

# Thermal Decomposition of Beta-Carotene in a Flow Reactor

This tutorial illustrates how to use the Uncertainty Quantification (UQ) functionality to answer questions regarding sensitivity and reliability of a flow reactor with thermal decomposition. The tutorial investigates which uncertainties in parameters dominate the survival of a nutrient ( $\beta$ -carotene) during a food-processing step. A fluid carrying the nutrient is injected into the reactor, and is subsequently heated by a downstream cylinder. Because  $\beta$ -carotene is heat sensitive it decomposes into fragments, reducing the nutrient level of the fluid. The model starts with a screening study, followed by a sensitivity analysis, then looks at error propagation, and finally performs a reliability analysis.

## Model Definition

The system models a plate reactor consisting of a channel in which carrot juice is being heated by a cylinder perpendicular to the juice's direction of flow. The model seeks to answer the question of how much of the nutrient  $\beta$ -carotene is lost due to its thermal decomposition. In this case the outflow rate of remaining  $\beta$ -carotene will be the *Quantity* of Interest (QoI).

The full 3D representation of the reactor geometry is shown in Figure 1.

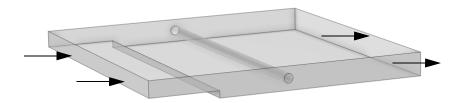


Figure 1: 3D geometry of a parallel plate reactor. The reacting fluid is heated as it passes the cylinder.

#### CHEMISTRY

A heat-sensitive chemical (β-carotene) undergoes thermal decomposition into fragments (byproduct) according to the following unimolecular reaction in water:

> kbetaCarotene ⇒ byprouduct

The reaction rate, r, is given by

$$r = kc_{\text{betaCarotene}}$$

where the rate coefficient k is temperature dependent according to the Arrhenius equation:

$$k = A \exp\left(-\frac{E}{R_g T}\right) \tag{1}$$

In Equation 1, A is the frequency factor  $(9.4 \times 10^{13} \text{ 1/s})$ , E the activation energy (110  $\times$  10  $^3$  J/mol),  $R_{\rm g}$  the gas constant (8.314 J/(mol·K)), and T the temperature (SI unit: K).

In addition, the decomposition reaction is endothermic, and the rate of energy expelled is given by

$$Q = -\text{rate} \cdot H$$

where H is the heat of reaction (8.4 kJ/mol).

The reaction kinetics are set up with the Chemistry interface. The conversion of the species β-carotene in the reactor is a function of the residence time; that is, it depends on the detailed fluid flow. Furthermore, the decomposition is influenced by the temperature distribution.

#### **PARAMETERIZATION**

To investigate the performance and reliability of the reactor, a number of parameters in the model setup need to be selected. The uncertainty quantification of the quantity of interest will be evaluated with respect to these inputs. The parameters can, for example, be physical properties of the fluid or the reactor material, such as the fluid viscosity or the heat conductivity of the reactor walls. It can also be parameters defining the reactor configuration, such as the position and radius of the cylinder or the height of the channel. Specific to the field of chemical engineering, parameters used to describe the reaction kinetics, such as reaction rate constants, activation energy, or the enthalpy of reaction, can also be used. Table 1 lists the parameters varied in this model, along with the default values and the respective statistical distributions describing their variation, in this example the distributions are arbitrary and simply used for demonstration purposes. Both the joint

effect and the individual sensitivity of the results on these uncertainties will be studied using the study steps provided by the Uncertainty Quantification Module.

TABLE I: INPUT PARAMETERS SUBJECT TO UNCERTAINTY QUANTIFICATION

Name	Description	Default value	Distribution
T_cyl	Temperature, heating cylinder	92[degC]	Uniform, ±5 K
Е	Activation energy	110[kJ/mol]	Normal, σ/E=0.3%
Α	Frequency factor	9.4e13[1/s]	Normal, σ/A=10%
Н	Enthalpy of reaction	8.4[kJ/mol]	Uniform, ±3%
D_BetaC_ref	Diffusion coefficient	2.07e-9[m^2/s]	Normal, σ/ D_BetaC_ref=20%
dDdT	Temperature derivative of diffusion coefficient	7.5e-11[m^2/s/K]	Uniform ±50%
xpos	Cylinder x-coordinate	6[cm]	Uniform, ±3 cm
ypos	Cylinder y-coordinate	5[mm]	Uniform, ±2.25 mm
RI	Cylinder radius	2[mm]	Uniform [I mm, 2.5 mm]

Here, the default values of the Arrhenius parameters are taken from Ref. 1.

#### THE UNCERTAINTY QUANTIFICATION STUDIES

The Uncertainty Quantification Module provides four different study types:

- Screening, MOAT
  - Identifies the most influential inputs, for each QoI
  - Is based on the Morris One-at-a-Time (MOAT) method
  - Outputs MOAT mean and MOAT standard deviation values for each input investigated
- Sensitivity Analysis
  - Computes the fraction of impact for the inputs, for each QoI
  - Outputs first-order and total Sobol indices
- Uncertainty Propagation
  - Computes the statistical variation of the QoI
  - Outputs a kernel density estimation (KDE) plot representing an estimate of the probability distribution of the QoI

- Reliability Analysis
  - Computes the probability for the fulfillment of a condition based on the QoI
  - For example, what is  $P(c < c_{min})$ , the probability that the outlet concentration is below some minimum value?

For more information, see the Uncertainty Quantification Module User's Guide.

#### SURROGATE MODELS

To get statistical data based on a physics model, a lot of simulation results are needed where input parameters are varied according to their probability distributions. For a 3D model, this might be computationally infeasible. To get around this problem, the Uncertainty Quantification Module trains a so-called surrogate model, which is used for sensitivity analysis, uncertainty propagation, and reliability analysis (but not for screening).

This process is typically adaptive and the surrogate model can approximate the original model to a user-defined degree of accuracy. The Uncertainty Quantification Module uses two different types of surrogate models:

- Sparse Polynomial Chaos Expansion (SPCE)
  - This surrogate model improves its accuracy by adaptively solving the full model and thereby adding new QoI data using sequential Latin hypercube sampling
- Gaussian Process (GP)
  - This surrogate model improves its accuracy, using information from the current Gaussian Process surrogate model, by adaptively solving the full model for new carefully selected sets of parameter values

## Results and Discussion

As a first step, a qualitative screening study is performed. Here the most influential parameters are highlighted, along with the information of what parameters have such a small effect that they can be excluded from further analysis. One of the main benefits of the screening study is that it is computationally cheap. The result of the screening study is a so-called MOAT plot shown in Figure 2. The three parameters with the least impact are found in the bottom left of the plot. These three parameters (H, D\_BetaC\_ref, and dDdT) were excluded from further analysis in order to reduce the overall computational cost.

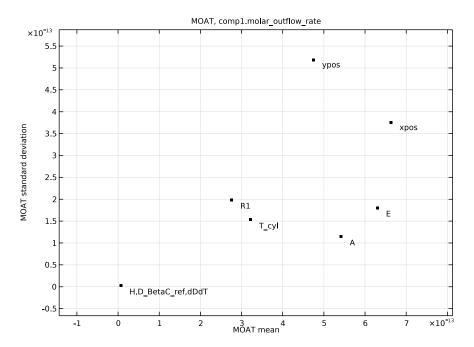


Figure 2: MOAT (Morris one at-a-time) plot from screening study, indicating general effect on x-axis and nonlinearity on y-axis.

After reducing the number of studied parameters, a more rigorous and quantitative Sensitivity Analysis is performed. In this step a surrogate model is trained automatically by COMSOL, and then used internally for a Monte Carlo simulation to perform a great number of evaluations. The output from this study step are the so-called Sobol sensitivity indices which are presented in a bar chart as seen in Figure 3.

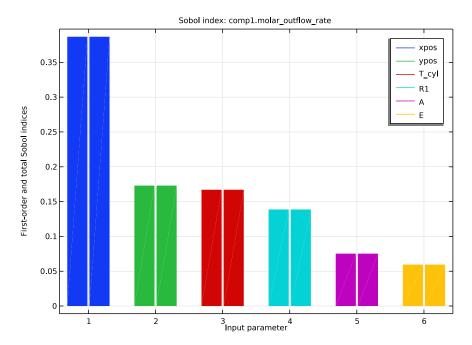


Figure 3: Bar chart of pairs of Sobol indices (first and total order) for each parameter from the sensitivity analysis.

The sum of all first-order Sobol indices (the left bar in each pair) is less than, or equal to, one; only for problems where there are no higher-order interactions between parameters would the value be one. The sum of all total-order Sobol indices (the right bar in each pair) is equal to, or greater than, one. And again, only for additive models would equality hold. The inclusion of higher-order effects in the total index makes differences within pairs useful as an indicator of what parameters interact with the QoI in a nonlinear manner.

The next step is to compute the probability density of the Quantity of Interest. This will provide information about the joint effect of the uncertainties in all varied parameters. Oftentimes, the probability density approximately takes the shape of the normal distribution (bell curve), but skewness and heavy tails are quite common, and it is therefore valuable to study a plot of the distribution. The probability density function is presented in Figure 4.

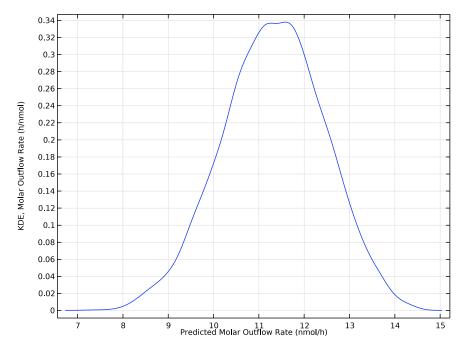


Figure 4: Probability density function, visualizing the distribution of the outflow rate of  $\beta$ -carotene from the reactor (the Quantity of Interest).

Finally a reliability analysis is performed, allowing us to quantify tail risks. In our case, we might want to know how big a risk there is of the  $\beta$ -carotene content dropping below a prescribed threshold. In this model, the risk of the outlet flux dropping below 33% of the inlet was found to be 2%. Note that even though we might be tempted to simply integrate the probability density function up to this threshold, we should refrain from doing so since the tails are not sufficiently sampled in the other study types. The reliability analysis intentionally refines the surrogate model by increasing the sampling in the regions of the multidimensional parameter window yielding these off chance conditions.

It is possible to keep intermediate solutions from the Uncertainty Quantification studies, and it can be quite instructive to compare a set of such results. An example of a plot with multiple such solutions is shown in Figure 5.

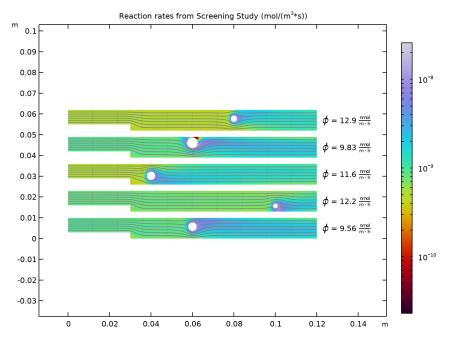


Figure 5: Rate of reaction from a subset of sampled parameters during the screening process.

## Reference

1. C. Dhuique-Mayer and others, "Thermal Degradation of Antioxidant Micronutrients in Citrus Juice: Kinetics and Newly Formed Compounds," J. Agric. Food Chem., vol. 55, pp. 4209-4216, 2007.

Application Library path: Chemical\_Reaction\_Engineering\_Module/ Reactors with Mass and Heat Transfer/thermal decomposition uq

# Modeling Instructions

From the File menu, choose New.

#### NEW

In the New window, click Model Wizard.

#### MODEL WIZARD

- I In the Model Wizard window, click **2** 2D.
- 2 In the Select Physics tree, select Chemical Species Transport>Reacting Flow>Laminar Flow, **Diluted Species.**
- 3 Click Add.
- 4 In the Added physics interfaces tree, select Transport of Diluted Species (tds).
- 5 Click + Add Concentration.
- 6 In the Concentrations (mol/m³) table, enter the following settings:

cBetaC cByprod

- 7 In the Select Physics tree, select Heat Transfer>Heat Transfer in Fluids (ht).
- 8 Click Add.
- 9 Click Study.
- 10 In the Select Study tree, select General Studies>Stationary.
- II Click Done.

#### **GLOBAL DEFINITIONS**

#### Parameters 1

- I In the Model Builder window, under Global Definitions click Parameters I.
- 2 In the Settings window for Parameters, locate the Parameters section.
- 3 Click Load from File.
- **4** Browse to the model's Application Libraries folder and double-click the file thermal decomposition ug parameters.txt.

#### **GEOMETRY I**

Rectangle I (rI)

- I In the Model Builder window, expand the Component I (compl)>Geometry I node.
- 2 Right-click Geometry I and choose Rectangle.
- 3 In the Settings window for Rectangle, locate the Size and Shape section.
- 4 In the Width text field, type W1.

5 In the **Height** text field, type H1.

## Rectangle 2 (r2)

- I In the Geometry toolbar, click Rectangle.
- 2 In the Settings window for Rectangle, locate the Size and Shape section.
- 3 In the Width text field, type W2.
- 4 In the Height text field, type H2.

## Circle I (c1)

- I In the Geometry toolbar, click Circle.
- 2 In the Settings window for Circle, locate the Size and Shape section.
- 3 In the Radius text field, type R1.
- 4 Locate the **Position** section. In the **x** text field, type xpos.
- 5 In the y text field, type ypos.

## Line Segment I (Is I)

- I In the Geometry toolbar, click More Primitives and choose Line Segment.
- 2 On the object c1, select Point 3 only.
- 3 In the Settings window for Line Segment, locate the Endpoint section.
- 4 From the Specify list, choose Coordinates.
- 5 In the x text field, type xpos+10\*R1.
- 6 In the y text field, type ypos.

#### Difference I (dif1)

- I In the Geometry toolbar, click Booleans and Partitions and choose Difference.
- 2 Select the object rl only.
- 3 In the Settings window for Difference, locate the Difference section.
- 4 Click to select the Activate Selection toggle button for Objects to subtract.
- 5 Select the objects cl and r2 only.

#### Mesh Control Edges I (mcel)

- I In the Geometry toolbar, click \times \text{Virtual Operations} and choose Mesh Control Edges.
- **2** On the object **fin**, select Boundary 6 only.
- 3 In the Geometry toolbar, click **Build All**.

#### DEFINITIONS

#### Inlet

- I In the Model Builder window, expand the Component I (compl)>Definitions node.
- 2 Right-click **Definitions** and choose **Selections>Explicit**.
- 3 In the Settings window for Explicit, locate the Input Entities section.
- 4 From the Geometric entity level list, choose Boundary.
- **5** Select Boundary 1 only.
- 6 In the Label text field, type Inlet.
- 7 Right-click Inlet and choose Duplicate.

#### Outlet

- I In the Model Builder window, under Component I (compl)>Definitions>Selections click Inlet I
- 2 In the Settings window for Explicit, type Outlet in the Label text field.
- **3** Select Boundary 6 only.
- 4 Right-click Outlet and choose Duplicate.

#### Heater

- I In the Model Builder window, under Component I (compl)>Definitions>Selections click Outlet I.
- 2 In the Settings window for Explicit, type Heater in the Label text field.
- **3** Select Boundaries 7–9 only.

#### MESH I

- I In the Model Builder window, under Component I (compl) right-click Mesh I and choose **Build All.**
- 2 Right-click Component I (compl)>Mesh I and choose Edit Physics-Induced Sequence.

#### Size 2

- I In the Model Builder window, right-click Mesh I and choose Size.
- 2 Drag and drop Size 2 below Size.
- 3 In the Settings window for Size, locate the Geometric Entity Selection section.
- 4 From the Geometric entity level list, choose Boundary.
- **5** Select Boundary 11 only.
- 6 Locate the Element Size section. From the Predefined list, choose Extra fine.

7 From the Calibrate for list, choose Fluid dynamics.

#### Boundary Layer Properties 1

- I In the Model Builder window, expand the Boundary Layers I node, then click Boundary Layer Properties I.
- 2 In the Settings window for Boundary Layer Properties, locate the Layers section.
- 3 In the Number of layers text field, type 3.
- 4 In the Thickness adjustment factor text field, type 4.
- 5 Right-click Boundary Layer Properties I and choose Duplicate.

## Boundary Layer Properties 2

- I In the Model Builder window, click Boundary Layer Properties 2.
- 2 In the Settings window for Boundary Layer Properties, locate the Boundary Selection section.
- 3 From the Selection list, choose Heater.
- 4 Locate the Layers section. In the Number of layers text field, type 5.
- 5 In the Thickness adjustment factor text field, type 2.

#### Size

- I In the Model Builder window, under Component I (compl)>Mesh I click Size.
- 2 In the Settings window for Size, click to expand the Element Size Parameters section.
- 3 In the Maximum element growth rate text field, type 1.1.
- 4 Click III Build All.

## **DEFINITIONS**

#### Variables 1

- I In the Model Builder window, under Component I (compl) right-click Definitions and choose Variables.
- 2 In the Settings window for Variables, locate the Variables section.
- **3** In the table, enter the following settings:

Name	Expression	Unit	Description
D_BetaC	D_BetaC_ref + dDdT*(T- 298.15[K])	m²/s	Diffusion coefficient

#### ADD MATERIAL

I In the Home toolbar, click **Add Material** to open the Add Material window.

- 2 Go to the Add Material window.
- 3 In the Search text field, type water.
- 4 Click Search.
- 5 In the tree, select Built-in>Water, liquid.
- 6 Click Add to Component in the window toolbar.
- 7 In the Home toolbar, click Radd Material to close the Add Material window.

## LAMINAR FLOW (SPF)

Inlet 1

- I In the Model Builder window, under Component I (compl) right-click Laminar Flow (spf) and choose Inlet.
- 2 Select Boundary 1 only.

Outlet I

- I In the Physics toolbar, click Boundaries and choose Outlet.
- 2 Select Boundary 6 only.

## HEAT TRANSFER IN FLUIDS (HT)

In the Model Builder window, under Component I (compl) click Heat Transfer in Fluids (ht).

Inflow I

- I In the Physics toolbar, click Boundaries and choose Inflow.
- 2 Select Boundary 1 only.

Outflow I

- I In the Physics toolbar, click Boundaries and choose Outflow.
- 2 Select Boundary 6 only.

Inflow I

- I In the Model Builder window, click Inflow I.
- 2 In the Settings window for Inflow, locate the Upstream Properties section.
- 3 In the  $T_{\rm ustr}$  text field, type T\_inlet.

Temperature I

- I In the Physics toolbar, click Boundaries and choose Temperature.
- **2** Select Boundaries 7–9 only.
- 3 In the Settings window for Temperature, locate the Temperature section.

**4** In the  $T_0$  text field, type T\_cyl.

#### ADD PHYSICS

- I In the Physics toolbar, click and Physics to open the Add Physics window.
- 2 Go to the Add Physics window.
- 3 In the tree, select Chemical Species Transport>Chemistry (chem).
- 4 Click Add to Component I in the window toolbar.
- 5 In the Physics toolbar, click and Physics to close the Add Physics window.

## CHEMISTRY (CHEM)

## Reaction I

- I Right-click Component I (compl)>Chemistry (chem) and choose Reaction.
- 2 In the Settings window for Reaction, locate the Reaction Formula section.
- 3 In the Formula text field, type betaCarotene=>byproduct.
- 4 Locate the Rate Constants section. Select the Use Arrhenius expressions check box.
- **5** In the  $A^f$  text field, type A.
- 6 In the  $E^{f}$  text field, type E.
- 7 Locate the Reaction Thermodynamic Properties section. From the Enthalpy of reaction list, choose User defined.
- **8** In the *H* text field, type H.

#### Species: betaCarotene

- I In the Model Builder window, click Species: betaCarotene.
- 2 In the Settings window for Species, locate the Chemical Formula section.
- 3 In the M text field, type Mn BetaC.

#### Species: byproduct

- I In the Model Builder window, click Species: byproduct.
- 2 In the Settings window for Species, locate the Chemical Formula section.
- 3 In the M text field, type Mn BetaC.
- 4 In the Model Builder window, click Chemistry (chem).
- 5 In the Settings window for Chemistry, locate the Species Matching section.
- 6 From the Species solved for list, choose Transport of Diluted Species.

7 Find the **Bulk species** subsection. In the table, enter the following settings:

Species	Туре	Molar concentration	Value (mol/m^3)
betaCarotene	Variable	cBetaC	Solved for
byproduct	Variable	cByprod	Solved for

## HEAT TRANSFER IN FLUIDS (HT)

In the Model Builder window, under Component I (compl) click Heat Transfer in Fluids (ht).

Heat Source I

- I In the Physics toolbar, click **Domains** and choose **Heat Source**.
- 2 Select Domain 1 only.
- 3 In the Settings window for Heat Source, locate the Material Type section.
- 4 From the Material type list, choose Nonsolid.
- 5 Locate the Heat Source section. From the  $Q_0$  list, choose Heat source of reactions (chem).

#### MULTIPHYSICS

Nonisothermal Flow I (nitf1)

In the Physics toolbar, click Multiphysics Couplings and choose Domain> Nonisothermal Flow.

#### TRANSPORT OF DILUTED SPECIES (TDS)

Transport Properties 1

- I In the Model Builder window, under Component I (compl)> Transport of Diluted Species (tds) click Transport Properties 1.
- 2 In the Settings window for Transport Properties, locate the Diffusion section.
- **3** In the  $D_{\mathrm{cBetaC}}$  text field, type D\_BetaC.

Inflow I

- I In the Physics toolbar, click Boundaries and choose Inflow.
- 2 Select Boundary 1 only.
- 3 In the Settings window for Inflow, locate the Concentration section.
- **4** In the  $c_{0,cBetaC}$  text field, type cBetaC\_inlet.

Outflow I

- I In the Physics toolbar, click Boundaries and choose Outflow.
- 2 Select Boundary 6 only.

#### Reactions 1

- I In the Physics toolbar, click **Domains** and choose **Reactions**.
- 2 Select Domain 1 only.
- 3 In the Settings window for Reactions, locate the Reaction Rates section.
- 4 From the  $R_{\rm cBetaC}$  list, choose Reaction rate for species betaCarotene (chem).
- 5 From the  $R_{
  m cBvprod}$  list, choose Reaction rate for species byproduct (chem).

## LAMINAR FLOW (SPF)

#### Inlet I

- I In the Model Builder window, under Component I (compl)>Laminar Flow (spf) click Inlet I.
- 2 In the Settings window for Inlet, locate the Boundary Condition section.
- 3 From the list, choose Fully developed flow.
- 4 Locate the Fully Developed Flow section. In the  $U_{
  m av}$  text field, type v\_inlet.

#### Outlet I

- I In the Model Builder window, click Outlet 1.
- 2 In the Settings window for Outlet, locate the Pressure Conditions section.
- 3 Select the Normal flow check box.

#### STUDY I

## Step 1: Stationary

- I In the Model Builder window, under Study I right-click Step I: Stationary and choose Duplicate.
- 2 In the Model Builder window, click Step 1: Stationary.
- 3 In the Settings window for Stationary, locate the Physics and Variables Selection section.
- **4** In the table, enter the following settings:

Physics interface	Solve for	Equation form
Laminar Flow (spf)		Automatic (Stationary)
Transport of Diluted Species (tds)		Automatic (Stationary)
Heat Transfer in Fluids (ht)		Automatic (Stationary)
Chemistry (chem)		Automatic (Stationary)

5 In the Home toolbar, click **Compute**.

#### DEFINITIONS

Integration over Exit

- I In the Definitions toolbar, click Nonlocal Couplings and choose Integration.
- 2 In the Settings window for Integration, locate the Source Selection section.
- 3 From the Geometric entity level list, choose Boundary.
- 4 From the Selection list, choose Outlet.
- 5 In the Label text field, type Integration over Exit.
- 6 In the Operator name text field, type intop exit.

Variables 1

- I In the Model Builder window, click Variables I.
- 2 In the Settings window for Variables, locate the Variables section.
- **3** In the table, enter the following settings:

Name	Expression	Unit	Description
molar_outflow_ rate	<pre>intop_exit(cBetaC *u)</pre>	mol/(m·s)	$\beta$ -carotene outflow

## STUDY I

In the Model Builder window, right-click Study I and choose Uncertainty Quantification> Add Uncertainty Quantification Study Using Study Reference.

## STUDY 2: UQ SCREENING

In the Settings window for Study, type Study 2: UQ Screening in the Label text field.

Uncertainty Quantification

- I In the Model Builder window, under Study 2: UQ Screening click **Uncertainty Quantification**.
- 2 In the Settings window for Uncertainty Quantification, locate the Quantities of Interest section.
- 3 Click + Add.
- 4 Click Replace Expression in the upper-right corner of the Quantities of Interest section. From the menu, choose Component I (compl)>Definitions>Variables> compl.molar\_outflow\_rate -  $\beta$ -carotene outflow - mol/(m·s).
- 5 Locate the Input Parameters section. Click + Add nine times.

- **6** In the table, click to select the cell at row number 1 and column number 3.
- 7 In the Lower bound text field, type T cyl-5[K].
- **8** In the **Upper bound** text field, type T\_cyl+5[K].
- **9** In the table, click to select the cell at row number 2 and column number 3.
- **10** From the **Distribution** list, choose **Normal**( $\mu$ , $\sigma$ ).
- II In the Mean text field, type E.
- 12 In the Standard deviation text field, type 0.003\*E.
- **13** In the table, click to select the cell at row number 3 and column number 3.
- 14 From the Distribution list, choose Normal( $\mu$ , $\sigma$ ).
- **I5** In the **Mean** text field, type A.
- 16 In the Standard deviation text field, type 0.1\*A.
- 17 In the table, click to select the cell at row number 4 and column number 3.
- **18** In the **Lower bound** text field, type 0.97\*H.
- 19 In the Upper bound text field, type 1.03\*H.
- **20** In the table, click to select the cell at row number 5 and column number 3.
- **21** From the **Distribution** list, choose **Normal**( $\mu$ , $\sigma$ ).
- 2 In the Mean text field, type D\_BetaC\_ref.
- 23 In the Standard deviation text field, type 0.2\*D BetaC ref.
- **24** In the table, click to select the cell at row number 6 and column number 3.
- **25** In the **Lower bound** text field, type 0.5\*dDdT.
- **26** In the **Upper bound** text field, type 1.5\*dDdT.
- **27** In the table, click to select the cell at row number 7 and column number 3.
- 28 In the Lower bound text field, type 4[cm].
- **29** In the **Upper bound** text field, type 10[cm].
- **30** In the table, click to select the cell at row number 8 and column number 3.
- 31 In the Lower bound text field, type 2.75[mm].
- 32 In the Upper bound text field, type 7.25[mm].
- **33** In the table, click to select the cell at row number 9 and column number 3.
- **34** In the **Lower bound** text field, type 1[mm].
- **35** In the **Upper bound** text field, type 2.5[mm].
- **36** Locate the **Output While Solving** section. Select the **Plot** check box.

- 37 From the Plot group list, choose Concentration, BetaC (tds).
- 38 Locate the Advanced Settings section. From the Keep model evaluations in memory list, choose All.
- **39** In the **Home** toolbar, click **Compute**.

This will produce Figure 2.

#### STUDY 2: UQ SCREENING

Uncertainty Quantification

- I In the Model Builder window, expand the Results>Uncertainty Quantification Graph> MOAT, compl.molar\_outflow\_rate node.
- 2 Right-click Study 2: UQ Screening>Uncertainty Quantification and choose Add New Uncertainty Quantification Study For>Sensitivity Analysis.

## STUDY 3: UQ SENSITIVITY ANALYSIS

- I In the Model Builder window, click Study 3.
- 2 In the Settings window for Study, type Study 3: UQ Sensitivity Analysis in the Label text field.
- I In the Model Builder window, under Study 3: UQ Sensitivity Analysis click **Uncertainty Quantification**.
- 2 In the Settings window for Uncertainty Quantification, locate the Input Parameters section.
- **3** In the table, click to select the cell at row number 4 and column number 3.
- 4 Click Delete.
- 5 Click Delete.
- 6 Click Delete.
- 7 In the table, click to select the cell at row number 6 and column number 3.
- **8** In the **Lower bound** text field, type 1 [mm].
- **9** In the **Upper bound** text field, type 2.5[mm].
- 10 Find the Input parameters sampling settings subsection. From the Number of input points type list, choose Manual.

Uncertainty Quantification 2

I In the **Home** toolbar, click **Compute**.

This will produce Figure 3.

#### STUDY 3: UQ SENSITIVITY ANALYSIS

Uncertainty Quantification

Right-click Uncertainty Quantification and choose

Add New Uncertainty Quantification Study For>Uncertainty Propagation.

### STUDY 4: UQ UNCERTAINTY PROPAGATION

- I In the Model Builder window, click Study 4.
- 2 In the Settings window for Study, type Study 4: UQ Uncertainty Propagation in the Label text field.

Uncertainty Quantification 3

In the **Home** toolbar, click **Compute**.

#### RESULTS

Line Graph 1

- I In the Model Builder window, expand the Results>Uncertainty Quantification Graph 2> Kernel Density Estimation, compl.molar\_outflow\_rate node, then click Line Graph 1.
- 2 In the Settings window for Line Graph, locate the y-Axis Data section.
- 3 In the Expression text field, type Kde/1e9/3600.
- 4 In the **Description** text field, type KDE, Molar Outflow Rate (h/nmol).
- 5 Locate the x-Axis Data section. In the Expression text field, type predicted\*1e9\*3600.
- 6 In the **Description** text field, type Predicted Molar Outflow Rate (nmol/h).

Kernel Density Estimation, compl.molar outflow rate

- I In the Model Builder window, click Kernel Density Estimation, compl.molar\_outflow\_rate.
- 2 In the Settings window for ID Plot Group, click to expand the Title section.
- **3** From the **Title type** list, choose **None**.
- This will produce Figure 4.

#### STUDY 4: UQ UNCERTAINTY PROPAGATION

Uncertainty Quantification

In the Model Builder window, under Study 4: UQ Uncertainty Propagation right-click Uncertainty Quantification and choose Add New Uncertainty Quantification Study For> Reliability Analysis.

#### STUDY 5: UQ RELIABILITY ANALYSIS

- I In the Model Builder window, under Study 5 click Uncertainty Quantification.
- 2 In the Settings window for Uncertainty Quantification, locate the Quantities of Interest section.
- **3** In the table, enter the following settings:

Expression	Description	Individual solution to use	True if	Threshold
comp1.molar _outflow_ra te		From "Solution to use"	Smaller than threshold	15e-13

- 4 In the Model Builder window, click Study 5.
- 5 In the Settings window for Study, type Study 5: UQ Reliability Analysis in the Label text field.

Uncertainty Quantification 4

In the **Home** toolbar, click **Compute**.

#### RESULTS

In the Model Builder window, expand the Results>Tables node.

#### Reaction Rate

- I In the Model Builder window, expand the Results>Tables>Reliability Analysis node.
- 2 Right-click Results and choose 2D Plot Group.
- 3 In the Settings window for 2D Plot Group, type Reaction Rate in the Label text field.
- 4 Locate the Data section. From the Dataset list, choose None.
- 5 Click to expand the Title section. From the Title type list, choose Manual.
- 6 In the Title text area, type Reaction rates from Screening Study (mol/(m<sup> 3</sup>\*s)).
- 7 Click to expand the Plot Array section. Select the Enable check box.
- 8 From the Array axis list, choose y.

## Surface I

- I Right-click Reaction Rate and choose Surface.
- 2 In the Settings window for Surface, locate the Data section.
- 3 From the Dataset list, choose Study 2: UQ Screening/Parametric Solutions I (sol4).

- 4 From the Parameter value (T\_cyl (K),E (J/mol),A (1/s),H (J/mol),...) list, choose I: T\_cyl=360.15 K, E=1.0898E5 J/mol, A=8.996E13 1/s, H=8652 J/mol, D\_BetaC\_ref=7.9064E-10 m^2/s, dDdT=3.75E-11 m^2/(s\*K), xpos=0.06 m, ypos=0.00575 m, R1=0.002 m.
- **5** Locate the **Expression** section. In the **Expression** text field, type -tds.R\_cBetaC.
- 6 Locate the Coloring and Style section. Click | Change Color Table.
- 7 In the Color Table dialog box, type prism in the text field.
- 8 Click **Filter**.
- 9 In the tree, select Rainbow>PrismDark.
- IO Click OK.
- II In the Settings window for Surface, locate the Coloring and Style section.
- 12 From the Color table transformation list, choose Reverse.
- 13 From the Scale list, choose Logarithmic.
- 14 Click to expand the Plot Array section. Select the Manual indexing check box.
- **I5** Right-click **Surface I** and choose **Duplicate**.

#### Surface 2

- I In the Model Builder window, click Surface 2.
- 2 In the Settings window for Surface, locate the Data section.
- 3 From the Parameter value (T\_cyl (K),E (J/mol),A (1/s),H (J/mol),...) list, choose 10: T\_cyl=366.82 K, E=1.1014E5 J/mol, A=1.2305E14 1/s, H=8316 J/mol, D\_BetaC\_ref=2.2479E-9 m^2/s, dDdT=8.75E-11 m^2/(s\*K), xpos=0.1 m, ypos=0.00275 m, R1=0.001 m.
- 4 Click to expand the Inherit Style section. From the Plot list, choose Surface 1.
- 5 Locate the Plot Array section. In the Index text field, type 1.
- 6 Right-click Surface 2 and choose Duplicate.

#### Surface 3

- I In the Model Builder window, click Surface 3.
- 2 In the Settings window for Surface, locate the Data section.
- 3 From the Parameter value (T\_cyl (K),E (J/mol),A (1/s),H (J/mol),...) list, choose 20: T\_cyl=360.15 K, E=1.1102E5 J/mol, A=9.804E13 1/s, H=8652 J/mol, D\_BetaC\_ref=7.9064E-10 m^2/s, dDdT=3.75E-11 m^2/(s\*K), xpos=0.04 m, ypos=0.00425 m, R1=0.002 m.
- 4 Locate the Plot Array section. In the Index text field, type 2.

**5** Right-click **Surface 3** and choose **Duplicate**.

#### Surface 4

- I In the Model Builder window, click Surface 4.
- 2 In the Settings window for Surface, locate the Data section.
- 3 From the Parameter value (T\_cyl (K),E (J/mol),A (1/s),H (J/mol),...) list, choose 30: T\_cyl=370.15 K, E=1.0898E5 J/mol, A=8.996E13 1/s, H=8484 J/mol, D\_BetaC\_ref=3.3494E-9 m^2/s, dDdT=6.25E-11 m^2/(s\*K), xpos=0.06 m, ypos=0.00725 m, RI=0.0025 m.
- 4 Locate the Plot Array section. In the Index text field, type 3.
- **5** Right-click **Surface 4** and choose **Duplicate**.

#### Surface 5

- I In the Model Builder window, click Surface 5.
- 2 In the Settings window for Surface, locate the Data section.
- 3 From the Parameter value (T\_cyl (K),E (J/mol),A (1/s),H (J/mol),...) list, choose 40: T\_cyl=363.48 K, E=1.0986E5 J/mol, A=6.4952E13 1/s, H=8148 J/mol, D\_BetaC\_ref=1.8921E-9 m^2/s, dDdT=1.125E-10 m^2/(s\*K), xpos=0.08 m, ypos=0.00575 m, RI=0.0015 m.
- 4 Locate the Plot Array section. In the Index text field, type 4.

#### Streamline 1

- I In the Model Builder window, right-click Reaction Rate and choose Streamline.
- 2 In the Settings window for Streamline, locate the Data section.
- 3 From the Dataset list, choose Study 2: UQ Screening/Parametric Solutions I (sol4).
- 4 From the Parameter value (T\_cyl (K),E (|/mol),A (1/s),H (|/mol),...) list, choose 1: T\_cyl=360.15 K, E=1.0898E5 J/mol, A=8.996E13 1/s, H=8652 J/mol, D\_BetaC\_ref=7.9064E-10 m^2/s, dDdT=3.75E-11 m^2/(s\*K), xpos=0.06 m, ypos=0.00575 m, RI=0.002 m.
- 5 Locate the Streamline Positioning section. From the Entry method list, choose Coordinates.
- **6** In the **x** text field, type 0 0 0 0 0.
- 7 In the y text field, type 0.004 0.005 0.006 0.007 0.008 0.009.
- 8 Locate the Coloring and Style section. Find the Point style subsection. From the Color list, choose **Custom**.

- 9 On Windows, click the colored bar underneath, or if you are running the crossplatform desktop — the **Color** button.
- 10 Click Define custom colors.
- II Set the RGB values to 128, 128, and 128, respectively.
- 12 Click Add to custom colors.
- 13 Click Show color palette only or OK on the cross-platform desktop.
- **14** From the **Type** list, choose **Arrow**.
- 15 Click to expand the Plot Array section. Select the Manual indexing check box.
- **16** Right-click **Streamline I** and choose **Duplicate**.

#### Streamline 2

- I In the Model Builder window, click Streamline 2.
- 2 In the Settings window for Streamline, locate the Data section.
- 3 From the Parameter value (T\_cyl (K),E (J/mol),A (1/s),H (J/mol),...) list, choose 10: T\_cyl=366.82 K, E=1.1014E5 J/mol, A=1.2305E14 1/s, H=8316 J/mol, D\_BetaC\_ref=2.2479E-9 m^2/s, dDdT=8.75E-11 m^2/(s\*K), xpos=0.1 m, ypos=0.00275 m, RI = 0.001 m.
- 4 Locate the Plot Array section. In the Index text field, type 1.
- 5 Right-click Streamline 2 and choose Duplicate.

#### Streamline 3

- I In the Model Builder window, click Streamline 3.
- 2 In the Settings window for Streamline, locate the Data section.
- 3 From the Parameter value (T\_cyl (K),E (J/mol),A (1/s),H (J/mol),...) list, choose 20: T\_cyl=360.15 K, E=1.1102E5 J/mol, A=9.804E13 1/s, H=8652 J/mol, D\_BetaC\_ref=7.9064E-10 m^2/s, dDdT=3.75E-11 m^2/(s\*K), xpos=0.04 m, ypos=0.00425 m, RI=0.002 m.
- 4 Locate the Plot Array section. In the Index text field, type 2.
- **5** Right-click **Streamline 3** and choose **Duplicate**.

#### Streamline 4

- I In the Model Builder window, click Streamline 4.
- 2 In the Settings window for Streamline, locate the Data section.
- 3 From the Parameter value (T\_cyl (K),E (|/mol),A (1/s),H (|/mol),...) list, choose 30: T\_cyl=370.15 K, E=1.0898E5 J/mol, A=8.996E13 1/s, H=8484 J/mol,

- D\_BetaC\_ref=3.3494E-9 m^2/s, dDdT=6.25E-11 m^2/(s\*K), xpos=0.06 m, ypos=0.00725 m, RI=0.0025 m.
- 4 Locate the Plot Array section. In the Index text field, type 3.
- **5** Right-click **Streamline 4** and choose **Duplicate**.

#### Streamline 5

- I In the Model Builder window, click Streamline 5.
- 2 In the Settings window for Streamline, locate the Data section.
- 3 From the Parameter value (T\_cyl (K),E (J/mol),A (1/s),H (J/mol),...) list, choose 40: T\_cyl=363.48 K, E=1.0986E5 J/mol, A=6.4952E13 1/s, H=8148 J/mol, D\_BetaC\_ref=1.8921E-9 m^2/s, dDdT=1.125E-10 m^2/(s\*K), xpos=0.08 m, ypos=0.00575 m, RI=0.0015 m.
- 4 Locate the Plot Array section. In the Index text field, type 4.

#### Annotation I

- I In the Model Builder window, right-click Reaction Rate and choose Annotation.
- 2 In the Settings window for Annotation, locate the Data section.
- 3 From the Dataset list, choose Study 2: UQ Screening/Parametric Solutions I (sol4).
- 4 From the Parameter value (T\_cyl (K),E (|/mol),A (1/s),H (|/mol),...) list, choose 1: T\_cyl=360.15 K, E=1.0898E5 J/mol, A=8.996E13 1/s, H=8652 J/mol, D\_BetaC\_ref=7.9064E-10 m^2/s, dDdT=3.75E-11 m^2/(s\*K), xpos=0.06 m, ypos=0.00575 m, RI=0.002 m.
- **5** Locate the **Annotation** section. In the **Text** text field, type \$\phi\$ = eval(comp1.molar outflow rate, nmol/m/h) \$\mathrm{\frac{nmol}{m\cdot} h}}\$.
- 6 Select the LaTeX markup check box.
- 7 Locate the **Position** section. In the x text field, type W1.
- 8 In the y text field, type H1/2.
- **9** Locate the Coloring and Style section. Clear the Show point check box.
- 10 From the Anchor point list, choose Middle left.
- II Click to expand the Advanced section. In the Expression precision text field, type 3.
- 12 Click to expand the Plot Array section. Select the Manual indexing check box.
- 13 Right-click Annotation I and choose Duplicate.

#### Annotation 2

I In the Model Builder window, click Annotation 2.

- 2 In the Settings window for Annotation, locate the Data section.
- 3 From the Parameter value (T\_cyl (K),E (|/mol),A (1/s),H (|/mol),...) list, choose 10: T cyl=366.82 K, E=1.1014E5 |/mol, A=1.2305E14 1/s, H=8316 |/mol, D\_BetaC\_ref=2.2479E-9 m^2/s, dDdT=8.75E-11 m^2/(s\*K), xpos=0.1 m, ypos=0.00275 m, RI=0.001 m.
- 4 Locate the Plot Array section. In the Index text field, type 1.
- 5 Right-click Annotation 2 and choose Duplicate.

#### Annotation 3

- I In the Model Builder window, click Annotation 3.
- 2 In the Settings window for Annotation, locate the Data section.
- 3 From the Parameter value (T cyl (K), E (I/mol), A (I/s), H (I/mol),...) list, choose 20: T\_cyl=360.15 K, E=1.1102E5 J/mol, A=9.804E13 1/s, H=8652 J/mol, D\_BetaC\_ref=7.9064E-10 m^2/s, dDdT=3.75E-11 m^2/(s\*K), xpos=0.04 m, ypos=0.00425 m, RI=0.002 m.
- 4 Locate the Plot Array section. In the Index text field, type 2.
- 5 Right-click Annotation 3 and choose Duplicate.

#### Annotation 4

- I In the Model Builder window, click Annotation 4.
- 2 In the Settings window for Annotation, locate the Data section.
- 3 From the Parameter value (T\_cyl (K),E (|/mol),A (1/s),H (|/mol),...) list, choose 30: T\_cyl=370.15 K, E=1.0898E5 |/mol, A=8.996E13 1/s, H=8484 |/mol, D BetaC ref=3.3494E-9 m^2/s, dDdT=6.25E-11 m^2/(s\*K), xpos=0.06 m, ypos=0.00725 m, RI=0.0025 m.
- 4 Locate the Plot Array section. In the Index text field, type 3.
- 5 Right-click Annotation 4 and choose Duplicate.

#### Annotation 5

- I In the Model Builder window, click Annotation 5.
- 2 In the Settings window for Annotation, locate the Data section.
- 3 From the Parameter value (T\_cyl (K),E (|/mol),A (1/s),H (|/mol),...) list, choose 40: T\_cyl=363.48 K, E=1.0986E5 J/mol, A=6.4952E13 1/s, H=8148 J/mol, D\_BetaC\_ref=1.8921E-9 m^2/s, dDdT=1.125E-10 m^2/(s\*K), xpos=0.08 m, ypos=0.00575 m, RI=0.0015 m.
- 4 Locate the Plot Array section. In the Index text field, type 4.

Reaction Rate This is Figure 5.