Supplement 1: Validated 2-compartment population pharmacokinetic models of vancomycin.

Validated 2-compartment population pharmacokinetic models of vancomycin for critically ill and non–critically ill patients were used to obtain pharmacokinetic parameters as described below.^{1,2} Critical illness was defined by admission to an intensive care unit (ICU).

Clearance

[1]
$$CL = (\theta_1 \times CL_{Cr}) + (\theta_2 \times TBW)$$
 [for critically ill patients]¹

[2] $CL = (0.0322 \times CL_{cr}) + 0.32$ [for non-critically ill patients]²

CL = vancomycin clearance (mL/min), CL_{Cr} = creatinine clearance (as determined using the Cockroft–Gault equation; mL/min), TBW = total body weight (kg), $\theta_1 = 0.034$, $\theta_2 = 0.015$.

Volume of distribution

 $[3] V_D = (\theta_3 + \theta_4) \times TBW \text{ [for critically ill patients]}$ $[4] V_D = (0.478 \times TBW) + 60.6 \text{ [for non-critically ill patients]}$ $V_D = \text{volume of distribution for both central and peripheral compartments (L), } \theta_3 = 0.414, \theta_4 = 1.32.$

Upon determining vancomycin CL and V_D , the k_e (elimination rate constant) was

calculated according to the following equation:

$$[5] k_e = \frac{CL}{V_D}$$

k_e = elimination rate constant.

$$[6] C_{max} = \frac{C_{trough}}{e^{-k_e(t-t_i)}}$$

 C_{trough} = measured trough level, t_i = time from the end of the infusion period, t = time of the measured trough level, C_{max} = calculated maximum serum vancomycin level following infusion.

[7]
$$C_{min} = C_{trough} \times e^{-k_e(t_d-t)}$$

 C_{trough} = measured trough level, t = time from the measured trough level, t_d = time at the end of the dosing interval, C_{min} = calculated minimum serum vancomycin level during the dosing interval.

Finally, the 24-h area under the curve (AUC₂₄) was calculated according to the following equation³:

$$[8] AUC = \frac{t_v(C_{max} + C_{min})}{2} + \frac{C_{max} - C_{min}}{k_e}$$

AUC = area under the curve for dosing interval, t_v = time over which vancomycin is infused.

Supplementary material for Marko R, Hajjar J, Nzeribe V, Pittman M, Deslandes V, Sant N, et al. Therapeutic drug monitoring of vancomycin in adult patients with methicillin-resistant *Staphylococcus aureus* bacteremia or pneumonia. *Can J Hosp Pharm.* 2021;74(4):334-43.

The AUC₂₄ was determined based on the vancomycin dosing interval. For example, if the dosing interval was scheduled as every 12 h, the AUC determined using equation 8 was multiplied by 2 to obtain the AUC₂₄.

AUC/MIC Determination Using 2 Levels

The AUC/MIC using 2 vancomycin levels was determined based on methods described by Pai and others.³ Cases with 2 levels within one dosing interval were identified, with one level at least 2 h after the end of the infusion (C₁) and a trough level (C₂). The k_e was calculated using a first-order rate equation, where Δt is the time (h) between the 2 levels:

$$[9] C_2 = C_1 \times e^{-k_e \times \Delta t}$$

The extrapolated C_{min} and C_{max} values were then calculated according to the following equation:

$$[10] C_{max} = \frac{C_2}{e^{-k_e \times \Delta t}}$$

(where Δt = time between the end of infusion and trough level)

$$[11] C_{min} = C_2 \times e^{-k_e \times \Delta t}$$

(where Δt = time between trough level and the end of the dosing interval)

Finally, the AUC was calculated according to the following equation:

$$[12] AUC = \frac{t_{infusion} \times (C_{max} + C_{min})}{2} + \frac{C_{max} - C_{min}}{k_e}$$

(where t_{infusion} = the duration of the vancomycin infusion)

The calculated AUC was then divided by the MIC. The AUC/MIC values calculated using

2 levels reflect various time intervals (i.e., the values were not all necessarily AUC_{24}/MIC). These

AUC/MIC values were not included in the discordance analysis.

References

1. Llopis-Salvia P, Jimenez-Torres NV. Population pharmacokinetic parameters of vancomycin in critically ill patients. *J Clin Pharm Ther*. 2006;31(5):447-54.

2. Yamamoto M, Kuzuya T, Baba H, Yamada K, Nabeshima T. Population pharmacokinetic analysis of vancomycin in patients with gram-positive infections and the influence of infectious disease type. *J Clin Pharm Ther*. 2009;34(4):473-83.

3. Pai MP, Neely M, Rodvold KA, Lodise TP. Innovative approaches to optimizing the delivery of vancomycin in individual patients. *Adv Drug Deliv Rev.* 2014;77:50-7.