

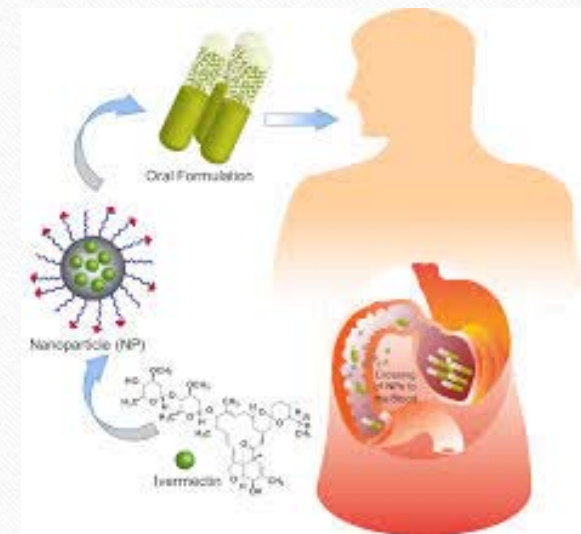
Lab 5

**Determination of Absorption Rate
Constants from Oral Absorption Data**

Absorption Rate Constant



- The overall **rate of systemic drug absorption** from an orally administered solid dosage form encompasses many individual rate processes, including
 1. **Dissolution of the drug,**
 2. **GI motility,**
 3. **Blood flow,**
 4. **Transport of the drug across the capillary membranes and into the systemic circulation.**
- The rate of drug absorption represents **the net result of all these processes.**



Absorption Rate Constant



- For **many immediate release dosage forms**, the absorption process is **first order** due to the **physical nature of drug diffusion**.
- For certain **controlled-release drug products**, the rate of drug absorption may be more appropriately described by a **zero-order rate constant**.



Importance of Absorption Rate Constant



1. The calculation of k_a is useful in designing a **multiple-dosage regimen**.
2. Knowledge of the k_a and k values allows for **the prediction of peak and trough plasma drug concentrations** following multiple dosing.

Importance of Absorption Rate Constant



3. In **bioequivalence studies**, drug products are given in **chemically equivalent** (ie, pharmaceutical equivalents) **doses**, and the respective rates of systemic absorption may **not differ markedly**. Therefore, for these studies, **t_{max}**, or **time of peak drug** concentration, can be very useful in **comparing** the respective rates of absorption of a drug from chemically equivalent drug products.

Method of Residuals



- Assuming $k_a \gg k$ in Equation (1) below:

- $$C_p = \left(\frac{Fk_a D_0}{v_d(k_a - k)} \right) * (e^{-kt} - e^{-k_a t}) \dots\dots 1$$

- The value for the second exponential will become **insignificantly small with time** (ie, $e^{-k_a t} \approx 0$) and can **therefore be omitted**.

- When this is the case, drug absorption is **virtually complete**. Equation (1) then reduces to:

- $$C_p = \left(\frac{Fk_a D_0}{v_d(k_a - k)} \right) * e^{-kt} \dots\dots 2$$

Method of Residuals

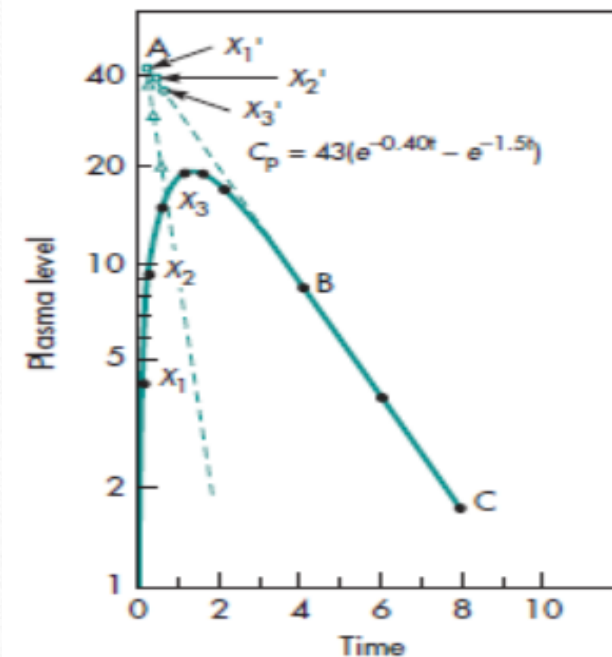


- From equation 1, one may also obtain **the intercept of the y axis**
- $A = \left(\frac{Fk_a D_0}{vd(k_a - k)} \right)$
- where **A is a constant**. Thus, Equation 2 becomes
- $C_p = Ae^{-kt}$ 3
- This equation, which represents **first-order drug** elimination, will yield a **linear plot on semilog paper**. In which the **slope is equal to - k/2.3**.

Method of Residuals



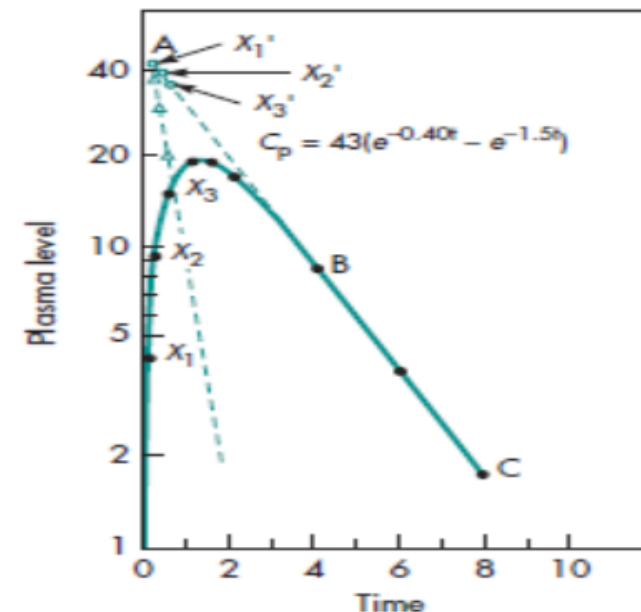
- The **value for k_a** can be obtained by **using the method of residuals** or a **feathering technique**, by the following procedure:
 1. **Plot the drug concentration versus time** on **semilog paper** with the concentration values on the logarithmic axis.
 2. **Obtain the slope of the terminal phase** (line BC,) by extrapolation.



Method of Residuals



3. Take any points on the upper part of line BC (eg, x_1' , x_2' , x_3' , ...) and drop vertically to obtain corresponding points on the curve (eg, x_1 , x_2 , x_3 , ...).
4. Read the concentration values at x_1 and x_1' , x_2 and x_2' , x_3 and x_3' , and so on.
5. Plot the values of the differences at the corresponding time points Δ_1 , Δ_2 , Δ_3 ,
6. A straight line will be obtained with a slope of $-ka/2.3$.



Notes on Method of Residuals



- When using the method of residuals, a **minimum of three points** should be used to **define the straight line**.
- Data points occurring **shortly after t_{max} may not be accurate**, Because this portion of the curve represents the **postabsorption phase**,

Notes on Method of Residuals



- If drug **absorption begins immediately after oral administration**, the residual lines obtained by feathering the plasma level–time curve will **intersect on the y axis at point A**.
- The value of this **y intercept, A**, The value of A has **no direct physiologic meaning**.

Example

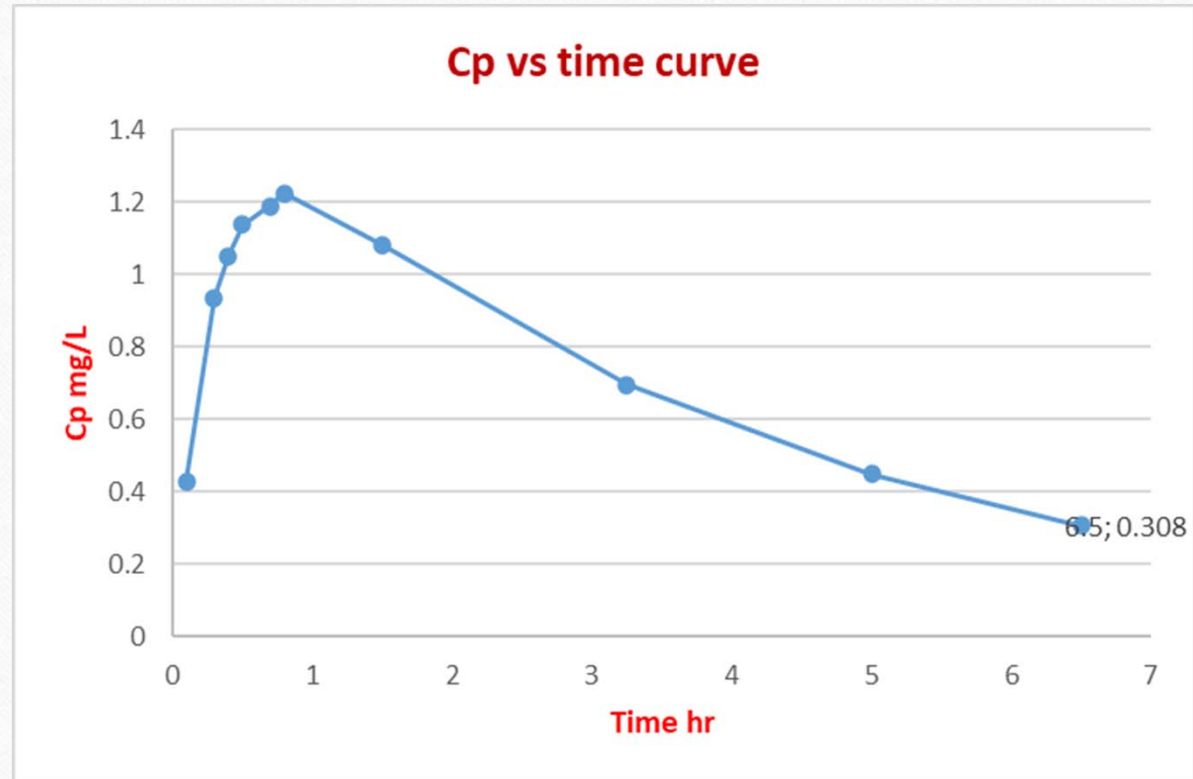


Drug concentrations in the blood at various times are listed in table. Assuming the drug follows a one compartment model.

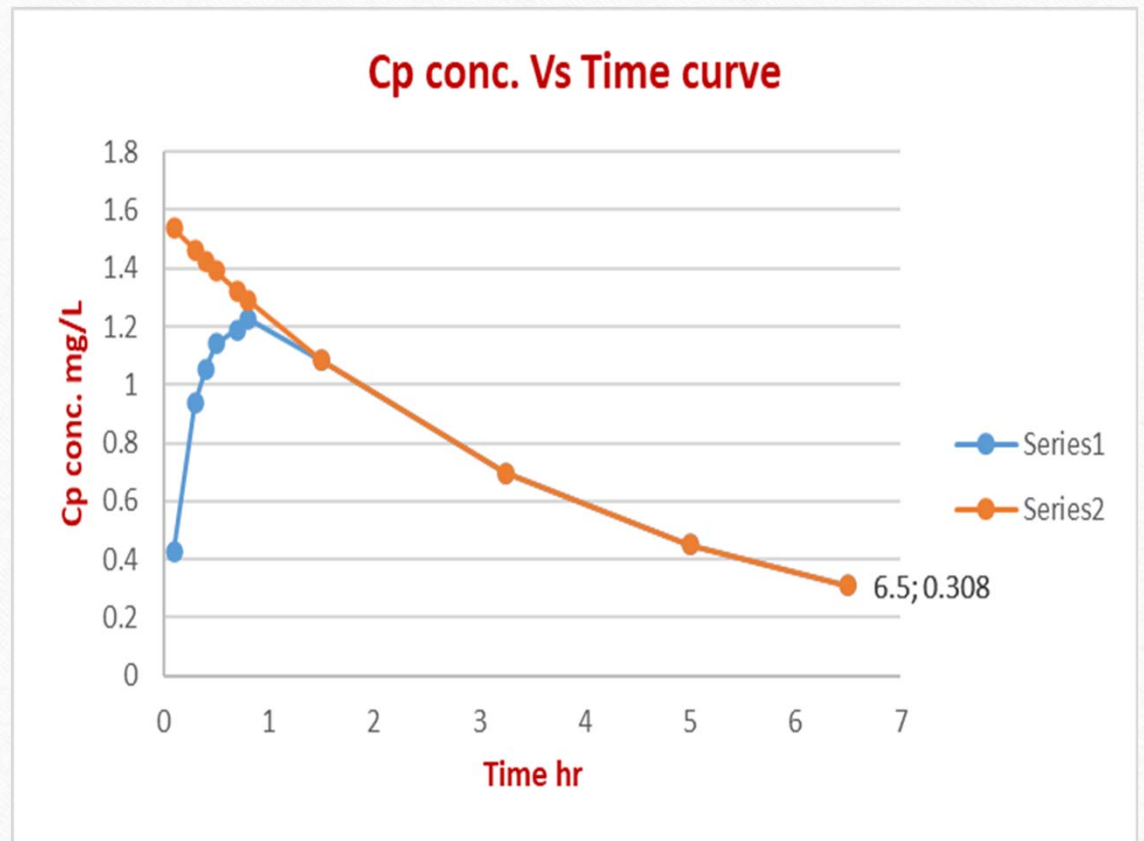
Find the k_a value by the method of residuals.

Time (hr)	Cp (mg/L)
0.1	0.429
0.3	0.936
0.4	1.051
0.5	1.139
0.7	1.189
0.8	1.226
1.5	1.082
3.25	0.697
5	0.449
6.5	0.308

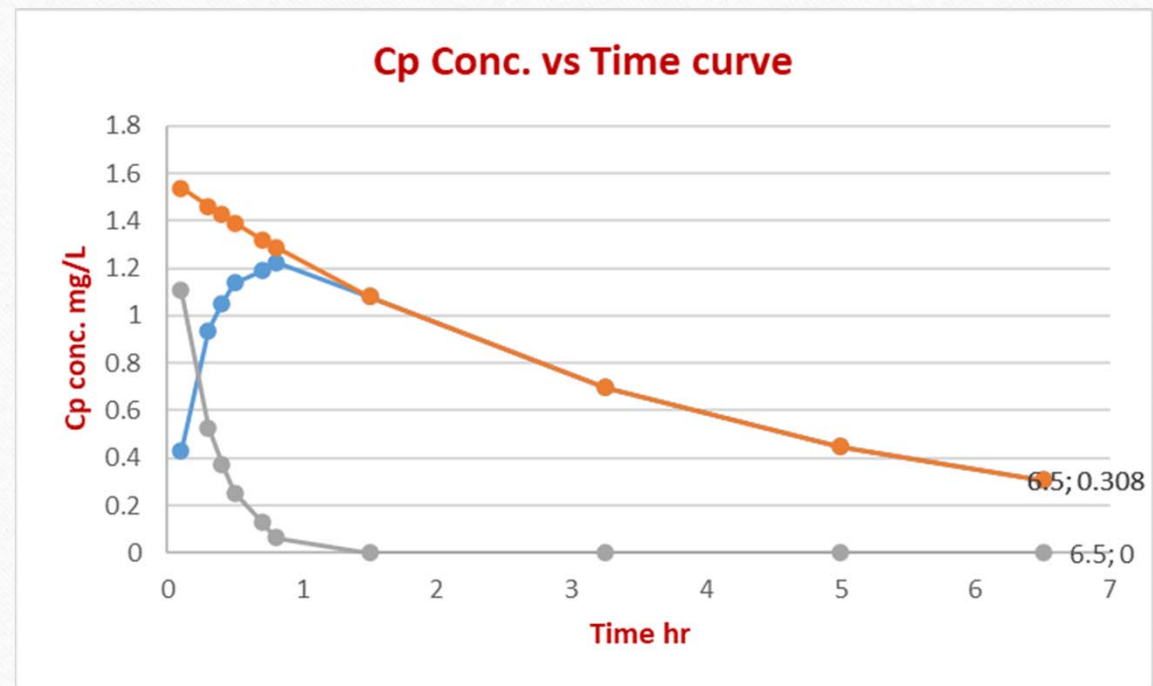
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0.7	1.189
0.8	1.226
1.5	1.082
3.25	0.697
5	0.449
6.5	0.308



Time (hr)	Cp (mg/L)	Cp _{late} (mg/L)
0.1	0.429	1.538
0.3	0.936	1.462
0.4	1.051	1.426
0.5	1.139	1.391
0.7	1.189	1.322
0.8	1.226	1.29
1.5	1.082	1.082
3.25	0.697	0.697
5	0.449	0.449
6.5	0.308	0.308



Time (hr)	Cp (mg/L)	Cp _{plate} (mg/L)	Residual (mg/L)
0.1	0.429	1.538	1.108
0.3	0.936	1.462	0.527
0.4	1.051	1.426	0.375
0.5	1.139	1.391	0.252
0.7	1.189	1.322	0.133
0.8	1.226	1.29	0.064
1.5	1.082	1.082	-
3.25	0.697	0.697	-
5	0.449	0.449	-
6.5	0.308	0.308	-



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$$K_a = -2.303 \text{ slope}$$

$$\text{slope} = (\log cp_2 - \log cp_1)/(t_2 - t_1)$$

$$= (\log 0.53 - \log 1.1)/(0.3 - 0.1)$$

$$= -0.322/0.2$$

$$= -1.61$$

$$\text{So } k_a = -2.303 * -1.61$$

$$= 3.71$$

THANK YOU FOR
YOUR ATTENTION