

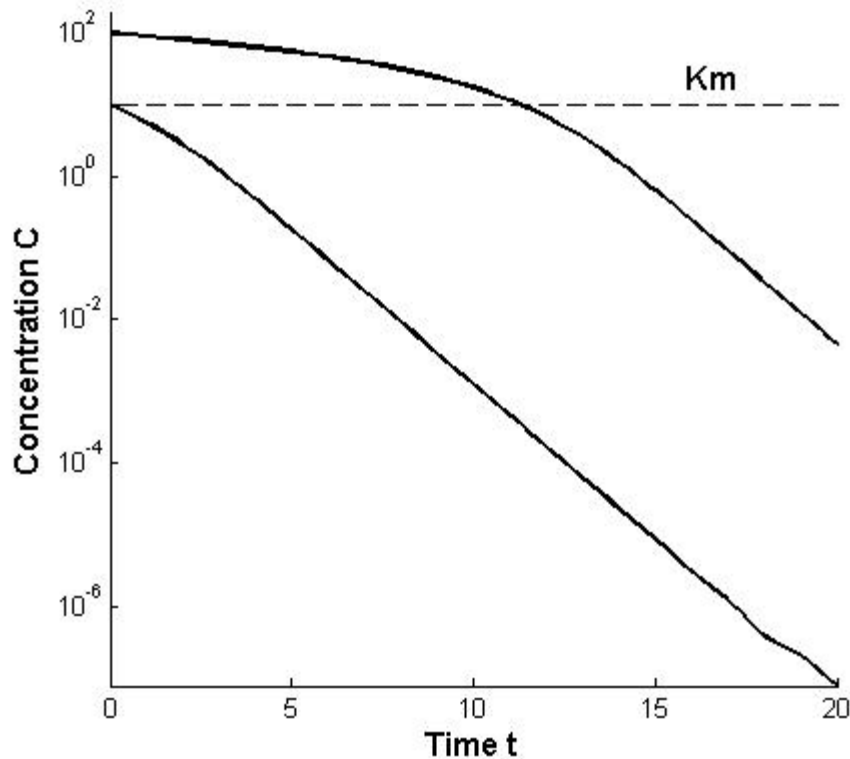
Dose Correction for the Michaelis-Menten Approximation of the Target - mediated Drug Disposition Model

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Michaelis-Menten Elimination Rate



$$V \frac{dC}{dt} = - \frac{V_{\max} C}{K_m + C}$$

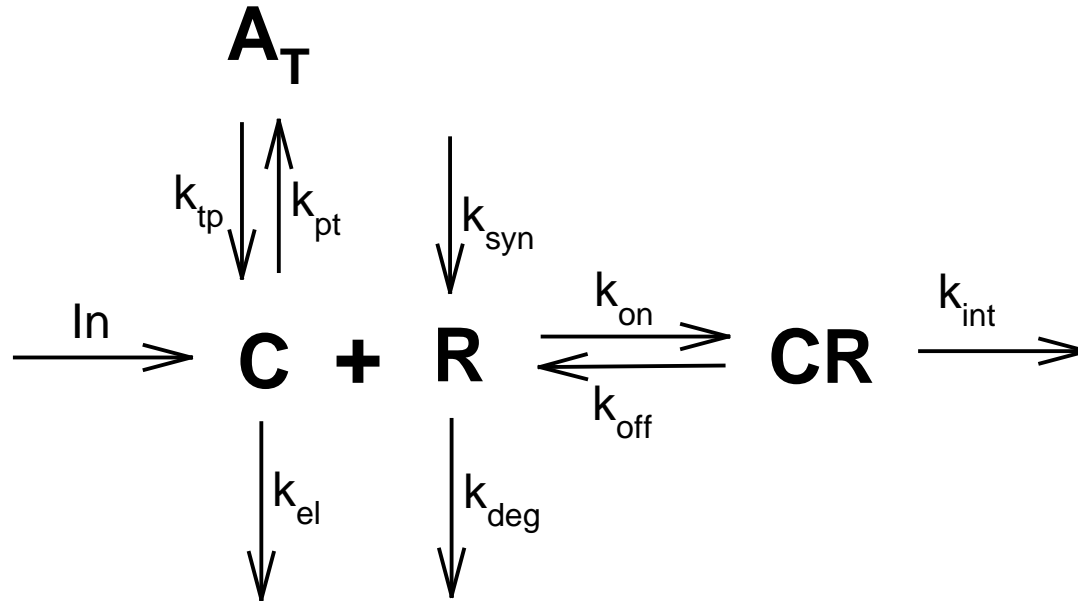
$$C(0) = \frac{\text{Dose}}{V}$$

V_{\max} = maximum elimination rate constant.

K_m = Michealis-Menten constant

Clearance saturation: If $C \gg K_m$ the concentration vs. time plateaus

Target Mediated Drug Disposition PK Model



$$dC/dt = In(t) - k_{on} \cdot R \cdot C + k_{off} \cdot CR - (k_{el} + k_{pt}) \cdot C + k_{tp} \cdot A_T / V_c$$

$$dA_T/dt = k_{pt} \cdot C \cdot V_c - k_{tp} \cdot A_T$$

$$dR/dt = k_{syn} - k_{on} \cdot R \cdot C + k_{off} \cdot CR - k_{deg} \cdot R$$

$$dCR/dt = k_{on} \cdot R \cdot C - (k_{off} + k_{int}) \cdot CR$$

Rapid Binding TMDD PK Model

Equilibrium assumption: Binding of free drug to its target is rapid, so that drug and receptor are in dynamic equilibrium

$$\frac{R \cdot C}{CR} = \frac{k_{\text{off}}}{k_{\text{on}}} \equiv K_D$$

$$C_{\text{tot}} = C + CR \quad \text{and} \quad R_{\text{tot}} = R + CR$$

$$dC_{\text{tot}}/dt = \text{In}(t) - k_{\text{int}} \cdot C_{\text{tot}} - (k_{\text{el}} + k_{\text{pt}} - k_{\text{int}}) \cdot C + k_{\text{tp}} \cdot A_T / V_c$$

$$dA_T/dt = k_{\text{pt}} \cdot C \cdot V_c - k_{\text{tp}} \cdot A_T$$

$$dR_{\text{tot}}/dt = k_{\text{syn}} - (k_{\text{int}} - k_{\text{deg}}) \cdot (C_{\text{tot}} - C) - k_{\text{deg}} \cdot R_{\text{tot}}$$

$$C = 1/2 \left[(C_{\text{tot}} - R_{\text{tot}} - K_D) + \sqrt{(C_{\text{tot}} - R_{\text{tot}} - K_D)^2 + 4 \cdot K_D \cdot C_{\text{tot}}} \right]$$

where C is a solution of the equilibrium equation

Michaelis-Menten Approximation of RB TMDD Model

If $R_{tot} = \text{constant}$:

$$\frac{dC}{dt} = \frac{\text{In}(t) - \frac{k_{int} R_{tot} C}{K_D + C} - (k_{el} + k_{pt})C + k_{tp} A_T / V_c}{1 + \frac{R_{tot} K_D}{(K_D + C)^2}}$$

Wagner, 1971

If $R_{tot} \ll K_D$ or $C \ll K_D$:

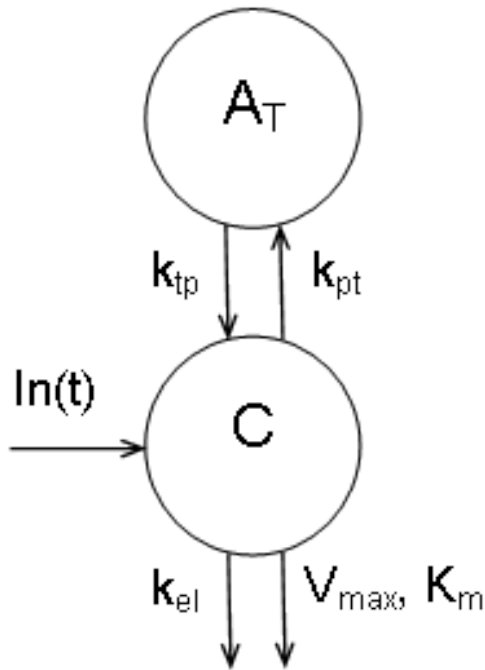
$$\frac{R_{tot} K_D}{(K_D + C)^2} \ll 1$$

Michaelis-Menten approximation:

$$\frac{dC}{dt} = \text{In}(t) - \frac{k_{int} R_{tot} C}{K_D + C} - (k_{el} + k_{pt})C + k_{tp} A_T / V_c$$

$$V_{max} = k_{int} R_{tot} V_c \quad K_m = K_D$$

M-M TMDD Model



$$\frac{dC}{dt} = -\frac{(V_{max} / V)C}{K_m + C} - (k_{pt} + k_{el})C + k_{tp}A_T / V$$

$$\frac{dA_T}{dt} = k_{pt}CV - k_{tp}A_T$$

$$C(0) = \frac{Dose}{V}$$

$$A_T(0) = 0$$

The initial condition for IV bolus dose is Dose/V.

The initial concentration for RB TMDD model $C(0) \neq Dose/V$

Correct Initial Condition for M-M Model

Initial condition based on RB TMDD model:

$$C_{\text{corr}}(0) = \frac{1}{2} \left[\frac{\text{Dose}}{V} - R_{\text{tot}0} - K_D + \sqrt{\left(\frac{\text{Dose}}{V} - R_{\text{tot}0} - K_D \right)^2 + 4K_D \frac{\text{Dose}}{V}} \right]$$

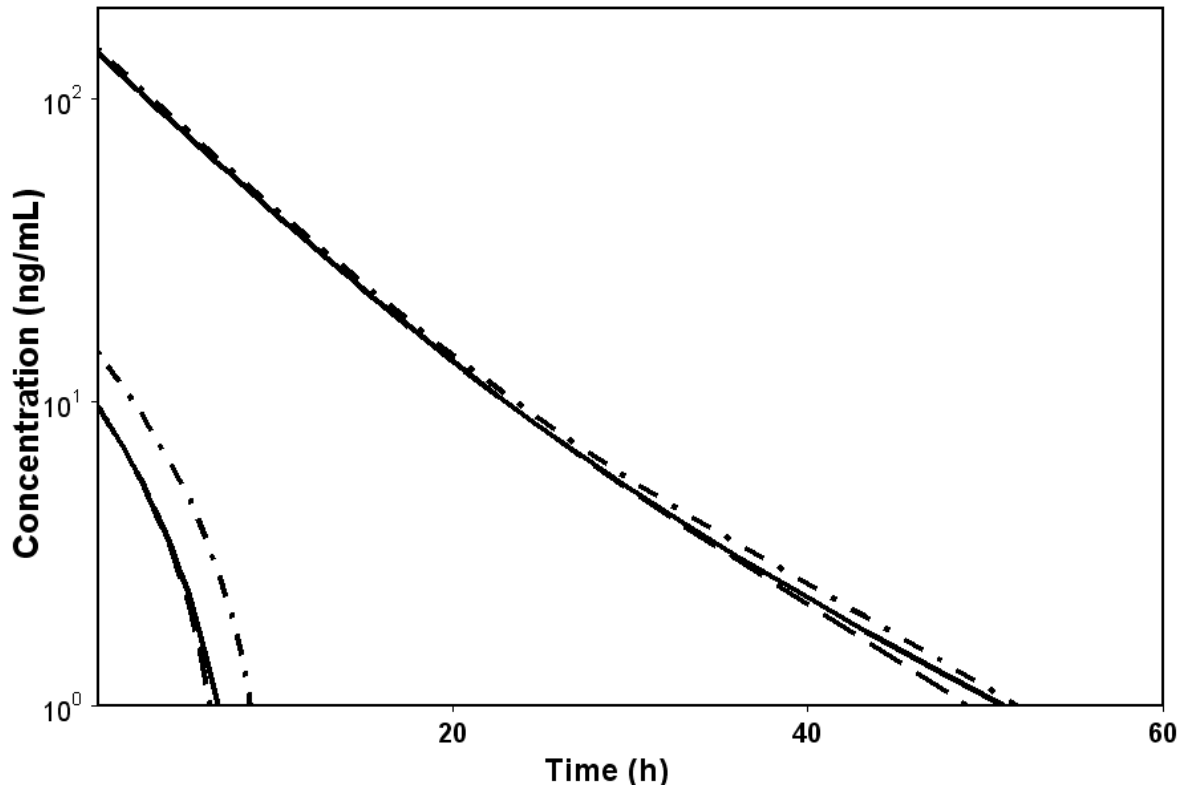
IV bolus dose needs to be corrected by amount bound to receptors:

$$C_{\text{corr}}(0) = \frac{\text{Dose}}{V} - \frac{R_{\text{tot}0} C_{\text{corr}}(0)}{K_D + C_{\text{corr}}(0)}$$

Correction introduces k_{int} into M-M model:

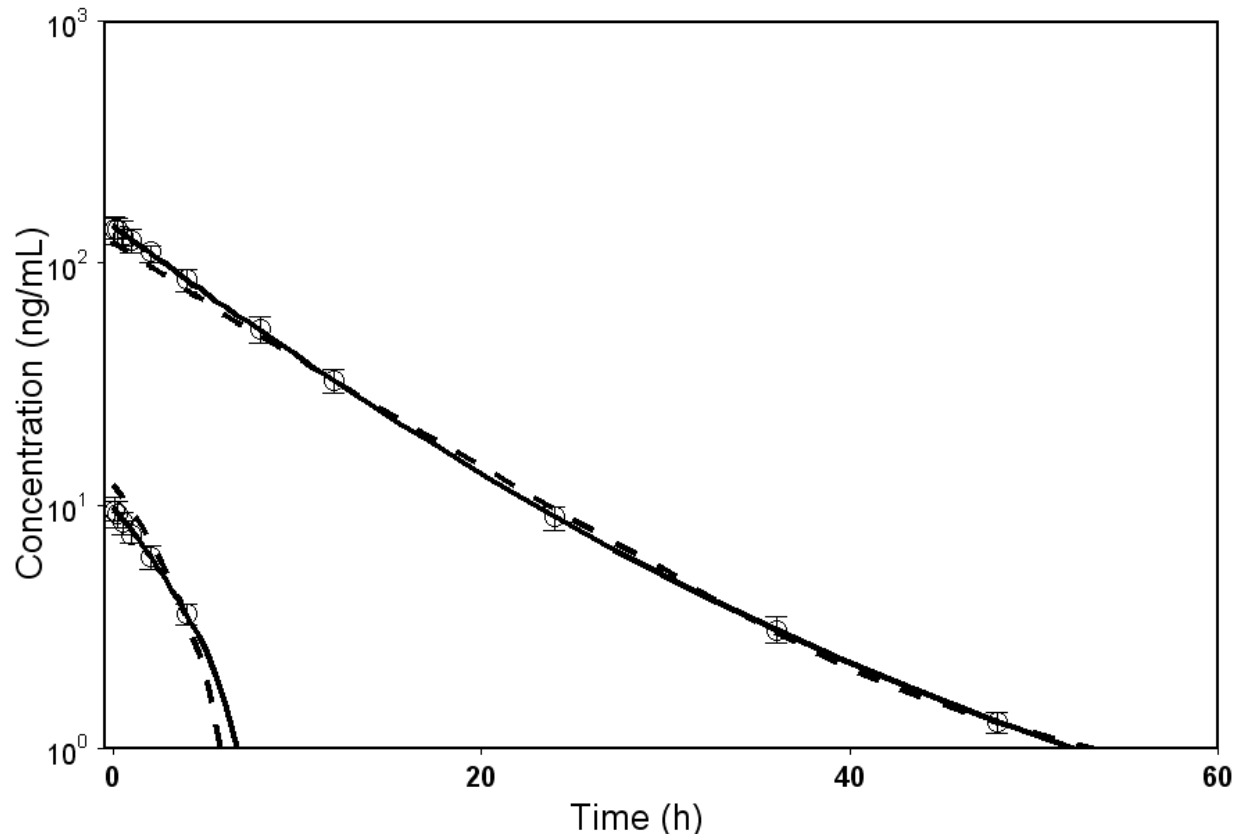
$$C_{\text{corr}}(0) = \frac{1}{2} \left[\frac{\text{Dose}}{V} - \frac{V_{\text{max}}}{k_{\text{int}} V} - K_m + \sqrt{\left(\frac{\text{Dose}}{V} - \frac{V_{\text{max}}}{k_{\text{int}} V} - K_m \right)^2 + 4K_m \frac{\text{Dose}}{V}} \right]$$

Impact of Dose Correction on PK Profile



Simulation of the pharmacokinetic profile with IV bolus doses using the RB TMDD model (solid line), M-M model with corrected initials (dash line), and M-M model with uncorrected initials (dash-dot line).

Bias of Parameter Estimates of M-M TMDD Model



Simulated PK data fitted with the original and corrected M-M models. Open circle represents simulated median data from 100 replicates. Error bar represents the interquartile range. Model predictions are shown for the corrected M-M model (solid lines) and original M-M model (dashed lines).

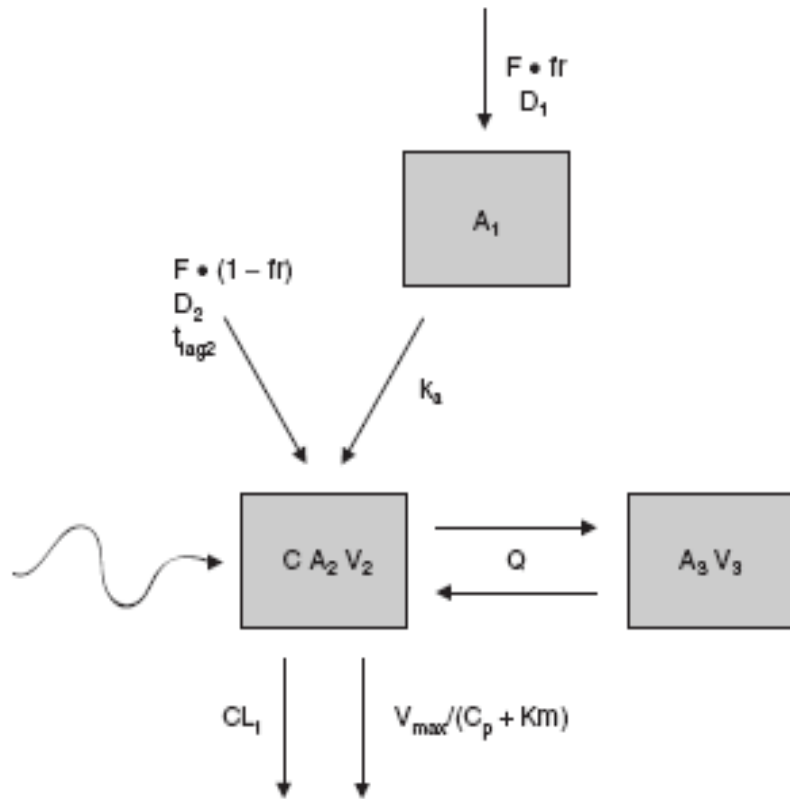
Parameter Estimates

Parameter	Estimate (IQR)			Parameter bias (%)	
	True value	Original M-M	Corrected M-M	Original M-M	Corrected M-M
k_{el} (h^{-1})	0.0382	0.00143 (0.0003, 0.00386)	0.0423 (0.0147, 0.0725)	-96.2	10.6
V_c ($mL\ kg^{-1}$)	68.3	81.9 (78.7, 84.8)	68.3 (64.6, 72.3)	20	0.027
k_{cp} (h^{-1})	0.0806	0.0979 (0.0918, 0.105)	0.078 (0.0594, 0.0962)	21.4	-3.24
k_{pc} (h^{-1})	0.0148	0.0224 (0.0194, 0.0258)	0.0157 (0.0115, 0.0254)	51	5.75
k_{int} (h^{-1})	0.173	N/A	0.167 (0.107, 0.223)	N/A	-3.64
k_{deg} (h^{-1})	0.173 ^a	N/A	N/A	N/A	N/A
K_{eq} ($ng\ mL^{-1}$)	0.131	N/A	N/A	N/A	N/A
R_{tot0} ($ng\ mL^{-1}$)	5	N/A	N/A	N/A	N/A
V_{max} ($ng\ kg^{-1}\ h^{-1}$)	59.1 ^b	144.5 (122.2, 157.7)	56.9 (43.8, 72.1)	145	-3.7
K_m ($ng\ mL^{-1}$)	0.131 ^c	1.06 (0.616, 1.3)	0.29 (0.0685, 0.664)	711	121
WSSR	N/A	0.567 (0.436, 0.748)	0.243 (0.173, 0.321)	N/A	N/A
AIC	N/A	2.63 (-2.11, 7.07)	-10.0 (-15.8, -5.32)	N/A	N/A

^a Fixed at k_{int} value. ^b Calculated as $R_{tot0}/k_{int} \cdot V$. ^c Same to K_D . IQR, interquartile range. N/A, not applicable.

Population Pharmacokinetics Meta-Analysis of Recombinant Human Erythropoietin in Healthy Subjects

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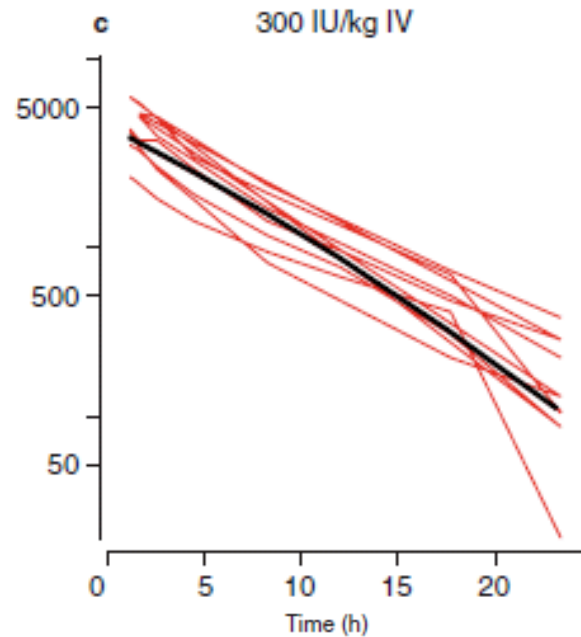
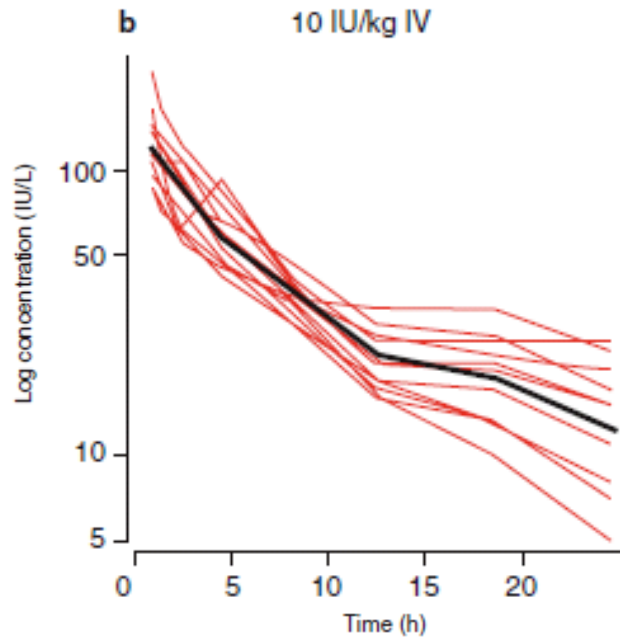
$$\frac{dA_1}{dt} = \begin{cases} \frac{\text{Dose} \cdot \text{fr} \cdot F}{D_1} - k_a A_1, & \text{if } t \leq D_1 \\ - k_a A_1, & \text{if } t > D_1 \end{cases}$$

$$\frac{dA_2}{dt} = \begin{cases} \frac{\text{Dose} \cdot (1 - \text{fr}) \cdot F}{D_2} + k_a A_1 + k_{32} A_3 - k_{23} A_2 - k_{20} A_2 - \frac{V_{\max} A_2 / V_2}{K_m + A_2 / V_2}, & \text{if } t_{\text{lag}2} \leq t \leq D_2 \\ k_a A_1 + k_{32} A_3 - k_{23} A_2 - k_{20} A_2 - \frac{V_{\max} A_2 / V_2}{K_m + A_2 / V_2}, & \text{if } t < t_{\text{lag}2} \text{ or } t > D_2 \end{cases}$$

Correction Factor for Lower Dose EPO Concentrations

$$C = F_{iv} \frac{A_2}{V_2}$$

$$C = \frac{A_2}{V_2}$$



Relative exposure at the two lowest intravenous dose levels, 10 IU/kg and 20 IU/kg, was found to be significantly lower than at higher dose levels. This was best fitted in the model by using a correction factor for these doses.

Parameter	Estimate	RSE (%)
F_{iv} , dose ≤ 20 IU/kg (%)	60.0	7

Conclusions

- Correction of the initial dose the M-M approximation of the TMDD model is necessary due to fraction of dose bound to the receptors.
- Correction is significant for lower doses and negligible for larger doses.
- Inclusion of the correction introduces k_{int} parameter into M-M model.
- Lack of correction biases parameter estimates.
- Further studies are necessary to investigate the impact of the correction on multiple IV bolus data .
- Correcting doses for other routes of administration remains unclear.

Acknowledgments

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