



# Use of Cefiderocol in Adult Patients: Descriptive Analysis from a Prospective, Multicenter, Cohort Study

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## ABSTRACT

**Introduction:** Cefiderocol is a siderophore cephalosporin showing activity against various carbapenem-resistant Gram-negative bacteria

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(CR-GNB). No data currently exist about real-world use of cefiderocol in terms of types of therapy (e.g., empirical or targeted, monotherapy or combined regimens), indications, and patient characteristics.

**Methods:** In this multicenter, prospective study, we aimed at describing the use of cefiderocol in terms of types of therapy, indications, and patient characteristics.

**Results:** Cefiderocol was administered as empirical and targeted therapy in 27.5% (55/200) and

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72.5% (145/200) of cases, respectively. Overall, it was administered as monotherapy in 101/200 cases (50.5%) and as part of a combined regimen for CR-GNB infections in the remaining 99/200 cases (49.5%). In multivariable analysis, previous isolation of carbapenem-resistant *Acinetobacter baumannii* odds ratio (OR) 2.56, with 95% confidence interval (95% CI) 1.01–6.46,  $p = 0.047$ ] and previous hematopoietic stem cell transplantation (OR 8.73, 95% CI 1.05–72.54,  $p = 0.045$ ) were associated with administration of cefiderocol as part of a combined regimen, whereas chronic kidney disease was associated with cefiderocol monotherapy (OR 0.38 for combined regimen, 95% CI 0.16–0.91,  $p = 0.029$ ). Cumulative 30-day

mortality was 19.8%, 45.0%, 20.7%, and 22.7% in patients receiving targeted cefiderocol for infections by Enterobacterales, *A. baumannii*, *Pseudomonas aeruginosa*, and any metallo- $\beta$ -lactamase producers, respectively.

**Conclusions:** Cefiderocol is mainly used for targeted treatment, although empirical therapies account for more than 25% of prescriptions, thus requiring dedicated standardization and guidance. The almost equal distribution of cefiderocol monotherapy and cefiderocol-based combination therapies underlines the need for further study to ascertain possible differences in efficacy between the two approaches.

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**Keywords:** Cefiderocol; Antimicrobial resistance; Clinical practice; Carbapenem resistance; Carbapenemases

### Key Summary Points

In a real-life, observational, multicenter study, cefiderocol was mostly administered as targeted therapy (72.5%), although the proportion of empirical therapy was non-negligible (27.5%).

Cefiderocol was mainly administered for lower respiratory tract infections and bloodstream infections caused by *Acinetobacter baumannii*, followed by *Pseudomonas aeruginosa*.

Notably, in a real-life scenario, cefiderocol was administered almost equally as monotherapy or as combination therapy (50.5% vs. 49.5%).

The almost equal distribution of cefiderocol monotherapy and cefiderocol-based combination therapies underlines the need for further study to ascertain possible differences in efficacy between the two approaches.

## INTRODUCTION

Cefiderocol is a catechol-substituted siderophore cephalosporin showing in vitro rapid bactericidal activity against various carbapenem-resistant Gram-negative bacteria (CR-GNB) [1, 2].

Based on the results of phase-3 randomized controlled trials (RCT), cefiderocol was approved in Europe for the treatment of GNB infections with limited therapeutic options [3–5]. Various studies have described the use of cefiderocol for treatment of CR-GNB infections, including those caused by carbapenem-resistant *Acinetobacter baumannii* and CR-GNB producing metallo- $\beta$ -lactamases (MBL) for which cefiderocol is usually among the very few active options [6–35]. However, such studies were designed mostly to evaluate cefiderocol for targeted treatment of specific infections and/or pathogens, while no data currently exist about the real-world use of cefiderocol in terms of types of therapy (e.g., empirical or targeted, monotherapy or in combination with other agents), indications, and

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characteristics of the treated patients. These data could be useful to improve the development of empirical and targeted therapeutic algorithms of cefiderocol in real life, ultimately aiming at better patient care.

In this multicenter, observational, prospective study, we primarily aimed at describing how cefiderocol is used in Italian hospitals, in terms of types of therapy, indications, and patient characteristics.

## METHODS

### Setting and Objectives

The MULTI-SITA project is a novel platform developed by the Italian Society of Anti-Infective Therapy (SITA) and dedicated to conducting observational studies on invasive bacterial and fungal diseases. CEFI-SITA is an ongoing observational, prospective, multicenter study conducted in Italian hospitals within the MULTI-SITA project, registering use of cefiderocol in consecutive adult patients according to clinical practice. The prospective study period of the CEFI-SITA study is from 1 August 2022 to 31 December 2025. Here, in a pre-planned analysis, we report the preliminary descriptive results of the CEFI-SITA study. The primary objective of this preliminary analysis was to describe the characteristics of patients and infections treated with cefiderocol in the first 200 enrolled patients (see sample size calculation below) from 17 hospitals. Secondary objectives were: (1) to exploratorily describe

factors associated with use of cefiderocol as part of a combined regimen versus monotherapy; and (2) to describe clinical cure rates and 30-day mortality in patients with GNB infections receiving targeted cefiderocol therapy. Patients receiving at least one dose of cefiderocol for any reason according to local practice were included in the study, in line with its observational nature. Exclusion criteria were (1) age less than 18 years and (2) already included in the study for a previous cefiderocol administration.

### Microbiological Procedures

Identification of bacterial isolates from clinical specimens was performed by means of matrix-assisted laser desorption ionization time of flight mass spectrometry (Vitek MS MALDI-TOF mass spectrometer, bioMérieux, Craponne, France; or MALDI Biotyper, Bruker Daltonics, Billerica, MA, USA) or automated systems, depending on standard local procedures. Antimicrobial susceptibility testing (AST) for antibiotics other than cefiderocol was also performed by means of automated systems (Vitek 2, bioMérieux; MicroScan, Beckman Coulter, Brea, CA, USA; or Phoenix, Becton Dickinson Diagnostics, Sparks, MD, USA) according to local standard procedures. Cefiderocol AST was performed by means of disk diffusion or broth microdilution methods, including either the reference broth microdilution MIC determination using iron-depleted cation-adjusted Mueller Hinton Medium [36] or commercial broth microdilution tests, according to local practice. Cefiderocol AST by gradient test was reported for one isolate. Results of susceptibility testing were interpreted in accordance with the European Committee on Antimicrobial Susceptibility Testing (EUCAST) clinical breakpoints, version 14.0 (<http://www.eucast.org/>), also for cefiderocol. In more detail, resistance to cefiderocol was defined as: (1) for Enterobacterales, minimum inhibitory concentration (MIC) > 2 mg/L for broth microdilution and zone diameter < 23 mm for disk diffusion; and (2) for *P. aeruginosa*, MIC > 2 mg/L for broth microdilution and zone diameter < 22 mm for disk diffusion). For other GNB currently lacking sufficient evidence for defining cefiderocol breakpoints

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according to EUCAST, isolates were defined as non-wild-type based on EUCAST ECOFF breakpoints for ceftiderocol (MIC > 0.5 mg/L for broth microdilution). For disk diffusion, zone diameters < 17 mm (for *A. baumannii*) and < 19 mm (for *S. maltophilia*), corresponding to MIC values below the EUCAST PK-PD breakpoint of ceftiderocol, were also deemed as defining non-wild-type isolates in this study. An MIC > 2 mg/L and an MIC > 0.5 mg/L were also considered as the cut-off for defining resistant and non-wild-type isolates by gradient test, respectively. For the purpose of the study, carbapenem resistance was defined as resistance to any of the carbapenems not intrinsically inactive against the given GNB species. Presence of carbapenemase-encoding gene/s was detected using Verigene BC-GN (Nanosphere, Northbrook, IL, USA) or Xpert Carba-R (Cepheid, Sunnyvale, CA, USA), or inferred by means of the BioFire BCID2 or Pneumonia Plus syndromic panels (bioMérieux), whereas the production of carbapenemase/s was assessed by NG-test Carba 5 (NG Biotech, Guipry, France), depending on standard local procedures at the time the study was conducted.

### Definitions and Data Collected for the Study

The following demographic and clinical variables were collected at the time of ceftiderocol initiation: age in years; sex; previous hospitalization (within 6 months); admission from a long-term care facility (LTCF); diabetes mellitus; chronic obstructive pulmonary disease (COPD); previous myocardial injury; New York Heart Academy (NYHA) score; chronic liver disease (defined histologically as liver cirrhosis or in presence of a clinical diagnosis supported by laboratory, endoscopy, and radiologic findings [37]); chronic kidney disease (defined as estimated glomerular filtration rate < 60 mL/min/1.73 m<sup>2</sup>); chronic intermittent hemodialysis; solid neoplasm; metastatic solid neoplasm; hematological malignancy; previous hematopoietic stem cell transplantation (HSCT); previous solid organ transplantation (SOT); human immunodeficiency virus (HIV) infection; autoimmune disease; age-adjusted Charlson Comorbidity Index [38]; previous therapy

with ceftiderocol (within 6 months); previous antibiotic therapy with agents other than ceftiderocol (within 6 months); previous chemotherapy (within 6 months); previous steroid therapy (within 6 months); previous therapy with other immunosuppressants (within 6 months); previous major surgery (within 3 months); previous isolation of carbapenem-resistant GNB (within 6 months); days from admission to ceftiderocol initiation; intensive care unit (ICU) stay; sequential organ failure assessment (SOFA) score [39]; presence of central venous catheter (CVC); presence of urinary catheter; presence of septic shock [40]; presence of at least mild acute respiratory distress syndrome (ARDS) [41]; estimated creatine clearance (CL<sub>Cr</sub>) and presence of at least stage 1 of acute kidney injury (AKI) according to the Kidney Disease: Improving Global Outcome (KDIGO) criteria [42]; concomitant coronavirus disease 2019 (COVID-19); total parenteral nutrition; neutropenia (defined as blood absolute neutrophil count < 500 cell/mm<sup>3</sup>); continuous renal replacement therapy (CRRT); extracorporeal membrane oxygenation (ECMO); absolute blood white cell count; serum C-reactive protein; serum procalcitonin; type of ceftiderocol therapy, defined as monotherapy or combined therapy based on the anti-carbapenem-resistant GNB (CR-GNB) activity of agents administered concomitantly with ceftiderocol (more in detail, combination of ceftiderocol with at least one of the following agents was considered as a combined anti-CR-GNB therapy: aminoglycosides; fosfomycin; tigecycline, with the exception of targeted therapy of *Pseudomonas aeruginosa* infections; polymyxins; sulbactam or ampicillin/sulbactam, as empirical treatment or as targeted therapy for *Acinetobacter baumannii* infections); timing of ceftiderocol therapy (empirical vs. targeted after identification of the causative agent); initial ceftiderocol dosage according to estimated CL<sub>Cr</sub> or hemodialytic treatment; type of infection treated with ceftiderocol (according to the US Centers for Disease Control and Prevention/National Healthcare Safety Network [CDC/NHSN] surveillance definitions [43]; if CDC/NHSN criteria were not satisfied, local investigators indicated the diagnosis registered in the clinical chart, e.g., sepsis provided Sepsis-3

criteria were satisfied [40]); causative agent/s of the infection treated with targeted ceftiderocol therapy. The following information was also collected during follow-up: causative agent/s of the infection initially treated with empirical ceftiderocol, if subsequently identified; change of ceftiderocol dosage during therapy due to changes in  $CL_{Cr}$ ; clinical cure, defined as resolution of clinical signs and symptoms of infection, at day 7, 14, and 28 after ceftiderocol initiation; cumulative mortality at day 30 after ceftiderocol initiation; adverse events (AE), including serious AE (SAE), occurring during ceftiderocol therapy, if any.

### Sample Size Calculation

A sample size of 200 patients was considered an acceptable compromise regarding the external validity of study results when considering the primary descriptive endpoint (i.e., use of ceftiderocol in terms of indications and characteristics of treated patients). In more detail, by assuming normal distribution of the estimates of population parameters measured in the study sample (e.g., proportion of patients receiving ceftiderocol as empirical therapy/all patients receiving ceftiderocol), a sample size of 200 patients would guarantee a maximum error margin (confidence interval) of  $\pm 7\%$  with  $\alpha = 0.05$ .

### Statistical Analysis

The characteristics of patients and infections treated with ceftiderocol were described through median (interquartile range, IQR) and absolute frequency (relative frequency, %) for continuous and categorical variables, respectively. The 95% confidence interval (CI) was calculated for both proportions [44] and median values [45] estimates. To assess factors associated with ceftiderocol combination therapy, we performed Rubin's multiple imputation as a preliminary step [46]. Then, the association of demographic/clinical variables with ceftiderocol combination therapy (vs. ceftiderocol monotherapy as reference) was first assessed through univariable

logistic regression (LR) models. Subsequently, all variables showing a  $p$  value  $< 0.10$  in univariable comparisons were included in an initial multivariable LR model, and further selected for inclusion in the final multivariable LR model (model A) by means of a stepwise backward procedure. Finally, as sensitivity analysis, variables included in model A were also included in an additional multivariable LR model that also included center as random effect (model B). The crude 30-day mortality from the initiation of ceftiderocol therapy was summarized graphically using the Kaplan–Meier method in patients receiving (1) targeted ceftiderocol therapy for Enterobacterales infection, (2) targeted ceftiderocol therapy for *Acinetobacter baumannii* infection, (3) targeted ceftiderocol therapy for *Pseudomonas aeruginosa* infection, and (4) targeted ceftiderocol therapy for MBL-producing Gram-negative bacteria.

The analyses were conducted using SAS software (version 9.4; SAS Institute, Cary, NC, USA) and R Statistical Software (version 4.2.1; R Foundation for Statistical Computing, Vienna, Austria).

### Ethical Considerations

The MULTI-SITA project was approved by the ethics committee of the coordinating center (Liguria Region Ethics Committee, registry number 390/2020). The amendment authorizing the conduct of the CEFI-SITA study within the MULTI-SITA project was approved by the Liguria Region Ethics Committee on 12 April 2022. The other participating centers followed the local ethical committees' requirements and started to prospectively enroll patients once activated. All conscious patients at time of enrollment signed an informed consent to participate in the study. A waiver of informed consent for data collection from unconscious patients at the time of enrollment due to severe clinical conditions was obtained within the ethics committee approval, in line with the observational nature of the analyses and in order not to bias research results towards high cure rates and low mortality prejudicing scientific validity.

**Table 1** Demographic and clinical characteristics of adult patients treated with cefiderocol

Variables	No. of patients	%	95% CI
Demographics			
Age in years, median (IQR)	66 (52–75)	–	50–68
Female sex	48/200	24	18.4–30.4
Comorbidities and medical history			
Previous hospitalization	94/193	48.7	41.7–56.0
Admission from LTCF	14/198	7.0	4.1–11.4
Diabetes mellitus	35/199	17.6	12.6–23.5
COPD	23/197	11.7	7.7–16.9
Previous myocardial injury	27/192	14.1	9.7–19.6
NYHA score, median (IQR)	1 (1–2)	–	1–1
Chronic liver disease	13/200	6.5	3.5–10.8
Chronic kidney disease	32/198	16.2	11.4–21.8
Chronic intermittent hemodialysis	7/197	3.6	1.6–7.2
Solid neoplasm	29/197	14.7	10.2–20.4
Metastatic solid neoplasm	10/196	5.1	2.6–9.0
Hematological malignancy	29/198	14.6	10.2–20.3
Previous HSCT	9/198	4.5	2.3–8.4
Previous SOT	12/198	6.1	3.4–10.2
HIV infection	3/187	1.6	0.4–4.6
Autoimmune disease	21/198	10.6	6.9–15.7
Age-adjusted Charlson Comorbidity Index, median (IQR)	4 (2–6)	–	2–4
Previous therapy with cefiderocol			
Monotherapy	0/194	0.0	0.0–1.8
Combination therapy	4/194	2.1	0.7–5.2
Previous antibiotic therapy other than cefiderocol			
Previous piperacillin/tazobactam	129/186	69.4	62.4–75.7
Previous piperacillin/tazobactam	85/186	45.7	38.6–53.0
Previous ceftazidime/cefepime	11/186	5.9	3.1–10.3
Previous ceftolozane/tazobactam	16/186	8.6	5.2–13.5
Previous carbapenems	56/186	30.1	23.8–37.0
Previous ceftazidime/avibactam	14/186	7.5	4.4–12.2
Previous meropenem/vaborbactam	1/186	0.5	0.0–2.7
Previous imipenem/relebactam	0/186	0.0	0.0–1.9

Table 1 continued

Variables	No. of patients	%	95% CI
Previous polymyxins	5/186	2.7	1.1–6.0
Previous chemotherapy	23/197	11.7	7.7–16.9
Previous steroid therapy	65/187	34.8	28.0–41.9
Previous therapy with immunosuppressants	26/193	13.5	9.2–19.0
Previous major surgery	75/199	37.7	31.1–44.7
Previous isolation of CR-GNB	67/186	36.0	29.2–43.2
Previous CRE	37/186	19.9	14.6–26.2
Previous CRPA	9/186	4.8	2.4–8.9
Previous CRAB	27/186	14.5	10.0–20.3
Previous MBL-producing CR-GNB	17/186	9.1	5.4–14.1
Variables at cefiderocol initiation			
Days from admission to cefiderocol initiation, median (IQR)	22 (10–40)	–	9–25
ICU stay	102/200	51.0	44.0–58.0
SOFA score, median (IQR)	5 (3–7)	–	2–5
Presence of CVC	140/194	72.2	65.5–78.2
Presence of urinary catheter	155/195	79.5	73.2–84.8
Presence of septic shock	51/198	25.8	20.0–32.2
Presence of ARDS	21/192	10.9	7.1–16.0
Presence of AKI	68/199	34.2	27.8–41.2
Concomitant COVID-19	16/197	8.1	4.9–12.8
Total parenteral nutrition	55/191	28.8	22.7–35.5
Neutropenia	10/200	5.0	2.6–8.8
CRRT	26/198	13.1	8.9–18.5
ECMO	5/198	2.5	1.0–5.6
White blood cell $\times 10^{-3}/\text{mm}^3$ , median (IQR)	10.26 (5.58–15.34)	–	4.9–12.0
Serum C-reactive protein in mg/L, median (IQR)	84.3 (22.8–154.0)	–	69–104
Serum procalcitonin in ng/mL, median (IQR)	1.4 (0.3–5.1)	–	0.9–1.9

Results are presented as No. of patients/Total of patients unless otherwise indicated. Number of missing values per variable were as follows: SOT ( $n = 2/200$ ); hematological malignancy ( $n = 2/200$ ); HSCT ( $n = 2/200$ ); solid neoplasm ( $n = 3/200$ ); metastatic solid neoplasm ( $n = 4/200$ ); HIV infection ( $n = 13/200$ ); previous chemotherapy ( $n = 3/200$ ); COPD ( $n = 3/200$ ); autoimmune disease ( $n = 2/200$ ); diabetes mellitus ( $n = 1/200$ ); chronic kidney disease ( $n = 2/200$ ); previous myocardial injury ( $n = 8/200$ ); previous hospitalization ( $n = 7/200$ ); previous cefiderocol therapy ( $n = 6/200$ ); previous antibiotic therapy other than cefiderocol ( $n = 14/200$ ); previous steroid therapy ( $n = 13/200$ ); previous therapy with immunosuppressants ( $n = 7/200$ ); previous major surgery ( $n = 1/200$ ); Previous isolation of CR-GNB ( $n = 14/200$ ); admission from LTCF ( $n = 2/200$ ); presence of CVC ( $n = 6/200$ ); presence of urinary catheter ( $n = 5/200$ ); ECMO



**Table 1** continued

( $n = 2/200$ ); chronic intermittent hemodialysis ( $n = 3/200$ ); CRRT ( $n = 2/200$ ); presence of septic shock ( $n = 2/200$ ); presence of ARDS ( $n = 8/200$ ); presence of AKI ( $n = 1/200$ ); concomitant COVID-19 ( $n = 3/200$ ); total parenteral nutrition ( $n = 9/200$ ); serum C-reactive protein ( $n = 12/200$ ); serum procalcitonin ( $n = 43/200$ ); HIV infection ( $n = 13/200$ ). No missing values were registered for all other remaining variables

*AKI* acute kidney injury, *ARDS* acute respiratory distress syndrome, *CI* confidence interval, *COPD* chronic obstructive pulmonary disease, *COVID-19* coronavirus disease 2019, *CR-GNB* carbapenem-resistant gram-negative bacteria, *CRAB* carbapenem-resistant *Acinetobacter baumannii*, *CRE* carbapenem-resistant Enterobacterales, *CRPA* carbapenem-resistant *Pseudomonas aeruginosa*, *CRRT* continuous renal replacement therapy, *CVC* central venous catheter, *ECMO* extracorporeal membrane oxygenation, *HIV* human immunodeficiency virus, *HSCT* hematopoietic stem cell transplantation, *ICU* intensive care unit, *IQR* interquartile range, *MBL* metallo  $\beta$ -lactamases, *NYHA* New York Heart Association, *LTCF* long-term care facility, *SOFA* sequential organ failure assessment, *SOT* solid organ transplantation

## RESULTS

The demographic and clinical characteristics at time of cefiderocol initiation of the 200 included patients (enrolled from August 2022 to September 2023) are displayed in Table 1. Their median age was 66 years (IQR 52–75) and 76% were male (152/200). The characteristics of cefiderocol treatments are reported in Table 2. Overall, cefiderocol was started as empirical and targeted therapy in 27.5% (55/200) and 72.5% (145/200) of cases, respectively.

### Empirical Therapy

As shown in Supplementary Table S1, the most frequently reported indications for empirical cefiderocol therapy were sepsis (33/55, 65.5%) and lower respiratory tract infection (10/55, 18.2%). In 28/55 cases of empirical therapy (50.9%), GNB grew from cultures collected at the time of treatment initiation, of which 14/26 (54%, missing = 2/28) were CR-GNB, mostly *A. baumannii*, *P. aeruginosa*, and *Klebsiella* spp. (all 4/14 each, 28.6%). Among identified CR-GNB, production of carbapenemase/presence of carbapenemase-encoding genes was assessed in 12/14 cases (85.7%), resulting positive in 5/12 cases (41.7%), of which 3/5 were positive for MBL (60.0%). Identified MBL-producers were 1 VIM-producing *Escherichia* spp., 1 NDM-producing *Escherichia* spp., and 1 NDM-producing

*K. pneumoniae*. The other identified carbapenemase-producing organisms were 1 KPC-producing *K. pneumoniae* and 1 OXA-producing *A. baumannii*. Cefiderocol AST was performed among 8/14 identified CR-GNB (including 2/3 MBL-producers), with one *A. baumannii* isolate being non-wild-type (production of carbapenemase/presence of carbapenemase-encoding genes investigated but not detected). The other 7 isolates were all wild-type/susceptible to cefiderocol.

### Targeted Therapy

Regarding targeted cefiderocol therapy (Supplementary Table S1), the most frequently reported indications were lower respiratory tract infection (63/145, 43.4%) and bloodstream infection (56/145, 38.6%). Overall, 170 GNB causative agents were retrieved from 145 infections treated with targeted cefiderocol therapy (see details in the legend of Supplementary Table S1). Infections treated with targeted cefiderocol therapy were most frequently monomicrobial (122/145, 84.1%), mainly caused by *A. baumannii* (67/122, 54.9%), *P. aeruginosa* (25/122, 20.5%), and *Klebsiella* spp. (18/122, 14.8%). Rates of carbapenem resistance among these infections were 91.9% (57/62 tested isolates), 95.5% (21/22 tested isolates), and 93.8% (15/16 tested isolates) for *A. baumannii*, *P. aeruginosa*, and *Klebsiella* spp., respectively. Production of carbapenemase/presence of carbapenemase-encoding genes was

**Table 2** Characteristics of cefiderocol therapy

Variables <sup>a</sup>	No. of patients	%	95% CI
Type of anti-CR-GNB therapy <sup>b</sup>			
Cefiderocol monotherapy	101/200	50.5	43.5–57.5
Combination therapy <sup>c,d,e,f,g</sup>	99/200	49.5	42.5–56.5
Timing of cefiderocol therapy			
Empirical therapy <sup>h</sup>	55/200	27.5	21.6–34.2
Empirical cefiderocol monotherapy	23/55	41.8	28.7–55.5
Empirical combination therapy	32/55	58.2	44.5–71.3
Targeted therapy <sup>l</sup>	145/200	72.5	65.8–78.4
Targeted cefiderocol monotherapy	69/145	47.6	39.2–55.9
Targeted combination therapy	76/145	52.4	44.1–60.8
Targeted therapy for Enterobacterales infection <sup>i</sup>			
Cefiderocol monotherapy	14/26	53.8	34.1–71.8
Combination therapy	12/26	46.2	28.2–65.9
Targeted therapy for <i>Pseudomonas aeruginosa</i> infection <sup>i</sup>			
Cefiderocol monotherapy	14/25	56.0	35.5–74.8
Combination therapy	11/25	44.0	25.2–64.5
Targeted therapy for <i>Acinetobacter baumannii</i> infection <sup>i</sup>			
Cefiderocol monotherapy	29/67	43.3	31.9–55.3
Combination therapy	38/67	56.7	44.7–68.1
Targeted therapy for <i>Stenotrophomonas maltophilia</i> infection <sup>i</sup>			
Cefiderocol monotherapy	3/4	75.5	24.9–98.7
Combination therapy	1/4	25.5	1.3–75.1
Targeted therapy for MBL-producing Gram-negative infection <sup>i,k,l</sup>			
Cefiderocol monotherapy	13/22	59.1	38.3–78.3
Combination therapy	9/22	40.9	21.7–61.7
Initial cefiderocol dosage according to estimated CL <sub>Cr</sub> or hemodialytic treatment			
CL <sub>Cr</sub> > = 120 mL/min	34/200	17.0	12.1–22.9
CL <sub>Cr</sub> 60 to 119 mL/min	97/200	48.5	41.5–55.5
CL <sub>Cr</sub> 30 to 59 mL/min	37/200	18.5	13.6–24.4
CL <sub>Cr</sub> 15 to 29 mL/min	18/200	9.0	5.6–13.8
CL <sub>Cr</sub> < 15 mL/min	6/200	3.0	1.3–6.3
IHD	3/200	1.5	0.4–4.3

Table 2 continued

Variables <sup>a</sup>	No. of patients	%	95% CI
CRRT	5/200	2.5	1.0–5.6
Change of dosage during therapy due to changes in $CL_{Cr}$ <sup>m</sup>	11/196	5.6	2.9–9.8

CI confidence interval,  $CL_{Cr}$  creatinine clearance, CR-GNB carbapenem-resistant gram-negative bacteria, IHD intermittent haemodialysis, MBL metallo  $\beta$ -lactamases, CRRT continuous renal replacement therapy

<sup>a</sup>Results are presented as No. of patients/Total of patients unless otherwise indicated

<sup>b</sup>Anti-CR-GNB combination was defined as treatment with ceftiderocol in combination with at least one of the following agents: aminoglycosides; fosfomycin; tigecycline (with the exception of targeted therapy of *P. aeruginosa* infections); polymyxins; sulbactam or ampicillin/sulbactam (as empirical treatment of as targeted therapy for *A. baumannii* infections)

<sup>c</sup>Agents combined with ceftiderocol for empirical therapy: fosfomycin ( $n = 9$ ); tigecycline ( $n = 6$ ); colistin ( $n = 3$ ); colistin plus tigecycline ( $n = 2$ ); aminoglycoside ( $n = 1$ ); aminoglycoside plus fosfomycin ( $n = 1$ ); ampicillin/sulbactam plus tigecycline ( $n = 1$ )

<sup>d</sup>Agents combined with ceftiderocol for targeted therapy of Enterobacterales infection: fosfomycin ( $n = 5$ ); tigecycline ( $n = 4$ ); aminoglycoside ( $n = 2$ ); aminoglycoside plus tigecycline ( $n = 1$ )

<sup>e</sup>Agents combined with ceftiderocol for targeted therapy of *Pseudomonas aeruginosa* infections: fosfomycin ( $n = 8$ ); aminoglycosides ( $n = 2$ ); colistin ( $n = 1$ )

<sup>f</sup>Agents combined with ceftiderocol for targeted therapy of *Acinetobacter baumannii* infections: fosfomycin ( $n = 13$ ); ampicillin/sulbactam ( $n = 8$ ); colistin ( $n = 8$ ); tigecycline ( $n = 5$ ); aminoglycosides ( $n = 1$ ); aminoglycosides plus tigecycline ( $n = 1$ ); ampicillin/sulbactam plus tigecycline ( $n = 1$ ); colistin plus tigecycline ( $n = 1$ )

<sup>g</sup>Agents combined with ceftiderocol for targeted therapy of *Stenotrophomonas maltophilia* infections: colistin plus tigecycline ( $n = 1$ )

<sup>h</sup>In 28/55 cases of empirical therapy (51%), a Gram-negative etiological agent grew from cultures collected at the time of treatment initiation, of which 14/26 (54%, missing = 2/28) were carbapenem-resistant (of them 3/12, 25%, were MBL producers, missing = 2/14)

<sup>i</sup>Ceftiderocol therapy started after identification of the causative agent

<sup>j</sup>Analyses limited to Infection by only one Gram negative genus (with the exception of Enterobacterales infection, for which concomitant infection by more than one member of the Enterobacterales order was also considered)

<sup>k</sup>Type of MBL enzyme ( $n = 20$ ): NDM ( $n = 12$ ); VIM ( $n = 19$ ); NDM ( $n = 3$ )

<sup>l</sup>Type of MBL-producing causative agent: *P. aeruginosa* ( $n = 12$ ); Enterobacterales ( $n = 10$ )

<sup>m</sup>Changes in  $CL_{Cr}$  deemed as not related to ceftiderocol therapy. Presence of missing values ( $n = 4/200$ )

assessed in 121/170 isolates (71.2%), resulting positive in 51/121 cases (42%), of which 26/51 (51.0%) were MBL-producing (21.5%), mostly VIM-producing *P. aeruginosa* (12/26, 46.2%), followed by NDM-producing *Klebsiella* spp. (5/26, 19.2%) and VIM-producing *Enterobacter* spp. (4/26, 15.4%). Other detected carbapenemases were KPC-type ( $n = 11$ ), OXA-type ( $n = 11$ ), and not specified ( $n = 3$ ). Ceftiderocol AST was performed in 74/170 isolates (43.5%) from infections receiving targeted ceftiderocol therapy (including 13/26 MBL producers, 50.0%), with 3/57 tested *A. baumannii* isolates (5.3%) being non-wild-type (production of carbapenemase/presence of carbapenemase-encoding genes was

investigated but not detected in all three isolates). No other isolates were resistant/non-wild-type. The demographic and clinical characteristics at time of targeted ceftiderocol initiation, stratified according to the different causative agents of infection, are available in supplementary Table S2.

### Monotherapy and Combined Therapy

Ceftiderocol was administered as monotherapy and as combined therapy for CR- GNB infections in 101/200 (50.5%) and 99/200 (49.5%) cases, respectively (Table 2). Supplementary Table S3

reports the results of univariable analyses of factors associated with administration of ceftiderocol as a combined therapy for CR-GNB infections, whereas factors retaining an independent association with combined therapy for CR-GNB infections in the final multivariable models are presented in Table 3. In model A, previous isolation of carbapenem-resistant *Acinetobacter baumannii* [odds ratio (OR) 2.56, 95% CI 1.01–6.46,  $p = 0.047$ ] and previous hematopoietic stem cell transplantation (OR 8.73, 95% CI 1.05–72.54,  $p = 0.045$ ) were associated with administration of ceftiderocol within a combined regimen, whereas chronic kidney disease was associated with ceftiderocol monotherapy (OR 0.38 for combined regimen, 95% CI 0.16–0.91,  $p = 0.029$ ). In model B, including the same variables of model A plus center as a random effect, the direction of fixed effects was the same registered in model A.

### Cure Rates and Tolerability

Cure rates at days 14, 21, and 28 are descriptively summarized in supplementary Table S4, divided by pathogen and combination or monotherapy. As shown in Fig. 1, the cumulative 30-day mortality was 19.8% (95% CI 7.0–37.4) in patients receiving targeted ceftiderocol therapy for Enterobacterales infection (panel A), 45.0% (95% CI 32.4–56.8) in those receiving targeted ceftiderocol therapy for *A. baumannii* infection (panel B), 20.7% (95% CI 7.3–38.7) in patients receiving targeted ceftiderocol therapy for *P. aeruginosa* infection (panel C), and 22.7% (95% CI 8.0–41.9) in patients receiving targeted ceftiderocol therapy for MBL-producing GNB (panel D). Overall, 4/200 patients (2.0%) experienced a suspected drug-related AE during ceftiderocol administration. Two patients developed mild skin rash, one patient experienced hyperchromic urine and moderate increase in liver enzymes values, and one patient developed status epilepticus categorized as SAE. Ceftiderocol was discontinued in 2/4 patients (50.0%) experiencing AE.

**Table 3** Multivariable analysis of factors associated with use of ceftiderocol in combination with other anti-CR-GNB agents<sup>a</sup>

Model A (AIC 265.28)	OR (95% CI)	<i>P</i>
Chronic kidney disease	0.38 (0.16–0.91)	<b>0.029</b>
Previous HSCT	8.73 (1.05–72.54)	<b>0.045</b>
Previous CRAB	2.56 (1.01–6.46)	<b>0.047</b>
ICU stay	1.57 (0.83–2.98)	0.168
Presence of septic shock	1.77 (0.86–3.62)	0.120
Urinary tract infection	0.21 (0.02–1.99)	0.174
Model B <sup>b</sup> (AIC 289.36)	OR (95% CI)	<i>P</i>
Chronic kidney disease	0.81 (0.68–0.97)	<b>0.024</b>
Previous HSCT	1.50 (1.09–2.07)	<b>0.013</b>
Previous CRAB	1.21 (0.99–1.48)	0.062
ICU stay	1.12 (0.97–1.30)	0.124
Presence of septic shock	1.13 (0.96–1.33)	0.137
Urinary tract infection	0.77 (0.53–1.11)	0.156

Analyses conducted after multiple imputation (see study methods). Values in bold are significant,  $p < 0.05$

*AIC* Akaike information criterion, *CI* confidence interval, *CR-GNB* carbapenem-resistant Gram-negative bacteria, *OR* odds ratio, *HSCT* hematopoietic stem cell transplantation, *CRAB* carbapenem-resistant *Acinetobacter baumannii*, *COVID-19* coronavirus disease 2019

<sup>a</sup>Anti-CR-GNB combination was defined as treatment with ceftiderocol in combination with at least one of the following agents: aminoglycosides; fosfomycin; tigecycline (with the exception of targeted therapy of *P. aeruginosa* infections); polymyxins; sulbactam or ampicillin/sulbactam (as empirical treatment of as targeted therapy for *A. baumannii* infections)

<sup>b</sup>Model B also included center as a random effect. For complete details, see Methods

## DISCUSSION

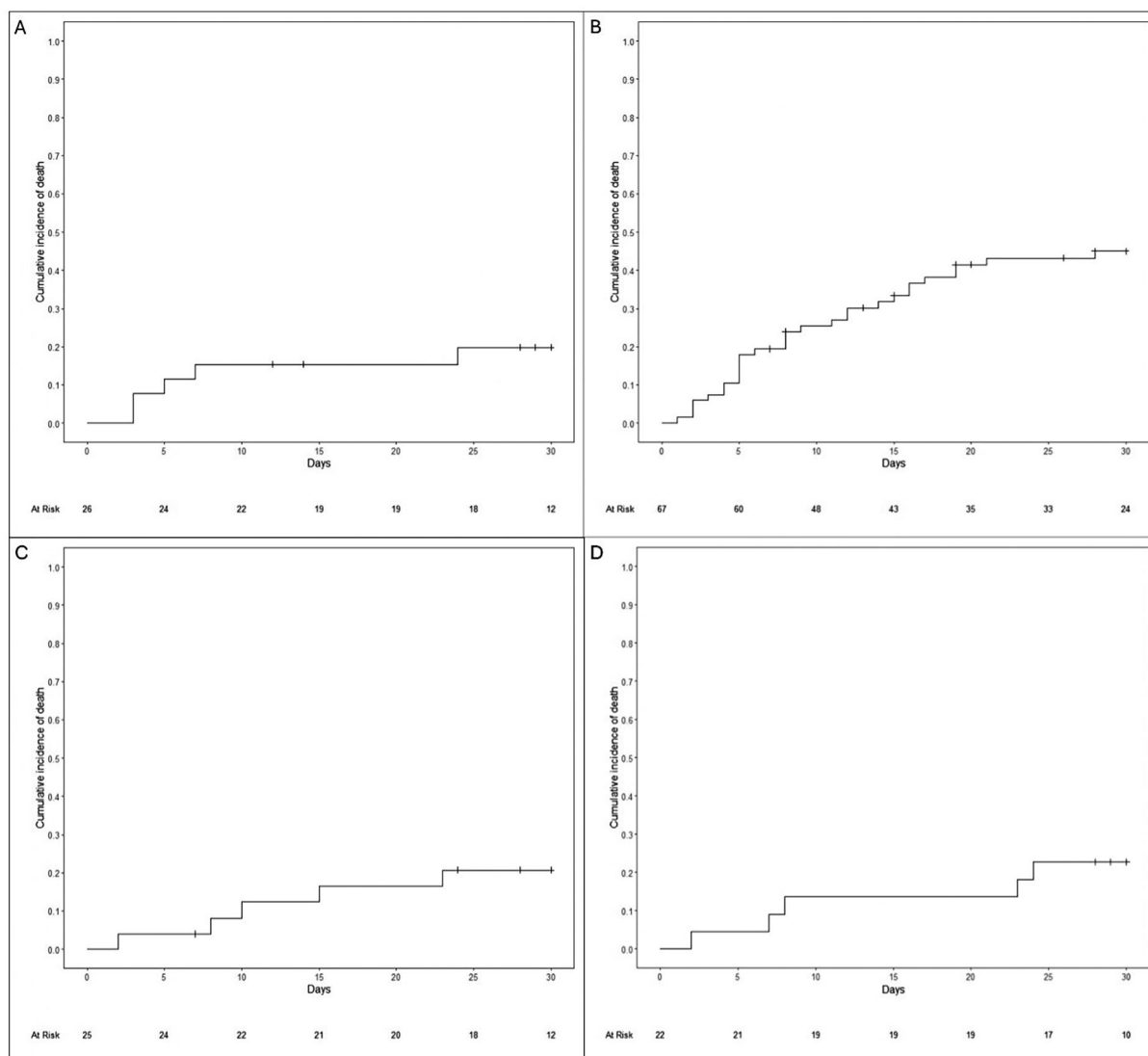
In this multicenter study, ceftiderocol was mostly administered as targeted therapy (72.5% vs. 27.5% as empirical therapy), mainly for lower respiratory tract infections and bloodstream infections caused by *A. baumannii*, followed by

*P. aeruginosa*. Cefiderocol was almost equally administered as monotherapy or as combination therapy (50.5% vs. 49.5%).

Most available studies on the use of cefiderocol in real life are focused on the targeted treatment of infections caused by various or specific CR-GNB [7, 10–15, 17, 18, 21–24, 26–35]. While useful for reporting cure rates in specific infections, these studies were unable to generally depict how cefiderocol is currently used by physicians, having been designed for other purposes. In our opinion, this topic is of interest considering the following: (1) current international and national guidelines/guidance documents provide recommendations on the use of cefiderocol for targeted treatment, while there is still no clear guidance regarding its empirical use [47–51]; and (2) various studies have reported use of cefiderocol either in combination or as monotherapy, thus we deemed it of interest to descriptively explore their relative frequency [30, 52]. Regarding the first point, in our study, cefiderocol was mainly administered as targeted therapy. However, the proportion of empirical therapies was not negligible, representing more than one-quarter of all cefiderocol prescriptions. This could reflect the physicians' willingness to prescribe an active therapy without delay in patients with severe infections and risk factors for CR-GNB, in areas where the prevalence rate of CR-GNB, especially MBL producers, is high. Overall, these results suggest the need for standardized consensus regarding the use of cefiderocol within empirical therapeutic algorithms, in order to maximize proper indications in line with antimicrobial stewardship principles, but also to not perilously delay treatment in patients with infections caused by cefiderocol-susceptible CR-GNB.

Regarding the use of cefiderocol as monotherapy or within combined regimens, our results could reflect the current lack of solid evidence on this aspect. Against this background, the possible perception of some clinicians of potential reduced activity of cefiderocol in patients with severe *A. baumannii* infections could have prompted to consider combination therapy in some cases. This is in line with the independent association we found in multi-variable models between previous isolation of

carbapenem-resistant *A. baumannii* and use of cefiderocol in combined regimens, although it is also of note that no association was found between targeted treatment of *A. baumannii* infections and use of cefiderocol within combined regimens. Furthermore, in our opinion, this controversial point merits some additional considerations. First, it should be considered that the apparently higher mortality rates (50% [21/42] vs. 18% [3/17] for best available therapy, as 49-day mortality) in patients with *A. baumannii* infections treated with cefiderocol in the CREDIBLE-CR RCT were possibly confounded by factors such as severity of presentation and baseline diseases (e.g., severe renal dysfunction, ongoing shock, shock 31 days before randomization, and ICU stay at randomization were more frequent in patients treated with cefiderocol than in those receiving best available therapy) [5]. A similar situation was present in our study, in which mortality of *A. baumannii* infection was higher than mortality of infections caused by other organisms. Indeed, severe clinical presentation was more frequent in patients with *A. baumannii* infections than in those with infections caused by other organisms (e.g., septic shock was present in 29% of patients receiving cefiderocol for the targeted treatment of *A. baumannii* infections vs. 15.4% and 16.0% in patients receiving cefiderocol for the targeted treatment of Enterobacterales infections and *P. aeruginosa* infections, respectively, as shown in supplementary Table S2). Second, it should also be noted that, although mortality rates of cefiderocol-treated *A. baumannii* infections in some other studies were in line with our results [13, 15, 18, 23, 24, 33], in many other studies, including the APEKS-NP randomized controlled trial, they were far lower, ranging from 18 to 37% [3, 7, 12, 14, 16, 27, 29, 32, 53]. Third, lower mortality of cefiderocol-treated *A. baumannii* infections in comparison with non-cefiderocol-based regimens was suggested by recent meta-analyses including data from both CREDIBLE-CR and observational studies [54–56]. Lastly, among isolates subjected to cefiderocol AST, non-susceptibility (i.e., non-wild-type) was exclusively detected in *A. baumannii*, a finding consistent with that of a recent meta-analysis reporting the highest prevalence of cefiderocol non-susceptibility for



*A. baumannii*, compared to other major Gram-negative pathogens [57]. Overall, considering all the above, in our opinion, the relevant research question for future studies should regard which subgroups of patients with *A. baumannii* infections could benefit the most from ceftiderocol monotherapy versus combination therapy (in turn possibly influencing the choice of combined regimens). Regarding other results, the significant association between previous HSCT and administration of ceftiderocol within a combined regimen deserves further investigations. Indeed, while the perception of severe baseline conditions and increased mortality due to immunosuppression connected to baseline disease

and/or its treatment could have played a role in influencing the choice of combination, also for targeted treatment, the subgroup of patients who underwent HSCT was small in this preliminary analysis ( $n = 9$ ), thus a spurious association due to chance alone cannot be definitely ruled out pending further dedicated data. By contrast, the association between chronic kidney disease and ceftiderocol monotherapy could reflect the physicians' decision, at least in some cases, not to administer ceftiderocol together with potentially nephrotoxic agents (e.g., polymyxins, aminoglycosides) in patients with already impaired renal function, or the fact that these patients are at increased risk of urinary tract infections

◀**Fig. 1** Cumulative mortality up to day 30 in patients receiving targeted ceftiderocol therapy for Enterobacterales infection (panel A), *Acinetobacter baumannii* infection (panel B), *Pseudomonas aeruginosa* infection (panel C), and MBL-producing Gram-negative bacteria (panel D). MBL metallo- $\beta$ -lactamases. Analyses limited to Infection by only one Gram negative genus (with the exception of Enterobacterales infection, for which concomitant infection by more than one member of the Enterobacterales order was also considered). The time of origin was set at the day of ceftiderocol initiation. Death was the event of interest and right-censoring was applied at the end of follow-up (hospital discharge or day 30, whichever came first). Site/s of Enterobacterales infection: bloodstream infection ( $n = 12$ ); lower respiratory tract infection ( $n = 7$ ); urinary tract infection ( $n = 2$ ); skin and soft tissue infection ( $n = 1$ ); intra-abdominal infection ( $n = 1$ ); intra-abdominal infection plus bloodstream infection ( $n = 1$ ); lower respiratory tract infection plus bloodstream infection ( $n = 1$ ); urinary tract infection plus bloodstream infection ( $n = 1$ ). Site/s of *P. aeruginosa* infection ( $n = 25$ ): lower respiratory tract infection ( $n = 12$ ); bloodstream infection ( $n = 6$ ); urinary tract infection ( $n = 2$ ); bone and joint infection plus bloodstream infection ( $n = 1$ ); intra-abdominal infection ( $n = 1$ ); lower respiratory tract infection plus bloodstream infection ( $n = 1$ ); skin and soft tissue infection ( $n = 1$ ); skin and soft tissue infection plus bloodstream infection ( $n = 1$ ). Site/s of *A. baumannii* infection ( $n = 67$ ): lower respiratory tract infection ( $n = 30$ ); bloodstream infection ( $n = 29$ ); bone and joint infection ( $n = 2$ ); urinary tract infection ( $n = 2$ ); intra-abdominal infection ( $n = 1$ ); lower respiratory tract infection plus bloodstream infection ( $n = 1$ ); skin and soft tissue infection ( $n = 1$ ); site/s not reported ( $n = 1$ ). Site/s of MBL-producing Gram-negative infection ( $n = 22$ ): lower respiratory tract infection ( $n = 8$ ); bloodstream infection ( $n = 5$ ); urinary tract infection ( $n = 3$ ); intra-abdominal infection ( $n = 2$ ); skin and soft tissue infection ( $n = 2$ ); lower respiratory tract infection plus bloodstream infection ( $n = 1$ ); skin and soft tissue infection plus bloodstream infection ( $n = 1$ ). Type of MBL enzyme ( $n = 20$ ): NDM ( $n = 12$ ); VIM ( $n = 19$ ); NDM ( $n = 3$ ). Type of MBL-producing causative agent: *P. aeruginosa* ( $n = 12$ ); Enterobacterales ( $n = 10$ )

(for which a trend towards preference of monotherapy was observed in our study, albeit not statistically significant, as reported in Supplementary Table S3).

This present preliminary analysis of the CEFI-SITA study has some limitations to be acknowledged. The first is that, while we reported

cumulative mortality in patients receiving ceftiderocol in subgroups according to different causative organisms, the analysis was not primarily designed with this aim, thus the resulting unadjusted estimates (e.g., not adjusted for appropriateness of targeted therapy based on in vitro activity) should be interpreted with caution pending further data. However, the low cumulative mortality registered in infections by Enterobacterales, *P. aeruginosa*, and MBL producers is worth mentioning, and is in line with our previous findings of the possibly changing landscape in the treatment of CR-GNB registered in the past few years [58]. A second limitation is connected to the small sample size of patients treated with empirical ceftiderocol. Indeed, although the registered 27.5% proportion of patients receiving empirical ceftiderocol is solidly based on sample size estimates, subgroup proportions (within empirical therapy) have a larger degree of uncertainty and may require confirmation in further dedicated studies. Of note, this also includes the analysis of either crude or adjusted mortality in patients receiving empirical ceftiderocol with subsequent isolation of CR-GNB as etiological agents, which has been deferred to a later phase of CEFI-SITA due to the limited sample size of this subgroup in this preliminary analysis. Third, no standardized microbiological approach was used for ceftiderocol AST across participating centers, due to the observational, descriptive representation of daily routine practice. Fourth, owing to the expression of genes not commonly included among targets of rapid molecular tests, production of carbapenemases might have not been thoroughly evaluated in some cases (e.g., OXA-23 in *A. baumannii*). Additional studies employing WGS are therefore needed to decipher the molecular bases of carbapenem resistance in study isolates.

## CONCLUSIONS

Ceftiderocol is mainly used for targeted treatment in Italian hospitals, although empirical therapies account for more than 25% of prescriptions and should require dedicated standardization and guidance. The almost equal distribution of

cefiderocol monotherapy and cefiderocol-based combination therapies underlines the need for further study to ascertain possible differences in efficacy between the two approaches.

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**Data Availability.** The data presented in this study will be available from the corresponding author on reasonable request and provided



all regulatory and privacy requirements are fulfilled.

### Declarations

**Conflict of Interest.** Outside the submitted work, Daniele Roberto Giacobbe reports investigator-initiated grants from Pfizer, BioMérieux, and Gilead Italia, and speaker/advisory board fees from Pfizer, Menarini, and Tillotts Pharma. Outside the submitted work, Matteo Bassetti has received funding for scientific advisory boards, travel, and speaker honoraria from Angelini, Astellas, Bayer, bioMérieux, Cidara, Cipla, Gilead, Menarini, MSD, Pfizer, Shionogi, Tetrphase, and Nabriva. Outside the submitted work, Vincenzo di Pilato reports travel grants from Arrow Diagnostic, and speaker honoraria from A.D.A. Outside the submitted work, Emanuele Pontali has received funding for scientific advisory boards, travel, and/or speaker honoraria from Abbvie, Angelini, Gilead, Janssen, MSD, and Viiv. Outside the submitted work, Andrea Lombardi reports travel grants from Shionogi. Outside the submitted work, Rosario Cultrera has received funding for scientific advisory boards, travel, and speaker honoraria from Angelini, Menarini, MSD, Pfizer, Shionogi, and TRX Italy. Outside the submitted work, Andrea Cortegiani reports fees for lectures/ advisory board membership from Gilead, MSD, Mundipharma, Pfizer, Shionogi. The other authors have no conflicts of interests to disclose.

**Ethical Approval.** The MULTI-SITA project was approved by the ethics committee of the coordinating center (Liguria Region Ethics Committee, registry number 390/2020). The amendment authorizing the conduct of the CEFI-SITA study within the MULTI-SITA project was approved by the Liguria Region Ethics Committee on 12 April 2022. The other participating centers followed the local ethical committees requirements and started to enroll patients prospectively once activated. All conscious patients at time of enrollment signed an informed consent to participate in the study. A waiver of informed consent for data collection from unconscious patients at the time of enrollment due to severe clinical

conditions was obtained within the ethics committee approval, in line with the observational nature of the analyses and in order not to bias research results towards high cure rates and low mortality prejudicing scientific validity.

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