NOROVIRUS GASTROENTERITIS CAUSES SEVERE AND LETHAL COMPLICATIONS AFTER CHEMOTHERAPY AND HEMATOPOIETIC STEM CELL TRANSPLANTATION

Short title: Norovirus enteritis in immunocompromised patients

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scientific heading: Clinical Observations, Interventions, and Therapeutic Trials
Abstract

Norovirus (NV) infections are a frequent cause of gastroenteritis (GE), but data on this disease in immunocompromised patients are limited. We analysed an NV-outbreak, which affected immunosuppressed patients in the context of chemotherapy or hematopoietic stem cell transplantation (HSCT). Upon recognition, 7 days post admission of the index patient, preventive measures were implemented. Attack rates were only 3% (11/334) and 10% (11/105) among patients and staff members, respectively. The median duration of symptoms was 7 days in patients compared to only 3 days in staff members (p=0.02). Three patients died of the NV infection. Commonly used clinical diagnostic criteria (Kaplan-criteria) were unsuitable, because they applied to 11 patients with proven NV-GE but also to 15 patients without NV-GE. With respect to the therapeutic management, it is important to differentiate intestinal graft-versus-host disease (GvHD) from NV-GE. Therefore, we analysed the histopathological patterns in duodenal biopsies, which were distinctive in both conditions. Stool specimens in patients remained positive for NV-RNA for a median of 30 days, but no transmission was observed beyond an asymptomatic interval of 48 hours. NV-GE is a major threat to patients with chemotherapy or HSCT and meticulous measures are warranted to prevent transmission of NV to these patients.
Introduction

Infections with norovirus (NV) are a dominating cause of gastroenteritis (GE) worldwide with great impact on public health. The clinical disease and its association with the winter season has been first described in 1929. In 1945, the transmissibility of NV-GE from patients to healthy volunteers was recognised, but it was not until 1972 when virus particles were discovered in fecal specimens obtained during an outbreak in Norwalk, Ohio. Until now, the NV is found to be refractory to replication in cell cultures or animal models, except for a study using a sophisticated, highly differentiated 3-D cell culture model and another study investigating gnotobiotic animals. Thus, the understanding about NV infections is largely derived from volunteer studies and outbreak investigations. Traditionally, outbreaks or cases of NV-GE are diagnosed by clinical criteria (Kaplan-criteria), which include a short 24 - 48 hour incubation period, ≥50% of affected individuals with vomiting, an average duration of illness of only 12 - 60 hours, negative stool cultures for bacteria and parasites for outbreaks or ≥2 episodes of vomiting and/or ≥3 episodes of diarrhea for case defining criteria. Detection of NV-specific RNA in stool specimens by PCR is currently considered the standard to confirm the diagnosis of NV-GE.

Outbreaks of NV-GE display notoriously high attack rates, which is at least in part explained by a very low infectious dose (<20 viral particles) and easy transmission not only by contaminated food or fecal-oral route, but also through
infectious aerosols, which could be inhaled during nearby vomiting.\textsuperscript{8,9} Attack rates exceeding 50\% have been reported during NV-GE outbreaks, resulting in ward closures and major disruption of hospital services.\textsuperscript{10,11} Symptoms of NV-GE typically appear with abrupt onset and are mostly limited to 1 - 2 days in healthy individuals, but surveillance data indicate that the duration of symptoms is prolonged in hospitalized patients.\textsuperscript{6,12} In addition, a growing number of reports suggest that NV-GE could result in severe and, occasionally, life-threatening complications in immunocompromised patients associated with prolonged viral shedding.\textsuperscript{13-17} However, clinical and virological characteristics of otherwise healthy individuals and immunocompromised patients have yet not been compared during a single outbreak caused by a particular NV strain.

During an outbreak of GE in our hematological and transplantation unit, which was caused by the predominant NV genotype II.4 variant strain, we recorded the pattern of disease among patients and staff members, the disease duration and severity, virological surveillance data, the attack rates and outbreak management. Histo-blood group antigens on gastroduodenal epithelium have been identified as receptors for NV.\textsuperscript{18} A low to absent susceptibility to NV infections has been recognised among individuals with particular histo-blood group characteristics, i.e. reduced susceptibility to NV genogroup I in patients with histo-blood group B and resistance against NV genogroup II among nonsecretors.\textsuperscript{19,20} Thus, we evaluated
histo-blood group characteristics of all infected patients, including those with allogeneic hematopoietic stem cell transplantation (HSCT). A subtle onset or prolonged illness in patients with NV infection after allogeneic HSCT may be undistinguishable from gastrointestinal graft-versus-host disease (GvHD). Since both conditions trigger completely different therapeutic consequences (increase of immunosuppression in patients with GvHD versus decrease of immunosuppression in NV infection), we compared the histopathological patterns in biopsies from the gastrointestinal tract in patients with GvHD or NV-GE after allogeneic HSCT.
Patients and Methods

**Hospital structure and staffing patterns.** The outbreak occurred in the department of hematology and oncology of a 1057-bed tertiary center, which consisted in a closed transplantation unit and a 50-bed ward for patients with cancer or hematological diseases, both separated from facilities for general medical patients. None of the patients nursed in the transplant unit were affected, but some patients with autologous HSCT and patients with stable engraftment and without intensified immunosuppressive therapies after allogeneic HSCT, who were treated outside the transplant unit, together with patients with cancer or hematological diseases, acquired NV-GE. Outside the transplant unit, lavatories were separately available in two single rooms but were otherwise shared by patients from two neighboring double rooms.

Except for a single physician during night shift duty and trained cleaning staff, there were no common staff responsibilities in- and outside the transplant unit. However, staff member duties outside the transplant unit included care for patients with cancer, hematological diseases or autologous HSCT as well as stable patients after allogeneic HSCT.

**Hygiene measures and case definitions.** Before this outbreak, standard hygiene practises included reverse isolation of patients with neutropenia or intensified immunosuppressive therapies (e.g., high dose steroids) with the use of
face masks by any visitor, strict avoidance of handshaking, hand disinfection before entering and after leaving any patient room and limited access of visitors, excluding individuals younger than 16 years and visitors with symptoms of infection or with recent contact to another individual with symptoms of infection. In addition, extra gowns, shoes and face masks were used in the transplant unit with a limitation of visitors (usually less than 2) per patient.

Upon recognition of the NV outbreak, any patient or staff member with nausea, vomiting or diarrhea was considered as potentially harboring a NV infection. Patients with suspected NV infection were immediately transferred to an isolation ward and nursed in single rooms. Preventive measures always included the use of surgical face masks, single-use coats, gloves and hand disinfectants with enhanced (75.1%) ethanol content. In individual patients, preventive measures were continued until negative results from stool specimen testing for NV-specific RNA were available. In patients with NV-GE confirmed by PCR testing, isolation and preventive measures were continued for at least 48 hours beyond complete resolution of symptoms of GE. All contact patients were also transferred to the isolation ward and observed in a similar fashion for at least 48 hours with screening for NV RNA in stool specimens. Staff members with symptoms of GE were immediately set off duty until they were without symptoms for at least 48 hours. All individuals with symptoms of GE were classified according to established outbreak and case-defining criteria.\textsuperscript{5,6}
**Data collection and statistical analyses.** Patient and staff member data were collected prospectively. Staff members were requested to complete questionnaire forms about symptoms and clinical signs. In all patients with suspected NV infection, clinical signs and symptoms of GE were recorded. In addition, patient files were reviewed in order to obtain clinical (e.g., underlying diseases, ongoing and previous therapeutic interventions, immunosuppressive therapy or medication) and laboratory data (e.g., neutrophil count, histo-blood group typing, microbiological findings in stool specimens). Patient and staff member data were anonymised and filed after Charité ethical committee approval was obtained.

Categorical and continuous variables were compared by Fisher’s exact and Mann-Whitney test, respectively. Kaplan-Meier plots were also produced and time-dependent variables were analysed using the Log-rank test. Calculations were performed using Prism statistical software (GraphPad Software Inc., San Diego, CA).

**Norovirus detection.** Stool samples were screened for NV-specific RNA by real-time PCR located in the junction between ORF 1 and ORF 2 of the NV genome as previously described. Identification of the NV genotype was done by a nested RT-PCR approach targeting the RNA polymerase gene within the open reading frame 1 with subsequent sequencing of PCR products. Stool specimens from all symptomatic individuals were screened at least once for NV-specific RNA by real-time PCR. To investigate the duration of viral shedding in patients with confirmed
NV-GE, sequential PCR testing of stool specimens was scheduled at 2 week intervals.

**Histo-blood group and secretor status typing.** Phenotyping of ABO histo-blood group characteristics in patients were available from previously performed standard serological testing using red blood cells and serum. In patients with allogeneic HSCT, full information on ABO histo-blood group characteristics of donors and recipients were available. In staff members without prior information on ABO histo-blood group characteristics (e.g., from previous blood donation), ABO phenotyping was done with informed consent using red blood cells and bed-side testing cards containing anti-A and anti-B monoclonal antibodies (Meditro GmbH, Leimen-Gau, Germany).

The secretor gene status in patients was identified by using stored peripheral blood, bone marrow cells or isolated DNA, which were in patients with allogeneic HSCT always obtained prior transplantation. The embryonic epithelial cell line INT-407 harboring FUT2 polymorphism associated with a nonsecretor phenotype served as a control. *FUT2* genotyping that predicts the secretor status was done by PCR with sequence specific priming as described previously.\(^{23}\)
Results

Outbreak description. The index patient was a 55 year-old female with high-risk B-CLL, carrying a del17p13, who had received an allogeneic HSCT seven month prior to admission. Post allogeneic HSCT, this patient experienced acute stage 4 intestinal GvHD and subsequently, protracted intestinal GvHD (1-2 bowel movements per day) requiring chronic immunosuppressive therapy with cyclosporine and methylprednisolone (8mg per day). The current hospital admission was required due to disorientation, which was caused by dehydration and cyclosporine accumulation. Three days after admission, symptoms of GE, i.e. emesis and diarrhea, occurred. Stool specimens were negative for bacterial and parasitic agents and symptoms were first considered to be caused by intestinal GvHD. Within 6 days after admission of the index patient and before the NV outbreak was confirmed, 4 other patients developed vomiting and diarrhea with abrupt onset, which prompted laboratory screening for NV-GE. On day 5 after admission of the index patient, her roommate (secondary patient) developed emesis and diarrhea, after being referred on day 4 to another room. The roommate of this secondary patient (tertiary patient) developed symptoms of GE on day 6. Another two patients nursed in the adjacent room and who shared the lavatory with the secondary and tertiary patient, became symptomatic also on day 6. On day 7, results from PCR testing were available and confirmed presence of a NV outbreak. Overall, the likely modes of transmission were
inhalation of infectious aerosols during nearby vomiting in 5 patients and 5 staff members and fecal-oral through lavatory sharing in 2 patients. The routes of transmission in the remaining 4 patients and 6 staff members remained less clear, but person-to-person transmission via fecal-oral route appeared likely. We were unable to identify obvious breaks in hygiene practices. However, 6 of 11 staff members with NV-GE developed their first symptoms 3 days or later after NV was identified by PCR, which makes at least occasional breaks or less strict adherence to the modified hygiene policies likely.

Further investigations revealed that the index patient was previously treated in a rehabilitation center where an outbreak of NV-GE was recorded, indicating the likely source of the infection. During this outbreak no other hospital facilities recorded cases of NV-GE making a food- or water-borne source of this outbreak from our hospital facilities unlikely.

Upon recognition that this cluster of GE was caused by NV, additional hygiene measures were implemented. Patients with nausea, vomiting or diarrhea and contact patients (roommates or those who had shared lavatories with symptomatic patients) were immediately transferred to an isolation ward and nursed in single rooms until negative results from stool specimen testing for NV-specific RNA were available. In patients with NV-GE confirmed by PCR, isolation and preventive measures were continued for at least 48 hours beyond complete resolution of symptoms. The total
duration of this outbreak, defined by admission of the index patient to the occurrence of the last documented infection, was 54 days (10\textsuperscript{th} April to 3\textsuperscript{rd} June, 2005). During this period, a total of 334 inpatients were treated and 105 staff members were involved into patient care. Thus, the attack rates for patients and staff members were 3\% (11 of 334) and 10\% (11 of 105), respectively (p=0.008).

**Clinical characteristics of patients and staff members.** Overall, 26 patients and 11 staff members experienced vomiting, diarrhea or both during the outbreak. Of these, NV-GE was confirmed by PCR in 11 patients and 11 staff members. Affected staff members were younger and otherwise healthy compared to patients (Table 1). The underlying diseases among the 11 patients with confirmed NV-GE were acute leukemia (n=5) and malignant lymphoma (n=6: B-CLL 3, diffuse large cell lymphoma 2, lymphoblastic B-cell lymphoma 1). A single patient with B-CLL had been treated only with rituximab monotherapy until 4 month prior to the occurrence of NV-GE (Table 1). Polychemotherapy had been administered to 5 patients 4 to 56 days (median: 19 days) prior to the onset of NV-GE. Two patients with autologous HSCT were roommates and experienced vomiting and diarrhea during conditioning BEAM chemotherapy or shortly thereafter (days -1 and -7). Another 3 patients had received allogeneic HSCT 71, 72 and 210 days prior to onset of NV-GE. Overall, 7 patients had neutropenia (neutrophil count <0.5/nL) at the onset of symptoms (Table 1).
There was no difference in ABO types between patients and donors (O 10, A 9, B 2, AB 1). Genotyping for the Secretor status in the 11 patients with confirmed NV-GE revealed always a homozygous Se/Se (n=4) or heterozygous Se/se (n=7) FUT2 polymorphism predicting a Secretor positive phenotype (Table 1).

**Outbreak and case defining criteria.** Clinical outbreak criteria were fulfilled with a ≥50% frequency of vomiting not only in patients with confirmed NV-GE, but also in patients without NV-GE (73% of patients with vomiting in both groups; Table 2). The median duration of illness was longer in patients with confirmed NV-GE (7 days) than the median 12 - 60 hour disease duration proposed by commonly accepted outbreak-defining criteria. Microbiological testing of stool specimens was done in 8 patients with confirmed NV-GE and 11 patients without NV-GE, which gave always negative results except in 4 patients without NV-GE (Table 2). Case-defining criteria, designed for hospital patients, were always fulfilled in patients with confirmed NV-GE. However, these case-defining criteria were also fulfilled in up to 10 of 15 (67%) patients with symptoms of GE not caused by NV (Table 2).

**Symptoms and complications.** Diarrhea and vomiting occurred in the majority of patients and staff members with confirmed NV-GE. However, in 6 patients compared to only 1 staff member fever was recorded (Table 1). The median duration of symptoms in affected patients was significantly longer compared to staff members (7 versus 3 days, p=0.02; Table 1 and Figure 1).
The severity of symptoms in patients varied largely and ranged from subtle and slowly progressive in the index patient to severe in 2 patients with BEAM chemotherapy and autologous HSCT. In one of these latter 2 patients, bowel perforation was suspected due to severe abdominal pain and tenderness. In this particular patient, an abdominal CT-scan disclosed extensive and pronounced edema of the small bowel sparing other parts of the gastrointestinal tract (Figure 2). A similar small bowel edema was seen in CT-series in another patient who developed NV-GE with acute abdomen after high-dose methotrexate-based polychemotherapy. The other patient with BEAM chemotherapy and autologous HSCT suffered from continuous retching, which was refractory to intensified antiemetic therapy. Unfortunately, this patient experienced severe aspiration resulting in hypoxic brain damage and death 11 days after onset of NV-GE. Another 2 patients experienced fatal sepsis 17 and 49 days after onset of symptoms of NV-GE. The first of these latter two patients showed profound neutropenia (neutrophil count: 0.03/nL) after intensified chemotherapy for acute myeloid leukemia and at the onset of NV-GE. Subsequently, this patient experienced fatal sepsis despite broad-spectrum antibacterial and antifungal therapy (repetitive blood cultures were negative). The other patient, who had received allogeneic HSCT 2 month prior to the NV infection, *E. coli* was repeatedly recovered from the blood stream without any
clinical site of infection, suggesting that bacterial translocation from the intestine was promoted by the NV infection.

Serial RT-PCR analyses of stool specimens from patients with confirmed NV-GE was scheduled. Three patients already had negative RT-PCR test results at the first follow-up investigation, 2 patients showed positive RT-PCR test results in serial stool specimens without further follow-up specimens and in 1 patient only a single stool specimen was available. The minimal duration of viral shedding (onset of symptoms to last positive RT-PCR) ranged from 11 to 87 days (median: 30 days). The median interval between onset of symptoms until documented cessation of viral shedding (first negative RT-PCR) was 36 days (range: 9 – 118 days).

**Endoscopic findings.** Endoscopy of the upper gastrointestinal tract was done in 3 patients with allogeneic HSCT and showed a normal appearance of the gastric mucosa in 2 patients and a pangastric erythema in the remaining patient. In all patients, the endoscopic appearance of the duodenal mucosa was unremarkable. Two patients with allogeneic HSCT underwent additional sigmoidoscopy, which showed petechial lesions in one patient (low platelet counts, no biopsy taken) and a normal appearance of the mucosa in the other patient.

**Histopathology findings.** Duodenal biopsies obtained during intestinal GvHD and after onset of the NV infection were available from 3 patients with allogeneic HSCT. A detailed analysis revealed a distinct histopathological pattern during GvHD
and NV-GE. Intestinal GvHD was characterised by a partial loss of epithelial cells, increased numbers of apoptotic crypt cells pronounced at the base of the crypts and increased numbers of CD8+ T-cells within the lamina propria with only focal infiltration of the crypt epithelium (Figure 3 A and B). In contrast, biopsies during NV-GE showed villous blunting, only slightly elevated numbers of apoptotic epithelial cells predominantly at the luminal surface and a massive increase in the numbers of intraepithelial lymphocytes consisting of CD8+ T-cells (Figure 3 C and D).

The gastric biopsies from 2 of these patients showed a mild chronic gastritis with no signs of GvHD and were normal in the remaining patient. The rectosigmoidal biopsies obtained from one patient showed increased numbers of apoptotic crypt cells with an edematous lamina propria consistent with stage I GvHD (data not shown).
Discussion

Norovirus infection is among the most frequent causes of infectious gastroenteritis worldwide and outbreaks could result in major disruption of hospital services. The disease in otherwise healthy individuals is self-limiting and of short duration, but growing numbers of case reports and retrospective analyses indicate that immunocompromised patients with NV-GE experience a prolonged and more severe illness.\textsuperscript{13-17} However, a study documenting the disease severity in adult patients after chemotherapy or allogeneic HSCT in comparison to healthy individuals was missing.

Norovirus GE is very easily transmitted with a very low infectious dose (<20 viral particles), including transmission through inhalation of infectious aerosols during nearby vomiting.\textsuperscript{8} Noteworthy, we recorded a low overall attack rate, which was significantly lower among patients compared to staff members (3\% versus 10\%). These low attack rates contrast results from a recently published analysis, which demonstrated a median >32\% patient attack rate recorded during 21 hospital outbreaks.\textsuperscript{24} This large difference might be explained by the early identification of NV in this study and the more carefully applied standard hygiene procedures established in the care of severely immunocompromised patients. We recorded prolonged viral shedding in patients beyond the resolution of symptoms, which is in line with previous observations.\textsuperscript{13,14,17} However, we did not observe further transmission of NV
from any patient beyond a 48-hour asymptomatic interval, which suggests that transmission from immunocompromised hosts with prolonged shedding is uncommon. All NV infected patients had a Secretor positive phenotype, which expresses the carbohydrate H type 1 known as a target structure for NV binding. Thus, it remains unclear whether severely immunocompromised patients with Secretor negative phenotype display any enhanced susceptibility to infection by NV genogroup II.

In a recently published analysis of 12 patients with allogeneic HSCT and epidemiologically unrelated NV-GE, the median duration of illness was remarkably longer compared to our study (3 months versus 7 days). In this previously published series of patients with allogeneic HSCT, one patient died due to malnutrition after more than one year with unresolved NV-GE compared to 3 patients in our study with NV-related deaths, which occurred within 49 days after onset of symptoms. In our study, only 3 patients had received allogeneic HSCT and the majority of patients acquired the NV infection during chemotherapy-induced neutropenia. It appears therefore likely that T-cell directed immunosuppressive therapies after allogeneic HSCT promote chronic NV infections with prolonged diarrhea and wasting, whereas recent chemotherapy and neutropenia enhance the disease severity and the likelihood of life-threatening complications. In the previously published series of patients with NV-GE after allogeneic HSCT, the median time from
onset of symptoms until the diagnosis was established widely ranged from 0.25 to 6 months\textsuperscript{14}, which illustrates to some extent the diagnostic dilemma in patients with diarrhea after allogeneic HSCT. Diarrhea is a frequent complication after HSCT and might be caused by the conditioning therapy, intestinal GvHD or infectious agents, or could also be drug-related (e.g., mycophenolate mofetil). In a previously published study with HSCT patients, 22 (79\%) of 28 patients with allogeneic HSCT experienced diarrhea sometime after the transplantation procedure. Most diarrheal episodes occurred early (i.e. within 20 days after transplantation) and in none of these episodes an infectious agent was identified.\textsuperscript{25} Four of these 22 allogeneic HSCT recipients experienced diarrhea later than 20 days after transplantation, which was always caused by viral infections (rotavirus 3, adenovirus 1). However, screening did not include NV in this study.\textsuperscript{25}

We found a significantly prolonged duration of symptoms in patients compared to staff members, which clearly confirms findings from previously published, non-comparative reports on patients with various immunocompromising conditions and NV-GE.\textsuperscript{13-17} In this outbreak, the maximum duration of symptoms among patients was limited to 36 days, but profuse diarrhea with wasting lasting up to more than one year has been previously observed in allogeneic HSCT patients with delayed diagnosis of NV-GE.\textsuperscript{14} Importantly, we observed severe to fatal complications, which were clearly attributable or likely caused by the NV infection. Two patients with
BEAM chemotherapy developed symptoms of an acute abdomen with peritonitis, which prompted abdominal CT-scans showing extensive and pronounced bowel wall edema restricted to the small intestine. This radiological pattern is identical to previous observations in a small number of non-immunocompromised patients with NV-GE.\(^{26,27}\) However, BEAM chemotherapy, gastrointestinal CMV infection or GvHD might also cause small bowel edema, but exclusive involvement of the small intestine has been observed infrequently in intestinal CMV infection or GvHD and radiological studies in patients with gastrointestinal complications after BEAM chemotherapy (e.g., colitis and stomatitis) are missing.\(^{28,29}\) Coexistence of NV infection and intestinal GvHD could represent a diagnostic and therapeutic challenge. We found evidence of simultaneous GvHD in large bowel biopsies in a single patient with NV-GE. It appears less likely that the GvHD-like pattern in this particular patient was directly caused by the NV as studies in healthy volunteers with experimental NV-GE showed unremarkable rectal biopsies.\(^{30}\) However, we found no evidence of small intestine GvHD coexisting with NV infection in any of our patients.

Three patients finally died in close association with the NV-GE. One patient experienced fatal aspiration. The other two patients died as a consequence of sepsis, which was likely promoted by the mucosal barrier defect induced by the NV infection and consecutive translocation of intestinal bacteria.
Infections with widespread enteric viral pathogens other than NV have been repeatedly reported in patients with hematological diseases or HSCT patients. Two larger studies evaluating rotavirus or astrovirus infections in patients with HSCT or hematological diseases demonstrated a pronounced illness and a prolonged duration of illness (median >10 days).\textsuperscript{31,32} In contrast to NV, rotavirus and astrovirus infections appear to some extent less severe as lethal complications have not been reported from these studies. Interestingly, asymptomatic shedding of astrovirus was found in 8 of 21 pediatric patients investigated during an outbreak in a transplant unit,\textsuperscript{32} which further indicates that this small round structured virus is less virulent than NV.

As shown in this study, established clinical outbreak and case-defining criteria are unsuitable to recognize NV infections in immunocompromised patients, because side effects of chemotherapy or HSCT may mimic symptoms of NV infection. In addition, symptoms of intestinal GvHD may be undistinguishable from NV-GE and the therapeutic strategies in these two conditions are greatly divergent. Comparison of duodenal biopsies from patients with GvHD prior to and after onset of the NV infection exhibited distinct histopathological patterns, which allow segregation of GvHD from NV infection. During the NV infection, histopathological findings in immunocompromised patients were similar to those recently described in non-immunocompromised hosts with NV-GE, which clearly contrasts the characteristic
histopathological pattern in the same patients during acute GvHD and prior to the onset of NV-GE.\textsuperscript{33}

Our data demonstrate that NV infections cause severe disease and fatal outcomes in patients after chemotherapy and HSCT. Meticulous isolation may reduce transmission of NV and such enhanced hygiene measures should always be implemented at first suspicion of an NV outbreak in health care facilities with immunocompromised patients, while the laboratory screening for NV in risk patients is pending. Based on these experiences we maintained our modified hygiene strategy in suspicious cases and incorporated routine testing for NV and other enteric viral pathogens into the diagnostic algorithm.
Acknowledgements

The authors are grateful to M. Höhne and K. Stanossek for NV genotyping and skillful technical assistance. Maria Vergoulidou has received a research grant from the Berliner Krebsgesellschaft.

Authorship Contributions

Stefan Schwartz, Thomas Schneider and Eckhard Thiel designed the study, analyzed/interpreted the data and wrote the paper. Maria Vergoulidou, Mark Reinwald and Martin Schmidt-Hieber collected data and performed data analyses. Eckart Schreier and Willy Flegel contributed vital analytical tools (Norovirus genotyping and detection; molecular Histo-blood group typing), analyzed data and wrote the paper. Christoph Loddenkemper performed and analyzed histopathology and wrote the paper.

Conflict of Interest Disclosures

None of the authors have relevant conflicts of interest to declare.
Reference List


Table 1. Characteristics and symptoms of patients and staff members with Norovirus gastroenteritis.

<table>
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<th>Patients (11)</th>
<th>Staff members (11)</th>
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<tbody>
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<td><strong>Age (years)</strong></td>
<td>Median (range)</td>
<td>56 (17 – 71)</td>
<td>38 (20 – 56)</td>
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<td><strong>Sex</strong></td>
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<td>4 / 7</td>
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<td><strong>Secretor status</strong></td>
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<td>n.d.</td>
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<tr>
<td></td>
<td>Se/Se</td>
<td>4</td>
<td>n.d.</td>
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<tr>
<td><strong>Underlying diseases</strong></td>
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<tr>
<td>Median (range)</td>
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<td><strong>1st symptoms relative to HSCT (n=5)</strong></td>
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<tr>
<td>Median (range)</td>
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<td><strong>neutrophils &lt; 500/µL</strong></td>
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<td><strong>Symptoms</strong></td>
<td>Median (range)</td>
<td>7 (2 – 36) days</td>
<td>3 (1 – 13) days</td>
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<tr>
<td>Vomiting</td>
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<td>7</td>
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<tr>
<td>Diarrhea</td>
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<tr>
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<td>Septicemia</td>
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Numbers of individuals are given, if not stated otherwise; HSCT = hematopoietic stem cell transplantation; 1 at onset of symptoms
Table 2. Outbreak and case-defining criteria in all patients who experienced diarrhea and/or vomiting during the outbreak with Norovirus gastroenteritis.

<table>
<thead>
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<th>Norovirus PCR</th>
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<th>Negative (n=15)</th>
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<td><strong>Outbreak criteria (ref. 5)</strong></td>
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<tr>
<td>Duration of symptoms (days)</td>
<td>7 (2 – 36)</td>
<td>6 (1 – 58)</td>
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<tr>
<td>Median (range) *</td>
<td></td>
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<tr>
<td>Patients with vomiting</td>
<td>8/11 (73%)</td>
<td>11/15 (73%)</td>
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<tr>
<td>Patients with stool specimens negative for bacterial pathogens</td>
<td>8/11 (73%) 1</td>
<td>7/15 (47%) 2</td>
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<td><strong>Case-defining criteria (ref. 6)</strong></td>
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</tr>
<tr>
<td>≥ 2 episodes of vomiting +</td>
<td>11/11 (100%)</td>
<td>6/15 (40%)</td>
</tr>
<tr>
<td>≥ 3 episodes of diarrhea +</td>
<td>11/11 (100%)</td>
<td>10/15 (67%)</td>
</tr>
<tr>
<td>Both criteria fulfilled during a 24-hour period +</td>
<td>7/11 (64%)</td>
<td>3/15 (20%)</td>
</tr>
</tbody>
</table>

* not significant (p = 0.47), + p ≤ 0.05, 1 3 patients without microbiological testing, 2 4 patients without microbiological testing, 3 patients with positive test result for C. difficile toxin, 1 patient with enhanced number of yeasts in stool specimen
Figure legends

Figure 1. Kaplan-Meier curves showing the disease duration in norovirus gastroenteritis.

The symptom durations in 11 patients and 11 staff members with norovirus gastroenteritis are displayed.

Figure 2. Abdominal CT scan of a patient with norovirus gastroenteritis.

Extensive bowel edema in a patient with norovirus gastroenteritis acquired during BEAM conditioning chemotherapy for autologous hematopoietic stem cell transplantation. The bowel wall edema was restricted to the small intestine and the colon showed only a moderate dilatation.

Figure 3. Sequential duodenal biopsies from a patient with allogeneic HSCT.

(A) Duodenal biopsy with severe GvHD showing partial loss of surface epithelium (H&E) and increased numbers of apoptotic crypt epithelial cells (inset: cleaved caspase-3 staining showing apoptotic epithelial cells at the base of crypts), (B) predominance of CD8⁺ T-cells in the lamina propria with focal infiltration of the crypt epithelium (immunohistochemistry, APAAP-method). (C) In contrast, duodenal biopsy in the same patient during Norovirus infection shows villous blunting (H&E) and only a
slight increase of apoptotic epithelial cells at the tip of the villi (inset: cleaved caspase-3) and (D) a massive increase of intraepithelial CD8+ T-cells pronounced in the surface epithelium. Original magnifications, x 200; and x 600 (inset). Microscope: Olympus AX70 (Olympus, Melville, NY); numeric aperture of objective lenses: x 20, 0.70 mm; x 60, 1.40. Stains: hematoxylin and eosin (H&E, A and C); APAAP and strepavidinAP (B and D, inset). Camera: JVC KY-F70 (JVC, Yokohama, Japan); acquisition software: DISKUS (Koenigswinter, Germany); and software used for image processing: Adobe Photoshop 7.0 (Adobe Systems, San Jose, CA).
Figure 1

- **patients (n=11)**
- **staff members (n=11)**

\[ p = 0.02 \]
Figure 3

A

B

C

D
Norovirus gastroenteritis causes severe and lethal complications after chemotherapy and hematopoietic stem cell transplantation

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