A Collaborative Effort to Identify the Causative Agents of Two Waterborne Outbreaks of Gastroenteritis in Wyoming

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Background

- Caliciviruses are included on EPA's Contaminant Candidate List because they are a major cause of viral gastroenteritis in the U.S. and worldwide (Table 1).
- The caliciviruses most commonly associated with human disease are the noroviruses (Fig 1).
- The potential for widespread outbreaks caused by the large, diverse group of noroviruses in drinking water is a public health concern as shown by numerous outbreaks that have occurred in municipal and private water systems.
- This collaborative study illustrates the usefulness of the methods used to isolate and identify noroviruses from outbreak associated water.

Methods

- Epidemiologic and Environmental Investigation by CDC, EPA, and the Wyoming Department of Health
 - To determine sources of illness and to examine food processing and groundwater well conditions.
- Water Sample and Stool Sample Processing
 - To examine for the presence of coliforms and viruses.
- Molecular Methods
 - To use RT-PCR analysis and sequencing for the detection of a norovirus gene.

Table 1. Drinking Water Contaminate Candidate List Microbiological Contaminants

Acanthamoeba Adenoviruses Aeromonas hydrophila Caliciviruses Coxsackieviruses Cyanobacteria (blue-green algae), other freshwater algae, and their toxins Echoviruses Helicobacter pylori Microsporidia (Enterocyotzoon and Encephalitozoon) Mycobacterium avium intracellulare (MAC)



Figure 1: Norwalk Virus

Results

Wyoming Snowmobile Lodges

- 41% (22/54) of persons from Lodge A and 48% (13/27) of persons from Lodge B developed gastroenteritis symptoms between January and March, 2001 (Fig. 2).
- For lodge A, illness was significantly associated with drinking water.
- For lodge B, guests who ate or drank at Lodge A had a greater risk of disease than those who did not.
- Environmental investigation of the groundwater wells at Lodge A revealed three on-site wells that were within 92-115 feet of a septic tank or outhouse.
- Environmental investigation of the septic system revealed that the system was not designed to effectively treat the volume of wastewater that it received.
- 7/8 well water samples collected at Lodge A were positive for fecal coliforms. Well water samples from Lodge B were negative.
- RT-PCR analysis showed the lodge A well water sample (Figure 3) and stool specimens had caliciviruses.

Wyoming Tourist Saloon

- 76% (84/111) of persons from the tourist saloon developed gastroenteritis symptoms during September to October, 2001 (Fig. 4).
- Illness was significantly associated with drinking water and/or consuming ice.
- Environmental investigation of the groundwater well revealed that it was situated 50 feet from the saloon's septic tank, 50 feet from the leach field of a separate septic system, and 100 feet from an effluent disposal of an RV park.
- Environmental investigation of the tourist saloon's septic system revealed perforations in the septic tank.
- 5/6 well water samples were positive for fecal coliforms; all six samples were positive for total coliforms.
- RT-PCR analysis showed that the saloon well water sample (Figure 3) and three stool samples had caliciviruses.
- Gene sequencing analysis revealed that the isolates from the lodge well water sample and stool specimens were a strain that belongs to a common genetic group of noroviruses.
- Gene sequencing analysis revealed that the isolates from the saloon water sample and one of three stool samples were a strain that belonged to a common genetic group of noroviruses. One stool sample differed genetically by three point mutations, and the other belonged to a different genetic group of noroviruses.



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