Pharmacokinetics and intrapulmonary concentrations of high-dose tigecycline in critically ill patients with severe infections

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Abstract

Background

In critically ill patients, the use of high tigecycline dosages (HD TGC) (200 mg/day) has been recently increasing but few pharmacokinetic/pharmacodynamic (PK/PD) data are available. We investigated plasmatic and pulmonary concentrations of HD TGC in the treatment of severe infections.

Methods

This was a single centre, prospective, observational study that was conducted in the twenty-bed mixed ICU of a 1,500- bed teaching hospital in Rome, Italy. In all patients admitted to the ICU between 2015 and 2018, who received TGC (200 mg loading dose, then 100 mg q12) for the treatment of documented infections, serial blood samples were collected to measure TGC concentrations. Moreover, epithelial lining fluid (ELF) concentrations were determined in patients with nosocomial pneumonia.

Results

Among the 32 non-obese patients included, 11 had a treatment failure, while the other 21 subjects successfully eradicated the infection. There were no between-group differences in terms of demographic aspects and main comorbidities. In nosocomial pneumonia, for a target AUC0-24/MIC of 4.5, 75% of the patients would be successfully treated in presence of 0.5 mg/L MIC value and all the patients obtained the PK target with MIC≤0.12 mg/mL. In intra-abdominal infections, for a target AUC0-24/MIC of 6.96, at least 50% of the patients would be adequately treated against bacteria with MIC≤0.5 mcg/mL. Finally, in skin and soft-tissue infections, for a target AUC0-24/MIC of 17.9 only 25% of the patients obtained the PK target at MIC values of 0.5 mg/L and less than 10% were adequately treated against germs with MIC value ≥1 mcg/ml. HD TGC showed a relevant pulmonary penetration with a median and IQR ELF/plasma ratio (%) of 152.9 [73.5-386.8].

Conclusions

The use of HD TGC is associated with satisfactory plasmatic and pulmonary concentrations for the treatment of severe infections due to fully susceptible bacteria (MIC<0.5 mg/L). Even higher dosages and combination strategies are required in presence of difficult to treat pathogens.
Background

Tigecycline (TGC), the first antimicrobial of glycyclcycline class, has shown an expanded-spectrum activity against gram-positive, gram-negative, aerobic, anaerobic and atypical bacterial species, including antibiotic-resistant strains [1]. Indeed, it has been demonstrated that methicillin-resistant Staphylococcus aureus (MRSA), penicillin-resistant Streptococcus pneumoniae, vancomycin-resistant enterococci (VRE), extended-spectrum β-lactamase (ESBL)-producing Gram-negative bacilli and extensively drug-resistant (XDR) Acinetobacter baumannii are susceptible to TGC [2–4].

TGC is currently approved by the U.S. food and Drug Administration (FDA) for complicated skin and skin-structure infections, complicated intra-abdominal infections, community-acquired pneumonia with an initial dose of 100 mg, followed by 50 mg every 12 hours. Nevertheless, due to an increased risk of death compared to other antimicrobials, its use has recently been restricted in situations when alternative treatments are not suitable [5].

However, the alarming increase in antimicrobial resistance among the nosocomial pathogens is leading the clinicians to consider the use of TGC as an important therapy in the management of difficult to treat infection, particularly in critically ill patients. This is also supported by recent studies suggesting that previous failures of TGC therapy in critically ill patients were likely due to a drug underdosage [6, 7] and that standard doses provide serum concentrations that are below the minimum-inhibitory concentrations (MICs) of most MDR pathogens. Moreover, it has been reported an increased effectiveness of high-dose TGC (HD TGC) regimen to improve the clinical outcome, without safety issues [8–11].

Therefore, we designed this prospective observational study to describe the pharmacokinetic/pharmacodynamic (PK/PD) profile of HD TGC in a cohort of critically ill patients with severe infections.

Methods

Patients and study design

This was a prospective, observational study that was performed between 2015 and 2018 in the 20-bed ICU of a 1,500-bed teaching hospital in Rome, Italy. The protocol was approved by the Catholic University’s Ethical Committee (approval number Prot.sf 8431/13). Written informed consent was
obtained from the patients’ legally authorized representative. Critically ill adult patients were considered eligible for the study when the attending physician prescribed TGC as empirical treatment (within 12 h from microbiological sampling) of a possible MDR infection, or as targeted therapy based on definitive results, in the absence of any exclusion criteria: known TGC allergy, creatinine clearance less than 40 mL/min (calculated according to the Cockcroft-Gault formula) apart from those ones who were anuric and on continuous renal replacement therapy (CRRT), hyperbilirubinemia (bilirubin level higher than 3 mg/dL), severe hepatic failure (Child-Pugh C), little chance of survival as defined by the Simplified Acute Physiology 2 (SAPS 2) score > 80, concomitant treatment with other drugs that can potentially interfere with TGC. TGC was administered intravenously at loading dose (LD) of 200 mg over 30-min, followed by 100 mg over 30-min bid. On day four after the commencement of the HD TGC, at steady state, pharmacokinetic analyses of the study group were performed. Clinical and demographic data were recorded upon enrolment. Safety and adverse events were determined through the observed biochemical abnormalities, documented according to the Department of Health and Human Services–Common Terminology Criteria for Adverse Events (DHHS-CTCAE v.3.0) classification [12].

Clinical cure was defined as the complete resolution of all signs and symptoms of the infection by the end of TGC therapy. In case of ventilator associated pneumonia (VAP), improvement or lack of progression of all abnormalities on chest radiographs was also required [13]. Clinical outcomes were independently evaluated by two physicians (GDP, MSV) who were blinded to the treatment. When judgments were discordant (about 5% of patients), the reviewers reassessed the data and reached a consensus decision. Adequate source control included drainage of infected fluid collections, debridement of infected solid tissue, removal of devices/foreign bodies, and definitive measures to correct anatomic derangements resulting in on-going microbial contamination and to restore optimal function within 48 h after diagnosis [14].

Sample collection
Blood samples were collected after the seventh dose (on day 4 of treatment) at T0 (immediately before the initiation of the infusion) and 1, 1.5, 2, 4, 6, 8, 10, 12 after the start of infusion. According
to patients’ respiratory status, one mini-bronchoalveolar lavage (BAL) (40 mL sterile 0.9% saline solution was blindly instilled through a telescopic catheter and immediately aspirated in a trap) was performed at steady state.

**Preparation of stock solution and calibration standard**

Stock solution of TGC and the internal standard (IS), propranolol hydrochloride, were prepared by dissolving accurately-weighed amounts of each compound in MeOH to obtain a final concentration of 0.1 mg/ml. Calibration standards were prepared by diluting stock solutions of TGC in drug-free human plasma to yield TGC concentrations of 5000, 2500, 1250, 625, 312.5, 156.25, 78.125, 39.1, 19.5 and 9.76 ng/ml.

**Sample preparation**

**Tigecycline liquid/liquid extraction from plasma samples**

Liquid/liquid extraction was employed for purification and concentration of TGC from plasma samples. Briefly 25 µl of 0.1 mg/ml IS and then 50 µl of mobile phase A (H₂O 0.1% HCOOH) were added to 250 µl of the plasma sample. After a rapid vortex mixing, 2.5 ml of Methyl-tert butyl ether was added to all samples, the mixture was vortexed for 5 min and then centrifuged at 3200 rpm for 10 min at 10 °C. The upper organic layer was separated and evaporated to dryness at 40 °C. The residue was then reconstituted in 100 µl of mobile phase B [ACN (+ 0.1%HCOOH)] solution and transferred to an auto-sampler vial.

**Tigecycline solid-phase extraction from BAL samples**

Purification and concentration of TGC from BAL samples were carried out by solid-phase extraction (SPE). Briefly, 500 µL of BAL samples were mixed to 500 µL of 1%TFA in water, and 6 µL of concentrated TFA were subsequently added. After 5 min of vortex, the samples were centrifuged at 2500 rpm for 10 min; thereafter, the sur natants were collected and loaded onto SPE cartridges (OASIS HLB 1 cc; Waters) arranged on an extraction manifold (Vac-Elut, Analytichem International). The SPE cartridges were sequentially pre-conditioned with 1 ml of MeOH given twice, and subsequently with 1 ml of water given twice. The loaded cartridges were washed with 1 ml of 1%TFA in water, and eluted with 1 ml of MEOH. The eluted solutions were evaporated to dryness at 40 °C. The residues were reconstituted in 100 µl of solution and then transferred to an auto-sampler vial.
Chromatographic and Mass-Spectrometric Conditions
The chromatographic separation was performed on an AQUITY UPLC system (Waters Corp., Milford, MA, USA) with cooled auto-sampler and column oven for temperature control. An AQUITY UPLC BEH C18 column (1.7 um; 2.1 × 50 mm; Waters Corp., Milford, MA, USA) was employed and the column temperature was maintained at 40 °C. The elution was performed at a flow rate of 0.6 mL/min with a run time of 3 min. The UPLC was connected to a triple quadrupole tandem mass detector (TQD) (Waters Corp., Milford, MA, USA) with an electrospray ionization (ESI) source for the mass spectrometric detection. The ESI source was set in positive ion mode and the quantification was performed using multi reaction monitoring (MRM) mode for the most suitable mass transitions. The optimal mass spectroscopy (MS) parameters were showed in Table 1. Data were processed using MassLynx software with a QuanLynx program version 4.1 (Waters Corp., Milford, MA, USA).
## Table 1
Baseline patients characteristics

| Demographics and comorbidities                      | Total cohort (n = 32) | Treatment Failure (n = 11) | Treatment Success (n = 21) | p value |
|------------------------------------------------------|-----------------------|---------------------------|---------------------------|---------|
| **Age, years**                                       | 56 [46-68.5]          | 55 [49.75-71]             | 56 [45-68.25]             | 0.75    |
| **Male sex, N (%)**                                  | 17 (53.1)             | 5 (45.5)                  | 12 (57.1)                 | 0.8     |
| **Weight, (Kg)**                                     | 76.5 [60-90]          | 75 [67.8-80]              | 90 [60-100]               | 0.45    |
| **Albumin, (g/dL)**                                  | 23 [21.5-26.5]        | 22 [19.25-26.25]          | 24 [22.75-26.5]           | 0.17    |
| **Fluid balance, (mL)**                              | +762.9 [393-1124.2]   | +3332 [-358.5-4112]       | 616.3 [-2592.7]           | 0.5     |
| **SAPS II score**                                    | 53.5 [44.5-67.5]      | 61 [44.7-66.5]            | 52 [43.5-67.5]            | 0.92    |
| **COPD, N (%)**                                      | 6 (18.75)             | 3 (27.3)                  | 3 (14.3)                  | 0.39    |
| **Chronic renal failure, N (%)**                     | 7 (21.9)              | 4 (19.1)                  | 4 (19.1)                  | 0.4     |
| **Diabetes, N (%)**                                  | 3 (9.4)               | 0                         | 3 (14.3)                  | 0.53    |
| **Neoplasm, N (%)**                                  | 7 (21.9)              | 4 (14.3)                  | 3 (14.3)                  | 0.2     |
| **ICU LOS before TGC, (days)**                       | 7.5 [2.5-16]          | 5 [0.5-11.25]             | 12 [3.75-18.25]           | 0.13    |
| **MV duration before TGC, (days)**                   | 8 [3-12]              | 5 [0.5-11.25]             | 8 [3.75-14.75]            | 0.19    |
| **Vasopressors duration before TGC, (days)**         | 4.5 [0-8.5]           | 5 [0.25-8.25]             | 4 [0-8.25]                | 0.89    |
| **SOFA score**                                       | 7 [4-10]              | 8 [4.75-12]               | 6 [4-9]                   | 0.2     |
| **Septic shock, N (%)**                              | 18 (56.3)             | 7 [63.6]                  | 11 (52.4)                 | 0.71    |
| **ARF requiring MV, N (%)**                          | 28 (87.5)             | 10 (90.9)                 | 18 (85.7)                 | 1       |
| **AKI requiring CRRT, N (%)**                        | 11 (34.4)             | 3 [27.3]                  | 8 (38.1)                  | 0.7     |
| **Creatinine clearance, (ml/min)**                   | 97.3 [32-150.8]       | 63.2 [32-155]             | 104 [30-142]              | 0.85    |
| **VAP, N (%)**                                       | 19 (59.4)             | 3 [27.3]                  | 16 (76.2)                 | 0.02    |
| **Intra-abdominal infection, N (%)**                 | 10 (31.3)             | 6 [54.5]                  | 4 [19.1]                  | 0.06    |
| **Skin and soft-tissue infection, N (%)**            | 3 (9.4)               | 2 [18.2]                  | 1 [4.8]                   | 0.27    |
| **Secondary bacteraemia, N (%)**                     | 13 (40.6)             | 4 [36.4]                  | 9 [42.9]                  | 1       |
| **Source control, N (%)**                            | 13 (40.6)             | 7 [63.6]                  | 6 [28.6]                  | 0.07    |
| **TGC therapy duration,(days)**                      | 12 [9-15]             | 12 [10-15]                | 11 [8-17]                 | 0.69    |
| **TGC empirical therapy, N (%)**                     | 17 (53.1)             | 7 [63.6]                  | 10 [47.6]                 | 0.47    |
| **Gram-positive bacteria N (%)**                     | 11 (34.4)             | 4 [36.4]                  | 7 [33.3]                  | 1       |
| **Gram-negative bacteria N (%)**                     | 29 (90.6)             | 10 (90.9)                 | 19 (90.5)                 | 1       |
| **ICU LOS after TGC, (days)**                        | 15 [10.5-27]          | 14.5 [12-19]              | 16 [10-31.4]              | 0.42    |
| **MV duration after TGC, (days)**                    | 10 [5-15]             | 14 [9.75-15.75]           | 8 [2-13.5]                | 0.04    |
| **Vasopressors duration after TGC, (days)**          | 3[1.5-13]             | 8 [2.25-13]               | 3 [0-10.75]               | 0.12    |
| **30-day mortality**                                 | 9 [28.1]              | 8 [72.7]                  | 1 [4.8]                   | < 0.001 |

Data are presented as median [IQR], unless otherwise indicated

Pts: patients; VAP: ventilator-associated pneumonia; TGC: tigecycline; SAPS II: Simplified Acute Physiology Score; COPD: chronic obstructive pulmonary disease; LOS: length of stay; ICU: Intensive Care Unit; MV: mechanical ventilation; SOFA: Sequential Organ Failure Assessment; AKI: acute kidney injury; CRRT: continuous renal replacement therapy; ARF: acute respiratory failure; MV: mechanical ventilation; Kg: kilograms; IQR: interquartile range

*Evaluated at TGC starting day

** i.e. Streptococcus spp., staphylococcus aureus, enterococci

*** i.e. Acinetobacter baumannii, Klebsiella pneumonia, Enterobacteriaceae, Bacteroides spp.
Urea assay
Determination of urea in plasma and BAL samples
Urea levels were detected by the QuantiChrom Urea Assay kit (BIOassay System, Hayward, CA, USA), which was used according to the manufacturer's instructions.

Pharmacokinetic/pharmacodynamics analysis
A one compartment model with first-order elimination determined pharmacokinetic parameters. The 0–12 h area under the time concentration curve \( \text{AUC}_{0-12} \) was determined by the linear trapezoidal rule. TGC \( \text{AUC}_{0-24} \) was calculated as \( \text{AUC}_{0-12} \times 2 \). TGC maximum and minimum concentrations \( (\text{C}_{\text{max}}, \text{C}_{\text{min}}) \) were directly obtained from observed peak and trough concentrations. Epithelial lining fluid (ELF) tigecycline \( \text{TGC}_{\text{ELF}} \) concentration was calculated from BAL concentration \( \text{TGC}_{\text{BAL}} \) using urea as dilution marker: \( \text{TGC}_{\text{ELF}} = \text{TGC}_{\text{BAL}} \times \text{urea dilution index (plasma urea concentration/BAL urea concentration)} \) [15]. In all patients, distribution volume \( (\text{Vd}) \), drug clearance \( (\text{CL}) \), and elimination half-life \( (t_{1/2}) \) were calculated after a single 100-mg intravenous dose at steady state.

Area under the concentration curve \( \text{AUC}_{0-24}/\text{MIC} \) ratio \( \geq 4.5, 6.96 \) and 17.9 were used as PD targets for VAP, intra-abdominal infections (IAI) and skin-soft tissue infections (SSTI), respectively [6].

Graphing of data was undertaken using Prism version 6.0 for Windows (graphPad Software, San Diego, CA).

Microbiological analysis
Isolates were identified by matrix-assisted laser desorption ionization-time-of-flight (MALDI-TOF) mass spectrometry (MALDI Biotyper, Bruker Daltonics GmbH, Leipzig, Germany). The in vitro susceptibility of the isolates was assessed with the Vitek 2 system (bioMérieux, Marcy l’Etoile, France) or with panels manufactured by MERLIN Diagnostica GmbH (Bornheim, Germany). Results were interpreted in accordance with the European Committee on Antimicrobial Susceptibility Testing (EUCAST) clinical breakpoints. The presence of carbapenemase genes of \( \text{blaKPC}, \text{blaNDM}, \text{blaVIM}, \text{blaOXA-48}, \text{blaOXA-23}, \) and \( \text{blaOXA-58} \) types was determined by polymerase chain reaction and DNA sequencing analysis using previously described protocols (Endemiani, Poirel, Woodford). [16]

Statistical analysis
All statistical analyses were performed using MedCalc software, version 12.2.1 (MedCalc®,
MariaKerke, Belgium). Kolmogorov-Smirnov test was used to value the variables distribution. The data with a non-Normal distribution were assessed with Mann-Whitney test and the median and selected centiles (25th-75th) value was given (interquartile range, IQR). The data with a Normal distribution were assessed with Student’s test. Categorical variables are presented as proportions and were analysed with the use of the chi-square test or Fisher’s exact test, as appropriate. A P value < 0.05 was considered significant. Due to the PK/PD design of the study, a sample size was not calculated, foreseeing the recruitment of at least 30 patients during the predefined study period (July 2015-July 2017).

Results

Patients characteristics

The clinical details of the 32 non-obese patients included in the study are listed in Table 1. Albumin levels were quite low with an overall positive fluid balance at enrolment. Median SAPS II score was 53.5 and the most relevant comorbidities were cardiovascular diseases, chronic obstructive pulmonary disease, chronic renal failure and neoplasm (Table 1). Median SOFA score was 7 and many patients were in septic shock or presented with acute respiratory failure (ARF) and acute kidney injury (AKI) requiring mechanical ventilation (MV) and CRRT, respectively. More than half of the patients had VAP, followed by intra-abdominal infections and skin and soft-tissue infections: in the microbiological case-mix Gram-negative bacteria were mostly represented (90.6%). Median duration of TGC therapy was 12 days and it was started empirically in half of the cases. The use of vasopressors and MV during TGC therapy was high and 30-day mortality rate was 28.1%. Eleven patients had a treatment failure, while the other 21 successfully eradicated the infection. There were no between-group differences in terms of demographic aspects and main comorbidities. Further the two groups were similar in terms of presenting features and outcomes with the exception of VAP rate which was higher in the treatment success group (76.2% vs. 27.3%, p = 0.02) and a trend to a higher percentage of skin and soft tissue infections and source control among patients who failed TGC treatment (p = 0.06 and p = 0.07, respectively).

Pharmacokinetic results

A one-compartment model with first-order disposition processes adequately described the
concentration-time curve, although significant inter-individual variability was observed. Vd, Cl, and t_{1/2} were 438.6 L, 42.1 L/h and 7.2 h, respectively. Median and IQR values of Cmax and Cmin were 0.34 [0.15–1.03] mg/L and 0.09 [0.05–0.26] mg/L (Table 1). Figure 1 shows the mean±SD time-concentration profile at different time points of plasma tigecycline concentrations, compared with most frequently observed MIC values (0.12–0.25–0.5 mcg/mL). AUC_{0–24} and IQR was calculated for each patient and the percentage of target attainment was also computed for nosocomial pneumonia (NP) (AUC_{0–24}/MIC breakpoint of 4.5), complicated intra-abdominal infections (cIAI) (AUC_{0–24}/MIC breakpoint of 6.96) and complicated skin and soft tissue infections (cSSTI) (AUC_{0–24}/MIC breakpoint of 17.9). Considering a target AUC_{0–24}/MIC of 4.5, 75% of the patients would be successfully treated in presence of 0.5 mg/L MIC value and all the patients obtained the PK target with MIC ≤ 0.12 mg/mL. Considering a target AUC_{0–24}/MIC of 6.96, at least 50% of the patients would be adequately treated against bugs with MIC ≤ 0.5 mcg/mL, while only 15.6% obtained the PK target with MIC of 2 mg/mL. Finally, with a target AUC_{0–24}/MIC of 17.9 only 25% of the patients obtained the PK target at MIC values of 0.5 mg/L and less than 10% were adequately treated against germs with MIC value ≥ 1 mcg/ml (Fig. 2).

Pulmonary concentrations
Tigecycline pulmonary concentrations were measured in 12 (1 h) and 7 (12 h) patients, respectively. Main reasons to exclude samples were presence of blood and excessive dilution, while five patients did not undergo BAL due to severe respiratory failure. Median and IQR ELF Cmax was 0.42 [0.15–1.2] and ELF Cmin was 0.32 [0.17–0.43]; median and IQR ELF/plasma ratio (%) was 152.9 [73.5–386.8] (Table 2). Mean±SE ELF concentrations were similar at 1 h and 12 h (0.78±0.2 mg/mL vs. 0.36±0.1 mg/mL; p = 0.19) (Fig. 3). Conversely, no significant differences were found comparing mean±SE ELF/plasma ratio at 1 h and 12 h (281±107.6 vs. 298.3±60.7; p = 0.9) (Fig. 4).
Table 2
Steady-state serum and alveolar TGC PK parameters in the 32 enrolled patients

| Parameter | Patients (n = 32) |
|-----------|------------------|
| Vd, L     | 438.6            |
| CL, L/h   | 42.1             |
| t\textsubscript{1/2}, h | 7.2 |
| C\textsubscript{max}, mg/L | 0.34 [0.15–1.03] |
| C\textsubscript{min}, mg/L | 0.09 [0.05–0.26] |
| ELF C\textsubscript{max}, mg/L* | 0.42 [0.15–1.2] |
| ELF C\textsubscript{min}, mg/L* | 0.32 [0.17–0.43] |
| ELF/plasma ratio (%), median [IQR]* | 152.9 [73.5–386.8] |
| AUC\textsubscript{0–24}, mg h/L | 3.61 [2.55–10.39] |
| AUC\textsubscript{0–24}/0.12 mg/L MIC ≥ 4.5, (%) | 100 |
| AUC\textsubscript{0–24}/0.25 mg/L MIC ≥ 4.5, (%) | 94 |
| AUC\textsubscript{0–24}/0.5 mg/L MIC ≥ 4.5, (%) | 75 |
| AUC\textsubscript{0–24}/1 mg/L MIC ≥ 4.5, (%) | 40.6 |
| AUC\textsubscript{0–24}/2 mg/L MIC ≥ 4.5, (%) | 28.1 |
| AUC\textsubscript{0–24}/0.12 mg/L MIC ≥ 6.96, (%) | 100 |
| AUC\textsubscript{0–24}/0.25 mg/L MIC ≥ 4.5, (%) | 91 |
| AUC\textsubscript{0–24}/0.5 mg/L MIC ≥ 6.96, (%) | 50 |
| AUC\textsubscript{0–24}/1 mg/L MIC ≥ 6.96, (%) | 34.4 |
| AUC\textsubscript{0–24}/2 mg/L MIC ≥ 6.96, (%) | 15.6 |
| AUC\textsubscript{0–24}/0.12 mg/L MIC ≥ 17.9, (%) | 78 |
| AUC\textsubscript{0–24}/0.25 mg/L MIC ≥ 17.9, (%) | 44 |
| AUC\textsubscript{0–24}/0.5 mg/L MIC ≥ 17.9, (%) | 25 |
| AUC\textsubscript{0–24}/1 mg/L MIC ≥ 17.9, (%) | 9.4 |
| AUC\textsubscript{0–24}/2 mg/L MIC ≥ 17.9, (%) | 3.1 |

Data are expressed as median [IQR] and N (%)

*TGC ELF concentrations were measured in 12 (1 h) and 7 (12 h) samples, respectively

Legend
TGC tigecycline; PK pharmacokinetic; Vd volume of drug distribution, IQR interquartile range; CL drug clearance; t\textsubscript{1/2} elimination half-life; Cmax peak plasmatic concentration; Cmin trough plasmatic concentration; ELF epithelial lining fluid; MIC minimum inhibitory concentration; AUC total drug area under the time-concentration curve

Discussion

Our study shows a HD TGC (200 mg LD, then 100 mg q12) time-curve concentration with mean peak and trough levels of 0.65 mg/mL and 0.25 mg/mL, respectively (Fig. 1). AUC\textsubscript{0–24}/MIC targets for nosocomial pneumonia (≥ 4.5) and complicated intra-abdominal infections (≥ 6.96) were obtained in the majority of cases in presence of bacteria with MIC values ≤ 0.25. Otherwise lower MIC values (≤ 0.12 mcg/mL) were required to have satisfactory AUC\textsubscript{0–24}/MIC results (78%), while treating a skin/soft tissue infection (Fig. 2, Table 2). Similar to plasma 1 h and 12 h, pulmonary concentrations (0.78 mg/L and 0.36 mg/L, respectively) were observed with a good median ELF/plasma ratio of 152.9% (Table 2, Fig. 3). This high-dose regimen was associated with a 65.6% of treatment success rate in a normal weight population including 60% of VAP, 31% of cIAI and 9% of SSTI. TGC was used in
half of the cases as targeted regimen for a median duration of 12 days. The rate of septic shock, acute respiratory failure requiring MV and acute kidney injury requiring CRRT was also high, with a mortality rate of 28.1% (Table 1).

The pharmacokinetics/pharmacodynamics and tissue penetration of tigecycline have been extensively studied in various in vitro and human models [17]. However, these studies were generally carried out in healthy volunteers, and few pharmacokinetic data concerning infected patients are available, which may present pathophysiologic conditions influencing the pharmacokinetic profile of this molecule. Additionally, the majority of available data in infected patients derive from studies where normal doses are used, although for severe nosocomial infections a double-dose regimen is warranted [18, 19].

Recently, standard dose TGC pharmacokinetics in ten critically ill patients have been studied [6]. The authors observed that a larger body mass index was associated with increased TGC Cl, but standard doses produced satisfactory plasmatic levels for VAP and cIAI treatment due to Enterobacter cloacae, Escherichia coli, Klebsiella pneumoniae and methicillin-resistant Staphylococcus aureus. However higher dosages were required for the treatment of SSTI, especially in obese patients.

Eleven out of 32 patients in our cohort were receiving CRRT while being treated with high-dose TGC. Interestingly, in a recent paper, Broeker and cow. [20] described the PK/PD of standard dose TGC in eleven patients on continuous veno-venous hemodialysis (CVVHD) or hemodiafiltration (CVVHDF). TGC dialysability, as expressed by saturation coefficients (0.79 and 0.9 for CVVHD and CVVHDF, respectively) was very high, but the contribution of CRRT TGC clearance was minimal (about 2 L/h), compared with the total body clearance (18.3L/h). Peak drug concentrations were below 1 mg/L and trough levels about 0.2 mg/L. The authors, considering the AUC0-24/MIC referral value for cIAI (6.96), observed that such target was accomplished in 88% of the case if MIC was ≤ 0.5.

Indeed, our results are in line with current available data, underlying the plus-value of increased dosages while treating critically ill patients especially with severe cIAI and SSTI. In addition, there is a high need of PK/PD data on TGC administered at higher than approved dosages, in light of the wide spread of increased resistance to TGC among Gram-negative rods and Acinetobacter spp. The first
investigation on PK/PD of HD TGC derives from Ramirez et al who conducted a randomized phase 2 trial to evaluate the clinical efficacy of two high-dosage regimen of TGC (75 mg bid and 100 mg bid) versus imipenem-cilastatin for the treatment of nosocomial pneumonia[8]. In the clinically evaluable population, clinical cure with TGC 100 mg bid was higher than with 75 bid and imipenem-cilastatin (85% vs. 69.6% vs. 75%). Mean peak TGC concentration was about 1 mg/mL, declining to less than 0.5 mg/ml after 8 hours, observing a safety profile comparable to that one known for the approved those. The only other study investigating the PK/PD of HD TGC profile was conducted by Borsuk-De Moor et al in 37 ICU patients with severe infections [21]. The time-concentration curve was similar to our data, displaying a peak concentration about 1 mg/L and 12 h level below 0.5 mg/L. Interestingly, the authors developed a model which showed that no individual covariates may influence target concentrations, advising to modify TGC daily dosage according to pathogens type, susceptibility pattern and PK targets.

Tissue concentrations of antibiotics at the target site contribute to therapeutic effects: using plasma concentrations may frequently overestimate the target site concentrations and therefore clinical efficacy. This is the first study to report steady-state ELF percentage penetration of TGC administered 100 mg q12 after 200 mg LD. Considering the AUC$_{0−24}$/MIC target of 4.96, our data shows satisfactory pulmonary concentrations with potential clinical success in 100% – 94% – 75% – 41% of the cases treating bacteria with MIC of 0.12–0.25 mg/L – 0.5–1 mg/L, respectively [Fig. 2]. These data confirm the results observed in healthy subjects by Conte et al., where the Cmax/MIC90, AUC/MIC90 ratios, T > MIC90 and extended serum and intrapulmonary half-lives following the standard regimen are favourable for the treatment of TGC-susceptible pulmonary infections [22]. Penetration ratio may be even higher when in presence of infected lungs. Crandon et al. demonstrated in infected and non-infected mice lungs that the baseline penetration ratio of 8.1 is incremented to 23.3 in case of Acinetobacter pneumonia [23]. Conversely, the majority of lung penetration occurs in alveolar cells, than in ELF, as suggested by Welte et al. in three cases of MDR lung infections [24]. Finally in a recent study on 58 healthy subjects treated with standard TGC dose, the ratio of ELF and AUC to total plasma concentration of tigecycline was 1.71 and 20.8, respectively [25].
Our study has several limitations. First we adopted a single high-dose of tigecycline and we do not know if even higher dosages may result in better PK/PD profiles. Second we measured only pulmonary tissue concentration through ELF collection and we can only postulate the real tissue/plasma ratio for cIAI and SSTI. Third, our analysis focused on total TGC concentration rather unbound $AUC_{0−24}$, due to the lack of clinical reliable breakpoint of $fAUC_{0−24}/MIC_{90}$. Finally the sample size may be likely responsible of an under-estimated interindividual variability in the observed PK/PD profile.

Conclusions
Our study is the first investigation where not only plasmatic but also pulmonary tigecycline concentrations are investigated during the treatment of severe infections in critically ill patients with high dose TGC. Observed plasmatic and pulmonary concentrations confirm the safety of this molecule for the treatment of susceptible pathogens, including pneumonia. Higher than 200 mg/day dosages and combination with other active molecules should be adopted while treating Enterobacteriaceae and Acinetobacter spp. with MIC values close to the clinical breakpoint.

Abbreviations
HD, high dosages; TGC, Tigecycline; PK/PD, pharmacokinetic/pharmacodynamic; ICU, Intensive Care Unit; ELF, epithelial lining fluid; AUC, area under the curve; MIC, minimum inhibitory concentration; IQR, interquartile range; MRSA, Staphylococcus aureus; VRE, vancomycin-resistant enterococci; ESBL, extended-spectrum β-lactamase; XDR, extensively drug-resistant; FDA, Food and Drug administration; CRRT, continuous renal replacement therapy; SAPS II, Simplified Acute Physiology Score; LD, loading dose; DHHS-CTCAE, Department of Health and Human Services–Common Terminology Criteria for Adverse Events; VAP, ventilator associated pneumonia; BAL, bronchoalveolar lavage; IS, internal standard; SPE, solid-phase extraction; TQD, triple quadrupole tandem; ESI, electrospray ionization; MRM, multi reaction monitoring; MS, mass spectroscopy; Cmax, maximum concentration; Cmin, minimum concentration; Vd, distribution volume; CL, drug clearance; t1/2, elimination half-life; IAI, intra-abdominal infection; SSTI, skin-soft tissue infection; MALDI-TOF, matrix-assisted laser desorption ionization-time-of-flight; EUCAST, European Committee on Antimicrobial Susceptibility Testing; SOFA, Sequential Organ Failure Assessment; ARF, acute respiratory failure; AKI, acute kidney injury; MV,
mechanical ventilation; cIAI, complicated intra-abdominal infection; cSSTI, complicated skin and soft tissue infection; CVVHD, continuous veno-venous haemodialysis; CVVHDF, continuous veno-venous hemodiafiltration.

Declarations

Ethics approval and consent to participate: The protocol was approved by the Catholic University’s Ethical Committee (approval number Prot.sf 8431/13). Written informed consent was obtained from the patients’ legally authorized representative.

Consent for publication: Written informed consent was obtained from the patients’ legally authorized representative. No individual person’s data in any form are presented in this manuscript.

Availability of data and materials: The datasets used and/or analysed during the current study are available from the corresponding author on reasonable request.

Competing interests: The authors declare that they have no competing interests

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Authors’ contributions: GDP and MA had full access to all the data in the study and take responsibility for the integrity and the accuracy of the data analysis. GDP, MS, PN and MA conceived the study, and participated in its design and coordination and helped to draft the manuscript. GDP was in charge of the statistical analysis, participated in analysis and interpretation of data, helped to draft the manuscript, and critically revised the manuscript for important intellectual content. LL, GMPC, MSV, SLC, LC, CG, GB, LM, SC and VDG collected the data for the study, recruited patients and did sample analyses. GDP, LL, GMPC, MT, MS, PN and MA participated in the conception, design and development of the database, helped in analysis and interpretation of data, helped in drafting of the manuscript and critically revised the manuscript for important intellectual content. All authors read and approved the final manuscript.

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Figure 1

Total Tigecycline plasma concentration (mean±SE) versus time of administration.
Figure 2

Probability of target attainment of pharmacodynamics indices in plasma, according to different infection types and MIC values
Boxplot showing Tigecycline ELF concentrations. Boxes represent interquartile ranges (lower border 25th percentile; upper border 75th percentile), and the horizontal lines within the boxes indicate the medians (50th percentile). Whiskers indicate minimum and maximum values.
Figure 4

Boxplot showing Tigecycline ELF to plasma ratio (%). Boxes represent interquartile ranges (lower border 25th percentile; upper border 75th percentile), and the horizontal lines within the boxes indicate the medians (50th percentile). Whiskers indicate minimum and maximum values.