

Noninvasive screening identifies patients at risk for spontaneous bacterial peritonitis caused by multidrug-resistant organisms

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Background and aims: Spontaneous bacterial peritonitis (SBP) is a severe complication of decompensated cirrhosis. The prevalence of multidrug-resistant organisms (MDROs) in patients with cirrhosis is increasing. Identification of patients at risk for SBP due to MDROs (ie, SBP with the evidence of MDROs or *Stenotrophomonas maltophilia* in ascitic culture, MDRO-SBP) is crucial to the early adaptation of antibiotic treatment in such patients. We therefore investigated whether MDROs found in ascitic cultures can also be found in specimens determined by noninvasive screening procedures.

Patients and methods: This retrospective study was conducted at the liver center of the University Hospital Frankfurt, Germany. Between 2011 and 2016, patients with cirrhosis were included upon diagnosis of SBP and sample collection of aerobic/anaerobic ascitic cultures. Furthermore, the performance of at least one complete MDRO screening was mandatory for study inclusion.

Results: Of 133 patients diagnosed with SBP, 75 (56.4%) had culture-positive SBP and 22 (16.5%) had MDRO-SBP. Multidrug-resistant *Escherichia coli* (10/22; 45.5%) and vancomycin-resistant enterococci (7/22; 36.4%) resembled the major causatives of MDRO-SBP. Rectal swabs identified MDROs in 17 of 22 patients (77.3%) who developed MDRO-SBP with a time-dependent sensitivity of 77% and 87% after 30 and 90 days upon testing, while negative predictive value was 83% and 76%, respectively. The majority of patients were included from intensive care unit or intermediate care unit.

Conclusion: MDRO screening may serve as a noninvasive diagnostic tool to identify patients at risk for MDRO-SBP. Patients with decompensated cirrhosis should be screened for MDROs from the first day of inpatient treatment onward.

Keywords: multidrug resistance, liver cirrhosis, ascites, screening routine, antibiotic therapy

Plain language summary

Patients with liver cirrhosis are prone to develop bacterial infections, which mitigate quality of life and survival. The occurrence of multidrug-resistant organisms (MDROs) puts these patients at additional risk and is therefore a severe life event. Since repetitive antibiotic therapies promote the occurrence of MDROs due to further selection pressure, the antibiotic regimen needs to be chosen carefully based on individual patient data. Therefore, the identification of patients at risk for infections caused by MDROs is vital. MDROs are mostly harbored in the large intestine or on the skin and can easily be diagnosed using screening swabs.

Spontaneous bacterial peritonitis (SBP) is the second most frequent bacterial infection in patients with cirrhosis and ascites and is caused by intestinal bacteria migrating from the gut to the abdomen. Due to the late onset of symptoms, diagnosis may be delayed, and special attention

is warranted in the care of these patients. So far, studies determining the outcome of SBP caused by MDROs are scarce. We retrospectively compiled a cohort of patients with SBP who underwent at least one complete MDRO screening (rectal, nasopharyngeal and cutaneous). In this study, we described the clinical impact of ascitic MDRO detection and aimed to assess the utility of such noninvasive screening to identify patients at risk for SBP caused by MDROs.

Introduction

The prevalence of multidrug-resistant organisms (MDROs) in patients with cirrhosis is rapidly increasing.^{1,2} Bacterial infections have a devastating clinical impact in patients with decompensated cirrhosis, including prolonged and repeated hospitalization as well as reduced survival.^{3–5} Spontaneous bacterial peritonitis (SBP) is the second most common bacterial infection in patients with cirrhosis and ascites.⁶ Studies from different continents have determined cirrhotic patients' outcome as particularly poor in the case of infection by MDROs.^{2,3,7–13}

MDRO-mediated infections in cirrhosis have mostly been attributed to multidrug-resistant gram-negative bacteria (MDRGN).^{14,15} Secondary SBP prophylaxis with fluoroquinolones has been recommended for more than 2 decades, but multidrug resistance narrows therapeutic options.^{15–18} In addition, non-fermenting gram-negative bacteria expressing multidrug resistance, such as *Stenotrophomonas maltophilia*, might bloom under recurrent administration of antimicrobials.^{4,19,20} In addition, gram-positive pathogens such as enterococci are increasingly contributing to the poor prognosis of SBP, since patients with cirrhosis are at high risk to acquire vancomycin-resistant enterococci (VRE).^{15,21–23} The burden and spread of VRE are associated with extensive and repetitive antibiotic therapies.^{24,25} The occurrence of these pathogens is being promoted by intestinal dysbiosis, putting patients with cirrhosis at high risk for acquiring SBP by MDROs (ie, SBP with the evidence of MDROs or *S. maltophilia* in ascitic culture, MDRO-SBP).^{19,20}

Among the entities of bacterial infection in cirrhosis, SBP is particularly harmful due to the late onset of symptoms, risk of recurrence and the wide spectrum of causative bacteria.^{4,10,15,21,22,26} To guarantee optimal patient care and, at the same time, minimize selective pressure on further MDRO development, an individualized antibiotic regimen based on concise patient-specific data may be warranted. Therefore, the anticipation of expectable pathogens would be vital. Since SBP is derived from the intestinal microflora, it seems feasible to investigate whether these pathogens may be detected earlier by noninvasive procedures. In this retrospective study,

we assessed the diagnostic utility of rectal, nasopharyngeal and cutaneous MDRO screening in a cohort of patients with decompensated cirrhosis, and we described the microbiological characteristics and clinical impact of MDRO-SBP in a tertiary liver transplant center in Germany.

Patients and methods

Study characterization

This study was conducted at the liver transplant center at University Hospital Frankfurt (UHF), Germany. From an electronic database, we retrieved all cases coded as “acute peritonitis” (ICD codes such as K65.0, K65.8 and K65.9) from 2012 to 2016. We retrieved 419 cases in total, which are represented by 299 individuals receiving inpatient treatment in the given period. Inclusion criteria were as follows: 1) diagnosis of SBP as defined by polymorphonuclear leukocyte count of $>250/\mu\text{L}$ in ascites in patients with cirrhosis. Patients were included upon the first SBP episode diagnosed at UHF. 2) Sample collection of at least one pair of aerobic/anaerobic ascitic cultures at the bedside at the time of diagnosis of SBP. 3) Complete MDRO screening (rectal, throat, skin, nose swabs). Exclusion criteria were as follows: patients with primary peritonitis or malignant ascites due to any abdominal metastases or peritoneal carcinosis. Biometric, clinical, laboratory and microbiological data as well as empiric antibiotic therapy were collected at the date of study inclusion. If more detailed patient characterization was deemed necessary, the corresponding data were retrieved at defined time points after study inclusion. All the patient data were retrieved from electronic patient charts. Importantly, further microbiological data were collected from study inclusion until death or lost to follow up, resulting in multiple microbiological test results for most patients.

Colonization, infection and case definition

If MDROs were detected only in rectal, throat or cutaneous swabs, patients were defined as “colonized”. If the MDRO strain was detected only or additionally in ascites, blood, urine, bronchial or pleural secretion, bile, pus, wounds or surgical sites, on medical devices or catheters, the patient was classified as “infected” by the strain. “Nosocomial SBP” was defined as SBP diagnosed at least 3 days after hospitalization according to the current definition^{27,28} or as SBP acquired up to 7 days after the last hospital discharge. Sepsis was defined as suspected or evident bacterial infection combined with two of the following clinical parameters according to the quick sequential organ failure assessment (qSOFA) score²⁹:

hemodynamic instability with systolic blood pressure below 100 mmHg (including patients receiving noradrenaline with systolic blood pressure >100 mmHg), respiratory rate >22/min or PaCO₂ <32 mmHg and/or altered mental status.

Screening for MDROs at UHF

Screening routine was performed for MDRGN, VRE and methicillin-resistant *Streptococcus aureus* (MRSA). No regular screening was scheduled for *S. maltophilia*. MDRO screening was performed according to the infection control protocol at UHF which again is according to the German infection protection law (§23, *Infektionsschutzgesetz*³⁰). Screening measures apply to, eg, patients arriving from hospitals in high prevalence countries/high prevalence local hospitals for MDRGN and/or MRSA as well as patients admitted to intensive care unit/intermediate care unit (ICU/IMC) on the day of admission. In addition, MDRO screening is scheduled weekly on ICU and IMC at UHF, and patients included may therefore have undergone repetitive MDRO screening. As of 2011–2016, no regular MDRO screening was scheduled on normal hepatologic wards.

Definition of MDRO-SBP

MDRO-SBP was defined as SBP due to MDRGN, VRE and MRSA as well as due to *S. maltophilia*. Although *S. maltophilia* intrinsically exhibits a broad spectrum of antibiotic resistance, eg, against antibiotic resistance, eg against carbapenems, it does not genuinely qualify to be included in the collective term “MDRO”. However, *S. maltophilia*, MDRGN, VRE and MRSA all require treatment with broad-spectrum antibiotics. We therefore decided to include *S. maltophilia* in the collective term “MDRO-SBP”.

Definition of MDRGN

MDRGN were defined as *Enterobacteriaceae* with extended-spectrum β-lactamase (ESBL) phenotype as well as *Enterobacteriaceae*, *Pseudomonas aeruginosa* and *Acinetobacter baumannii* resistant against piperacillin, any third- or fourth-generation cephalosporin and fluoroquinolone resistance (QR) ± CR.^{31,32}

Detection of MDROs and molecular resistance analysis

All laboratory procedures were performed under quality-controlled criteria (laboratory accreditation according to ISO 15189:2007 standards; certificate number D-ML-13102-01-00, valid through January 25, 2021). To identify MDRGN and VRE, rectal swabs were collected using culture swabs with

Amies collection and transport medium (Hain Lifescience, Nehren, Germany) and streaked onto selective CHROMagar ESBL plates (Mast Diagnostica, Paris, France) and ChromID VRE agar (bioMérieux, Nürtingen, Germany), respectively. Species identification was performed by matrix-assisted laser desorption ionization–time-of-flight analysis (VITEK MS; bioMérieux). Antibiotic susceptibility testing was performed according to Clinical and Laboratory Standards Institute guidelines using VITEK 2 and/or antibiotic gradient tests (bioMérieux). In the case of gram-negative CR isolates, the detection of genes encoding carbapenemases was routinely performed via PCR analysis and subsequent sequencing including the bla genes for carbapenemases NDM, VIM, IMP, OXA-48 like, and KPC as well as OXA-23, OXA-24 and OXA-58 for *Acinetobacter baumannii*.^{33,34} For the detection of MRSA, moistened nasal swabs were inoculated on Brilliance MRSA Agar (Oxoid, Wesel, Germany), and identification and antimicrobial susceptibility testing were performed as described earlier. The detection of MDROs from screening and microbiological culture of clinical specimens (eg, ascites) were performed according to German microbiological procedure quality standards (MiQ), for example, current versions such as MiQ 2, 3a/b (2007), MiQ 25 (2006), MiQ 32 (2012).

Ethics statement

For this retrospective study, approval from the clinical ethics committee of the Goethe University Frankfurt was obtained prior to research (file number: 391/15). The database was anonymized, and informed consent was waived due to the retrospective nature of the study.

Statistical analysis

BiAS software (v. 11.06; Epsilon-Verlag, Darmstadt, Germany) and R (v. 3.2.0, R Core Team [2015], R: A language and environment for statistical computing; R Foundation for Statistical Computing, Vienna, Austria) were used for statistical calculations. Fatal outcome due to sepsis, fatal outcome due to other causes and the appearance of MDRO-SBP were defined as end points, and patients who did not reach an end point were censored from the last day of inpatient treatment. Model for End-stage Liver Disease (MELD), Albumin–Bilirubin (ALBI), Child–Pugh–Turcotte and Chronic Liver Failure–Acute Decompensation (CLIF-AD) scores were calculated with the laboratory parameters taken at the day of SBP detection. Standard deviation (SD) was calculated for age, MELD and CLIF-AD scores and laboratory parameters. SE of the mean was calculated for all observational intervals given in days. Clinical scores were compared using the

Wilcoxon–Mann–Whitney *U*-test. Kaplan–Maier curve was employed for the display of comparative survival analysis, and Cox–Mantel chi-squared test was used for the calculation of HR and *P*-values. Considering the appearance of MDRO-SBP as the event of interest, fatal outcome due to sepsis and fatal outcome due to other causes than sepsis were defined as competing risks, using the Aalen–Johansen estimator. In case that death fatal due to sepsis was considered as the event of interest, fatal outcome due to other causes than sepsis was defined as competing risk. Independent-risk factors for death were calculated using the multivariate Cox regression model. For calculating unbiased sensitivity and specificity for censored data with competing risks, the method proposed by Blanche et al³⁵ was utilized.

Results

Clinical characterization of the cohort

One hundred and thirty-three patients diagnosed with SBP between December 2011 and August 2016 were included in the analysis (Figure S1). Of these, 72 (54.1%) were positive for MDRO colonization or infection in any location, while MDRO-SBP was diagnosed in 22 patients (16.5%). Biometric and laboratory parameters as well as liver scores (Child–Pugh–Turcotte, MELD, CLIF-AD, ALBI) did not differ between patients with or without the detection of MDROs or MDRO-SBP, respectively. Clinical features of hepatic decompensation were equally distributed (Table 1). MELD score and ICU admission at the time of SBP diagnosis were

independent predictors of survival, and MELD, CLIF-AD and ALBI scores were higher in patients who died than those in patients who were lost to follow up (Table S1). Patients with MDRO-SBP were at increased risk of death (HR = 2.17, *P* = 0.007) and were more likely to develop lethal sepsis (HR = 5.67, *P* < 0.001). While MELD and CLIF-AD scores were marginally better in patients with culture-negative SBP, severe clinical courses due to bacterial infections were more frequent in patients with MDRO-SBP (Figure 1). As calculated by multivariate regression, liver scores or clinical features were not associated with the occurrence of MDRO-SBP.

Microbiological characterization of the cohort

The first MDRO screening was performed before or upon study inclusion in 105/133 patients (78.9%) and in timely fashion in the rest of the patients, resulting in MDRO detection in the majority of patients (Figure 2A and B). Among 72/133 patients with MDROs, MDRGN were detected in 42/133 cases (31.6%), VRE in 50/133 cases (37.6%) and MRSA in 5/133 cases (3.8%). Colonization – defined as positivity for MDRO screening without detection in invasive materials – was detected in 16/42 patients (38.1%) with MDRGN, 39/50 patients (78%) with VRE and 2/5 patients (40%) with MRSA. MDRO-associated ascitic or extra-ascitic infection was found in the remaining patients within the body compartments as given in Table 2. Multiple MDRO strains were detected within 24 individuals, resulting

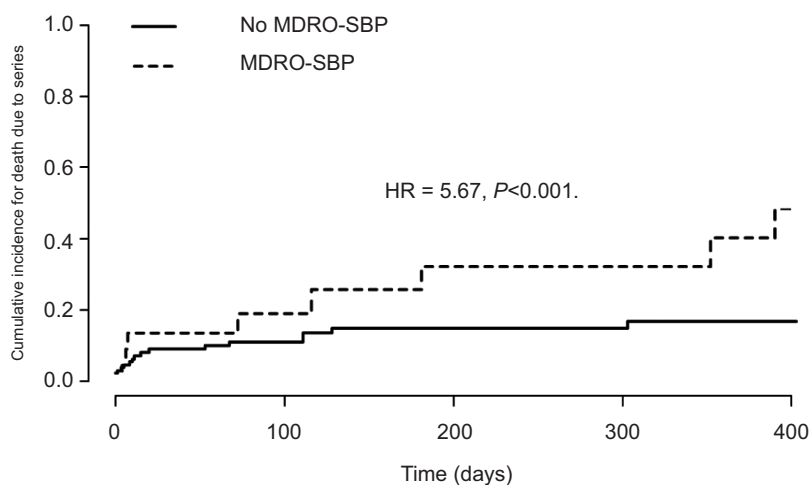


Figure 1 Competing risk analysis of 22 patients with MDRO-SBP compared to 111 patients without MDRO-SBP.

Notes: Patients were included from the day of first SBP detection, and death due to sepsis and death due to other reasons were defined as competing risks. Lethal sepsis was significantly increased in patients with MDRO-SBP (HR = 5.67, *P* < 0.001). Of note, MDRO-SBP was included as a time-dependent variable in this analysis. Therefore, the number at risk cannot be displayed in this plot, since the time-dependent calculation has been performed with censored data.³⁵ MDRO-SBP, SBP with the evidence of MDROs or *S. maltophilia* in ascitic culture.

Abbreviations: MDRO, multidrug-resistant organism; SBP, spontaneous bacterial peritonitis; *S. maltophilia*, *Stenotrophomonas maltophilia*.

Table 1 General characteristics of the included patients

	All patients	Patients with MDROs	Patients with MDRO-SBP	Patients with culture-negative SBP
Σ	133 (100%)	72 (54.1%)	22 (16.5%)	58 (43.6%)
Biometric characteristics				
Age (year), mean ± SD (range)	55.1±11.2 (28–86)	56.1±10.8 (28–74)	54.1±11.9 (46–74)	57.1±11.6 (30–86)
Male sex	98 (73.7%)	53 (73.6%)	20 (90.9%)	37 (63.8%) ^a
Mean observational time (d), mean (range)	230 (1–1,529)	206 (1–1,109)	174 (1–799)	264 (1–1,529)
Liver scores				
MELD	22.4 (7–40)	23.1 (11–40)	24.4 (13–40)	20.3 (7–40) ^a
Child A	1 (0.8%)	1 (1.4%)	0 (0%)	1 (1.7%)
Child B	35 (26.3%)	14 (19.4%)	3 (13.6%)	21 (36.2%)
Child C	97 (72.9%)	57 (79.2%)	19 (86.4%)	36 (62.1%)
CLIF-AD	61.9 (32–89)	63.3 (43–88)	64.4 (43–88)	59.2 (32–85) ^a
ALBI	1.15 (0.35–1.9)	1.17 (0.55–1.89)	1.21 (0.55–1.65)	1.09 (0.35–1.9)
Etiology of cirrhosis^b				
Alcoholic	80 (60.2%)	44 (61.1%)	15 (68.2%)	37 (63.8%)
HBV/HCV	45 (33.8%)	23 (31.9%)	7 (31.8%)	18 (31%)
Others ^c	19 (14.3%)	11 (15.3%)	4 (18.2%)	10 (17.2%)
Laboratory parameters				
Sodium (mmol/L)	134.1 (110–159)	133.1 (118–149)	131.5 (118–145) ^a	136.2 (122–159) ^a
Creatinine (mg/dL)	2.01 (0.45–7.66)	2.08 (0.60–7.66)	2.34 (0.60–7.66)	1.83 (0.45–4.61)
Bilirubin (mg/dL)	6.4 (0.2–44.6)	6.4 (0.4–43.1)	5.9 (0.4–18.6)	6.1 (0.2–44.6)
Albumin (g/dL)	2.8 (1.5–4.4)	2.8 (1.5–4.3)	2.7 (1.5–4)	2.9 (1.8–4.4)
INR	1.73 (1.04–3.75)	1.77 (1.04–3.1)	1.86 (1.04–3.1)	1.59 (1.05–2.79) ^a
Leukocytes/μL	11.6 (2–96.1)	12.6 (2.7–96.1)	11.8 (3.1–23.8)	12.7 (2.4–96.1)
CRP (mg/dL)	7.1 (0.2–30.4)	7.4 (0.2–30.4)	6.8 (0.2–28.2)	7.2 (0.5–26)
Hemoglobin (g/dL)	9.9 (5.5–15.3)	9.6 (5.5–14.6)	9.9 (6.3–14.6)	9.7 (5.8–14.8)
TSH	3.9 (<0.01–126)	3.27 (0.03–77)	2 (0.2–5.1)	5.18 (<0.01–126) ^a
Neutrophils/μL ascites	3.6 (0.25–31.4)	3.9 (0.26–31.4)	4.3 (0.26–16.4)	1.81 (0.25–9.94) ^a
Clinical courses				
Hepatic encephalopathy	88 (66.2%)	50 (69.4%)	15 (68.2%)	35 (60.3%)
Variceal bleeding	49 (36.8%)	30 (41.7%)	9 (40.9%)	18 (31%)
Hepatorenal syndrome	29 (21.8%)	16 (22.2%)	6 (27.3%)	11 (20%)
Portal vein thrombosis	25 (18.8%)	17 (23.6%)	5 (22.7%)	9 (15.5%)
HCC	25 (18.8%)	7 (9.7%) ^a	1 (4.5%)	14 (24.1%)
Liver transplantation ^d	9 (6.8%)	9 (12.5%) ^a	3 (13.6%)	1 (1.7%)
Sepsis	9 (6.8%)	7 (9.7%)	5 (22.7%)	1 (1.7%)
At SBP diagnosis				
Overall	29 (21.8%)	21 (29.2%) ^a	12 (54.5%) ^a	6 (10.3%)
ICU	67 (50.4%)	39 (54.2%)	10 (45.5%)	27 (46.6%)
At SBP diagnosis				
Overall	104 (78.2%)	62 (86.1%) ^a	20 (90.9%)	38 (65.5%)
Death	58 (43.6%)	33 (45.8%)	15 (68.2%) ^a	22 (38%)
Overall				
Due to sepsis	18 (13.5%)	12 (16.7%)	8 (36.4%) ^a	3 (5.2%) ^a
Due to others ^e	40 (30.1%)	21 (29.2%)	7 (31.2%)	19 (32.8%)

(Continued)

Table 1 (Continued)

	All patients	Patients with MDROs	Patients with MDRO-SBP	Patients with culture-negative SBP
Death after 30 days	33 (24.8%)	19 (26.4%)	9 (40.9%)	12 (20.7%)
Death after 90 days	37 (27.8%)	22 (30.6%)	10 (45.5%)	15 (25.9%)

Notes: ^aSignificant difference ($P < 0.05$) was detected in these patients compared to patients without the detection of MDROs, without the detection of MDRO-SBP and with culture-positive SBP. ^bCirrhosis was both alcoholic and due to viral hepatitis in eleven patients, resulting in 146 etiologies in total in this line. ^cCirrhosis due to other reasons composed of primary sclerosing cholangitis, autoimmune hepatitis, alpha 1-antitrypsin deficiency, and cirrhosis due to both HBV and cardiac cirrhosis, with $n=1$ each. ^dTwo patients suffered from newly occurring cirrhosis of the liver transplant eleven and 19 years after transplantation, respectively. ^eDeath due to other reasons composed of liver failure (ten patients), multi-organ failure (six patients), lactic acidosis (five patients), gastrointestinal bleeding (four patients), bleeding other than gastrointestinal (six patients), respiratory failure (four patients), kidney failure (two patients), liver transplant failure and heart failure (one patient each). MDRO-SBP, SBP with the evidence of MDROs or *S. maltophilia* in ascitic culture.

Abbreviations: ALBI, Albumin–Bilirubin; CLIF-AD, Chronic Liver Failure–Acute Decompensation; CRP, C-reactive protein; HCC, hepatocellular carcinoma; ICU, intensive care unit; MDRO, multidrug-resistant organism; MELD, Model for End-stage Liver Disease; SBP, spontaneous bacterial peritonitis; *S. maltophilia*, *Stenotrophomonas maltophilia*; TSH, thyroid-stimulating hormone.

in 22 patients tested positive for MDRGN and VRE, one patient for MRSA and VRE and one patient with MDRGN, MRSA and VRE.

Microbiological characterization of SBP and MDRO-SBP

Ascitic culture was positive in 60/133 patients (45.1%) at first paracentesis and in 75/133 patients (56.4%) altogether. Gram-positive bacteria were detected in 29/75 patients (38.7%), followed by gram-negative bacteria in 25/75 patients (33.3%). Both gram-positive and gram-negative bacteria were found in 21/75 patients (28%), which were due to sequential detection in 10/21 patients (47.6%) and simultaneous, ie, polymicrobial SBP, in 11/21 patients (52.4%). In these patients, abdominal perforation as a cause for peritonitis had been ruled out by computed tomography. Appropriate initial antibiotic therapy according to ascitic resistance profile was administered in 41/60 patients (68.3%) with positive first ascitic culture. Patients carrying MDROs yielded positive ascitic cultures more frequently (52/72 cases, 72.2%; HR =14.6, $P < 0.001$), demonstrating that in these patients ascitic MDRO presence was also more likely to be detected.

In 22/133 patients, MDROs or *S. maltophilia* were detected in ascitic cultures during at least one episode. This was the case upon the first diagnosis of SBP in 15/22 patients (68.2%), while MDRO-SBP as a recurring episode of SBP occurred in 7/22 patients (31.8%) with an estimated time-dependent probability of 14% (standard error [SE] =1.4%) after 60 days and 28% (SE =2%) after 90 days, respectively. In the cases of MDRO-SBP after study inclusion, mean MELD score did not differ from baseline MELD. MDRO-SBP was nosocomial in 15/22 cases (68.2%). In 13/22 patients with MDRO-SBP (59.1%),

further MDROs were detected in other compartments than ascites (Table S2).

Assessing the occurrence of MDRO-SBP upon rectal screening

Since SBP originates from the intestinal microflora, we reasoned that rectal MDRO screening might reveal causative to pathogens in patients with MDRO-SBP. Among all patients with MDRO-positive ascitic culture, the identical strain could be detected in other body compartments; rectal swabs being the most frequent localization with 17/22 cases (77.3%; Table 3). Rectal, cutaneous or nasopharyngeal screening identified the corresponding pathogens in 20/22 cases (90.9%). Of note, one of these did not undergo rectal MDRO screening. The estimated time-dependent probability for the occurrence of MDRO-SBP upon the first positive rectal MDRO screening was 11.5% (SE =2.8%) after 30 days and 12.5% (SE =3%) after 60 days, respectively. Time-dependent sensitivity of MDRO screening before the occurrence of MDRO-SBP was 77% after 30 days and 87% after 90 days, while negative predictive value (NPV) was 83% and 76%, respectively (Table S3; Figure S2).

Antibiotic treatment and secondary prophylaxis

Standard empirical therapy for community-acquired SBP at UHF is intravenous ceftriaxone, while the standard of care for nosocomial SBP is a carbapenem such as meropenem or imipenem. Based on the individual microbial data, diverging antibiotic regimens may have been chosen, and inappropriate initial therapy resulted in decreased survival in 19/60 (31.7%) patients with culture-positive SBP (HR =1.92, $P=0.014$). Initial treatment of MDRO-SBP comprised of carbapenems

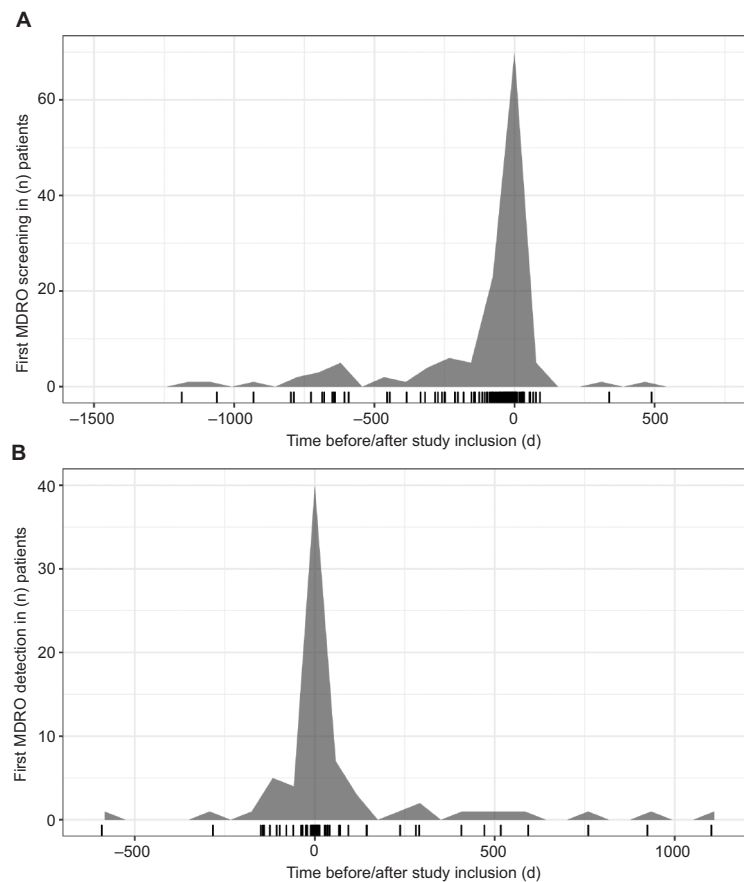


Figure 2 (A) Time of first in-house MDRO screening in relation to time of study inclusion. Ticks at the bottom represent first rectal swab. Screening was performed before or within 28 days upon study inclusion in 123 patients (92.5%), and later than 28 days upon inclusion in 10 patients (7.5%). **(B)** Time of first MDRO detection in relation to time of study inclusion. First overall diagnosis of MDRO had been made before study inclusion in 24/72 patients (33.3%) and was made upon or after study inclusion in 48/72 patients (66.7%).

Abbreviations: *E. coli*, *Escherichia coli*; ESBL, extended-spectrum β -lactamase; ICU, intensive care unit; IMU, intermediate care unit; MDRO, multidrug-resistant organism; QR, fluoroquinolone resistance; SBP, spontaneous bacterial peritonitis; *S. maltophilia*, *Stenotrophomonas maltophilia*.

Table 2 Cumulative evidence of MDROs and *S. maltophilia* collected during the observational period, arranged by sample type and strain

MDROs; $\Sigma=72$ patients ^a		Material						
MDRGN (42 patients, 31.6%)		Ascites	Blood	Rectal	Throat/skin/nose	Urine	Wound	Others
<i>E. coli</i>	ESBL + QR	8 (11%)	3 (4%)	16 (22%)	2 (3%)	2 (3%)	–	1 (1%)
	ESBL	2 (3%)	–	3 (4%)	–	1 (1%)	–	–
<i>K. pneumoniae</i>	ESBL + QR	–	2 (3%)	7 (10%)	2 (3%)	1 (1%)	–	–
	CR + QR	–	–	2 (3%)	–	–	–	–
	ESBL	2 (3%)	2 (3%)	4 (6%)	4 (6%)	–	1 (1%)	1 (1%)
<i>P. aeruginosa</i>	CR + QR	–	–	1 (1%)	1 (1%)	1 (1%)	–	–
<i>E. cloacae</i>	CephR	–	–	–	–	1 (1%)	–	–
	CephR/QR	–	–	2 (3%)	–	2 (3%)	–	–
<i>C. freundii</i>	CephR	–	–	2 (3%)	–	–	–	–
	CephR + QR	–	–	1 (1%)	–	–	–	–
<i>S. maltophilia</i>	No MDROs	2 (3%)	–	1 (1%)	–	–	–	–
VRE (50 patients, 37.6%)		8 (11%)	1 (1%)	47 (65%)	2 (3%)	3 (4%)	2 (3%)	1 (1%)
MRSA (5 patients, 3.8%)		1 (1%)	–	1 (1%)	7 (10%)	1 (1%)	2 (3%)	–

Notes: One patient was positive for both ESBL + QR *E. coli* and VRE in ascitic cultures, resulting in the detection of 23 ascitic MDROs in total. Of note, *S. maltophilia* was not genuinely classified as MDRO, but was nevertheless considered a relevant pathogen due to its intrinsic resistance to CRs. *Enterobacteriaceae* with ESBL phenotype as well as *Enterobacteriaceae*, *P. aeruginosa* and *A. baumannii* resistant to piperacillin, any third-/fourth-generation cephalosporins and QR \pm CR. ^aMultiple MDRO strains were detected in 24 individuals, resulting in 22 patients testing positive for MDRGN and VRE, one patient for MRSA and VRE and one patient with MDRGN, MRSA and VRE.

Abbreviations: *A. baumannii*, *Acinetobacter baumannii*; CephR, cephalosporine resistance; *C. freundii*, *Citrobacter freundii*; CR, carbapenem resistance; *E. cloacae*, *Enterobacter cloacae*; *E. coli*, *Escherichia coli*; ESBL, extended-spectrum β -lactamase; *K. pneumoniae*, *Klebsiella pneumoniae*; MDRGN, multidrug-resistant gram-negative bacteria; MDRO, multidrug-resistant organism; MRSA, methicillin-resistant *Streptococcus aureus*; *P. aeruginosa*, *Pseudomonas aeruginosa*; QR, fluoroquinolone resistance; *S. maltophilia*, *Stenotrophomonas maltophilia*; VRE, vancomycin-resistant enterococci.

Table 3 Colocalizations of MDROs and *S. maltophilia* in other body compartments in patients with MDRO-SBP

MDRO-SBP (n=21)	Material					
	Rectal (n=18)	Throat/ skin (n=2)	Blood (n=5)	Body fluid ^a (n=3)	Wound/ surgical (n=3)	Device ^b (n=2)
Gram-positive pathogens (n=8)						
VRE (n=7)	6	–	1	–	1	–
MRSA (n=1)	1	1	–	–	1	1
MDRGN and <i>S. maltophilia</i> (n=13)						
<i>K. pneumoniae</i> ESBL (n=2)	1	–	2	–	–	1
<i>E. coli</i> (n=9 ^c)	7	1	2	1	1	–
<i>S. maltophilia</i> (n=2)	1	–	–	1	–	–
VRE + MDRGN (n=1)						
<i>E. coli</i> ESBL + QR	1	–	–	–	–	–
VRE	1	–	–	1	–	–

Notes: Of note, strains were detected only in ascitic cultures, and the abovementioned sites are given. One patient was excluded from the chart due to incomplete MDRO screening, resulting in 21 patients. ^aBody fluid composed of tracheal secretion (*S. maltophilia*), bile (VRE) and pleural secretion (*E. coli*). ^bMedical devices with positivity for MDROs composed of left ventricular assistant device (MRSA) and Shaldon catheter (*K. pneumoniae*). ^cPatients with MDRO *E. coli* composed of n=2 expressing ESBL and n=7 expressing ESBL + QR. MDRO-SBP, SBP with the evidence of MDROs or *S. maltophilia* in ascitic culture.

Abbreviations: *E. coli*, *Escherichia coli*; ESBL, extended-spectrum β -lactamase; *K. pneumoniae*, *Klebsiella pneumoniae*; MDRGN, multidrug-resistant gram-negative bacteria; MDRO, multidrug-resistant organism; MRSA, methicillin-resistant *Streptococcus aureus*; QR, fluoroquinolone resistance; SBP, spontaneous bacterial peritonitis; *S. maltophilia*, *Stenotrophomonas maltophilia*; VRE, vancomycin-resistant enterococci.

in 16/22 patients (72.3%), cephalosporins and piperacillin/tazobactam in 2/22 patients (9.1% each), and a fluoroquinolone in one patient (4.5%). Initial treatment furthermore composed of glycopeptides, oxazolidinones or tigecycline in four of eight patients with the diagnosis of VRE-associated MDRO-SBP. Among these cases, resistance against one or more of these agents was detected in four of eight patients (50%; Table 4), while all cases of MDRO-SBP associated with VRE displayed QR expression. If necessary, antibiotic regimens were adapted immediately according to microbiological findings. Nevertheless, inadequate initial antibiotic therapy was associated with highly increased risk of death in MDRO-SBP patients (14/22 patients, 63.6%, HR =11.4, $P=0.002$). Clinical courses did not differ notably between different causatives of MDRO-SBP (Table S4).

One hundred and five patients (78.9%) could be discharged after first inpatient treatment. Secondary antibiotic prophylaxis as recommended by the current hepatologic guidelines³⁴ was prescribed in 86/105 patients (81.9%; Table S5). Recurrent SBP occurred in 35/105 patients (33.3%), and ascitic evidence of bacteria expressing QR was found in 13 of these patients (37.1%) and in 36/133 patients in total (27.1%). This was due to *Enterococcus* spp. or VRE in 16/36 cases (44.4%), MDRGN in 8/36 cases (22.2%), MRSA in one case (2.8%) and due to others (other pathogens expressing QR composed of coagulase-negative *Staphylococcus* spp. [four patients], *K. pneumoniae* [two patients] and *Clostridium tertium*, *E. coli*, *S. aureus*, *Streptococcus*

mitis and *S. maltophilia* [one patient each]) in 11/36 cases (30.6%). The estimated time-dependent probabilities for recurring SBP were 9.6% (SE =2.8%) after 30 days, 15.7% (SE =3.5%) after 60 days and 17.9% (SE =3.8%) after 90 days, and MDRO-SBP was more likely to occur in patients with recurrent SBP (HR =2.82, $P=0.001$).

Discussion

The global incidence of MDRO-SBP, especially caused by pathogens expressing resistance to cephalosporins and fluoroquinolones, is a growing health threat.³⁶ We showed that accurate microbiological screening measures helped identify patients who developed MDRO-SBP. These patients are at high risk of death due to sepsis. Although the study was designed in a retrospective fashion, our findings are consistent and clinically substantial due to the following arguments.

First, patients with and without MDRO detection had similar baseline parameters, thus resembling a homogenous population in terms of the severity of disease. Second, survival was impaired if the initial empiric antibiotic therapy did not cover the indicated pathogens, which is in line with previous findings and shows that lethality was mainly associated with bacterial infections.^{37,38} Third, while liver scores and survival in patients with culture-negative SBP were comparable to culture-positive patients, lethal sepsis was less frequent (Table 1).

Regarding patients with culture-negative SBP, one might argue that cases of ascitic MDRO presence might

Table 4 Distribution of resistance patterns among causatives of MDRO-SBP

Pathogen	Resistance pattern				
	ESBL	CR	QR	Amikacin/colistin	Teicoplanin (\pm tigecycline/linezolid ^a)
<i>E. coli</i> (n=10)	10	0	8	0	–
<i>K. pneumoniae</i> (n=2)	2	0	0	0	–
VRE (n=8)	–	–	8	–	4

Notes: The abovementioned pathogens were detected in both MDRO screening and ascites of patients. ^aVRE strains expressing teicoplanin resistance, 2/4 were additionally resistant to tigecycline and 2/4 to linezolid. Of note, cases of *S. maltophilia* are not included in this table. MDRO-SBP, SBP with the evidence of MDROs or *S. maltophilia* in ascitic culture.

Abbreviations: CR, carbapenem resistance; *E. coli*, *Escherichia coli*; ESBL, extended-spectrum β -lactamase; *K. pneumoniae*, *Klebsiella pneumoniae*; MDRO, multidrug-resistant organism; QR, fluoroquinolone resistance; SBP, spontaneous bacterial peritonitis; *S. maltophilia*, *Stenotrophomonas maltophilia*; VRE, vancomycin-resistant enterococci.

have been missed in some of these patients. However, it is well known that the sensitivity of ascitic cultures is only around 50%,¹⁸ a fact that is reflected by our findings and has to be dealt with throughout the management of SBP. Nevertheless, MDRO-SBP was exclusively detected in patients with carriage of MDROs in other body compartments as well, while patients negative for MDROs also had significantly less positive ascitic cultures (HR =0.07, $P<0.001$). Considering the significantly better outcome in regard of septic complications in these patients, who are regularly treated with ceftriaxone, these SBP cases have to be considered as nonresistant rather than resistant.

Time-dependent sensitivity of MDRO screening for developing MDRO-SBP was between 77% and 95% including a high NPV, while specificity of the test was only around 50% (Table S3). Thus, negative MDRO screening might rule out MDRO-SBP with a high likelihood. Due to its low specificity and positive predictive value (PPV), though, repetitive diagnostic paracentesis including microbiological sample collection remains mandatory. Patients with nosocomial SBP should receive empiric antibiotic therapy according to local resistance patterns.²⁸ A more “narrow” antibiotic therapy might be justified if MDRO screening was negative, but prospective data are warranted. Notably, the first MDRO evidence was made by the time of diagnosis of MDRO-SBP in almost half of the patients, since no screening results had been available beforehand.

Based on these observations, the ascitic detection of MDROs resembles a major prognostic factor in patients with decompensated cirrhosis.^{11,36} Because these patients are afflicted by rapid deterioration and death, early MDRO detection is vital.^{37,38} Since directional spread of MDROs goes from the intestine or skin into the abdomen (and not vice versa), it would be appropriate to screen for MDRO

colonization in patients with risk or even history of SBP. Our data give the first evidence that this approach might be valid in the clinical setting and suggest that MDRO screening should be conducted on the day of patients’ admission. However, a study including a larger sample size of patients with MDRO-SBP is warranted.

Conclusion

Due to the retrospective design of the study, it must be stressed that prospective data are warranted to validate the accuracy of MDRO screening in association with MDRO-SBP. Therefore, general treatment recommendations cannot be made based on our findings. Furthermore, patients were mainly recruited from ICU/IMC. This translates into comparably high liver scores upon inclusion and an overall reduced survival during inpatient treatment as well as under secondary prophylaxis.³⁹ Yet, differences between patients with and without MDRO-SBP were significant and comprehensible. Importantly, microbiological resistance data might depend on local resistance patterns, which is reflected by a high proportion of VRE, and may therefore not be transferable unrestrictedly. However, our microbiological data are comparable to reports from other monocenter studies.^{1,2,11–14,21,22,40} On a more general basis, we suggest a prospective multicenter setting to determine the long-term clinical impact of MDRO-SBP and the benefit of MDRO screening in patients with decompensated cirrhosis.

Author contributions

Compiling data, data interpretation and drafting and writing the manuscript: PGF, MM and OW; Biostatistical analysis and interpretation: PGF and NF. Data interpretation and critical review of the manuscript: SZ, VAJK, MH, TAW, CML, JV, CR and OW. All the authors approved the final version of the manuscript and agreed to be accountable for all aspects of

the work in ensuring that questions related to the accuracy or integrity of any part of the work are appropriately investigated and resolved. All authors contributed toward data analysis, drafting and critically revising the paper and agree to be accountable for all aspects of the work.

Disclosure

SZ has provided consultancy for Abbvie, BMS, Gilead, Intercept, Janssen and Merck (all outside the scope of this study). OW has provided consultancy for Amgen, Bayer, BMS, Celgene, Eisai, Merck, Novartis, Roche, Servier and Shire (all outside the scope of this study). VAJK was partially supported by a grant from the Deutsche Forschungsgemeinschaft (DFG, research unit 2251). The authors report no other conflicts of interest in this work.

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Supplementary materials

Impact of multidrug-resistant organisms in spontaneous bacterial peritonitis

Table S1 Liver scores at study inclusion, arranged by clinical end points

Score	Death (n=58)			Lost to follow up (n=75)			P-value
	Mean	Median	SD	Mean	Median	SD	
MELD	25.6	25	8.57	19.9	19	7.23	0.0001
CLIF-AD	64.8	67	10.20	59.7	58	10.66	0.007
ALBI	1.249	1.23	0.395	1.070	1.050	0.301	0.005

Abbreviations: ALBI, Albumin–Bilirubin; CLIF-AD, Chronic Liver Failure–Acute Decompensation; MELD, Model for End-stage Liver Disease.

Table S2 Timeline and additional microbiological findings in patients with MDRO-SBP

Patient	MDRO-SBP	Further MDROs	Further ascitic detections	Time in study (days)	Time to MDRO-SBP (days)
1	<i>E. coli</i> ESBL + QR	–	<i>S. aureus</i>	23	0
2	<i>K. pneumoniae</i> ESBL	–	–	116	111
3	<i>E. coli</i> ESBL + QR, VRE	<i>K. pneumoniae</i> ESBL+QR (rectal)	<i>E. cloacae</i> CephR/QR	697	593
4	VRE	–	<i>C. koseri</i> , <i>A. baumannii</i> <i>P. putida</i> , <i>Candida famata</i>	181	53
5	<i>S. maltophilia</i>	–	<i>E. faecium</i> , <i>Candida albicans</i>	6	5
6	VRE	–	–	7	0
7	VRE	–	<i>S. haemolyticus</i> , <i>Candida albicans</i>	7	0
8	<i>E. coli</i> ESBL + QR	VRE (rectal)	–	1	0
9	<i>E. coli</i> ESBL	–	–	4	0
10	MRSA	<i>E. coli</i> (rectal) VRE (rectal, wound) <i>K. pneumoniae</i> (rectal, medical device)	–	390	0
11	<i>E. coli</i> ESBL + QR	VRE (rectal) <i>E. cloacae</i> (urine)	<i>Ent. hirae</i> , <i>B. amyloliquefaciens</i>	799	94
12	VRE	<i>K. pneumoniae</i> (rectal, throat, blood) <i>P. aeruginosa</i> (tracheal secretion)	<i>S. mitis</i> , <i>E. coli</i>	671	297
13	<i>E. coli</i> ESBL	–	–	7	2
14	VRE	–	<i>P. rettgeri</i> , <i>S. aureus</i> , <i>B. cereus</i> , <i>S. epidermidis</i> , <i>S. haemolyticus</i>	14	8
15	VRE	–	<i>E. coli</i>	14	10
16	<i>K. pneumoniae</i> ESBL	–	<i>S. pneumoniae</i>	30	14
17	<i>E. coli</i> ESBL + QR	VRE	–	72	67
18	<i>E. coli</i> ESBL + QR	–	–	15	0
19	<i>S. maltophilia</i>	VRE (rectal)	<i>S. parasanguinis</i> , <i>E. faecalis</i> , <i>S. epidermidis</i>	352	303
20	<i>E. coli</i> ESBL + QR	–	<i>E. faecium</i>	96	0
21	VRE	<i>E. coli</i> ESBL+QR (rectal)	<i>E. gallinarum</i> , <i>E. faecium</i> , <i>S. epidermidis</i>	73	8
22	<i>E. coli</i> ESBL + QR	VRE (rectal)	–	265	0

Note: MDRO-SBP, SBP with the evidence of MDROs or *S. maltophilia* in ascitic culture.

Abbreviations: CephR, cephalosporine resistance; *E. cloacae*, *Enterobacter cloacae*; *E. coli*, *Escherichia coli*; ESBL, extended-spectrum β -lactamase; *K. pneumoniae*, *Klebsiella pneumoniae*; MDRO, multidrug-resistant organism; MRSA, methicillin-resistant *Streptococcus aureus*; *P. aeruginosa*, *Pseudomonas aeruginosa*; QR, fluoroquinolone resistance; *S. aureus*, *Streptococcus aureus*; SBP, spontaneous bacterial peritonitis; *S. maltophilia*, *Stenotrophomonas maltophilia*; *S. mitis*, *Streptococcus mitis*; VRE, vancomycin-resistant enterococci; *C. koseri*, *Citrobacter koseri*; *A. baumannii*, *Acinetobacter baumannii*; *P. putida*, *Pseudomonas putida*; *E. faecium*, *S. haemolyticus*, *Ent. Hirae*, *B. amyloliquefaciens*, *P. rettgeri*, *B. cereus*, *S. epidermidis*, *S. parasanguinis*, *E. faecalis*, *E. gallinarum*.

Table S3 Time-dependent sensitivity and specificity of rectal MDRO screening for the prediction of developing MDRO-SBP

Time after first positive screening	Sensitivity (%)	Specificity (%)	PPV (%)	NPV (%)
30 days	77	50	10	83
90 days	87	50	19	76
1 year	92	62	34	62
2 years	95	71	52	48

Notes: The values resemble the sensitivity and specificity of the test after the time given in the first column (time after the first positive screening). MDRO-SBP, SBP with evidence of MDROs or *S. maltophilia* in ascitic culture.

Abbreviations: MDRO, multidrug-resistant organism; NPV, negative predictive value; PPV, positive predictive value; SBP, spontaneous bacterial peritonitis; *S. maltophilia*, *Stenotrophomonas maltophilia*.

Table S4 Clinical courses of patients with MDRO-SBP, arranged by causative pathogens

	22 patients with MDRO-SBP				
	<i>E. coli</i> ESBL ± QR, n=10	<i>K. pneumoniae</i> ESBL, n=2	<i>S. maltophilia</i> , n=2	VRE, n=8	MRSA, n=1
ICU	9	6	2	8	1
Sepsis at SBP diagnosis	2	1	1	2	0
Sepsis overall	7	6	2	2	1
Death	7	5	2	6	1
Death due to sepsis	3	2	1	2	1
Death due to other reasons ^a	4	3	1	4	0

Notes: Since in one patient MDRO-SBP due to both *E. coli* and VRE was detected, cumulative number of causative pathogens is 23. ^aDeath due to other reasons composed of lactic acidosis (n=3), hemorrhagic shock due to variceal bleeding (n=2), end-stage kidney failure (n=1) and multi-organ failure (n=1). MDRO-SBP, SBP with evidence of MDROs or *S. maltophilia* in ascitic culture.

Abbreviations: *E. coli*, *Escherichia coli*; ESBL, extended-spectrum β-lactamase; ICU, intensive care unit; *K. pneumoniae*, *Klebsiella pneumoniae*; MDRO, multidrug-resistant organism; MRSA, methicillin-resistant *Streptococcus aureus*; QR, fluoroquinolone resistance; SBP, spontaneous bacterial peritonitis; *S. maltophilia*, *Stenotrophomonas maltophilia*; VRE, vancomycin-resistant enterococci.

Table S5 Antibiotic prophylaxis or curative treatment in patients after SBP

	Death (n=30)	Lost to follow up (n=75)
Quinolones (n=65)	23 (35.4%)	42 (64.6%)
Rifaximin (n=13)	2 (15.4%)	11 (84.6%)
TIPS (n=9)	2 (22.2%)	7 (77.8%)
Liver transplantation ^a (n=6)	3 (50%)	3 (50%)
No prophylaxis ^b (n=12)	0	12 (100%)

Notes: Patients were successfully discharged after SBP episode (n=105). ^aFrom 9/133 patients undergoing liver transplantation in total, six patients underwent liver transplantation after study inclusion. ^bReasons for no prophylaxis in these patients were as follows: discharge to other hospital or rehabilitation clinic with ongoing intravenous antibiotic therapy (n=3), transfer to palliative care (n=3), recurring colitis from *C. difficile* (n=1) and unidentifiable (n=5).

Abbreviations: *C. difficile*, *Clostridium difficile*; SBP, spontaneous bacterial peritonitis; TIPS, transjugular intrahepatic portosystemic shunt.

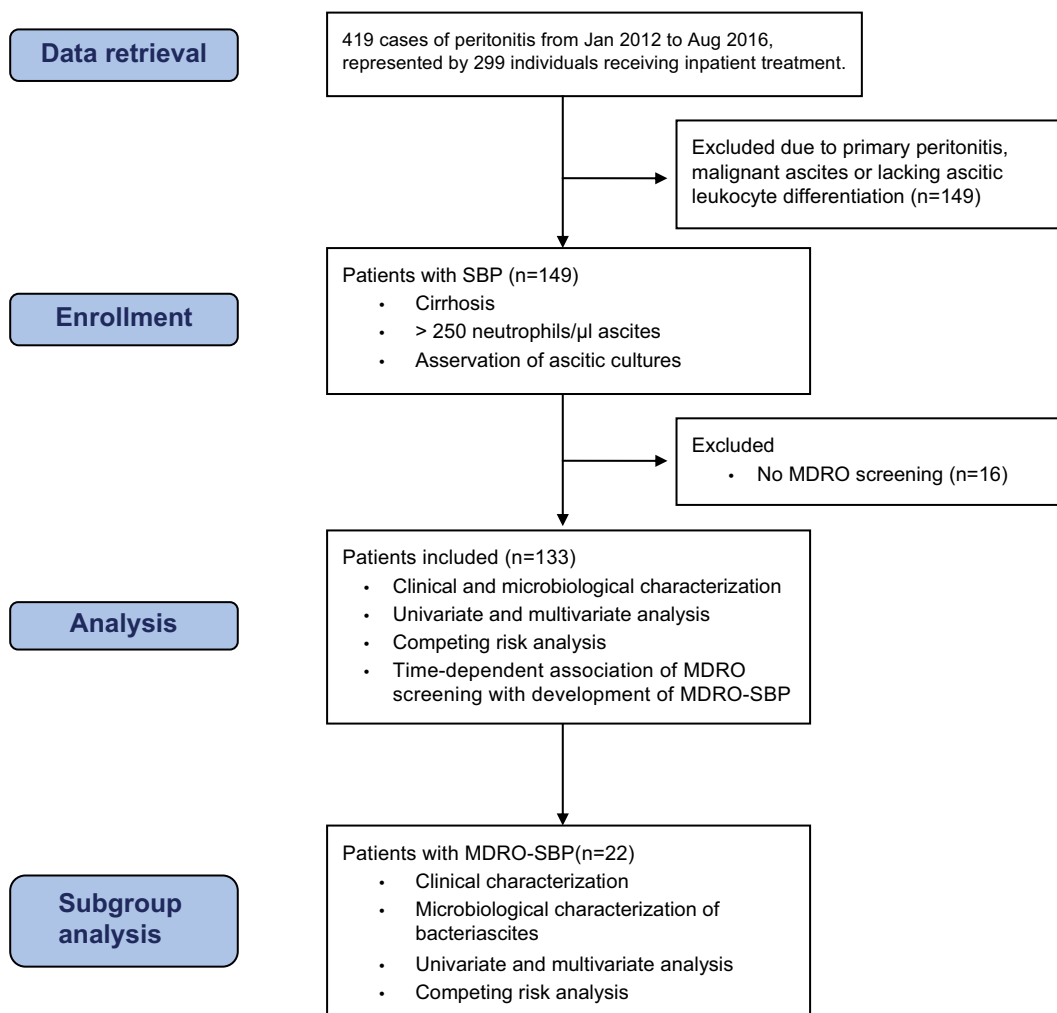


Figure S1 CONSORT diagram of patients included in the study, analysis and subgroup analysis.

Notes: One patient who admitted with MDRO-SBP caused by *E. coli* expressing ESBL and QR did not undergo rectal MDRO swabs, since he had not been admitted to ICU/IMC and no regular MDRO screening was scheduled on normal wards. The patient was included due to ascitic MDRO detection and completion of cutaneous MDRO screening and was defined as MDRO negative in rectal screening. MDRO-SBP, SBP with evidence of MDROs or *S. maltophilia* in ascitic culture.

Abbreviations: *E. coli*, *Escherichia coli*; ESBL, extended-spectrum β-lactamase; ICU, intensive care unit; IMC, intermediate care unit; MDRO, multidrug-resistant organism; QR, fluoroquinolone resistance; SBP, spontaneous bacterial peritonitis; *S. maltophilia*, *Stenotrophomonas maltophilia*.

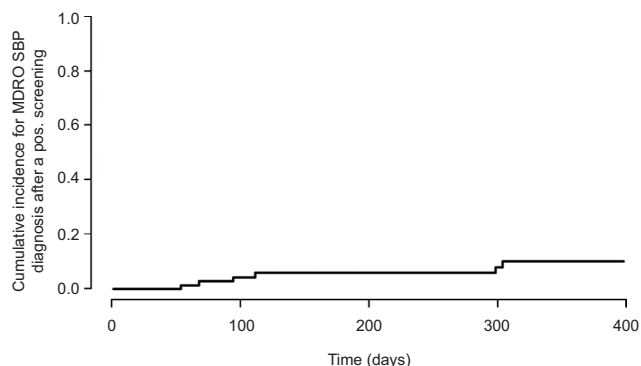


Figure S2 Cumulative incidence for MDRO-SBP diagnosis after a positive rectal SBP screening.

Note: MDRO-SBP, SBP with evidence of MDROs or *S. maltophilia* in ascitic culture.

Abbreviations: MDRO, multidrug-resistant organism; SBP, spontaneous bacterial peritonitis; *S. maltophilia*, *Stenotrophomonas maltophilia*.

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