PHAS0030: Computational Physics  
Session 5: More Differential Equations

David Bowler

In this session, we will continue our examination of differential equations, introducing matrix methods for their solution, and beginning to look at partial differential equations (PDEs). We will recall how PDEs are characterised, and investigate how two of these types are solved.

# Objectives

The objectives of this session are to:

* Understand matrix approaches to boundary value problems for ordinary differential equations
* Characterise partial differential equations into hyperbolic, elliptic and parabolic
* Demonstrate how parabolic equations can be solved, first with explicit finite differences and then with implicit methods
* Consider elliptic equations, and explore the use of matrix approaches in their solution, investigating inversion and alternatives to inversion

# Review of Session 4

In the fourth session, we looked at ordinary differential equations (ODEs), seeing how Euler’s method can be used as a simple but often unreliable approach. We discussed how second order ODEs can be broken down into coupled first order ODEs, and investigated more accurate approaches to their solution. We introduced boundary value problems and the shooting method to solve them, and considered the SciPy functions that can be used in place of hand-coded functions.

## Implicit methods

In Session 4, we saw that the Euler method is inherently unstable for large step sizes. For the simple differential equation:

we showed that this instability[[1]](#footnote-1) comes directly from the use of the forward difference approximation for . What would happen if we used the *backward* difference instead?

This formula is stable, and does not suffer from the problems of the simple Euler method. In most cases, the form of the function is complicated, and we have the difficulty that the update for the step depends on the value . This type of problem is a known as an *implicit* problem, and requires some form of iteration, sometimes using a root-finder to solve each step. In the next section, we will see an alternative way of solving implicit problems.

For the simple case above, however, we can write:

Notice that this will be stable for all values of , and will converge to the correct result for large values of . Notice also that the update step (from *n-1* to *n*) is the same *to first order* as the update step from the forward difference formula.

# Matrix approach to boundary-value problems

We will now introduce an alternate approach to boundary-value problems which uses matrices. We’ll consider a different first-order differential equation, which is concerned with the one-dimensional flow of heat in a bar with cross-sectional area , and when solved gives the temperature distribution in the bar:

In this case, is the heat flow, which will be a constant in the steady state, is the temperature in the bar at a point , is the thermal conductivity and we will fix the temperature at the ends of the bar. This gives a boundary value problem, and is illustrated in the diagram below.



One dimensional heat flow in a bar, in the steady state. Heat flow, Q, temperature θ.

To solve this with the shooting method, you would choose a value of , and then integrate to find values of from the start of the bar to the end. We would then adjust until the value of at the end of the bar matched our set point.

There is an alternative approach to the shooting method, using matrices, that is a standard approach to diffusion problems (which are *partial* differential equations) that we can also use for this simple ODE when is a constant.

We discretise the problem[[2]](#footnote-2), so that we will evaluate the temperature at a set of points along the bar, . Then we note that, since is constant, we must have:

Now, substituting the differentials with finite differences, and assuming that is constant for simplicity, we find:

This is a general relationship that links points that are adjacent along the bar. (You can see how we could write the integration point-by-point in terms of this kind of equation.) However, if we write out all the equations in one block, you should see that it would form a matrix. We will have a series of equations like:

where , etc. Notice that we have equations for points, but we will specify the values of two points: and , giving us the right number of equations.

We can then re-arrange these equations into a single matrix equation:

The only real manipulation needed to make these equations is for the first and last lines, where we move the terms involving and to the right-hand side of the equation.

We now have to solve a problem that can be written[[3]](#footnote-3) as , which is a classic problem in linear algebra. The obvious solution involves the inverse of the matrix: .

There are two main Numpy approaches that you can take: np.linalg.solve(M,b) will solve for, and return, ; and np.linalg.inv(M) will calculate the inverse matrix to M and return it (and it can then be applied to any boundary condition vectors). (Confusingly, Scipy also offers a module linalg and solvers linalg.inv and linalg.solve though they are not identical to the NumPy solvers; we’ll stick with NumPy for now.) You perform the matrix-vector multiply using np.dot(A,b) where A is the matrix and b is the vector.

## Exercises

### In class

1. You will construct the matrix for the bar with uniform cross-section, using the np.diag function. If you pass a 1D array to np.diag then it will return a 2D array with the 1D array along the diagonal. You can also specify a shifted diagonal (consider the diagonals in the matrix with 1 instead of -2) using np.diag(array,k=n) where n can be positive or negative.
   1. Define a variable for your problem size, , and create an array of length with every element set to -2.
   2. Create an array of length with every element set to 1 (remember the command np.ones)
   3. Create the matrix by summing together the result of *three* calls to np.diag and print it out

* Your matrix should look something like this (example for ):
* [[-2. 1. 0. 0. 0. 0. 0.]  
   [ 1. -2. 1. 0. 0. 0. 0.]  
   [ 0. 1. -2. 1. 0. 0. 0.]  
   [ 0. 0. 1. -2. 1. 0. 0.]  
   [ 0. 0. 0. 1. -2. 1. 0.]  
   [ 0. 0. 0. 0. 1. -2. 1.]  
   [ 0. 0. 0. 0. 0. 1. -2.]]

1. Use np.linalg.inv to invert the matrix and solve for the temperature, with boundary conditions for and , using np.dot(Minv,b). Plot the result (including the end-points). Note that we assume that the bar is 1m long.

### Further work

1. Now consider a bar which has cross-sectional area . Write a function that returns the area for an input of .
2. Construct a matrix for the varying cross-section, following these steps, which are an adaptation of the method we used above:
   1. Start as you did before with a size,
   2. Define for a bar of length 1m.
   3. For the main diagonal, you will need to create an array whose entries are , where and starts at zero.
   4. For the shifted diagonals you can use *one array*, with starting from one (note that this will still give the correct entries)
   5. Now build the matrix using np.diag. Print your matrix to check that it is symmetrical (you may find np.set\_printoptions(precision=5) helpful)
   6. Now solve for the temperature, setting the boundary conditions as and . Plot your result, with the boundary conditions, and ensure that there are no discontinuities.

# Partial Differential Equations

Partial differential equations (PDEs) involve solving for a function of more than one independent variable, which are most commonly position and time in physics. While it is perfectly possible to write a first-order PDE, the most common forms are second order. If we consider a function of and for simplicity[[4]](#footnote-4) then a general linear, second-order PDE can be written as:

The three functions , and are often constants, and determine the behaviour of the equation, and the approach to solving required. If the right-hand side of the equation is zero, or is linear in and its first-order derivatives, then the equation is called *homogeneous*: any constant multiple of a solution is also a solution.

Most PDEs in physics are linear, second-order PDEs, with examples including:

which are the one-dimensional wave equation (easily extended to three dimensions), Laplace’s equation, the diffusion or heat equation and the time-dependent Schrödinger equation respectively. These are all homogeneous; by adding a charge density to Laplace’s equation (to get Poisson’s equation) or a heat source to the heat equation we would obtain *inhomogeneous* equations, which are also extremely important.

These equations all behave differently, and require different boundary conditions and approaches to solving. They can be characterised by considering the discriminant function:

where there are three possibilities, named after conic sections (which are defined by the same equation):

1. Hyperbolic: propagating oscillations, such as waves (A=1, B=0, C=-1/c2)
2. Parabolic: transport processes, such as heat flow and diffusion (A=1, B=0, C=0)
3. Elliptic: stationary systems, such as electrostatic fields and steady-state temperature distributions (A=1, B=0, C=1)

We will consider the solution of each of these types separately (with hyperbolic considered in Session 6). Of course, as with any broad classification scheme, care must be taken: notice that the heat equation is both parabolic (when the temperature varies with time) and elliptic (in a steady-state situation).

## Boundary Conditions

For ordinary differential equations, the specification of boundary conditions was rather simple, and required only a number of pieces of information equivalent to the order of the equation. It is considerably more complex for second-order PDEs, with the information required and the domain over which it is specified varying with the nature of the problem.

Each equation type requires boundary conditions named after famous mathematicians who worked in the area. The boundary conditions are specified in terms of the value of the function, , or its derivative along the normal to the boundary, .

1. Hyperbolic: Cauchy conditions ( and on an open boundary)
2. Parabolic: Dirichlet () or von Neumann () along an open boundary
3. Elliptic: Dirichlet or von Neumann on a closed boundary

Note that both hyperbolic and parabolic are time-dependent, while elliptic is not. A helpful example of Dirichlet and von Neumann boundary conditions comes from electrostatics: we would specify either the potential (Dirichlet) or (von Neumann) along a boundary (where the normal vector is ).

# Parabolic equations

If we are interested in heat flow away from the steady state, then we need to consider a new equation for the temperature, :

where is the specific heat capacity of the material, is the (mass) density and is the thermal conductivity.

We will return to our example of one-dimensional heat flow in a bar, but now consider time dependence: the whole bar will start at 300K, but we will raise the end at to 500K at . For a typical metal, the parameters would have values of WmK, JkgK and kgm.

How do we go about solving this? The simplest approach would be to take finite differences; for the second derivative in we can use a centred formula, which we know is accurate. The time component is more difficult: we will start with a simple forward difference, and investigate its behaviour[[5]](#footnote-5). We will then introduce a better approach. We will discretise space and time, so that a point . Keep track of these indices: we will write the temperature as a function of space and time as: .

The spatial derivative is given by:

while we will, for now, write the time derivative as:

How do we proceed? We should have a starting temperature distribution (say for all values of ), and we want to know how it evolves over time. So we want to find for all in our bar, for various times indexed by . We can substitute and re-arrange:

where (this Greek letter is zeta). Notice that the value of the temperature at any point is given by its value and its neighbours’ values at the previous time step: a simple, explicit scheme. You will construct a simple implementation of this in the exercises.

You should find during the testing of your implementation that there is a critical value of the parameter . Notice that this combines both physical parameters of the system, and the timestep and grid spacing; there is a key ratio for stability when the timestep becomes too large (just as we saw for the Euler method). You may notice that the update for each timestep is rather similar to what we saw in Section [4](#sec:part-diff-equat). Indeed, we might imagine that we could write the overall process in a similar matrix equation—and indeed we can. It turns out that we can show that the stability is related to the eigenvalues of the update matrix, which need to be smaller than one[[6]](#footnote-6)

## Implicit methods

We know from our work on finite differences that centred formulae are more accurate and stable; can we use this here? We can indeed. If we think of the formula we’ve been using not as a forward difference, but as a *centred* difference at the timestep then, so long as we can update the right-hand side of the equation for the spatial derivative, we should be more stable.

We will approximate the spatial derivative at time by averaging:

If we substitute these into the heat-flow equation, we find:

This is an *implicit* method: the values of at a point at each timestep depend on the values of its neighbours in the previous timestep *and* at the present timestep. We cannot solve this by direct iteration, as before, but we can use a matrix approach, exactly paralleling what we did in Section [3](#sec:matr-appr-bound). As we have to incorporate values of from neighbours at two timesteps, and include boundary conditions, the overall result is a little more complex than before:

where the matrices and are defined as:

and the vector **b** contains the boundary conditions. This is more complicated than the example in Section 3 because we have time dependence. The boundary conditions for our bar would be a vector of zeros except at the first and last entries, which would be and . If we solve for once at the start and also calculate and then the propagation becomes rather simple. This implicit method is known as the Crank-Nicolson method, which can be shown to be stable for all values of (though of course large values of will not give accurate results).

## Exercises

### In class

1. Write a function to perform the step-by-step update (*not* the matrix version) based on forward differences in time. You should pass as parameters an array of temperatures at timestep only (i.e. a 1D array) and the constant zeta. Return an array of temperatures at timestep . When you iterate along the bar, remember to *exclude* the end-points. [You may find np.size useful to determine the iteration.]
2. Create a 2D array to store the temperature; don’t make the spatial domain too large (I chose 7 points) and use 10 time points. Set the boundary conditions ( at for all points, except ). Define a parameter zeta (start with 0.1) and loop over time, calling the update routine to evolve the differential equation forward in time. Ensure that you set the boundary conditions at each step if necessary. [Remember that, if you have an array temperature[N,I] with I spatial points and N time points then temperature[n] will give a 1D array with all points along the bar at timestep n. You can also store a 1D array in your 2D array using something like temperature[n] = array.]
3. Plot the resulting evolution of the temperature distribution (you could loop over time steps and put them all on the same graph using plt.plot or you create an array of plots using the figure approach).
4. Now repeat the calculation for zeta=0.7. What happens? If you have time, identify a critical value of zeta.

### Further work

1. Write two functions to create the matrices and . You should pass as parameters the dimensions of the problem (remember that for points your matrices should have size ) and (zeta) and return the matrix. You can use the same approach using np.diag as we did above.
2. Now solve the same problem as we did in Question 2 of the in-class work, and experiment with the value of . You should ask yourself two questions: 1. Is the approach stable for all values of ? 2. How does the long-time solution vary with ? You may need to change the number of steps you use with smaller values of .

# Elliptic equations: iterative approaches

We now turn to elliptic equations, which have two spatial dimensions, instead of one spatial and one temporal dimension as we have seen just now. The simplest equation is Laplace’s equation in two dimensions, which is written:

We can apply a centred finite-difference scheme, where we now have points and and we will notate as . We can then write:

If we set , then we can make a considerable simplification:

This can be solved using a matrix approach, but the creation of the matrix involved is a little complex, and involves non-trivial indexing; we will discuss this in Sec. [7](#sec:elliptic-equations) for information. As an alternative, we can use a set of methods generally known as *relaxation* methods to give an *iterative* procedure for the update of each point in the domain where we are solving the equation, *without* the need for matrices (and the associated issues involved in converting from a 2D (or 3D) grid to a 1D vector). We will see below in Sec. [7.1](#sec:altern-invers) that these relaxation approaches can also be used with matrix methods.

At the simplest level, we need an update for the potential at each step of the iteration, ; if we re-write the equation above, we find:

where in the second equation we have indicated how the update works. In this case, after creating an initial set of values for , we simply set the boundary values of at each step and then update. This is known as the Jacobi method; a sample implementation of the update is:

def update\_phi(phi, N):  
 *"""Update NxN grid of phi using Jacobi method"""  
 # Copy rather than equate to avoid update issues* phiout = np.copy(phi)  
 *# Avoid boundaries in update*  
 for i in range(1,N-1):  
 for j in range(1,N-1):  
 phiout[i,j] = 0.25\*(phi[i-1,j] + phi[i+1,j]   
 + phi[i,j-1] + phi[i,j+1])  
 return phiout

Note that we have not specified how to choose the initial value: this is inevitably a form of educated guess, and can have a significant effect on the speed and stability of the iteration.

An alternative method, known as the Gauss-Seidel method, can be implemented in almost the same way; the only change is to use the updated values of instead of the current values[[7]](#footnote-7). This is a more efficient, and importantly a more stable, method to use, particularly when introducing a source term. In that case, the equation we solve would be something like , and the update equation would need an extra term on the right-hand side: , where the grid spacing is .

A further refinement, called the successive over-relaxation (SOR) approach, can also be used here, and can significantly improve the efficiency. Essentially it mixes the present and updated values of via a parameter :

There is no general way to find the optimum value of , and some experimentation on small, simple problems is often needed. You should note that guarantees stability (and if then we have the Jacobi method).

## Exercises

### In-class exercises

1. Adapt the basic Jacobi solver above to implement first the Gauss-Seidel solver, and then the SOR solver with Gauss-Seidel. For the SOR method you should pass as a parameter.
2. We will solve for the electrostatic potential in a square, with the potential fixed at on the boundaries and on the boundaries. Set up an grid for (where is a variable that you can adjust) which will *include* the boundaries. Set the initial values of to include the boundary conditions (3V for and 4V for ) with elsewhere. Using a while loop, iterate using the Gauss-Seidel method to find a solution for (you should calculate the maximum change in any element of from step to step using np.max and np.abs). Keep a record of the number of iterations, and output it at the end.
3. Plot the resulting potential using plt.imshow or plt.contourf. With imshow you might experiment with interpolation, and compare to a solution using larger numbers of points. With contourf, remember that you can pass an array of contour values to use.

### Further work exercises

1. Now use the SOR method for values of between 0.1 and 0.9, and find the most efficient value.

# Supplementary: Elliptic equations: matrix approaches

[Note: this year we will not cover this section in detail in class; it is an important concept to grasp if you intend to go further in computational physics, and is included for completeness.] How do we go about solving elliptic equations with matrices? We need to do what we have done before: write a set of equations for each grid point, which can then be combined into a single matrix-vector operation. To do this, we need to map the points on the grid into a vector (so that the index in the vector, for an grid with and starting from 0). Two routines which convert from to and *vice versa* are:

def ij\_to\_index(i,j,N):  
 """Convert i,j pair in (NxN) grid to index"""  
 return i+j\*N

from math import floor  
def index\_to\_ij(index,N):  
 """Convert index to i,j pair in (NxN) grid"""  
 j = floor(index/N)  
 i = index - j\*N  
 return i,j

You will need these in the exercises below. We can then write the differential equation as a matrix acting on a vector; to calculate the matrix elements we will use the equation from before:

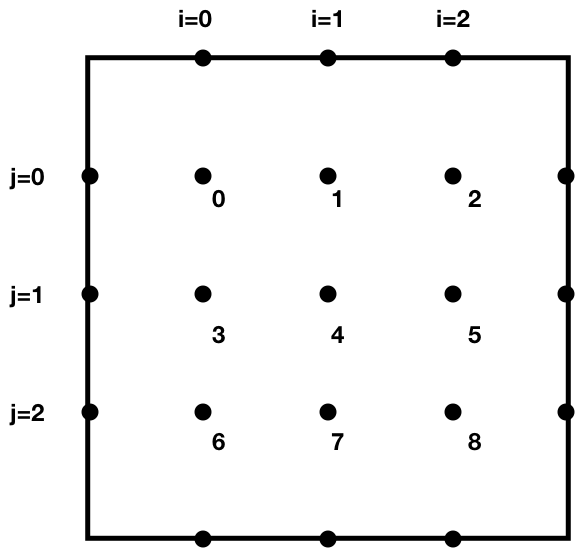
We can create a single, composite index *a=i+Nj* and rewrite the equation above to make it clearer how this maps to a matrix-vector problem:

Note that if or is on the boundary (i.e. out of the normal range) we replace the value of with the appropriate boundary condition, and move them to the right hand side of the equation.

We then form a matrix equation for the vector form of the potential, with the boundary conditions as a vector forming the right-hand side. The final form is:

The mapping from physical problem to computer code is a little complicated: we go from a 2D grid of points to a 1D vector for the potential.

To make this concrete, we will consider a small, simple problem, illustrated in the figure below: a mesh with the potential fixed at on the boundaries and on the boundaries. Note that you will need to use both (or ) and . In this case, and . There are 9 sites corresponding to 3 *x* coordinates and 3 *y* coordinates. The matrix is . The rest of the derivation is in the exercises.



The mapping from a simple grid to a length 9 vector

## Alternatives to inversion

Matrix inversion becomes expensive as matrices get large (it scales with the cube of the size: scaling). As the matrix that we are using scales as the square of the number of grid points in or , this can rapidly become very expensive. As our matrices are rather structured (not quite tridiagonal, but certainly dominated by the diagonal) then other methods are worth exploring.

We saw some of these methods briefly above, and in Session 2 when considering the optimisation of functions: we are again trying to solve the standard linear algebra problem . The Jacobi method writes where is just the diagonal of the matrix, and defines the iteration:

and the solution is iterated until it converges. This is often not a fast converging method, and the Gauss-Seidel method is used instead. Here, formally, we write the original matrix as the sum of a lower-diagonal matrix and an upper-diagonal matrix (though note that the diagonal entries of does not include the diagonal entries) and iterate:

This looks, at first sight, as if it will require inversion of the matrix , but it can be shown that the following element-by-element update solves without inversion, and without decomposing the original matrix:

This is easy to implement, and is iterated until convergence is reached. Notice that it is almost the same as Jacobi iteration, but with the vector split into the old (iteration *k*) and new (iteration *k+1*) parts. It is the decomposition into lower and upper that allows this to be done.

You should be aware that this approach may converge slowly, and a further refinement is sometime used, called the successive over-relaxation (SOR) method. In this case, at each step in the iteration, we mix a proportion of the previous solution with a proportion of the Gauss-Seidel solution:

where ; note that this is a reformulation of the SOR method above (with a different definition of ). To derive this, we have to decompose the matrix as , though the implementation is essentially exactly as above. The SOR method can be effective, but requires some study of the system to find the correct value of . If is chosen poorly, then the convergence can be very slow.

## Exercises

Note that the in-class exercise is basically a complicated indexing problem; do not worry if you cannot finish it – it is unusually hard.

### In class

1. Implement the functions given above to convert between the 2D and 1D representations. Now write a function to calculate the Laplacian matrix and the boundary condition vector, using the following steps:
   1. Pass as parameters *either* *or* and the boundary conditions for both and (in this case for and for ). You will need to calculate whichever of the two sizes you did not pass.
   2. The matrix can be built using np.diag. The main diagonal (where ) should be made from an array of length , containing entries of -4
   3. There are then *four* other sub-diagonals (based on the terms etc). For the and sub-diagonals you need arrays with entries 1 with a location of (use the parameter k when calling np.diag) and a length of . For the and sub-diagonals you need a location of and a length of ; the array should be set to 1 *except* for the elements where , which should be zero
   4. Now make the boundary condition vector: loop over , use the functions above to find and and *accumulate* the boundary conditions if or is or . (You need to accumulate for and for ; check you understand why.)
2. Now solve for the potential, using np.linalg.inv and np.dot, or np.linalg.solve to solve the appropriate equation. You will need to reshape the resulting 1D vector into a grid (use np.reshape(potential,(Nx,Ny))) and plot using plt.imshow. You may find the optional parameter interpolation=’bicubic’ makes the very blocky result clearer.

### Further work

1. Create a Jacobi or Gauss-Seidel solver for the electrostatic grid problem above, and check that the results match the exact inversion. You will need to define a tolerance and stop when the change in the solution (defined in some way - maybe the RMS change between iterations) is smaller than this. You might like to increase the size of the grid, and see whether the numerical cost becomes noticeable.

# Assignment

You will investigate the heat equation in two forms: first, a 2D steady state (actually equivalent to the Laplace equation); second, a 2D time varying solution (using the forward difference approach). You will consider a bar of length 21m and width 15m. The end of the bar at will be maintained at 400K, while the other boundaries will be maintained at 300K.

First, we seek a steady-state solution using an iterative approach to an elliptic equation. We have:

1. Write a function to implement the successive over-relaxation (SOR) update (using the approach from Section [6](#sec:ellipt-equat-an)). You may need to be careful to ensure that the boundary conditions are maintained.
2. Set up an array for a guess initial temperature of the bar (300K except where boundary conditions change it is a good guess). Don’t make the array too large. Plot the initial guess.
3. Choose a value of and a tolerance, and iterate the SOR function until the largest absolute difference between iterations is less than the tolerance (you may find np.abs and np.max useful). Plot the final temperature distribution.
4. If you want to investigate further, you could: experiment with values of to find how it affects convergence; experiment with the initial guess; see how the time required for the solution scales with the array size

Second, we will solve the parabolic equation for the time evolution of the bar (again in 2D - the extension is easy).

1. Write a function to perform the step-by-step update as you did in Section [5.2](#sec:exercises-2) for 1D. Note that the update equation will be:
2. As before, exclude the edges (the boundary conditions).
3. Set up an array for the temperature of the bar as before, and plot the initial temperature distribution.
4. Set and run the time evolution for 400 steps.
5. Plot the final temperature distribution and compare it to the answer from the first part.

This second approach is equivalent to finding the steady state solution for a bar which starts at 300K throughout, and then has one end raised to 400K at . If you want to explore further, you could experiment with different boundary conditions (e.g. one end oscillates in temperature).

# Progress Review

Once you have finished *all the material* associated with this session (both in-class and extra material), you should be able to:

* Use matrix approaches to solve boundary value problems for ODEs (where appropriate)
* Understand how to classify PDEs
* Solve simple examples of both elliptic and parabolic PDEs given appropriate boundary conditions

1. We showed that at the timestep, , we could write . [↑](#footnote-ref-1)
2. Of course, we discretise the problem with the shooting method, but in that case we solve for each point sequentially, whereas here we solve for all points at once. [↑](#footnote-ref-2)
3. I am using to indicate the vector of temperatures at points along the bar [↑](#footnote-ref-3)
4. We could just as easily have chosen and , and these arguments extend to three dimensions as well. [↑](#footnote-ref-4)
5. We may well suspect that it will not be stable beyond a certain time step, on the basis of our experience with Euler’s method. [↑](#footnote-ref-5)
6. We will not pursue this further, but it’s important to be aware of this kind of analysis. [↑](#footnote-ref-6)
7. In the example code, we would not need the variable phiout, but would instead just update phi: phi[i,j] = 0.25\*(phi[i-1,j] + …). [↑](#footnote-ref-7)