

Pharmacokinetic/Pharmacodynamic Target Attainment Based on Measured Versus Predicted Unbound Ceftriaxone Concentrations in Critically Ill Patients with Pneumonia: An Observational Cohort Study

Supplementary Materials

File S2: Linear and nonlinear protein binding equations

Linear protein binding

$$CEF_t = CEF_u + CEF_t \times b \quad (\text{Eq. S1})$$

where CEF_t stands for total ceftriaxone concentration (mmol/L), CEF_u stands for unbound ceftriaxone concentration (mmol/L), b stands for the linear protein binding constant.

Nonlinear protein binding

$$EF_t = CEF_u + CEF_u \times \frac{(B_{max} \times (\frac{ALB}{median\ ALB})^h)}{(B_{50} + CEF_u)} \quad (\text{Eq. S2})$$

where CEF_t stands for total ceftriaxone concentration (mmol/L), CEF_u stands for unbound ceftriaxone concentration (mmol/L), B_{max} stands for maximum binding capacity (mmol/L), ALB stands for albumin concentration (mmol/L), $median\ ALB$ stands for the median albumin concentration (mmol/L) in the study population, h stands for Hill coefficient, and B_{50} stands for dissociation constant (mmol/L).