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Title

Population pharmacokinetics of **single-dose** amikacin in critically-ill patients with suspected ventilator-associated pneumonia

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Keywords

Amikacin, Population pharmacokinetics, Intensive-care unit, Ventilator-associated pneumonia

Abstract

Aims

Modifications of antimicrobials' pharmacokinetic parameters have been reported in critically-ill patients, resulting in a risk of treatment failure. We characterized amikacin pharmacokinetic variability in critically-ill patients with ventilator-associated pneumonia (VAP) and evaluated several dosing regimens.

Methods

We conducted a prospective multicenter study in critically-ill patients with presumptive diagnosis of Gram-negative bacilli (GNB) VAP. Patients empirically received imipenem and a single-dose of amikacin, which was administered as a 30-minute infusion (20mg/kg). Concentrations were measured 0.5, 1, 8, 16 and 24 hours after beginning of infusion. Pharmacokinetic parameters were estimated using a population approach. Main pharmacodynamic target was a ratio ≥ 10 between the concentration achieved 1 hour after beginning of infusion (C_{1h}) and the minimal inhibitory concentration of the liable bacteria (MIC). We simulated individual C_{1h} for several dosing regimens by Monte Carlo method and computed C_{1h}/MIC ratios for MICs from 0.5 to 64mg/L.

Results

Sixty patients (47 males), median (range) age and bodyweight: 61.5 years (28-84) and 78kg (45-126), respectively were included. Amikacin median C_{1h} was 45mg/L (22-87). Mean value (between-patients variability) for CL, V1, Q and V2 were 4.3L/h (31%), 15.9L (22%), 12.1L/h (27%) and 21.4L (47%), respectively. CL increased with CrCL ($p < 0.001$) and V1 with body weight ($p < 0.001$) and PaO_2/FIO_2 ratio ($p < 0.001$). With a 25mg/kg regimen, the pharmacodynamic target was achieved in 20% and 96% for aMICs of 8 and 4mg/L, respectively.

Conclusion

Amikacin clearance was decreased and its volume of distribution was increased as previously reported. A ≥ 25 mg/kg single-dose is needed for empirical treatment of GNB-VAP.

Introduction

Ventilator-associated pneumonia (VAP), defined as pneumonia occurring more than 48 h after the initiation of mechanical ventilation, is the most common hospital-acquired infection in the intensive care unit (ICU) [1]. Many previously published studies have shown that early and appropriate antibiotic therapy is associated with better outcome in critically-ill patients with severe infections[2].The latest guidelines of the American Thoracic Society and Infectious Diseases Society of America recommend that the empirical antibiotic therapy for late-onset pneumonia or in patients with risk factors for infection by multidrug-resistant pathogens should include a combination of an anti-pseudomonal β -lactam and an anti-pseudomonal fluoroquinolone or an aminoglycoside such as amikacin[3].The superiority of once-daily dosing of aminoglycosides over multiple-daily dosing has been established[4]. However, no data supports a clinical benefit of a single-dose of aminoglycosides over multiple administrations. The rationale for an initial combination therapy is to broaden the antimicrobial spectrum of the empirical therapy; however some data suggest adverse effects linked to repeated administrations, such as adaptive resistance or nephrotoxicity.

Amikacin, like other aminoglycosides, exhibits a concentration-dependent killing and produces a prolonged post-antibiotic effect. Previous clinical studies have shown that a ratio of 10 or more between the concentration achieved 1 hour after the beginning of a 30-minute infusion (C_{1h}) and the minimal inhibitory concentration (MIC) of the bacteria responsible of the infection was predictive of therapeutic success [5, 6]. Some authors recently suggested that a ratio between the area under the curve of concentration over time (AUC) and MIC (AUC/MIC) greater than 90 also has some predictive value for therapeutic success, with no difference in predictive capacities when compared to C_{1h} / MIC ratio [7]. Those pharmacodynamic targets may be difficult to reach in critically-ill patients for several reasons. First, large interindividual variations of amikacin pharmacokinetic parameters have been reported [8-10], with variations of C_{1h} and AUC as a consequence. This variability is partially explained by total body weight and creatinine clearance, which are the most

frequently reported covariates of amikacin pharmacokinetic parameters[8-11]. Second, the current breakpoint of MICs defined by the European Committee on Antimicrobial Susceptibility Testing (EUCAST) for Enterobacteriaceae species and *Pseudomonas aeruginosa* is 8 mg/L [12].

Few authors investigated the amikacin dosage needed in a single infusion setting to achieve the defined pharmacodynamic target in a homogenous group of patients hospitalised in ICU with suspected VAP.

Using data from a prospective clinical trial, we aimed (i) to characterize amikacin pharmacokinetic parameters and their variability in ICU patients with suspected VAP, (ii) to study covariates of amikacin pharmacokinetic parameters, and (iii) to evaluate several amikacin dosing regimens using Monte Carlo simulations.

Material and methods

Patients and sampling

This study is part of the IMPACT trial, a prospective multicenter clinical trial conducted between 2009 and 2011 in three ICUs, at Bichat university hospital, Paris, France, and Victor Dupouy hospital, Argenteuil, France (ClinicalTrials #NCT00950222).

Patients were included in the trial if they presented the following criteria: mechanical ventilation for more than 48 hours, a clinical suspicion of Gram-negative bacilli VAP, risk factors for multidrug resistant bacteria, and if a microbiologic sample was obtained before initiation of antimicrobial therapy using blinded protected telescoping catheter or bronchoalveolar lavage. Clinical suspicion of VAP was defined by the onset of new lung infiltrates on chest radiography, fever greater than 38.3°C, purulent tracheal aspirates or a leukocytosis > 10000/mL. Risk factors for multidrug-resistant bacteria were antimicrobial therapy in the preceding 15 days or late-onset VAP (≥ 6 days)[3]. Patients were not eligible if they were younger than 18-year-old, if they had renal failure requiring renal replacement therapy or received a treatment by imipenem or amikacin at the time of inclusion.

Written consent was obtained from patients or their legal representative. The study was approved by the Ethics Committee of the Hôtel-Dieu university hospital (APHP, Paris).

All patients were treated with a combination of imipenem and amikacin. Imipenem was administered every 8 hours over a 30-minute infusion and should not be changed for the first 48 hours. Doses ranged from 500 mg to 1000 mg, according to creatinine clearance (CrCL). Amikacin treatment consisted in one single 30-minute infusion of a suggested dose of 20 mg/kg administered the first day of antimicrobial therapy. This dose was recommended by guidelines at the time of this study was designed [3].

De-escalation therapy was encouraged after isolation of the Gram-negative pathogen and obtaining the results of susceptibility tests.

Usual clinical and demographic characteristics were recorded at ICU admission, as well as ventilator parameters, two scores for assessment of severity (SOFA [13] and SAPS-II [14]) and routine biologic markers.

Blood samples were collected 0.5, 1, 8, 16 and 24 hours after the onset of infusion. Exact times of beginning and end of infusion were recorded, as well as exact sampling times. Amikacin concentrations were determined by fluorescence polarization immunoassay using the amikacin Innofluor® kit [15]. The limit of quantification of the technique is 0.5 mg/L, and coefficients of variation for intra- and inter-assay were 4.1% and 5.8%, respectively.

Population pharmacokinetic analysis

Basic model

Population pharmacokinetic analysis was performed using the Stochastic Approximation Expectation Minimization (SAEM) algorithm in Monolix v4.2 (Lixoft, Orsay, France, available at <http://www.lixoft.com>). This algorithm handles concentrations below the limit of quantification to

improve parameters estimation [16]. We used the total dose administered to each patient for model building.

Both one- and two-compartment(s) models with first order elimination were tested[8, 10, 11, 17].

We used an exponential random effects model for each pharmacokinetic parameter. We assumed the random effects to have a normal distribution with a mean of 0 and a variance of ω^2 . Correlation (ρ) between individual random effects was kept in the model if the estimated correlation coefficient was ≥ 0.2 . The residual error model was supposed to be additive, proportional or combined, with a being the standard deviation of the additive component and b the standard deviation of the proportional component.

The best model was chosen using the Bayesian information criteria (BIC), derived for each model from the computation of likelihood by importance sampling[18].

Covariate model building

The influence of the following covariates at initiation of treatment on amikacin pharmacokinetic parameters was tested: 3 demographic variables (age, gender and total body weight), 2 clinical variables (shock and edema score[19]), 2 severity scores (SOFA[13] andSAPS-II[14]), 2 ventilator-related parameters (PEEP and PaO₂/FIO₂ ratio) and 3 biochemical markers (serum albumin, total bilirubin and 4-hour creatinine clearance). Four-hour creatinine clearance was calculated using serum creatinine level and a 4-hour urine collection at the day of inclusion. This approach has been validated for monitoring renal function in critically-ill patients[20]. Missing values for tested covariates were imputed to the median value observed in the analysis population. Steps for selection of covariate model are described in Online Resource 1.

Final model determination

We graphically studied the influence of covariates on their related pharmacokinetic parameters.

Outliers were studied and excluded from analysis when there was a suspicion of incomplete data collection, in particular for creatinine clearance which was estimated using a 4-hour urine collection.

Patients for whom covariates had been imputed were also excluded. The covariate model was adjusted to the reduced population obtained after exclusion of these patients, and a backward selection method was used in order to obtain a final model in which all covariates had a p-value < 0.05 using the likelihood-ratio test.

The coefficient estimated for creatinine clearance was compared to 1 using the likelihood ratio test.

Model evaluation

Evaluation of the final model was conducted using graphical methods. Basic goodness-of-fit plots were used, as well as the individual weighted residuals (IWRES) and the normalized prediction distribution errors (NPDE) over time and the visual predictive check (VPC). NPDE and VPC were generated using 500 Monte Carlo simulations.

Evaluation of doses by simulation

Using the estimated distribution of amikacin pharmacokinetic parameters in the final model with covariates, we simulated amikacin concentration obtained 1 hour after the start of a 30-minute infusion and AUC for 1000 patients by Monte Carlo simulation for several dosing regimen. This timing for amikacin sampling is commonly used for studying amikacin efficacy[17]. For pharmacokinetic simulations, we randomly re-sampled 1000 vectors of covariates among those observed in the patients included in the analysis and simulated individual pharmacokinetic parameters from their estimated distribution in the final model with covariates. Simulated dosing regimens were the following: 20, 25, 30, 35 and 40 mg/kg. We computed the probability to achieve a C_{1h}/MIC ratio ≥ 10 and an AUC/MIC ratio ≥ 90 for MICs ranging from 0.25mg/L to 64mg/L for each of these dosing regimens. These values are usually observed in clinical practice

(http://www.eucast.org/mic_distributions/), with an 8 mg/l susceptibility breakpoint for Enterobacteriaceae species and *Pseudomonas aeruginosa*.

Results

Patients' characteristics

Of the 61 patients included in the IMPACT trial, one had a kinetic profile which was not compatible with a unique injection and was withdrawn from analysis. Thus, data from 60 patients were available for modelling. Patients' characteristics are presented in Table 1. Median (min – max) age and total body weight were respectively 61.5 years (28 – 84) and 78 kg (45 – 126), and most patients were males (n=47, 78%). At inclusion, 26 patients had septic shock (43%). Median SAPS-II and SOFA scores were 42 (19 – 90) and 7 (2 – 17), respectively, and median 4-hour creatinine clearance was 82 mL/min (4 – 412).

Pharmacokinetic data and modelling

A total of 291 values of amikacin concentrations were available, with a median of 5 per patient (3 – 5). Median amikacin dose administered was 20 mg/kg (11 – 28). Median serum concentration observed 1 hour after the beginning of infusion was 45 mg/L (22 – 87). Individual observed concentrations are presented in Figure 1. Fourteen values (4.8%) of amikacin concentration were below the limit of quantification.

Serum concentrations were best described by a 2-compartment model with a combined residual error model (Online Resource 2). Correlations between estimates in the basic model were ≥ 0.2 between all individual random effects, and thus kept in the analysis. In this model, amikacin clearance (CL) was estimated to 4.0 L/h, central volume of distribution (V1) to 15.3 L, peripheral volume of distribution (V2) to 22.1 L and inter-compartmental clearance (Q) to 12.2 L/h (Table 2). Their relative standard errors were satisfactory, all being $<10\%$. The inter-individual variability of pharmacokinetic parameters ranged from 30% for Q, to 60% for CL. Estimates of inter-individual

variability of pharmacokinetic parameters were also satisfactory (Table 2). The goodness of fit plots of this model without covariate did not show any model deficiency (data not shown).

Among the 12 studied covariates, 8 were significantly associated with the individual Bayes estimates of pharmacokinetic parameters (Online Resource 1). The best model included 4 covariates (Online resource 1): creatinine clearance for CL, total body weight and PaO₂/FIO₂ ratio for V1, and creatinine clearance for V2. Two patients presented a low creatinine clearance but a high amikacin clearance. Their individual fits were satisfactory, and we hypothesised that their urine collection was incomplete. They were excluded from the analysis. In another patient, amikacin clearance was estimated to 11 mL/min, but creatinine clearance was missing and had been imputed to the median value. The 5 patients in whom creatinine clearance was not available were excluded. The 4-covariate model was adjusted to the 53 remaining patients.

Creatinine clearance was no longer a significant covariate for V2 ($p=0.4$), and the final model included 3 covariates: creatinine clearance ($p<0.001$) for CL, total body weight ($p<0.001$) and PaO₂/FIO₂ ratio ($p<0.001$) for V1 (Table 2). The coefficient for creatinine clearance on CL was significantly different from 1 ($p<0.001$). The variations of pharmacokinetic parameters according to covariates are presented in Figure 2. All three of them were positively correlated with the pharmacokinetic parameters. Amikacin clearance was estimated in the final model at 4.3 L/h (72 mL/min), V1 at 15.9 L and V2 at 21.4 L (Table 2). Variance – covariance matrix between the individual random effects is presented in Online Resource 3.

Goodness-of-fit plots did not show any model misspecification. The IWRES and NPDE were centered to zero and did not show any trend over time (Online Resource 4). The VPC did not show any model deficiency (Figure 3).

Pharmacokinetic simulations

Probabilities to achieve a C_{1h}/MIC ratio ≥ 10 according to the single-dose of amikacin simulated and MICs are presented in Figure 4. A 20 mg/kg dose was sufficient to achieve the pharmacokinetic target in 100% of patients for a MIC of 2 mg/L or less. With this dose only 80% and 4% of patients achieved the target for a MIC of 4 and 8 mg/L, respectively. With a 25 mg/kg simulated infusion, the probability to reach the target was 96% for a MIC of 4 mg/L and 20% for a MIC of 8 mg/L. With the 40 mg/kg regimen, the probability to achieve a C_{1h}/MIC ratio ≥ 10 was 80% for a MIC of 8 mg/L (Figure 5). Online Resource 5 presents the boxplots of the 1000 simulated C_{1h}/MIC ratio for a MIC of 8 mg/L, for doses from 20 to 40 mg/kg.

Results were similar when considering the target of $AUC/MIC \geq 90$ (Online Resource 6). With a dose of 20 mg/kg, 90% of patients achieved the target for a MIC of 2 mg/L or less, but this proportion was below 20% for a MIC value of 8 mg/L. Simulations for a dose of 25 mg/kg showed that 69% of patients achieved the pharmacodynamic target for a MIC of 4 mg/L, but only 22% did for a MIC of 8 mg/L. An increase to 40 mg/kg led to an increase of the probability to reach the pharmacodynamic target to 90% and 52% for a MIC of 4 mg/L or 8 mg/L, respectively.

Discussion

The present study confirms that the 20 mg/kg single-dose regimen traditionally used is not adequate based on the analysed pharmacokinetic and pharmacodynamic indices and that amikacin dosing regimen should be increased to at least 25 mg/kg for critically-ill patients when initiating empirical amikacin therapy for VAP caused by a Gram-negative rod. This increase has been recently suggested by Taccone et al [17]. To our knowledge, few data supported this recommendation [21, 22]. Those studies did not use a population approach and were performed in patients with severe sepsis or septic shock, thereby gathering patients with different varieties of infection. Although the influence of the nature of infection has not been thoroughly examined, some authors suggested that the clinical diagnosis might influence the pharmacokinetic parameters of aminoglycosides [10]. We focused our analysis on an homogenous group of patients. Our simulation-based analysis, using data

from a prospective multicenter trial and a solid methodology, confirms the results of Taccone et al [17], and suggest that a higher initial dose is necessary, at least in patients requiring mechanical ventilation.

A comparative study of different initial amikacin dosing regimen would confirm our conclusions. It would also allow to study the impact of higher doses on renal function. Nephrotoxicity is a well-known adverse effect of aminoglycoside therapy. However, this effect has been associated with duration of therapy [23]. Available studies suggest that short-course treatments would allow to minimize toxicity while keeping maximal efficacy [24]. In the majority of severe infections, aminoglycosides are administered as a single infusion in combined antimicrobial therapy. Higher single doses would therefore have potentially limited or no effect on renal function, as suggested by recent studies [21, 22]. It should also be noticed that neuromuscular blockade might have an increased frequency when using higher doses of aminoglycosides [25]. Its main manifestation is respiratory weakness, and this side effect is reversible. It would have a limited impact on patients with mechanical ventilation.

Another result is that amikacin clearance and volume of distribution were estimated to approximately 70 mL/min and 37 L, respectively. These values were in accordance with published data on ICU patients. This confirms the issues observed in antimicrobial pharmacokinetic parameters in critically-ill patients [26, 27], *i.e.* a decrease of drug clearance and an increase of volume of distribution [28-32]. Significant covariates were 4-hour creatinine clearance for amikacin clearance, and total body weight and PaO₂/FIO₂ ratio for central volume.

The association between total body weight and volume of distribution has already been reported [9, 10, 33], and administered amikacin doses are commonly adjusted to total body weight. It is still not clear which measure of body size best describes pharmacokinetic parameters. A recent meta-analysis explored the relationship between drug clearance and body size metrics [34]. The authors found that lean body weight to the exponent 2/3 was more suitable for describing the relationship between

drug clearance and body size. However, they did not question the link between the volume of distribution and body size, and this conclusion may not apply for all pharmacokinetic parameters. As body size was not collected in the IMPACT trial, we could not investigate the effect of this weight metrics or of body mass index on amikacin pharmacokinetic parameters. For hydrophilic drugs such as aminoglycosides, data on the influence of lean body weight and drugs volume of distribution are lacking and would be of interest.

The BIC was not improved by adding total body weight as a covariate for V2. This is quite unexpected, and V2 might represent a weight-independent compartment in which amikacin accumulates. It has previously been shown that total urine recovery of amikacin is not complete 24 hours after its administration, and that aminoglycosides including amikacin accumulates in the kidney [35, 36].

We found a high interindividual variability in amikacin pharmacokinetic parameters as previously reported in studies performed among critically-ill patients [8, 37]. Significant covariates reduced this variability, with a 50% maximal decrease for amikacin CL in our study. To our knowledge, no published covariate model could fully explain the observed interindividual variability. It is therefore highly probable that inside diagnosis-homogenous groups of patients, subpopulations exist that are not individualized by usual covariates. Such variability makes the choice of the optimal dosing strategy extremely challenging from an individual perspective. Considering the absence of severe adverse effect expected in a single-dose setting, mechanically-ventilated patients in intensive care units should receive a high amikacin dose in order to maximize the probability of pharmacodynamic target attainment. Further studies should furthermore focus on identification of covariates allowing for reducing the high unexplained interindividual variability.

The main limitation of this study is that it was restricted to a limited number of patients who had a suspicion on VAP. Therefore the results may not apply to patients who do not require mechanical ventilation. Nevertheless, the French surveillance network of nosocomial infections

recently reported that about 60% of patients hospitalized in ICU would require mechanical ventilation during their stay[38]. Another limitation is the use of the MIC breakpoint for Monte Carlo simulations. To our knowledge, no published data report the distribution of MICs for Gram-negative bacilli responsible for infection in intensive care units. As bacteria involved in infection of critically-ill patients frequently have higher MICs than those isolated from patients hospitalised in other wards, we did not use published MICs distributions that gather strains obtained from all wards. A recent study of MIC breakpoints based on simulations suggested that pharmacokinetic – pharmacodynamic breakpoints are similar to those defined by the EUCAST or the CLSI [39]. In the empirical setting, the worst-case assumption should be preferred when initiating an antimicrobial therapy. This approach is currently used for pharmacokinetic-pharmacodynamic simulations [17, 40, 41]. However, it is likely that the majority of Gram-negative rods involved in critical infections have a MIC below 8 mg/L. For a MIC of 4 mg/L, 96% of patients achieved a C_{1h}/MIC ratio ≥ 10 after a dose of 25 mg/kg in our simulations.

In conclusion, amikacin pharmacokinetic parameters values were similar to those previously reported. This study confirms recent results for the determination of the initial amikacin dose required in critically-ill patients. An empirical dose of 25mg/kg or more is needed to achieve the amikacin pharmacodynamic predictors of clinical efficacy in Gram-negative bacilli infections. Other studies are needed to prospectively evaluate these conclusions in order to improve the management of patients with severe infections.

Authors' contributions

FM, LAL, MW and OP conceived the study

AF, MW and OP provided medical care to trial participants

LM performed the pharmacological assays

CB, CC, CL and FM performed the statistical analysis

CB, FM, MW and OP wrote the first draft of the manuscript. All authors read and agreed with the manuscript in its published version.

Competing interests

All authors have completed the Unified Competing Interest form (available on request from the corresponding author) and declare no support from any organization for the submitted work; no financial relationship with any organisation that might have an interest in the submitted work in the previous 3 years; no other relationships or activity that could appear to have influenced the submitted work.

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