance for total coliforms, *Enterococci*, *Pseudomonas aeruginosa*, sulfite-reducing Clostridia or *Clostridium perfringens*, in a contaminated sample. This study followed the parameter regarding the number of 500 mL and 1.5 L mineral water samples collected from the same lot; total and thermotolerant coliforms were absent in the mineral water samples, although three samples were positive for heterotrophic bacteria (one 1.5 L bottle from brand F and two 20 L containers, locations 1 and 5). In previous studies in Brazil, Resende (2008) [14] and Ritter (2009) [15] analyzed bottled mineral water for the presence of coliforms and none of the samples were inconsistent with current regulations. Bacteriological analysis of bottled water has been evaluated in several studies nationwide and these findings indicate that contamination is above the levels established by regulations (Santana *et al.*, 2003 [3]). Silva *et al.* (2008) [16] compared the public water supply and mineral water collected from 20 L water containers and concluded that the public water supply had better microbiological quality than the bottled water sold in 20 L containers. Several studies worldwide have also demonstrated the presence of *Cryptosporidium*, *Pseudomonas*, *Aeromonas*, *Legionella*, *Penicillium*, enteric viruses and pesticides, among others in mineral water samples (Rosa *et al.* 2008 [17]; Zeenat *et al.* 2009 [18]; Pontara *et al.* 2011 [19]; Beuret *et al.* 2002 [20]). In this study, we demonstrated that enteric viruses were present in six commercialized brands in Brazil, even though Brazilian legislation does not require this evaluation. In our findings, at least one virus evaluated was present per brand of bottled mineral water, with HAdV being the most prevalent, reaching a positivity rate of 92.3% (qPCR; 7.39×10^3 to 5.78×10^4 gc/L). This virus was also present in 100% of the mineral water containers (20 L), with viral loads ranging from 1.34×10^3 to 9.94×10^4 gc/L. HAdV is one of the main etiological agents of gastroenteritis in children under 4 years (Lee and Kim 2002 [21]) and was included in the U.S. Environmental Protection Agency's (USEPA) "Contaminant Candidate List 2" for its sanitary significance and its elevated incidence in water sources and sewage samples (Xagorarakis *et al.* 2007 [22]). The frequency and viral loads found in the present study were higher when compared to previous studies. The occurrence of HAdV by molecular methods in finished drinking water and tap water has been reported worldwide: 39.1% in South Korea (Lee and Kim 2002 [21]), 52% in California (USA) (Jiang and Chu 2004 [23]), 12.9% in Benin (Verheyen *et al.* 2009 [24]) and 66% in South Brazil (Rigotto *et al.* 2010 [25]). The high rates of contaminated mineral water may be a reflection of the overall poor quality of Brazilian groundwater bodies, since the source of water for the commercial brands studied are all from wells and springs located in the southern region of Brazil. In a previous study from

our group, De Oliveira *et al.* (2012) [11] found HAdV in up to 23.2% of water samples from dairy farms, and the majority of the positive samples were from natural springs and groundwater. In Brazil, Rigotto *et al.* (2010) [25] evaluated chlorinated drinking water and ICC-PCR indicated that 50% (54/84) were found to be infected with HAdVs. Such high frequency of infectious adenovirus detection in drinking water from these regions is alarming. Studies based on risk analysis, using an adenovirus occurrence rate of 1/100–1/1000 L in drinking water, indicate an illness rate between 8.3/1000 and 8.3/10000 in drinking water (Jiang 2006 [26]).

In the samples analyzed, there was also incidence of GARV (25%, 10/40) in the six brands of bottled water used in the study, with the GARV genome present in three (50%) and absent in the 20 L containers. GARV has been considered to be the major etiological agent of childhood diarrhea in the world, and it has been related to 527,000 deaths annually, mostly in developing countries (Racaniello 2010 [27]). With regard to hEV detection, this virus was present in low frequency (17.5%) when compared to HAdV and GARV. Several authors have reported that simultaneous contamination of water by different enteric viruses is often found (Brassard *et al.* 2005 [28]; Gilgen *et al.* 1997 [29]; Fong and Lipp 2005 [5]). In this work, among the six brands surveyed, only one did not present simultaneous contamination (brand F). Brands A and B were contaminated with GARV and HAdV, brand C and E with hEV and HAdV and brand D presented the genome for the three enteric viruses (hEV, GARV and HAdV).

There is a large sanitation deficit in Brazil and only 48.7% of the population has sewage collection and only 37.9% of the sewage is treated, ranking Brazil 112th in Sanitation Development Index, which includes 200 countries (SNIS, 2014 [30]). This scenario generates a great impact on the environment due to the large amounts of sewage discharged into the environment without treatment, affecting surface and groundwater sources for public supply and mineral water exploitation. Studies performed in Brazil have already demonstrated the importance of monitoring bottled water quality, however, without any information regarding viral presence (Castro *et al.* 2010 [31]; Santana *et al.* 2003 [3]; Silva *et al.* 2008 [16]; Resende 2008 [14]). Another important aspect to be considered is the fact that the Brazilian regulations serve as references for published research, so the safety parameters must be expanded, thus preserving the health of the population. Until now, there have been very few studies worldwide evaluating viral contaminants in bottled water, even though the scientific community has made great efforts to highlight the significance of these agents regarding the risks populations are exposed to when consuming this product. Product storage is another issue to be discussed, since enteric viruses can easily adhere to polyethylene terephthalate (PET) bottle walls and glass, as shown in a study that intentionally contaminated mineral water in order to evaluate absorption of these agents in PET bottles and glass (Butot *et al.* 2007 [32]). Regarding water treatments applied to minimize contamination, those are based on chemical (chlorination, ozonation) or physical methods (high temperature). However, according to Brazilian legislation, mineral water cannot receive any disinfection treatment.

Given the above, we consider that bottled water may present potentially questionable microbiological qualities, thus exposing the public to health risks when consumed. We conclude that, in order to minimize disease outbreaks, monitoring mineral water quality is important and relevant, not only with regard to the presence of bacteria, but also the presence of enteric viruses. Due to the increased consumption of bottled water, new standards should be established in order to protect the consumer from waterborne diseases. For most people, mineral water is considered pure and safe and its consumption has grown, benefiting from the fact that there is mistrust of treated water provided by supply systems.

Silva *et al.* (2008) [16] has pointed out that there is a cultural tendency to opt for mineral water consumption, to the expense of public water system, because the population perceives groundwater to be of higher quality. Our findings demonstrate severe human fecal origin contamination. Nevertheless, it was not possible to trace the origin of the contamination within the scope of the current study, but we suggest that the contamination may have occurred at the source (water catchments) or from the product's processing or packaging (bottle).

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Author Contributions

VRS conducted water sampling, concentration and molecular detections, and wrote first draft of the manuscript, CR has compiled all data, wrote and prepared the final version of the manuscript, RS and ADV participated in standardization of molecular techniques and performed molecular assays, AH participated in writing of the manuscript, FRS was responsible of experimental design, general coordination and revised the manuscript.

Conflicts of Interest

The authors declare that there is no conflict of interest.

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