

# Fermentation in Beer Brewing

# Introduction

An important step in brewing beer is the fermentation process. Here, alcohol is formed together with various flavor substances from sugars in the presence of yeast. The initial sugar content, temperature, and yeast type dictate how the fermentation proceeds.

This model consists of two parts. In the first part the fermentation process is efficiently modeled using the Reaction Engineering interface, assuming that the reaction rate is neither mass- nor heat-transfer limited, that is, the system is perfectly mixed.

In the second part of this model, the perfectly mixed model in part 1 is expanded into a sphero-conical tank model accounting for mass transfer, heat transfer and natural convection. In both model setups, it is possible to evaluate several parameters affecting the final alcohol content and taste of the beer. The perfectly mixed example reproduces results in Ref. 1 and Ref. 2.

# Model Definition

When brewing beer, the fermentation step is subsequent of malting and mashing, and involves the conversion of sugars to alcohol. The previous steps cover wetting and drying of barley grains to form malt, followed by boiling and mixing the malt, to create a sugary liquid called wort. The fermentation starts as soon as the wort has been cooled down (<  $20^{\circ}$ C) and yeast has been added to it.

The fermentation usually takes place in a closed tank under anaerobic conditions. The time frame for the fermentation is weeks, but may vary considerably depending on the yeast type and fermentation temperature. The sugar content is mainly made up of three types of sugars: maltose, glucose, and maltotriose. Of these, the maltose content is predominant. Selecting the yeast type is sometimes a bit tricky, but most important is that it should be able to catalyze the fermentation reactions at the chosen process temperature. The type studied in this example thrives at temperatures near 12°C, which is ideal for brewing lager.

# REACTION KINETICS

The irreversible reactions taking place during the fermentation process can be written in the following simplified form:

$$G \xrightarrow{x} E + CO_2 + \text{flavors}$$
 (1)

$$M \xrightarrow{x} E + CO_2 + \text{flavors}$$
 (2)

$$N \xrightarrow{x} E + CO_2 + \text{flavors}$$
 (3)

where G, M, and N denote glucose, maltose, and maltotriose, respectively. Furthermore, E stands for alcohol and  $CO_2$  for the carbon dioxide dissolved in the wort. The X notation shows the presence of yeast. Aside from carbon dioxide and ethanol, different flavoring components are formed. This tutorial accounts for two types of flavors: Ethyl acetate (EtAc) and acetaldehyde (AcA). The former, an ester, gives a desirable taste, the latter, an aldehyde, gives a bad tasting beer.

The reaction kinetics are as follows (for reactions 1, 2, and 3). Note that as a consequence of the simplified reaction description, yield coefficients, Y, are used to compute product concentrations:

$$r_i = k_i c_x$$
, i=1,2,3

The fermentation mechanisms depend on the yeast concentration and the reaction rate constant,  $k_i$  (SI unit: s<sup>-1</sup>), can be described using Michaelis–Menten kinetics:

$$k_1 = \frac{k_G c_G}{K_G + c_G}$$

The last two reactions are also inhibited by high sugar concentrations:

$$\begin{aligned} k_2 &= \frac{k_M c_M}{K_M + c_M} \cdot \frac{K_G}{K_G + c_G} \\ k_3 &= \frac{k_N c_N}{K_N + c_N} \cdot \frac{K_G}{K_G + c_G} \cdot \frac{K_M}{K_M + c_M} \end{aligned}$$

 $k_G$ ,  $k_M$ , and  $k_N$  are the maximum velocities (SI unit: s<sup>-1</sup>), K the Michaelis–Menten constant, and K an inhibition constant for the fermentation reaction. These three properties are temperature dependent as defined by the Arrhenius equation:

$$k_{j} = A_{j}e^{\frac{-E}{RT}} , j=G,M,N$$

$$K_{j} = A_{Hj}e^{\frac{-E_{Hj}}{RT}}$$

$$K'_{j} = A'_{Hj}e^{\frac{-E'_{H'j}}{RT}}$$

Here, A is the frequency factor and E is the activation energy.

The yeast concentration is modeled as a free species, with the following reaction rate:

$$R_x = k_x c_x$$

where  $k_x$  is the reaction rate constant that depends on the reaction constant of the three governing reactions and the fact that a high yeast concentration inhibits its production:

$$k_{x} = (Y_{X1}k_{1} + Y_{X2}k_{2} + Y_{X3}k_{3}) \cdot \frac{K_{X}}{K_{X} + (c_{x} - c_{x0})^{2}}$$

where  $K_X$  is the yeast growth inhibition constant and  $c_{x0}$  the initial yeast concentration in the tank.

The alcohol production needs to be corrected with yield coefficients as well, giving the following total reaction rate:

$$R_E = (Y_{E1}k_1 + Y_{E2}k_2 + Y_{E3}k_3)c_x$$

In similar manner, the production of the ethyl acetate flavor compound can be written as:

$$R_{EtAc} = Y_{EtAc}(k_1 + k_2 + k_3)c_x$$

The actaldehyde flavor, on the other hand, also decomposes, as given by:

$$R_{AcA} = Y_{AcA}(k_1 + k_2 + k_3)c_x - k_{AcA}c_{AcA}c_x$$

where  $k_{AcA}$  is the rate constant for the decomposition of acetaldehyde and is defined with the Arrhenius equation.

Both the gaseous and dissolved carbon dioxide are computed in the example. The reaction rate of the gaseous species is described by:

$$R_{CO_2(\mathsf{g})} = (Y_{C1}k_1 + Y_{C2}k_2 + Y_{C3}k_3)c_x - K_{GL}(c_{co_2(\mathsf{sat})} - c_{co_2(\mathsf{l})})$$

where  $K_{GL}$  is the gas to liquid mass transfer coefficient of carbon dioxide and  $c_{CO2(sat)}$  the maximum solubility concentration of carbon dioxide in water.

For the dissolved species, the reaction rate becomes:

$$R_{CO_2(l)} = -K_{GL}(c_{co_2(sat)} - c_{co_2(l)})$$

#### 4 | FERMENTATION IN BEER BREWING

The reaction data required to simulated the fermentation reactions are tabulated in Table 1.

Parameters	Value	Parameters	Value
E <sub>G</sub>	9.46·10 <sup>4</sup> J/mol	A' <sub>HG</sub>	1.36·10 <sup>10</sup> mol/m <sup>3</sup>
$E_{\mathrm{M}}$	4.73·10 <sup>4</sup> J/mol	$A'_{\rm HM}$	1.42·10 <sup>24</sup> mol/m <sup>3</sup>
$E_{\rm N}$	3.00·10 <sup>4</sup> J/mol	$A_{ m AcA}$	9.13 m <sup>3</sup> /(s·mol)
$E_{ m HG}$	-2.87·10 <sup>5</sup> J/mol	$Y_{\rm X1}$	0.134
$E_{\rm HM}$	-6.03·10 <sup>4</sup> J/mol	$Y_{\rm X2}$	0.268
$E_{\rm HN}$	-8.33·10 <sup>4</sup> J/mol	$Y_{\rm X3}$	0.402
$E'_{\rm HG}$	4.27·10 <sup>4</sup> J/mol	$Y_{\rm E1}$	1.92
$E'_{\rm HM}$	1.10 <sup>.</sup> 10 <sup>5</sup> J/mol	$Y_{\rm E2}$	3.84
E <sub>AcA</sub>	4.64·10 <sup>4</sup> J/mol	$Y_{\rm E3}$	5.76
$A_{ m G}$	9.51·10 <sup>11</sup> 1/s	$Y_{\rm EtAc}$	9.92·10 <sup>-4</sup>
$A_{\mathrm{M}}$	3.68·10 <sup>3</sup> 1/s	Y <sub>AcA</sub>	1.00·10 <sup>-2</sup>
$A_{ m N}$	1.10 <sup>.</sup> 10 <sup>1</sup> 1/s	K <sub>X</sub>	3.65·10 <sup>5</sup> mol <sup>2</sup> /m <sup>6</sup>
$A_{ m HG}$	2.09·10 <sup>-53</sup> mol/m <sup>3</sup>	$K_{ m GL}$	1.94·10 <sup>-5</sup> 1/s
$A_{ m HM}$	3.40·10 <sup>-9</sup> mol/m <sup>3</sup>	$c_{\rm CO2(sat)}$	3.90·10 <sup>2</sup> mol/m <sup>3</sup>
$A_{ m HN}$	2.34·10 <sup>-12</sup> mol/m <sup>3</sup>		

TABLE I: REACTION PARAMETERS.

The perfectly mixed model is solved with the Reaction Engineering interface using the Batch, constant volume, reactor type at non-isothermal conditions.

For the three reactions, reactions heats are available:  $\Delta H_1 = -91.2 \text{ kJ/mol}, \Delta H_2 = -226.3 \text{ kJ/mol}, and \Delta H_2 = -361.3 \text{ kJ/mol}, that are entered into the energy balance settings in the interface. The wort mixture is assumed to have similar thermal properties as water, that is, water is included as solvent. A cooling medium, with a temperature, <math>\Delta T_C$ , lower than the intial tank temperature, cools the fermentation process with the rate,  $q_v$  (SI unit: W/ (m<sup>3</sup>·K)):

$$Q_{\text{ext}} = -q_v(T - (T_0 - \Delta T_C))$$

where  $Q_{\text{ext}}$  is the total heat removed from the reactor (SI unit: W).

Beer fermentation usually takes place in sphero-conical tanks (Figure 1). Such a design is suitable for easy separation of yeast from the liquid either at the top or the bottom, and it enables better temperature control. Modern beer brewing equipment often has a built-in cooling jacket, but sometimes the tank is just situated in a cooled environment.

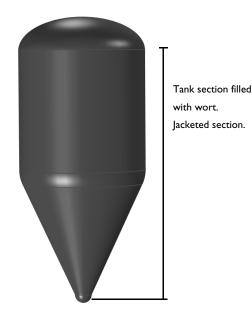


Figure 1: Sphero-conical tank for fermentation.

The space-dependent model is created using the Generate Space-Dependent Model feature. The kinetics are thus the same as in the perfectly mixed model. Due to rotational symmetry, you can model the whole system in 2D, using an axisymmetric geometry. The mass transport is modeled using the Transport of Diluted Species interface, the fluid motion with the Laminar Flow interface, and the heat transport using the Heat Transfer in Fluids interface. The wort takes up the part of the reactor as indicated in Figure 1.

The cooling is incorporated as a convective heat flux boundary condition driven by the temperature difference between the tank and the cooling media, as described by:

$$q = h(T_{ext} - T)$$

where h is an automatically defined heat transfer coefficient for external natural convection.

The only source of mixing is by natural convection, which is achieved by the coupling all three interfaces. As an assumption, the only property affecting the density of the mixture is the temperature. Also, the Boussinesq approximation is used, meaning that in the single-phase flow interface, the density is only varied in the volume force term.

The results from the perfectly mixed model are shown in Figure 2. The temperature in the cooling media and the initial tank temperature are both set to 12°C. The cooling rate is  $8 \text{ W/(m^3 \cdot K)}$ .

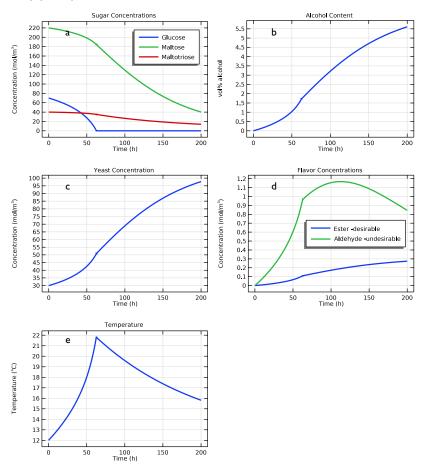


Figure 2: Plots displaying the results of the perfectly mixed model.

At these conditions, all three sugars decrease with time and the alcohol content reaches more than 5 vol%. Unfortunately, this beer will contain a considerable amount of aldehydes and probably taste bad. After reaching a maximum, the aldehyde concentration decreases and it is therefore important to continue the fermentation process long enough to allow the concentration to decrease to acceptable levels. A higher initial yeast concentration is one approach to decrease the aldehyde content more quickly. The temperature increase observed initially coincides with the quick consumption of glucose. After 60 h, all glucose has been consumed.

The results from the space-dependent model show a very even distribution in concentrations within the tank. In Figure 3, this is illustrated with the concentration of the undesired aldehyde after 3 hours. The same even concentration is seen throughout the whole simulation (24 h)

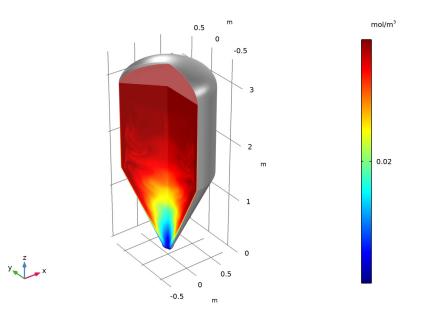


Figure 3: Concentration  $(mol/m^3)$  of acetaldebyde after 3 hours.

The well mixed behavior is also seen in Figure 4, showing the velocity magnitude, the temperature difference  $T - T_0$ , as well as the total energy flux (indicated by arrows), all at 3 h of fermenting. The velocity magnitude illustrates how natural convection creates a downward flow along the cold tank surface. In the temperature plot, the temperature difference is clearly seen along the fluid interface.

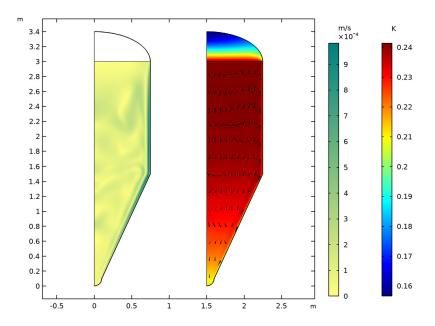
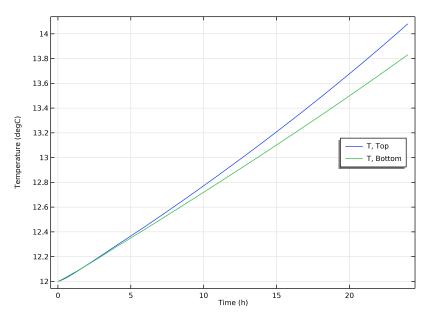


Figure 4: Velocity magnitude, temperature difference  $T - T_0$ , as well as the total energy flux (indicated by arrows). 3 h of fermentation.

Figure 5 shows the temperature in the top and at the bottom of the tank. During the simulated time frame the temperature is only increased a few degrees from the starting condition. Comparing the temperature reached at 24 h in the 0D model (Figure 2e) with



that in the space-dependent model (Figure 5), we see that the values are practically the same.

Figure 5: Temperature of the wort as a function of fermentation time. Temperatures at the top and in the bottom of the tank.

# References

1. D.A. Gee and W.F. Ramirez, "A Flavour Model for Beer Fermentation," *J. Inst. Brew.*, vol. 100, pp. 321–329, 1994.

2. W.F. Ramirez and J. Maciejowski, "Optimal Beer Fermentation," J. Inst. Brew., vol. 113, no. 3, pp. 325–333, 2007.

**Application Library path:** Chemical\_Reaction\_Engineering\_Module/ Reactors\_with\_Mass\_and\_Heat\_Transfer/beer\_fermentation

# Modeling Instructions

Setting up 0D (perfectly mixed) model using the Reaction Engineering interface.

From the File menu, choose New.

# NEW

In the New window, click 🙆 Model Wizard.

# MODEL WIZARD

- I In the Model Wizard window, click 0D.
- 2 In the Select Physics tree, select Chemical Species Transport>Reaction Engineering (re).
- 3 Click Add.
- 4 Click 🔿 Study.
- 5 In the Select Study tree, select General Studies>Time Dependent.
- 6 Click **M** Done.

# GLOBAL DEFINITIONS

Load model parameters and variables from text files.

# 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 **b** Load from File.
- **4** Browse to the model's Application Libraries folder and double-click the file beer\_fermentation\_parameters.txt.

# DEFINITIONS

Variables I

- 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 Click 📂 Load from File.
- **4** Browse to the model's Application Libraries folder and double-click the file beer\_fermentation\_variables1.txt.

# REACTION ENGINEERING (RE)

Use the **Batch**, **constant volume**, reactor type (the default) and model nonisothermal conditions by including the **Energy Balance**.

I In the Model Builder window, under Component I (compl) click Reaction Engineering (re).

- 2 In the Settings window for Reaction Engineering, locate the Energy Balance section.
- **3** From the list, choose **Include**.
- 4 In the  $Q_{\text{ext}}$  text field, type -qv\*(re.T-(T0-(dTc-273.15[K])))\*re.Vr.
- **5** Locate the **Mixture Properties** section. From the **Phase** list, choose **Liquid**.

Continue by entering free species, reactions, and a solvent.

#### Species 1

- I In the Reaction Engineering toolbar, click 🦂 Species.
- 2 In the Settings window for Species, locate the Species Name section.
- **3** In the text field, type X.

Reaction I

- I In the **Reaction Engineering** toolbar, click 👗 **Reaction**.
- 2 In the Settings window for Reaction, locate the Reaction Formula section.
- 3 In the Formula text field, type G=>E+C02+EtAc+AcA.
- 4 Click Apply.
- 5 Locate the Reaction Rate section. From the list, choose User defined.
- **6** In the  $r_i$  text field, type re.kf\_1\*re.c\_X.
- 7 Locate the Rate Constants section. In the k<sup>f</sup> text field, type kfG\_max\*re.c\_G/(KG1+ re.c\_G).
- 8 Locate the Reaction Thermodynamic Properties section. From the Heat source of reaction list, choose User defined.
- **9** In the Q text field, type -re.r\_1\*HG.

Reaction 2

- I In the Reaction Engineering toolbar, click 👗 Reaction.
- 2 In the Settings window for Reaction, locate the Reaction Formula section.
- **3** In the **Formula** text field, type M=>E+C02+EtAC+AcA.
- 4 Click Apply.
- 5 Locate the Reaction Rate section. From the list, choose User defined.
- 6 In the  $r_j$  text field, type re.kf\_2\*re.c\_X.
- 7 Locate the Rate Constants section. In the k<sup>f</sup> text field, type kfM\_max\*re.c\_M/(KM1+ re.c\_M)\*KG2/(KG2+re.c\_G).
- 8 Locate the Reaction Thermodynamic Properties section. From the Heat source of reaction list, choose User defined.

**9** In the Q text field, type -re.r\_1\*HM.

# Reaction 3

- I In the **Reaction Engineering** toolbar, click <u>A</u> **Reaction**.
- 2 In the Settings window for Reaction, locate the Reaction Formula section.
- 3 In the Formula text field, type N=>E+C02+EtAc+AcA.
- 4 Click Apply.
- 5 Locate the Reaction Rate section. From the list, choose User defined.
- 6 In the r<sub>i</sub> text field, type re.kf\_3\*re.c\_X.
- 7 Locate the Rate Constants section. In the k<sup>f</sup> text field, type kfN\_max\*re.c\_N/(KN1+ re.c\_N)\*KG2/(KG2+re.c\_G)\*KM2/(KM2+re.c\_M).
- 8 Locate the Reaction Thermodynamic Properties section. From the Heat source of reaction list, choose User defined.
- **9** In the Q text field, type -re.r\_1\*HN.

# Species 1

- I In the Reaction Engineering toolbar, click 📩 Species.
- 2 In the Settings window for Species, locate the Species Name section.
- **3** In the text field, type H20.
- 4 Locate the Species Type section. From the list, choose Solvent.
- 5 Click to expand the Species Thermodynamic Expressions section. From the list, choose User defined.
- **6** In the  $C_n$  text field, type CpH20.
- 7 In the Reaction Engineering toolbar, click 🦂 Species.

#### Species 1

- I In the Settings window for Species, locate the Species Name section.
- **2** In the text field, type CO2(g).

Most reaction products do not fully follow the reaction stoichiometry, therefore enter, where necessary, user defined reaction rates in their respective species nodes.

#### Species: X

- I In the Model Builder window, click Species: X.
- 2 In the Settings window for Species, click to expand the Reaction Rate section.
- 3 From the list, choose User defined.

4 In the R<sub>i</sub> text field, type (YXG\*re.kf\_1+YXM\*re.kf\_2+YXN\*re.kf\_3)\*re.c\_X\*KX/ (KX+(re.c\_X-cOX)^2).

#### Species: E

- I In the Model Builder window, click Species: E.
- 2 In the Settings window for Species, locate the Reaction Rate section.
- 3 From the list, choose User defined.
- **4** In the  $R_i$  text field, type (YEG\*re.kf\_1+YEM\*re.kf\_2+YEN\*re.kf\_3)\*re.c\_X.

#### Species: CO2

- I In the Model Builder window, click Species: CO2.
- 2 In the Settings window for Species, locate the Reaction Rate section.
- **3** From the list, choose **User defined**.
- 4 In the  $R_i$  text field, type max(hCO2\*(Csat\_CO2-re.c\_CO2),0).

# Species: CO2(g)

- I In the Model Builder window, click Species: CO2(g).
- 2 In the Settings window for Species, locate the Reaction Rate section.
- **3** From the list, choose **User defined**.
- 4 In the R<sub>i</sub> text field, type max(((YXG\*re.kf\_1+YXM\*re.kf\_2+YXN\*re.kf\_3)\* re.c\_X-max(hCO2\*(Csat\_CO2-re.c\_CO2),0)),0).

#### Species: EtAc

- I In the Model Builder window, click Species: EtAc.
- 2 In the Settings window for Species, locate the Reaction Rate section.
- 3 From the list, choose User defined.
- **4** In the  $R_i$  text field, type YEtAc\*(re.kf\_1+re.kf\_2+re.kf\_3)\*re.c\_X.

#### Species: AcA

- I In the Model Builder window, click Species: AcA.
- 2 In the Settings window for Species, locate the Reaction Rate section.
- 3 From the list, choose User defined.
- 4 In the R<sub>i</sub> text field, type YACA\* (re.kf\_1+re.kf\_2+re.kf\_3)\*re.c\_X-kACA\* re.c\_ACA\*re.c\_X.

Enter initial values.

Initial Values 1

- I In the Model Builder window, click Initial Values I.
- 2 In the Settings window for Initial Values, locate the General Parameters section.
- **3** In the  $T_0$  text field, type T0.
- **4** Locate the **Volumetric Species Initial Values** section. In the table, enter the following settings:

Species	Concentration (mol/m^3)	
E	cOE	
G	cOG	
H2O	rhoH20/MH20	
Μ	cOM	
N	CON	
х	cOX	

Solve the model for 200 h.

# STUDY I

Step 1: Time Dependent

- I In the Model Builder window, under Study I click Step I: Time Dependent.
- 2 In the Settings window for Time Dependent, locate the Study Settings section.
- 3 From the Time unit list, choose h.
- 4 In the **Output times** text field, type range(0,1,200).
- 5 From the Tolerance list, choose User controlled.
- 6 In the Relative tolerance text field, type 1e-6.

#### Solution 1 (soll)

- I In the Study toolbar, click **The Show Default Solver**.
- 2 In the Model Builder window, expand the Solution I (soll) node, then click Time-Dependent Solver I.
- **3** In the **Settings** window for **Time-Dependent Solver**, click to expand the **Absolute Tolerance** section.
- 4 From the Tolerance method list, choose Manual.
- 5 In the Absolute tolerance text field, type 1.0E-7.
- 6 Click **=** Compute.

# RESULTS

Sugars First, create 2a in Figure 2.

I In the Settings window for ID Plot Group, type Sugars in the Label text field.

# Global I

- I In the Model Builder window, expand the Sugars node, then click Global I.
- In the Settings window for Global, click Replace Expression in the upper-right corner of the y-Axis Data section. From the menu, choose Component I (compl)>
   Reaction Engineering>re.c\_G Concentration mol/m<sup>3</sup>.
- 3 Click Add Expression in the upper-right corner of the y-Axis Data section. From the menu, choose Component I (compl)>Reaction Engineering>re.c\_M Concentration mol/m<sup>3</sup>.
- 4 Click Add Expression in the upper-right corner of the y-Axis Data section. From the menu, choose Component I (compl)>Reaction Engineering>re.c\_N Concentration mol/m<sup>3</sup>.
- 5 Click to expand the Title section. From the Title type list, choose Manual.
- 6 In the Title text area, type Sugar Concentrations.
- 7 Click to expand the Coloring and Style section. In the Width text field, type 2.
- 8 Click to expand the Legends section. From the Legends list, choose Manual.
- **9** In the table, enter the following settings:

#### Legends

Glucose

Maltose

# Maltotriose

**IO** In the **Sugars** toolbar, click **IO Plot**.

Continue with 2b in Figure 2.

# Alcohol

- I In the Model Builder window, right-click Sugars and choose Duplicate.
- 2 In the Settings window for ID Plot Group, type Alcohol in the Label text field.
- 3 Locate the Plot Settings section. Select the y-axis label check box.
- **4** In the associated text field, type vol% alcohol.

#### Global I

- I In the Model Builder window, expand the Alcohol node, then click Global I.
- 2 In the Settings window for Global, click Replace Expression in the upper-right corner of the y-Axis Data section. From the menu, choose Component I (compl)>Definitions> Variables>Evol - vol% alcohol.
- 3 Locate the Title section. In the Title text area, type Alcohol Content.
- 4 Locate the Legends section. Clear the Show legends check box.
- **5** In the **Alcohol** toolbar, click **9 Plot**.

The yeast concentration plot (2c in Figure 2) is set up following these steps:

Yeast

- I In the Model Builder window, right-click Alcohol and choose Duplicate.
- 2 In the Settings window for ID Plot Group, type Yeast in the Label text field.
- 3 Locate the Plot Settings section. In the y-axis label text field, type Concentration (mol/m<sup>3</sup>).

Global I

- I In the Model Builder window, expand the Yeast node, then click Global I.
- In the Settings window for Global, click Replace Expression in the upper-right corner of the y-Axis Data section. From the menu, choose Component I (compl)>
   Reaction Engineering>re.c\_X Concentration mol/m<sup>3</sup>.
- 3 Locate the Title section. In the Title text area, type Yeast Concentration.
- **4** In the **Yeast** toolbar, click **O Plot**.

The flavors plot (2d) in Figure 2 is set up following these steps:

#### Flavors

- I In the Model Builder window, right-click Sugars and choose Duplicate.
- 2 In the Settings window for ID Plot Group, type Flavors in the Label text field.
- 3 Locate the Legend section. From the Position list, choose Middle right.

# Global I

- I In the Model Builder window, expand the Flavors node, then click Global I.
- In the Settings window for Global, click Replace Expression in the upper-right corner of the y-Axis Data section. From the menu, choose Component I (compl)>
   Reaction Engineering>re.c\_EtAc Concentration mol/m<sup>3</sup>.

- 3 Click Add Expression in the upper-right corner of the y-Axis Data section. From the menu, choose Component I (compl)>Reaction Engineering>re.c\_AcA - Concentration mol/m<sup>3</sup>.
- 4 Locate the Title section. In the Title text area, type Flavor Concentrations.
- **5** Locate the **Legends** section. In the table, enter the following settings:

Legends				
Ester	-desirable			

Aldehyde -undesirable

6 In the Flavors toolbar, click 💽 Plot.

Last, create the temperature plot 2e in Figure 2.

# Temperature (re)

- I In the Model Builder window, click Temperature (re).
- 2 In the Settings window for ID Plot Group, locate the Plot Settings section.
- 3 Select the y-axis label check box.
- **4** In the associated text field, type Temperature (<sup>o</sup>C).

Global I

- I In the Model Builder window, expand the Temperature (re) node, then click Global I.
- 2 In the Settings window for Global, locate the y-Axis Data section.
- **3** In the table, enter the following settings:

Expression	Unit	Description
re.T-273.15[K]	К	

- 4 Locate the Title section. From the Title type list, choose Manual.
- 5 In the Title text area, type Temperature.
- 6 Locate the Coloring and Style section. In the Width text field, type 2.
- 7 Locate the Legends section. Clear the Show legends check box.
- 8 In the Temperature (re) toolbar, click **D** Plot.

Now, create the space dependent model using the 0D model as a starting point. The Generate Space-Dependent Model feature makes this process straight forward.

# REACTION ENGINEERING (RE)

#### Generate Space-Dependent Model I

- I In the Reaction Engineering toolbar, click 🖙 Generate Space-Dependent Model.
- 2 In the Settings window for Generate Space-Dependent Model, locate the Component Settings section.
- 3 From the Component to use list, choose 2Daxi: New.
- **4** Locate the **Physics Interfaces** section. Find the **Fluid flow** subsection. From the list, choose **Laminar Flow**: **New**.
- 5 Find the Heat transfer subsection. From the list, choose Heat Transfer in Fluids: New.
- 6 Locate the Study Type section. From the Study type list, choose Time dependent.
- 7 Locate the Space-Dependent Model Generation section. Click Create/Refresh.

# MULTIPHYSICS

In the Model Builder window, expand the Component 2 (comp2) node, then click Multiphysics.

# Nonisothermal Flow 1 (nitf1)

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

# Reacting Flow, Diluted Species 1 (rfd1)

In the Physics toolbar, click An Multiphysics Couplings and choose Domain>Reacting Flow, Diluted Species.

Import the mesh from a file. The tank geometry is imported together with the mesh.

# MESH I

Import I

- I In the Mesh toolbar, click া Import.
- 2 In the Settings window for Import, locate the Import section.
- 3 In the Filename text field, type beer\_fermentation\_mesh.mphbin.
- 4 Click Import.

The imported geometry contains a number of selections. These are listed in the **Domain Selections** and **Boundary Selections** sections in the settings window.

Add materials to the geometry.

## MATERIALS

In the Home toolbar, click 📑 Windows and choose Add Material from Library.

# ADD MATERIAL

- I Go to the Add Material window.
- 2 In the tree, select Built-in>Air.
- 3 Click Add to Component in the window toolbar.
- 4 In the tree, select Built-in>Water, liquid.
- 5 Click Add to Component in the window toolbar.

# MATERIALS

Water, liquid (mat2)

- I In the Settings window for Material, locate the Geometric Entity Selection section.
- 2 From the Selection list, choose Wort.

# ADD MATERIAL

- I Go to the Add Material window.
- 2 In the tree, select Built-in>Copper.
- 3 Click Add to Component in the window toolbar.
- 4 In the Home toolbar, click 🙀 Add Material to close the Add Material window.

# MATERIALS

#### Copper (mat3)

- I In the Settings window for Material, locate the Geometric Entity Selection section.
- 2 From the Selection list, choose Tank.

Add variables for the space-dependent model.

# **DEFINITIONS (COMP2)**

#### Variables 2

- I In the Model Builder window, under Component 2 (comp2) right-click Definitions and choose Variables.
- 2 In the Settings window for Variables, locate the Variables section.
- 3 Click 📂 Load from File.

4 Browse to the model's Application Libraries folder and double-click the file beer\_fermentation\_variables2.txt.

Hide all the interior boundaries (used for meshing purposes) except the one that correspond to the vapor liquid interface.

# Hide for Physics 1

- I In the Model Builder window, right-click View I and choose Hide for Physics.
- 2 In the Settings window for Hide for Physics, locate the Geometric Entity Selection section.
- **3** From the **Geometric entity level** list, choose **Boundary**.
- 4 Select Boundaries 6, 8, 16, and 17 only.

# CHEMISTRY I (CHEM)

In this model, the transport properties used will be those defined from materials. The transport properties computed by the Chemistry interface can thus be disabled.

- I In the Model Builder window, under Component 2 (comp2) click Chemistry I (chem).
- **2** In the **Settings** window for **Chemistry**, click to expand the **Calculate Transport Properties** section.
- **3** Clear the **Calculate mixture properties** check box.

Add the molar masses, defined earlier, to the species where it is necessary. Species with names consisting only of chemical elements from the periodic table will get the molar mass calculated and added by default.

# Species: G

- I In the Model Builder window, expand the Chemistry I (chem) node, then click Species: G.
- 2 In the Settings window for Species, locate the General Parameters section.
- **3** In the M text field, type MG.

# Species: E

- I In the Model Builder window, click Species: E.
- 2 In the Settings window for Species, locate the General Parameters section.
- **3** In the M text field, type ME.

# Species: EtAc

- I In the Model Builder window, click Species: EtAc.
- 2 In the Settings window for Species, locate the General Parameters section.
- 3 In the *M* text field, type MEtAc.

#### Species: AcA

- I In the Model Builder window, click Species: AcA.
- 2 In the Settings window for Species, locate the General Parameters section.
- **3** In the *M* text field, type MACA.

# Species: M

- I In the Model Builder window, click Species: M.
- 2 In the Settings window for Species, locate the General Parameters section.
- **3** In the M text field, type MM.

#### Species: N

- I In the Model Builder window, click Species: N.
- 2 In the Settings window for Species, locate the General Parameters section.
- **3** In the M text field, type MN.

#### Species: X

- I In the Model Builder window, click Species: X.
- 2 In the Settings window for Species, locate the General Parameters section.
- **3** In the M text field, type MX.

In the next stage of the modeling process you will set up the physics interfaces describing the mass transport, heat transfer, and fluid flow.

#### TRANSPORT OF DILUTED SPECIES (TDS)

- I In the Model Builder window, under Component 2 (comp2) click Transport of Diluted Species (tds).
- **2** In the **Settings** window for **Transport of Diluted Species**, locate the **Domain Selection** section.
- 3 From the Selection list, choose Wort.

# HEAT TRANSFER IN FLUIDS I (HT)

- I In the Model Builder window, expand the Transport of Diluted Species (tds) node, then click Component 2 (comp2)>Heat Transfer in Fluids I (ht).
- 2 In the Settings window for Heat Transfer in Fluids, locate the Physical Model section.
- **3** In the  $T_{\rm ref}$  text field, type T0.

Heat Source 1

- I In the Model Builder window, under Component 2 (comp2)>Heat Transfer in Fluids I (ht) click Heat Source I.
- 2 In the Settings window for Heat Source, locate the Domain Selection section.
- 3 From the Selection list, choose Wort.

# Solid I

- I In the Physics toolbar, click **Domains** and choose Solid.
- 2 In the Settings window for Solid, locate the Domain Selection section.
- **3** From the **Selection** list, choose **Tank**.

# Fluid 2

- I In the Physics toolbar, click **Domains** and choose Fluid.
- 2 In the Settings window for Fluid, locate the Domain Selection section.
- **3** From the Selection list, choose Empty tank section.
- **4** Locate the **Model Input** section. Click **Make All Model Inputs Editable** in the upper-right corner of the section.
- 5 From the *T* list, choose **Temperature (ht)**.
- **6** From the  $p_A$  list, choose **User defined**.

#### Heat Flux 1

- I In the Physics toolbar, click Boundaries and choose Heat Flux.
- **2** Select Boundaries 2, 13, 15, 20, and 21 only.
- 3 In the Settings window for Heat Flux, locate the Heat Flux section.
- 4 Click the **Convective heat flux** button.
- 5 From the Heat transfer coefficient list, choose External natural convection.
- 6 In the *L* text field, type 3.
- 7 In the  $T_{\text{ext}}$  text field, type T0.

# LAMINAR FLOW I (SPF)

- I In the Model Builder window, under Component 2 (comp2) click Laminar Flow I (spf).
- 2 In the Settings window for Laminar Flow, locate the Domain Selection section.
- 3 From the Selection list, choose Wort.
- **4** Locate the **Physical Model** section. Select the **Include gravity** check box.

Initial Values 1

- I In the Model Builder window, under Component 2 (comp2)>Laminar Flow I (spf) click Initial Values I.
- 2 In the Settings window for Initial Values, locate the Initial Values section.
- 3 In the p text field, type spf.rhoref\*g\_const\*(3-z).

# Pressure Point Constraint I

- I In the Physics toolbar, click 💭 Points and choose Pressure Point Constraint.
- 2 Select Point 13 only.

# Symmetry I

- I In the Physics toolbar, click Boundaries and choose Symmetry.
- 2 Select Boundaries 10 and 18 only.

# STUDY 2

First generate a default solver for fluid and heat transfer only. This produces a default solver setting for natural convection problems, due to the presence of gravity.

## Step 1: Time Dependent

- I In the Model Builder window, expand the Study 2 node, then click Step I: Time Dependent.
- 2 In the Settings window for Time Dependent, locate the Study Settings section.
- **3** From the **Time unit** list, choose **h**.
- 4 In the **Output times** text field, type range(0,0.1,4) range(5,1,24).
- 5 Locate the Physics and Variables Selection section. In the table, clear the Solve for check boxes for Chemistry I (chem) and Transport of Diluted Species (tds).

#### Solution 2 (sol2)

I In the Study toolbar, click The Show Default Solver.

Now choose to include the mass transport as well.

# Step 1: Time Dependent

- I In the Model Builder window, click Step I: Time Dependent.
- **2** In the Settings window for Time Dependent, locate the Physics and Variables Selection section.
- 3 In the table, select the Solve for check boxes for Chemistry I (chem) and Transport of Diluted Species (tds).

#### Solution 2 (sol2)

- In the Model Builder window, expand the Study 2>Solver Configurations>
   Solution 2 (sol2)>Dependent Variables I node, then click Concentration (comp2.cAcA).
- 2 In the Settings window for Field, locate the Scaling section.
- **3** From the **Method** list, choose **Manual**.
- **4** In the Model Builder window, click Concentration (comp2.cEtAc).
- 5 In the Settings window for Field, locate the Scaling section.
- 6 From the Method list, choose Manual.
- 7 In the Model Builder window, click Concentration (comp2.cM).
- 8 In the Settings window for Field, locate the Scaling section.
- 9 From the Method list, choose Initial value based.
- **IO** In the **Model Builder** window, click **Concentration** (comp2.cN).
- II In the Settings window for Field, locate the Scaling section.
- 12 From the Method list, choose Initial value based.
- **I3** In the **Model Builder** window, click **Concentration** (comp2.cX).
- 14 In the Settings window for Field, locate the Scaling section.
- **I5** From the **Method** list, choose **Initial value based**.
- **I6** In the **Model Builder** window, click **Pressure (comp2.p)**.
- 17 In the Settings window for Field, locate the Scaling section.
- **18** From the **Method** list, choose **Initial value based**.
- **19** In the **Model Builder** window, click **Temperature (comp2.T)**.
- 20 In the Settings window for Field, locate the Scaling section.
- 21 From the Method list, choose Initial value based.

Natural convection is present in the wort due to the heat release from the reaction, and the cooling along the tank wall. Restrict the solver from taking too large time steps during the initial phase.

- **22** In the **Model Builder** window, click **Time-Dependent Solver I**.
- **23** In the **Settings** window for **Time-Dependent Solver**, click to expand the **Time Stepping** section.
- 24 Select the Initial step check box.
- **25** In the associated text field, type 1/3600.
- **26** From the Maximum step constraint list, choose Expression.

**27** In the **Maximum step** text field, type (60)\*(t<4\*3600)+(300)\*(t>4\*3600).

Use **Get initial condition** to initialize the datasets and to produce default plot groups. Continue by editing the plot groups before computing the solution.

**28** In the Study toolbar, click  $t_{=0}^{U}$  Get Initial Value.

Begin by creating the datasets to be used in the plot groups.

# RESULTS

Study 2/Wort

- I In the Model Builder window, expand the Results>Datasets node.
- 2 Right-click Results>Datasets>Study 2/Solution 2 (sol2) and choose Duplicate.
- 3 In the Settings window for Solution, type Study 2/Wort in the Label text field.

#### Selection

- I In the **Results** toolbar, click 🐐 **Attributes** and choose **Selection**.
- 2 In the Settings window for Selection, locate the Geometric Entity Selection section.
- **3** From the Geometric entity level list, choose Domain.
- **4** From the **Selection** list, choose **Wort**.

#### Revolution 2D - Wort

- I In the Model Builder window, under Results>Datasets right-click Revolution 2D I and choose Duplicate.
- 2 In the Settings window for Revolution 2D, type Revolution 2D Wort in the Label text field.
- 3 Locate the Data section. From the Dataset list, choose Study 2/Wort (sol2).

#### Revolution 2D - Wort, Study 2/Wort (sol2)

In the Model Builder window, under Results>Datasets, Ctrl-click to select Study 2/ Wort (sol2) and Revolution 2D - Wort.

# Study 2/Tank

- I Right-click and choose **Duplicate**.
- 2 In the Settings window for Solution, type Study 2/Tank in the Label text field.

#### Selection

- I In the Model Builder window, expand the Results>Datasets>Study 2/Tank (sol2) node, then click Selection.
- 2 In the Settings window for Selection, locate the Geometric Entity Selection section.

3 From the Selection list, choose Tank.

Revolution 2D - Tank

- I In the Model Builder window, under Results>Datasets click Revolution 2D Wort I.
- 2 In the Settings window for Revolution 2D, type Revolution 2D Tank in the Label text field.
- 3 Locate the Data section. From the Dataset list, choose Study 2/Tank (sol2).

#### Cut Plane 1

- I In the **Results** toolbar, click **Cut Plane**.
- 2 In the Settings window for Cut Plane, locate the Data section.
- 3 From the Dataset list, choose Revolution 2D Wort.
- 4 Locate the Plane Data section. From the Plane type list, choose General.
- **5** In row **Point 2**, set **x** to **0**.
- 6 In row **Point 2**, set **z** to **1**.
- 7 In row **Point 3**, set **x** to cos(-90[deg]).
- 8 In row Point 3, set y to sin(-90[deg]).

Cut Plane 2

- I Right-click Cut Plane I and choose Duplicate.
- 2 In the Settings window for Cut Plane, locate the Plane Data section.
- 3 In row Point 3, set x to cos(135[deg]).
- 4 In row Point 3, set y to sin(135[deg]).

Now edit the plot groups.

Surface 1

- I In the Model Builder window, expand the Concentration, AcA (tds) node, then click Surface 1.
- 2 In the Settings window for Surface, locate the Expression section.
- 3 In the Expression text field, type spf.U.
- 4 Locate the Coloring and Style section. From the Coloring list, choose Gradient.
- 5 From the Top color list, choose Custom.
- 6 On Windows, click the colored bar underneath, or if you are running the crossplatform desktop — the **Color** button.
- 7 Click Define custom colors.

- 8 Set the RGB values to 0, 128, and 128, respectively.
- 9 Click Add to custom colors.
- IO Click Show color palette only or OK on the cross-platform desktop.
- II From the Bottom color list, choose Custom.
- **12** On Windows, click the colored bar underneath, or if you are running the crossplatform desktop — the **Color** button.
- **I3** Click **Define custom colors**.
- 14 Set the RGB values to 255, 255, and 128, respectively.
- **I5** Click **Add to custom colors**.
- 16 Click Show color palette only or OK on the cross-platform desktop.

#### Concentration, AcA (tds)

- I In the Model Builder window, click Concentration, AcA (tds).
- 2 In the Settings window for 2D Plot Group, locate the Color Legend section.
- **3** Select the **Show units** check box.

# Surface 2

- I Right-click Concentration, AcA (tds) and choose Surface.
- 2 In the Settings window for Surface, locate the Expression section.
- **3** In the **Expression** text field, type T-TO.

#### Deformation 1

- I Right-click Surface 2 and choose Deformation.
- 2 In the Settings window for Deformation, locate the Expression section.
- 3 In the r component text field, type 1.5.
- **4** In the **z** component text field, type 0.
- 5 Locate the Scale section. Select the Scale factor check box.
- 6 In the associated text field, type 1.
- 7 In the Concentration, AcA (tds) toolbar, click **O** Plot.

## Line I

In the Model Builder window, right-click Concentration, AcA (tds) and choose Line.

# Deformation I

In the Model Builder window, right-click Deformation I and choose Copy.

#### Line I

- I In the Model Builder window, click Line I.
- 2 In the Settings window for Line, locate the Expression section.
- **3** In the **Expression** text field, type **1**.
- 4 Locate the Coloring and Style section. From the Coloring list, choose Uniform.
- 5 From the Color list, choose Black.

#### Deformation I

Right-click Line I and choose Paste Deformation.

# Arrow Surface 1

In the Model Builder window, right-click Concentration, AcA (tds) and choose Arrow Surface.

#### Deformation I

In the Model Builder window, right-click Arrow Surface I and choose Paste Deformation.

#### Arrow Surface 1

- I In the Settings window for Arrow Surface, click Replace Expression in the upper-right corner of the Expression section. From the menu, choose Component 2 (comp2)> Heat Transfer in Fluids 1>Domain fluxes>ht.tefluxr,ht.tefluxz Total energy flux.
- **2** Locate the **Arrow Positioning** section. Find the **r grid points** subsection. In the **Points** text field, type **10**.
- 3 Locate the Coloring and Style section. From the Arrow length list, choose Logarithmic.
- **4** From the **Arrow type** list, choose **Cone**.
- 5 In the Range quotient text field, type 1000.
- 6 From the Color list, choose Black.

#### Line 2

- I In the Model Builder window, right-click Concentration, AcA (tds) and choose Line.
- 2 In the Settings window for Line, locate the Expression section.
- **3** In the **Expression** text field, type **1**.
- 4 Locate the Coloring and Style section. From the Coloring list, choose Uniform.
- 5 From the Color list, choose Black.

#### Deformation I

- I Right-click Line 2 and choose Deformation.
- 2 In the Settings window for Deformation, locate the Expression section.
- **3** In the **r component** text field, type **0**.

- 4 In the **z component** text field, type w.
- **5** Locate the **Scale** section. Select the **Scale factor** check box.
- 6 In the associated text field, type 50.

#### Streamline 1

In the Model Builder window, right-click Streamline I and choose Delete.

#### Velocity Field and Temperature

- I In the Model Builder window, under Results click Concentration, AcA (tds).
- 2 In the Settings window for 2D Plot Group, type Velocity Field and Temperature in the Label text field.

#### Concentration, AcA, 3D (tds)

- I In the Model Builder window, expand the Results>Concentration, AcA, 3D (tds) node, then click Concentration, AcA, 3D (tds).
- 2 In the Settings window for 3D Plot Group, locate the Plot Settings section.
- **3** Clear the **Plot dataset edges** check box.

#### Surface 2

- I In the Model Builder window, under Results>Concentration, AcA, 3D (tds) right-click Surface I and choose Duplicate.
- 2 In the Settings window for Surface, locate the Data section.
- **3** From the **Dataset** list, choose **Revolution 2D Tank**.
- 4 Locate the Expression section. In the Expression text field, type 1.
- 5 Locate the Coloring and Style section. From the Coloring list, choose Uniform.
- 6 From the Color list, choose Gray.

#### Concentration, AcA (Acetaldehyde)

- I In the Model Builder window, under Results click Concentration, AcA, 3D (tds).
- 2 In the Settings window for 3D Plot Group, type Concentration, AcA (Acetaldehyde) in the Label text field.

# Temperature, 3D (htl)

- I In the Model Builder window, click Temperature, 3D (htl).
- 2 In the Settings window for 3D Plot Group, locate the Plot Settings section.
- **3** Clear the **Plot dataset edges** check box.

# Surface

I In the Model Builder window, expand the Temperature, 3D (htl) node, then click Surface.

- 2 In the Settings window for Surface, locate the Data section.
- 3 From the Dataset list, choose Revolution 2D Wort.

# Surface 2

- I Right-click Surface and choose Duplicate.
- 2 In the Settings window for Surface, locate the Data section.
- 3 From the Dataset list, choose Revolution 2D Tank.
- 4 Locate the Expression section. In the Expression text field, type 1.
- 5 Locate the Coloring and Style section. From the Coloring list, choose Uniform.
- 6 From the Color list, choose Gray.

Arrow Surface 1

- I In the Model Builder window, right-click Temperature, 3D (htl) and choose Arrow Surface.
- 2 In the Settings window for Arrow Surface, locate the Data section.
- 3 From the Dataset list, choose Cut Plane I.
- 4 Click Replace Expression in the upper-right corner of the Expression section. From the menu, choose Component 2 (comp2)>Heat Transfer in Fluids I>Domain fluxes>ht.tefluxr, ...,ht.tefluxz Total energy flux.
- 5 Locate the Coloring and Style section. From the Arrow type list, choose Cone.
- 6 From the Color list, choose Black.

#### Arrow Surface 2

- I Right-click Arrow Surface I and choose Duplicate.
- 2 In the Settings window for Arrow Surface, locate the Data section.
- 3 From the Dataset list, choose Cut Plane 2.

#### Temperature, 3D

- I In the Model Builder window, under Results click Temperature, 3D (htl).
- 2 In the Settings window for 3D Plot Group, type Temperature, 3D in the Label text field.

Delete the default plot groups that are not needed, namely all concentration plots for CO2, E, EtAc, G, M, N and X (in total 16), plus the two Velocity plots. Easiest way to delete them is to Ctrl-click to select all of them and then right-click and choose **Delete**.

# Temperature, Top and Bottom

I In the Home toolbar, click 🚛 Add Plot Group and choose ID Plot Group.

2 In the Settings window for ID Plot Group, type Temperature, Top and Bottom in the Label text field.

# Point Graph 1

- I Right-click Temperature, Top and Bottom and choose Point Graph.
- **2** Select Point 13 only.

#### Temperature, Top and Bottom

- I In the Model Builder window, click Temperature, Top and Bottom.
- 2 In the Settings window for ID Plot Group, locate the Data section.
- 3 From the Dataset list, choose Study 2/Solution 2 (sol2).

#### Point Graph 1

- I In the Model Builder window, click Point Graph I.
- 2 In the Settings window for Point Graph, locate the Selection section.
- **3** Select the **Image Activate Selection** toggle button.
- **4** Select Point 13 only.
- 5 Locate the y-Axis Data section. In the Expression text field, type T.
- 6 From the Unit list, choose degC.
- 7 Click to expand the Legends section. Select the Show legends check box.
- 8 Find the Include subsection. Select the Expression check box.
- 9 Clear the **Point** check box.
- **IO** In the **Suffix** text field, type , Top.

#### Point Graph 2

- I Right-click **Results>Temperature**, **Top and Bottom>Point Graph I** and choose **Duplicate**.
- 2 In the Settings window for Point Graph, locate the Selection section.
- **3** Select the **I** Activate Selection toggle button.
- 4 Click Clear Selection.
- **5** Click the  $\longleftrightarrow$  **Zoom Extents** button in the **Graphics** toolbar.
- **6** Select Point 2 only.
- 7 Locate the Legends section. Find the Include subsection. In the Suffix text field, type , Bottom.
- 8 In the Temperature, Top and Bottom toolbar, click 🗿 Plot.

With all the plot groups ready, compute the study and choose the setting to update the plots with results while solving.

# STUDY 2

Step 1: Time Dependent

- I In the Model Builder window, under Study 2 click Step I: Time Dependent.
- **2** In the **Settings** window for **Time Dependent**, click to expand the **Results While Solving** section.
- **3** Select the **Plot** check box.
- 4 From the Plot group list, choose Velocity Field and Temperature.
- 5 From the Update at list, choose Time steps taken by solver.
- 6 In the **Home** toolbar, click **= Compute**.

# RESULTS

Velocity Field and Temperature

Time to look at the results.

- I In the Settings window for 2D Plot Group, locate the Data section.
- 2 From the Time (h) list, choose 3.
- 3 In the Velocity Field and Temperature toolbar, click on Plot.

Concentration, AcA (Acetaldehyde)

- I In the Model Builder window, click Concentration, AcA (Acetaldehyde).
- 2 In the Settings window for 3D Plot Group, locate the Data section.
- 3 From the Time (h) list, choose 3.
- 4 In the Concentration, AcA (Acetaldehyde) toolbar, click 💽 Plot.

#### Temperature, 3D

- I In the Model Builder window, click Temperature, 3D.
- 2 In the Settings window for 3D Plot Group, locate the Data section.
- 3 From the Time (h) list, choose 3.
- **4** In the **Temperature**, **3D** toolbar, click **O Plot**.

# Surface

- I In the Model Builder window, click Surface.
- 2 In the Settings window for Surface, locate the Data section.

3 From the Solution parameters list, choose From parent.

# 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 Solution parameters list, choose From parent.

# Arrow Surface 1

- I In the Model Builder window, click Arrow Surface I.
- 2 In the Settings window for Arrow Surface, locate the Data section.
- 3 From the Solution parameters list, choose From parent.

# Arrow Surface 2

- I In the Model Builder window, click Arrow Surface 2.
- 2 In the Settings window for Arrow Surface, locate the Data section.
- 3 From the Solution parameters list, choose From parent.
- 4 Click to expand the Inherit Style section. From the Plot list, choose Arrow Surface 1.

# Temperature, Top and Bottom

- I In the Model Builder window, click Temperature, Top and Bottom.
- 2 In the Settings window for ID Plot Group, locate the Legend section.
- **3** From the **Position** list, choose **Middle right**.