SAAM II Version 2.2 Basic Tutorials

Working with Samples

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Working with Samples

Prerequisites

The prerequisite for this tutorial is having worked through the **SAAM II** introductory tutorial, "Getting Started with **SAAM II Compartmental**."

What you will learn in this tutorial

The purpose of this tutorial is to show you how to use the **Sample** tool beyond just linking your model to your data. Some examples of such linkage will be given in the Introduction. Other examples will be discussed in this tutorial. You will learn

- How to create PKPD models (Part 1)
- How to estimate the AUC of a sample using the SAAM II AUC option (Part 2)
- How to estimate the AUC using a sum of exponentials (Part 3)

Files Required

Study Files: The study file for this tutorial is

pkpd.stu study_0.stu

These files are included as part of this tutorial. **study_0.stu** is the same study file that is installed in the **SAAM II** program folder.

Introduction

This tutorial focuses on the different uses of the sample tool in the **SAAM II Toolbox**. Normally samples are used to link a compartment in your model to a set of data, and to specify the sample equation which makes the units of model (determined by the units of the input) consistent with the units of the data (determined by the experiment). The typical example is taken from **study_0** shown below:

q1 k(0,1) k(0,1)	1) 2)
Sample Attributes	×
Nam Uni Associate with Data Nam	e: s1 ts: e: plasma
Equations:	
Done Can	cel Help

In terms of reproducing experiments, however, samples can be more complex, and serve purposes other than linking the model to data.

Consider the following example where a drug is given as a bolus into plasma. The drug is known to convert to a metabolite, but when taking plasma samples, only the sum of the drug and metabolite can be measured. The model and sample is shown as follows:



However, even though the data are the sum of the drug and its metabolite, the model is able to predict the behavior of the drug and metabolite separately. This can be done as shown in the following figure:



Here separate samples are associated with Compartments **q1** and **q2**. The sample equation, as shown above, is in terms of concentration. Thus **s2** and **s3**, which are not linked to any data because separate data on the drug and its metabolite do not exist, predict the behavior of the drug and its metabolite. If you plot, as shown below, **plasma**, **s1**, **s2** and **s3**, you can see the model predicted fit, and the contribution of **s2** and **s3** to the solution:



Another similar example comes when radioiodinated low-density lipoprotein (LDL) is injected into plasma. The plasma model contains two compartments; this reflects the kinetic heterogeneity of LDL. In addition, urine data are also collected, and analyzed in terms of individual and cumulative urine samples. This is an example of a single-input three-output experiment. A typical model looks like the following:



The input, **ex1**, is split between Compartments **q1** and **q2**; the sample equation, as shown in the figure above, is a sum of the radioactivity in Compartments **q1** and **q2** divided by the plasma volume. Samples **s2** and **s3** are linked to the cumulative and individual data respectively.

The final example comes from an experiment in which radiolabeled zinc was injected into plasma as a bolus. Serial plasma samples were taken. Radioactivity in red blood cells was also collected. Radioactivity in urine and faeces was measured. Using an external counter, radioactivity in the liver was calculated. This is very similar to MRI or PET studies where images over regions of interest are collected and test substances quantitated. Thus developing models of regions of interest is almost identical to the model for liver radioactivity. The model of the experiment appears as follows:



This model is obviously complex because of the nature of the data collected. Radioactive zinc was injected into Compartment **q1** as indicated by **ex1**. The plasma, urine and faeces data are linked through samples **s1**, **s4** and **s3**.

The red cell system consists of two compartments, Compartment **q5** and **q6**. Since total radioactivity in red cell is counted, the **Sample Attributes** dialog box appears as follows:

Sample Attributes
Name: s5
Units:
Associate with Data Name: rbc
Equations:
s5 = q5 + q6
Done Cancel Help

The liver system consists of two compartments, Compartment **q8** and **q9**. However, these are hypothesized to represent labeled zinc in liver cells while the counter sees both the cells and plasma. The **Sample Attributes** dialog box for the measurement as follows:



s2 is the measurement equation for the liver data. It consists of the zinc in the liver cells, "q8 + q9". Total blood is given by "q1+q5+q6", i.e. blood in plasma and the red cells. However, this is total blood. Only a fraction of total blood is in the liver. Hence *frac* is a parameter that estimates the fraction of total blood that is in the liver. It is known that around 20% of total blood is in the liver, so when fitting the model to the data, *frac* should be around 0.2, or 20%.

The above give some examples of how samples can be used to describe experimental data. The use of samples depends on the particular experiments.

In the following tutorials, we will describe how to create pharmacokinetic-pharmacodynamic (PKPD) models using the **Sample** tool. We will also give two examples of how to estimate area under the curve. This is discussed here because, in SAAM II, the area under the curve is calculated only for samples.

Part 1. Creating PKPD Models

In this tutorial, you will create a PKPD model. The system model for the drug is two compartments. The drug is administered orally, and complete absorption is assumed. The PD model is a direct Emax model. The weighting scheme for the data is 1/y^2 as specified by (GEN 0 1 2); the variance model is data-relative.

- 1. Start the SAAM II Compartmental application. The SAAM II Compartmental main window will open.
- 2. Open the SAAM II Compartmental study file pkpd.
 - a. The file **pkpd.stu** should appear in the file list; if it does not, find the folder where you put this file.
 - b. In the **File** menu, click **Open**. The **SAAM II Compartmental** main window will appear as shown below:



The system model, i.e. the behavior of the drug once in the systemic circulation, is described by Compartments 1 and 2. Compartment 3 is the compartment into which the drug is introduced; the upper part of the GI tract. Delay 4 simulates the transit time through the GI tract as the drug is absorbed.

3. View the model and the experiment on the model. In the **SAAM II Toolbox**, click **Experiment**. The model of the experiment will appear on the **Drawing Canvas** as follows:



The experiment as been created to analyze the plasma data collected following a single dose of a tablet containing 400mg of the drug. The dose can be confirmed by opening the **Exogenous Input** dialog box. The **Sample Attributes** dialog box associated with s1 are shown below:

ample Attributes	
Name:	s1
Units:	mg/l
Associate with Data Name:	conc
s1 = q1/Vc	
	<u>×</u>

The sample is simply concentration in terms of mg/l, and the plasma data are named "conc". The **Parameters** dialog box appears as follows:

Name	Туре	Current	Low Limit	High Limit
Vc	Adj	50.0000	5.0000	100.0000
k(0,1)	Adj	0.5000	0.0500	5.0000
k(1,2)	Adj	0.0100	0.0010	0.1000
k(2,1)	Adj	0.0100	0.0010	0.1000
k(4,3)	Adj	1.0000	0.1000	10.0000
tlag	Adj	0.5000	0.1000	1.0000
Name: Vc	Valu	e: 50.000000		Edit - J
Name: Vc Type: C Fixed C Adjustable	Valu Low Lim High Lim	e: [\$0.0000000 it: [5.00000000 it: 100.000000		Edit 년 ave소

Click Cancel or Done to close the Parameters dialog box.

- 4. Solve the model, and view the solution.
 - a. In the **Compute** menu, click **Solve**, or alternatively, on the **SAAM II Toolbar**, click **Solve**
 - b. In the Show menu, click Plot, or alternatively, on the SAAM II Toolbar, click Plot
 Image: The Plot and Table Variables dialog box will open. Be sure the List All Variables check box is not selected.
 - c. Click **s1:conc** to move this to the **Current Selection** pane. The **Plot and Tables Variables** dialog box will appear as follows:



d. Click **Done**. Your plot will appear as shown below (in linear mode):



Leave the **Plot** window open.

5. Fit the model to the data.

a. Fit the model to your data. In the **Compute** menu, click **Fit**, or alternatively, on the



- b. Close the **Plot** window.
- 6. Create the PD model.

Besides collecting plasma data, an effect was also measured. The key in PKPD models is linking the PK model to the effect, or PD, data.

a. View the data. In the Show menu, click Data, or alternatively, on the SAAM II
 Toolbar click Data
 The Data window will open as follows:

d _{ij} Data - pkpd.dat	- O ×
Gij Data - pkpd.dat DATA #time in hours; conc in mg/ml (GEN 0 1 2) t conc .5 7.24 1 14.21 2 17.2 4 7.27 6 3.18 9 1.25 12 .8 24 .23 36 .11 48 .05 60 .03 END # DATA #time in hours; effect in % (GEN 0 1 2) t effect .5 85.69 1 93.90 2 94.43 4 85.20 6 74.62 9 50.48 12 42.31 24 20.85	
END	•

You can see there are two sets of data. One, "conc" is the plasma concentration of the drug. The other, "effect" is a measurement of an effect of the drug in terms of %. In the above, the weights assigned to both sets of data are the same. It is often the case that the "effect" data are known with less precision than the plasma data, and hence are assigned less weight.

Close the **Data** window.

- b. In the **SAAM II Toolbox**, click **Sample**.
- c. Click Compartment **q1** and then click the **Drawing Canvas**. The sample **s2** will appear associated with Compartment **q1**.



- d. Double-click s2 to open the Sample Attributes dialog box.
 - (1) Type "effect" in the Associate with Data Name box.
 - (2) Type "s2=Emax*s1/(EC50+s1)" in the **Equations** box. The **Samples Attributes** dialog box will appear as follows:

Sample Attributes		×
	Name: s2 Units:	
Associate with	Data Name: ef	fect
Equations:	145	
s2 = Emax*s1/(EC5	0+s1)	
		-
P	Transfer II	
Done	Cancel	Help

(3) Click **Done**. The model will appear as follows:



Notice s2 is no longer associated with any compartment in the model. This is because the equation for s2 does not contain any of the **qi**.



Samples and PKPD models. In SAAM II, the model is linked to the data via the **Sample** tool. When you create such samples, it does not matter which compartment you click. The reason is that the "link" equation involves a plasma measurement, and hence will contain, as illustrated above, "**s1**" and not "**q1**". If, on the other hand, you had written the sample equation

"s2=Emax*(q1/Vc)/(EC50 + (q1/Vc))",

s2 would be linked to Compartment q1 since "q1" appears in the equation; the equivalence is because "s1=q1/Vc".

Obviously there are several different effect models that can be used; they can be specified in this fashion in SAAM II.

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- 7. Enter the new parameters, solve the model, and view the solution.
 - a. Open the **Parameters** dialog box. Two new parameters, *Emax* and *EC50* will appear in the list. *Enter the values as shown below* (remember since you have previously Fitted the model to the plasma data, the system parameters are those from the best fit). The **Parameters** dialog box should appear as follows:

Name	Туре	Current	Low Limit	High Limit
EC50	Adj	10.0000	1.0000	20.0000
Emax	Adj	80.0000	50.0000	200.0000
Vc	Adj	15.3199	5.0000	100.0000
k(0,1)	Adj	0.3605	0.0500	5.0000
k(1,2)	Adj	0.0745	0.0010	0.1000
k(2,1)	Adj	0.0691	0.0010	0.1000
k(4,3)	Adj	1.6776	0.1000	10.0000
tlag	Adj	0.2902	0.1000	1.0000
	Valu	e: 80		Fditピー
Type: C Fixed	Low Limi	it: 50.0000000		ave 分
Type: O Fixed • Adjustable	Low Limi High Limi	it: 50.0000000		ave D
Type: O Fixed Adjustable Adjust value:	Low Lim High Limi	it: 50,0000000		ave D

Notice for both *Emax* and *EC50* the **Low** and **High Limits** were separately set. Click **Done** to close the **Parameters** dialog box.

b. Solve the model, and view the solution. In the **Plot and Tables Variables** dialog box, select both **s1:conc** and **s2:effect**. The plot of **s1:conc** (which will not change from the previous best-fit solution) and **s2:effect** will appear as follows:



Leave the **Plot** window open.



8. Fit the model to the data. The **Plot** window will be updated as follows:

The **Statistics** window will appear as follows:

Parameter/Variable	Value	Std.Dev.	Coef. of Var.	95% Confidence	Interval	
EC50	1.08787	1.12842e-001	1.03727e+001	0.84201	1.33373	
Emax	99.47768	4.00656e+000	4.02759e+000	90.74815	108.20720	1
Ve	14.71656	1.92542e+000	1.30834e+001	10.52143	18.91169	
k(0,1)	0.37365	3.71837e-002	9.95148e+000	0.29263	0.45467	
k(1,2)	0.08218	6.77301e-003	8.24187e+000	0.06742	0.09694	100
k(2,1)	0.08067	1.42906e-002	1.77139e+001	0.04954	0.11181	
k(4,3)	1.58733	5.38182e-001	3.39048e+001	0.41474	2.75993	
C Correlation Matrix	C Covarian	ce Matrix 🔎	Objective			
		Objective	Scaled Data V	ariance		
s2 : effect	1.37936	9e+000	5.152350e-00	3		
s1 : conc	-2.03375	1e+000	1.402546e-00	2		
Total objective	-6. <mark>54</mark> 382	8e-001				
AIC	1.09174	7e+000				L
BTC	1 34068	0e+000				

Close the **Plot** and **Statistics** windows.

Quit the SAAM II Compartmental application. Do not save the changes to pkpd.stu.

This is an example of how a PKPD model can be created in SAAM II. The link between the PK model and the effect data comes via the sample equation s2 containing the Emax model. There are many other options available to specify the link. Consider the following model:



This is a model in which a drug was introduced into plasma Compartment **q1**, and serial plasma samples were taken. The system model for the metabolism of the drug consists of Compartments **q1**, **q2** and **q3**. The sample **s1** is typical; it links the model with the plasma data. Compartment **q4** is the biophase or link compartment. It will be used to link the pharmacokinetics of the drug with the measured effect of the drug.

The input, **ex2**, into Compartment **q4** is defined "ex2=ki*s1". This links the pharmacokinetics of the drug with Compartment **q4**, and introduces a new parameter, *ki* which will be the compartment turnover. The sample **s2** is also typical; the sample equation is "s2 = q4/V4"; V4 is a new parameter in the model. The sample is not linked to any data.

The pharmacodynamic equations are written in the **Equations Defined Here** pane in the **Equations** dialog box as shown below:



The first equation describes the relationship between the observed effect, **E**, the initial measurement of the drug effect, *Eo*, the concentration of drug in the link compartment, **Ce**, and the constants of the Emax model, *Emax* and *IC50*. The second equation specifies **Ce** as equal to **s2**. The third equation specifies that the maximum drug effect, *Emax*, would simply consist of a complete reversal of the effect that was initially observed, *Eo*. In the last equation k(0,4) is defined as a fractional clearance that is equal to *ki*. In terms of Compartment **q4**, this is equivalent to an intercompartmental fractional clearance k(4,1). However, the system model is structured without actual mass transfer between the pharmacokinetic model and the effect compartment. This allows the effect kinetic data to be analyzed without perturbing the pharmacokinetic "Fit."

Part 2. Estimating the AUC for a Sample using the AUC Option in the Computational Settings Dialog Box.

There are two ways one can estimate area under the curve in the **SAAM II Compartmental** application. One way is to use the **AUC** command in the **Computational Settings** dialog box. This will be described in this tutorial.

- 1. Start the SAAM II Compartmental application. The SAAM II Compartmental main window will open.
- 2. Open the **SAAM II Compartmental** study file **study_0**.
 - a. The file **study_0.stu** should appear in the file list; if it does not, find the folder where you put this file.
 - b. In the **File** menu, click **Open**. The **SAAM II Compartmental** main window will appear as shown below:



3. View the model and the experiment on the model. In the **SAAM II Toolbox**, click **Experiment**. The model of the experiment will appear on the **Drawing Canvas** as follows:



This is the experimental model developed and used in the **Getting Started with Compartmental** tutorial.

- 4. Estimate the area under **s1** and the first-moment (mean) area under **s1** for the duration of the experiment.
 - a. In the **Compute** menu, click **Computational Settings**. The **Computational Settings** dialog box will open.
 - b. In the **Integrator** pane, select **Compute Sample AUC's**. The **Computational Settings** dialog box will appear as follows:

Rosenbrock Pade Runge-Kutta	Use Relative Error: 0.00100000 (1.0e-10 to 1.0) Use Absolute Error:
Max. Nr. of Fit Iterations	(greater than 0.0) 20 (0 to 50) Derivative
C Data C Abs C Model Convergence Criterion:	solute Forward C Central
Include Bayesian Term Lambdar 10,000000	(1.0e-7 to 1.0) 00 (1.0e-7 to 1.0e7
Save Results to Text F	File C Replace C Add



Sample AUC's. SAAM II will not estimate the sample area under the curve (AUC) unless this option is selected. SAAM II estimates AUC's only for samples. Thus, for example, if you wanted the AUC for Compartment q^2 in the model, you would need to put a sample on this compartment.



- c. Create a sample to estimate the first-moment (mean) area under s1.
 - (1) In the SAAM II Toolbox, click Sample.
 - (2) Click Compartment **q1** and then the **Drawing Canvas**. A sample **s2** will appear associated with Compartment **q1**.
 - (3) Double-click s2 to open the Sample Attributes dialog box.
 - (4) Type "s2=t*q1/vol" in the **Equation** box. The **Sample Attributes** dialog box will appear as follows:

Sample Attributes		2
	Name: s2	
	Units:	
Associate with Da	ata Name:	
-		
Equations:		
s2 = t*q1/vol		-
~		-
Done	Cancel	Help
	Sector Sector	

The model will appear as follows:



(5) Click Done.



Samples and AUC's. The sample for the first-moment (mean) s2 is written in terms of q1, and hence is associated with Compartment q1. The equation could have been written

"s2=t*s1"

In that case, since **q1** does not appear directly in the equation, **s2** would no longer be associated with Compartment **q1**.



- d. Fit **1** the model to the data. You can view the solution if you wish.
- e. View the AUC's.
 - (1) In the **Show** menu, click **AUC's**. The **Area Under Curve** window will open as follows:

🔜 Area U	<u>- 0 ×</u>		
Sample	AUC	Std.Dev.	<u>-</u>
s1	1.2225e+005	1.2756e+003	
s2	3.4048e+005	3.7761e+003	
			T
<			× //

Because you have Fitted the model to the data, you will obtain standard deviations for the AUC's. If you had Solved the model, the standard deviations would not appear. Remember the AUC for s1 is the area under the plasma curve, and the AUC for s2 is the area under the first moment.

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AUC's. When using the AUC option in the Computational Settings dialog box, the AUC will be calculated up to the end of the experiment. In this case, the numbers reported above are the AUC up to day 9. You can thus calculate the AUC over any time interval you wish (starting from time zero). To obtain an estimate of the AUC to time infinity, you need to set the time of the experiment appropriately long. This will be explained below.



- (2) Close the Area Under Curve window.
- 5. Estimate the AUC to "infinity".

To obtain an estimate of the AUC to infinity, you need to set the time of the experiment to an appropriately long value. While there is no set rule on how to do this, a reasonable approach is the following.

Calculate the inverse of the smallest rate constant in your model. By looking either in the **Parameters** dialog box or **Statistics** window, this value is approximately 0.175; the inverse is about 6.

Set the time of the experiment to 10 times this value, and Re-Fit the model to the data. In this case, the time will be set equal to 60. Then double this time, and Re-Fit the model to the data. The time will be 120. Keep doubling the time until there is no discernable difference between successive AUC calculations.

- a. In the Set menu, click Experimental Attributes.
- b. Change the length of the experiment from "9" to "60". The **Experiment Attributes** dialog box will appear as follows:

periment Attributes	
Independent Variable:	t
Units:	days
Start at:	0.0
End at:	60

- c. Click Done.
- d. Re-Fit the model to the data, and view the AUC's. The **Area Under Curve** window will appear as follows:

s1 1.5310e+005 6.4296e	+003
s2 8.4461e+005 1.3998e	+005

These numbers are obviously different from the previous estimates because the "integration" has gone from 9 to 60.

e. Set the time of the experiment equal to 120. Re-Fit the model to the data and view the AUC's. The **Area Under Curve** window will appear as follows:

🔜 Area U	_ 🗆 🗙		
Sample	AUC	Std.Dev.	<u>^</u>
91	1.5313e+005	6.4775e+003	
s2	8.4649e+005	1.4330e+005	
			T
4			► //.

You can see now there is little difference; it is in the third significant figure for s2. If you set the time of the experiment equal to 240 and Re-Fit the model to the data, the **Area Under Curve** window will appear as follows:

Sample	AUC	Std.Dev.	-
 s1	1.5313e+005	6.4775e+003	
s2	8.4655e+005	1.4314e+005	

Now there is no difference in s1, and the difference in s2 is in the fourth significant figure. It is probably safe to stop at this stage.

f. Close the Area Under Curve window.

Quit the SAAM II Compartmental application. Do not save the changes to study_0.stu.

Using this option may seem a little cumbersome if the area from zero to infinity is all that is desired. For a simple bolus injection into plasma, this can be the case. However, the option does allow you to estimate the area over different time intervals, something that can be important in observing the behavior of a substance in the body.

This option is really useful when you have inputs other than the bolus, especially multiple inputs. Writing equations to estimate areas in this case can be very tricky. Using other methods such as trapezoidal or log-trapezoidal rules with extrapolations can get even trickier.

For the simple bolus injection, sums of exponentials can be used in the **SAAM II Compartmental** application. This is explained in Part 3.

Part 3. Estimating the AUC for a Sample a Sum of Exponentials.

There are two ways one can estimate area under the curve in the **SAAM II Compartmental** application. One way is to use the **AUC** command in the **Computational Settings** dialog box as illustrated in Part 2. The other is to use sums of exponentials. This will be described in this tutorial.

- 1. Start the SAAM II Compartmental application. The SAAM II Compartmental main window will open.
- 2. Open the **SAAM II Compartmental** study file **study_0**.
 - a. The file **study_0.stu** should appear in the file list; if it does not, find the folder where you put this file.
 - b. In the **File** menu, click **Open**. The **SAAM II Compartmental** main window will appear as shown below:



3. View the model and the experiment on the model. In the **SAAM II Toolbox**, click **Experiment**. The model of the experiment will appear on the **Drawing Canvas** as follows:



This is the experimental model developed and used in the **Getting Started with Compartmental** tutorial.

4. Since you have previously worked with **study_0**, you can proceed to Fit the model to the data, and view the solution. The plot of **s1:plasma** will appear as follows (in linear mode):



The statistics will appear as follows:

Σ <mark>,</mark> Statistics					_	
Parameter/Variable	Value	Std.Dev.	Coef. of Var.	95% Confidence	e Interval	
k(0,1)	0.29227	1.46155e-002	5.00071e+000	0.26010	0.32444	A
k(1,2)	0.28311	7.51183e-002	2.65335e+001	0.11777	0.44844	
k(2,1)	0.17452	1.82636e-002	1.04652e+001	0.13432	0.21472	
vol	2323.77661	3.85619e+001	1.65945e+000	2238.90219	2408.65103	
						Ţ
C Correlation Matrix	C Covariant	e Matrix 📀	Objective			
		Objective	Scaled Data V	ariance		
s1 : plasma	1.33674	3e+001	1.113515e-00	1		
Total objective	1.33674	3e+001				
AIC	7.93598	7e+000				
BIC	8.05399	5e+000				-
1						▶

Close the **Plot** and **Statistics** windows.

- 5. Estimate the area under the plasma curve and first-moment (mean) using a sum of exponentials.
 - a. Open the **Parameters** dialog box, and fix all model parameters. Select the first variable, click **Edit**, select **Fix**, and then click the **Save** button. Repeat for all variables. The **Parameters** dialog box will appear as follows:

Turumeters				
Name	Туре	Current	Low Limit	High Limit
k(0,1)	Fix	0.2923		
k(1,2)	Fix	0.2831		
k(2,1)	Fix	0.1745		
vol	Fix	2323.7766		
Name: vol Type: • Fixed	Val Low Li	lue: 2323.776	61009	⊐Edit⊅
O Adjustable	High Lir	nit: 20000.00	000000	Save之
Adjust value:				
Adjust value:				
Adjust value:			_ _	Auto solve

The reason you fix the model parameters is that, when you write your sum of exponentials, you introduce a new "numerical" model. The parameters of this model, as you will see, are adjustable so the compartmental model parameters must be fixed.

Click **Done** to close the **Parameters** dialog box.

- b. Create a sample to describe the plasma data by a sum of exponentials.
 - (1) In the **SAAM II Toolbox**, click **Sample**.

- (2) Click Compartment **q1** and then the **Drawing Canvas**. A sample **s2** will be associated with Compartment **q1**.
- (3) Double-click s2 to open the Sample Attributes dialog box.
- (3) Type "plasma" in the Associate with Data Name box.
- (4) Type the equation " $s^2 = A1 \exp(-a1 t) + A2 \exp(-a2t)$ " in the **Equations** box. The **Sample Attributes** dialog box will appear as follows:

Sample Attributes
Name: s2 Units: Associate with Data Name: plasma
Equations: s2 = A1*exp(-a1*t)+A2*exp(-a2*t)
<u></u>
Done Cancel Help

(5) Click **Done**. The following error message will appear:





Associating samples and data. In SAAM II, you cannot associate the same data name with more than one sample. To solve this problem, since the compartmental model and its sample **s1** has already been fitted, you can remove "plasma" from **s1** so that you can use it with **s2**.

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- (6) Click **OK**.
- (7) In the **Sample Attributes** dialog box, click **Cancel**.

- (8) Double-click s1 to open the Sample Attributes dialog box.
- (9) Remove "plasma" in the Associate with Data Name box.
- (10) Click **Done**. Because the sample equation remains, and the best fit parameters remain in the **Parameters** dialog box, the solution for **s1** will remain the best fit solution.
- (11) Double-click s2 to open the Sample Attributes dialog box.
- (12) Type "plasma" in the Associate with Data Name box.
- (13) Type the equation " $s2 = A1 \exp(-a1 t) + A2 \exp(-a2 t)$ " in the **Equations** box.
- (14) Click **Done**. The model will appear as follows:



s1 is an open circle now because it is no longer associated with data. **s2** is a closed circle because it is associated with data. It is not connected to Compartment **q1** because **q1** does not appear in the equation for **s2**.

- c. Estimate the area under the curve AUC and the mean area under the curve MAUC.
 - In the Show menu, click Equations, or alternatively, on the SAAM II Toolbar, click Equations Eq. The Equations dialog box will open.
 - (2) Enter the following equations in the **Equations Defined Here** pane:

AUC = A1/a1 + A2/a2 $MAUC = A1/a1^2 + A2/a2^2$

The **Equations** dialog box will appear as follows:



(3) Close the **Equations** dialog box.



Sums of exponentials. Remember the number of exponential terms in your sum of exponentials equals the number of compartments in your model (in most cases). In this case, there are two compartments and hence two exponential terms. If there had been three compartments, there would have been three exponential terms. The formulas for AUC and AUMC are easy in this case because the mode of administration is a bolus. If the mode of administration were different, the number of exponential terms would still be two, but the expressions for AUC or AUMC could change depending upon the circumstances. In the case of multiple boluses or multiple infusions followed by a washout, the formulas can be very complex. For this reason, the scheme presented in Part 2 is more efficient.



d. Enter the parameters for the sum of exponentials. Open the **Parameters** dialog box. Notice all the model parameters remain and are fixed. The new parameters are those for the sum of exponentials -A1, A2, a1 and a2. Enter the values as shown in the **Parameters** dialog box below:

Parameters						
Name	Туре	Current	Low Limit	High Limit		
A1	Adj	40000.0000	4000.0000	4.000e+005		
A2	Adj	10000.0000	1000.0000	1.000e+005		
al	Adj	0.5000	0.0500	5.0000		
a2	Adj	0.0500	0.0050	0.5000		
k(0,1)	Fix	0.2923				
k(1,2)	Fix	0.2831				
k(2,1)	Fix	0.1745				
vol	Fix	2323.7766				
Name: a2 Value: 0.05						
 Adjustable 	Low Li High Li	mit: 0.0050000		Save		
Adjust value:				Auto solve		
Done	Cancel	Help				

e. Fit the model to the data (remember the model now is the sum of exponentials). In the **Plot and Tables Variables Dialog Box**, select **s2:plasma** and click **Done**. The plot of **s2:plasma** will appear as follows:



III Table			
t	plasma	s1	s2
0.000	-	44754.732	44749.022
7.00000e-003	46780.000	44608.789	44603.249
4.20000e-002	43522.000	43887.777	43883.067
0.125	42535.000	42234.517	42231.621
0.250	40125.000	39887.687	39887.149
0.375	36221.000	37701.625	37703.041
0.500	35562.000	35664.642	35667.660
0.750	-	31995.409	32000.745
1.000	28194.000	28802.604	28809.324
1.450	-	24050.218	24057.848
1.725	-	21664.652	21672.096
2.000	19573.000	19601.193	19608.130
2.450	-	16796.027	16801.837
2.725	-	15368.982	15373.936
3.000	14403.000	14121.006	14125.098
3.450	-	12397.968	12400.771
3.725	-	11506.245	11508.309
4.000	11278.000	10715.665	10717.068
4.450	-	9603.457	9603.980
4.725	-	9016.183	9016.260
5.000	8081.000	8487.392	8487.102
5.450	-	7728.050	7727.333
5.725	-	7318.556	7317.654
6.000	6999.000	6944.009	6942.981
6.450	-	6395.269	6394.150
6.725	-	6093.436	6092.311
7.000	5653.000	5813.411	5812.317
7.450	-	5395.888	5394.908
7.725	-	5162.366	5161.483
8.000	5139.000	4943.185	4942.417
8.450	-	4611.791	4611.240
8.725	-	4424.042	4423.634
9.000	-	4246.276	4246.018
9.000	4210.000	4246.276	4246.018

The plot, as expected, is essentially identical to the best fit of the two-compartment model to the data. You can actually verify this by plotting **s1**, **s2** and **plasma**. If you look at the table of values as shown below:

you will see only minor differences due simply to the numerics of the software.

f. View the statistics. The **Statistics** window will appear as follows:

Parameter/Variable	Value	Std.Dev.	Coef. of Var.	95% Confider	ice Interval	
A1	30953.79919	3.05931e+003	9.88348e+000	24220.27976	37687.31862	F
12	13795.22249	3.29157e+003	2.38602e+001	6550.50902	21039.93597	Ē
1	0.61431	7.72115e-002	1.25688e+001	0.44437	0.78425	
2	0.13419	2.61249e-002	1.94685e+001	0.07669	0.19169	
(0,1)	0.29227	** Fixed **	** Fixed **	** Fixed **	** Fixed **	-
k(1,2)	0.28311	** Fixed **	** Fixed **	** Fixed **	** Fixed **	
c(2,1)	0.17452	** Fixed **	** Fixed **	** Fixed **	** Fixed **	[
C Correlation Matrix	C Covarian	ce Matrix 📀 🧿	Objective			
		Objective	Scaled Data V	ariance		
s2 : plasma	1.33674	5e+001	1.113536e-00	1		Ī
Total objective	1.33674	5e+001				
AIC	7.93599	6e+000				
BIC	8.05400	4e+000				Ē

If you scroll in the **Parameter/Variable** pane, you will see the estimates for AUC and AUMC are:

Parameter/Variable	Value	Std.Dev.	Coef. of Var.	95% Confidence Interval	
k(1,2)	0.28311	** Fixed **	** Fixed **	** Fixed ** ** Fixed **	
k(2,1)	0.17452	** Fixed **	** Fixed **	** Fixed ** ** Fixed **	
vol	2323.77661	** Fixed **	** Fixed **	** Fixed **	
		- Derived Vari	ables		
AUC	153191.18396	6.51383e+003	4.25209e+000	138854.30653 167528.06140	
MAUC	848122.02456	1.44358e+005	1.70209e+001	530391.33253 1165852.71658	

g. Close the **Statistics** window.

Quit the SAAM II Compartmental application. Do not save the changes to study_0.stu.



AUC calculations. Two points can be made in terms of the AUC calculations. If you worked through Part 2 as well as Part 3, how do the AUC calculations compare? The results are shown below:

	AUC Option		Sum of Ex	kponential
Parameter	Value	SD	Value	SD
AUC	153130	6478	153191	6514
MAUC	846550	143140	848122	144358

You can see there is very good correspondence, especially for AUC. For AUMC, in Part 2, you may want to extend the time of the experiment a little longer; the difference is in the third significant figure.

The other point deals in this tutorial with the fact that "plasma" cannot be associated with two samples. Suppose you wanted to Fit the compartmental model and the sum of exponentials simultaneously. You can do this if you create another table for the plasma data, but give it a different name. Below is an example:

d _{ij} Data	- study_0.dat		
DATA			
(FSD 0.1	1)		
t	plasma		
0.007	46780		
0.042	43522		
0.125	42535		
0.25	40125		
0.3/5	36221		
0.5	35562		
1.0	10572		
2.0	19575		
4.0	11278		
5.0	8081		
6.0	6999		
7.0	5653		
8.0	5139		
9.0	4210		
END			
DATA			
(FSD 0.1	1)		
t	plasma 1		
0.007	46780		
0.042	43522		
0.125	42535		
0.25	40125		
0.375	36221		
0.5	35562		
1.0	20194		
3.0	14403		
4.0	11278		
5.0	8081		
6.0	6999		
7.0	5653		
8.0	5139		
9.0	4210		
END		_	
Data Fo	Data Format is okay Edite		

Here the plasma data have been duplicated, but in the second table, they are named "plasma1". In this situation, you can leave **s1** associated with "plasma" and associate **s2** with "plasma1". Now there will be no error message. When you Fit the two models, there are in fact two models – the compartmental and sum of exponential -, to the data, you will get approximately the same final parameter values, but the statistics will be a little different. The reason is that SAAM II creates only one objective function. Here with two models, there should be two. This is why we separated the models as we did.

