

Molecular Flow Through a Microcapillary

Introduction

Computing molecular flows in arbitrary geometries produces complex integral equations which are very difficult to compute analytically. Analytic solutions are therefore only available for simple geometries. One of the earliest problems solved was that of gas flow through tubes of arbitrary length, which was first treated correctly by Clausing (Ref. 1). Later the integral expressions he derived were computed more accurately by Cole (Ref. 2). These authors derived values for the transmission probability of molecules incident on a tube of arbitrary length, which is independent of the pressure, provided that the Knudsen number is much greater than one (that is in the molecular flow regime).

This model uses COMSOL Multiphysics to compute the transmission probability for molecular flow of molecules through a microcapillary with different ratios of the length to the radius. The results are compared with the accurate solution due to Cole (Ref. 2).

Model Definition

Nitrogen gas flows through a microcapillary tube of diameter 200 µm, from a reservoir at a fixed pressure of 10^{-3} mBar to a reservoir at high vacuum. The flux of molecules entering the tube from the high vacuum end is assumed to be negligible. At this pressure the mean free path of the nitrogen is approximately 100 mm, so the Knudsen number of the flow is much greater than 1. The ratio of the flux of molecules entering the tube to the flux of molecules leaving the tube is computed for tube lengths of 0.4 mm, 0.8 mm, 1.2 mm, 1.6 mm, and 2.0 mm. The model is axially symmetric and is built in the 2D axially symmetric interface. The geometry is shown in Figure 1 for the case of the 2.0 mm long tube. The lower boundary is connected to the reservoir at low vacuum, the upper boundary is connected to the reservoir at high vacuum and the nonaxial vertical boundary is the capillary wall.

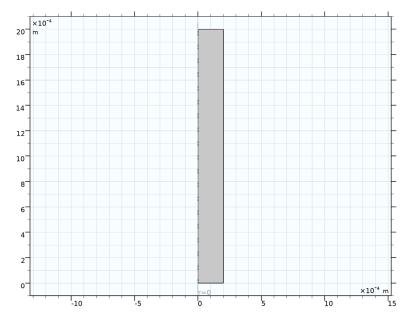


Figure 1: Model geometry.

The probability, χ , of molecules that strike the tube passing through the tube can be computed as the following integral:

$$\chi = \frac{\int 2\pi r G dr}{\int 2\pi r J dr}$$
 RS

where J is the outgoing flux, G is the incident flux, F is the radial distance, F is the boundary adjacent to the reservoir, and F is the boundary adjacent to high vacuum.

Results and Discussion

The incident flux on the surfaces of the tube is shown in Figure 2 for the case of the 2 mm long tube. A line plot of the pressure along the wall is shown in Figure 3. As expected, the incident flux decreases steadily from the reservoir end to the end at high vacuum, as does the pressure. Note that the incident flux (in contrast with the emitted flux) is not constant along the two reservoir boundaries, as it results from reflections within the geometry. The

transmission probability for the tube is plotted against length to radius ratio in Figure 4. Good agreement is obtained between the calculated results and the probabilities calculated by Cole (Ref. 2).

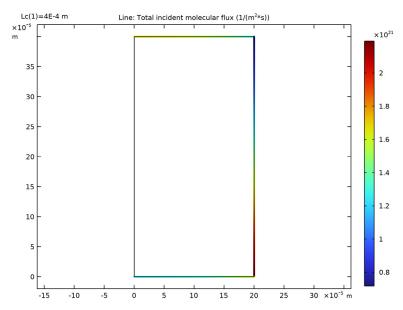


Figure 2: The incident flux inside the tube.

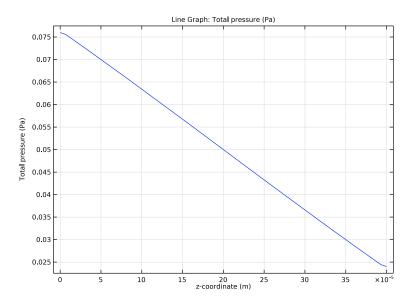


Figure 3: The pressure along the tube wall.

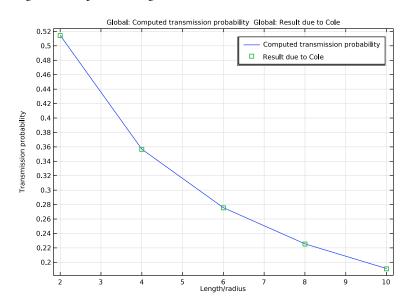


Figure 4: The transmission probability for the pipe as a function of the length to radius ratio. The results due to Cole (Ref. 2) are shown for comparison.

References

- 1. P. Clausing, "Über die Strömung sehr verdünnter Gase durch Röhren von beliebiger Länge", Ann. Physik, vol. 404, no. 8, pp. 961–989, 1932. English translation available as: "The Flow of Highly Rarefied Gases Through Tubes of Arbitrary Length", J. Vacuum Science and Technology, vol. 8. no. 5, pp. 636-646, 2009.
- 2. R.J. Cole, "Complementary Variational Principles for Knudsen Flow Rates", IMA J. Appl. Math., vol. 20, pp. 107-115, 1977.

Notes About the COMSOL Implementation

The model is straightforward to set up using the Molecular Flow interface. The capillary walls are assigned the wall boundary condition, the reservoir boundary condition is used for the bottom opening, and the top opening is assigned the total vacuum boundary condition. The geometry is parameterized, and a parametric solver is used to vary the capillary dimensions.

Application Library path: Molecular Flow Module/Benchmarks/ vacuum_capillary

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 2D Axisymmetric.
- 2 In the Select Physics tree, select Fluid Flow>Rarefied Flow>Free Molecular Flow (fmf).
- 3 Click Add.
- 4 Click 🔵 Study.
- 5 In the Select Study tree, select General Studies>Stationary.
- 6 Click M 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** In the table, enter the following settings:

Name	Expression	Value	Description
Lc	2[mm]	0.002 m	Capillary length
Rc	0.2[mm]	2E-4 m	Capillary radius
p0	1e-3[mbar]	0.1 Pa	Reservoir pressure

GEOMETRY I

Rectangle I (rI)

- 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 Rc.
- 4 In the Height text field, type Lc.
- 5 Click Build All Objects.
- **6** Click the **Zoom Extents** button in the **Graphics** toolbar.

FREE MOLECULAR FLOW (FMF)

Set boundary conditions for the tube entrance and exit.

Total Vacuum 1

- I In the Model Builder window, under Component I (compl) right-click Free Molecular Flow (fmf) and choose Total Vacuum.
- 2 Select Boundary 3 only.

Reservoir I

- I In the Physics toolbar, click Boundaries and choose Reservoir.
- 2 Select Boundary 2 only.
- 3 In the Settings window for Reservoir, locate the Reservoir section.
- **4** In the $p_{0,G}$ text field, type p0.

DEFINITIONS

Define an interpolation function for Cole's solution for the transmission probability as a function of the length-to-radius ratio. You will use it later for comparison with the model result.

Interpolation | (intl)

- I In the Home toolbar, click f(x) Functions and choose Local>Interpolation.
- 2 In the Settings window for Interpolation, locate the Definition section.
- **3** In the table, enter the following settings:

t	f(t)	
2	0.5142	
4	0.3566	
6	0.2755	
8	0.2253	
10	0.1910	

Set up integration nonlocal couplings to compute the total flux of molecules through the system.

Integration | (intob|)

- 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 Select Boundary 2 only.
- 5 Locate the Advanced section. Clear the Compute integral in revolved geometry check box.

Integration 2 (intob2)

- 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** Select Boundary 3 only.
- 5 Locate the Advanced section. Clear the Compute integral in revolved geometry check box.

Use a mapped mesh for this simple geometry.

MESH I

Mapped I

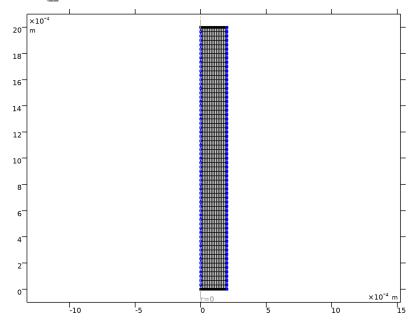
In the Mesh toolbar, click Mapped.

Distribution I

- I Right-click Mapped I and choose Distribution.
- **2** Select Boundaries 2 and 3 only.
- 3 In the Settings window for Distribution, locate the Distribution section.
- 4 In the Number of elements text field, type 15.

Distribution 2

- I In the Model Builder window, right-click Mapped I and choose Distribution.
- **2** Select Boundaries 1 and 4 only.
- 3 In the Settings window for Distribution, locate the Distribution section.
- 4 In the Number of elements text field, type 60.
- 5 Click **Build All**.



Set up a parametric sweep over the capillary length.

STUDY I

Parametric Sweep

- I In the Study toolbar, click Parametric Sweep.
- 2 In the Settings window for Parametric Sweep, locate the Study Settings section.
- 3 Click + Add.
- 4 From the list in the Parameter name column, choose Lc (Capillary length).
- 5 Click Range.
- 6 In the Range dialog box, type 0.4[mm] in the Start text field.
- 7 In the **Stop** text field, type 2[mm].
- 8 In the Step text field, type 0.4[mm].
- 9 Click Add.
- 10 In the Study toolbar, click **Compute**.

RESULTS

Incident Molecular Flux (fmf)

- I In the Settings window for 2D Plot Group, locate the Data section.
- 2 From the Parameter value (Lc (m)) list, choose 4E-4.
- 3 In the Incident Molecular Flux (fmf) toolbar, click Plot.
- 4 Click the Zoom Extents button in the Graphics toolbar.

The default plot shows the incident molecular flux inside the microcapillary tube. Compare with the plot in Figure 2.

Plot the pressure along the tube wall by following these steps.

Total Pressure

- I In the Home toolbar, click Add Plot Group and choose ID Plot Group.
- 2 In the Settings window for ID Plot Group, type Total Pressure in the Label text field.
- 3 Locate the Data section. From the Dataset list, choose Study 1/ Parametric Solutions I (sol2).
- 4 From the Parameter selection (Lc) list, choose First.

Line Graph 1

- I Right-click Total Pressure and choose Line Graph.
- 2 Select Boundary 4 only.

- 3 In the Settings window for Line Graph, click Replace Expression in the upper-right corner of the y-Axis Data section. From the menu, choose Component I (compl)>
 Free Molecular Flow>Pressure>fmf.ptot Total pressure Pa.
- **4** Click to expand the **Quality** section. From the **Resolution** list, choose **No refinement**. Use the z-coordinate for the x-axis data.
- 5 Locate the x-Axis Data section. From the Parameter list, choose Expression.
- **6** In the **Expression** text field, type z.
- 7 In the Total Pressure toolbar, click Plot.

Compare the resulting plot with that in Figure 3.

Finally, plot the result for the transmission probability versus the length-to-radius ratio and compare with Cole's solution.

ID Plot Group 5

- I In the Home toolbar, click Add Plot Group and choose ID Plot Group.
- 2 In the Settings window for ID Plot Group, locate the Data section.
- 3 From the Dataset list, choose Study I/Parametric Solutions I (sol2).

Global I

- I Right-click ID Plot Group 5 and choose Global.
- 2 In the Settings window for Global, locate the y-Axis Data section.
- **3** In the table, enter the following settings:

Expression	Unit	Description
<pre>intop2(2*pi*r*G)/intop1(2* pi*r*fmf.J_G)</pre>		Computed transmission probability

- 4 Locate the x-Axis Data section. From the Parameter list, choose Expression.
- 5 In the Expression text field, type Lc/Rc.

Transmission Probability

- I In the Model Builder window, under Results click ID Plot Group 5.
- 2 In the Settings window for ID Plot Group, type Transmission Probability in the Label text field.
- 3 Locate the Plot Settings section.
- 4 Select the x-axis label check box. In the associated text field, type Length/radius.
- **5** Select the **y-axis label** check box. In the associated text field, type Transmission probability.

Global 2

- I Right-click Transmission Probability and choose Global.
- 2 In the Settings window for Global, locate the y-Axis Data section.
- **3** In the table, enter the following settings:

Expression	Unit	Description
int1(Lc/Rc)		Result due to Cole

- 4 Locate the x-Axis Data section. From the Parameter list, choose Expression.
- 5 In the Expression text field, type Lc/Rc. Change the plot style to display only the data points.
- 6 Click to expand the Coloring and Style section. Find the Line style subsection. From the Line list, choose None.
- 7 Find the Line markers subsection. From the Marker list, choose Square.

Transmission Probability

- I In the Model Builder window, click Transmission Probability.
- 2 In the Transmission Probability toolbar, click **2** Plot.

Compare with Figure 4.