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Numerical Project

PLANAR POTENTIAL FLOW

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Part I

Functional requirement of the program

1 The project

The goal of this project is to simulate the movement of a fluid through different geometries. The program creates a box of a choosen size, builds a geometry inside it and simulates the movement of a given fluid.

2 Files

In order to increase readability, the project is made of several files. I made the choice to work with Object Oriented Programming.

- `main.py`: This file calls for the needed functions/class
- `matrices.py`: This file contains the class “Matrices”, it builds the geometry, the different matrices to plot and stores them
- `plot.py`: This file plots the matrices built in “matrices.py”
- `parameters.py`: This file contains all the variables that can be changed by the user
- `data_check.py`: This file checks the variables and makes sure that the program will run

3 Data

This project uses several piece of data set by the user to work.

- `Nx`: `uint` and `Ny`: `uint`, size of the domain
- `h`: `uint`, size of a cell
- `geometry`: `str`, choosen geometry
- `angle`: `uint`, angle of the widening/shrinkage geometry
- `vx`: `float`, initial fluid speed (Neuman’s condition)
- `phi_ref`: `float`, reference velocity potential (Dirichlet’s condition)
- `rho`: `float`, relative density of the fluid
- `pressure_init`: `float`, initial pressure of the fluid

Be careful in the case of a widening/shrinkage geometry! In order for the program to generate a domain from one end to another, there is a restriction on the angle, if the restriction is not met, the program will output a `ValueError`. The restriction is as follows:

$$|angle| < \arctan\left(\frac{0.5 \times N_y - 1}{N_x}\right)$$

The angle parameter should be set in degree, the program will convert it to radians for the computation.

4 Outputs

As of the alpha version, the program outputs 4 pdf files, one for each plot. The files are saved in a subfolder named **figures/** and the filenames are set with the following rule:

$$\langle data \rangle_ \langle geometry \rangle_ Nx = \langle Nx \rangle_ Ny = \langle Ny \rangle. pdf$$

`data` stands for the plotted data (potential, velocity, streamlines, pressure).

5 Concerning the running time

Due to the function `numpy.linalg.solve()` being slow for big matrices, the bigger the size of the domain, the higher the running time.

For a domain size of 3600 cells (60×60), it takes around 20 seconds to run, for a domain size of 14 400 cells (120×120), it increases to 23 minuts.

I searched for a faster method to solve the linear system in vain, thus I recommend to stay on relatively low values for N_x and N_y , the graphs are easily readable for a value of 60 each.

By default, python only uses one CPU core when executing a program, the solution to that problem is parallel programming, however I am not enough experienced in that field to develop an optimized program.

Part II

Internal structure of the program

6 Physical model

In order to build the model, the program uses a squared structured meshe model (matrix). The values are computed at each point of the matrix.

6.1 Velocity potential field

The velocity potential field is computed by solving a system of linear equations. The result is computed using the matrix method, by solving the equation $Ax = b$ we can find the potential flow value in each cell.

The matrix A , of size $N_{\text{fluid}} \times N_{\text{fluid}}$ stores 2 information per cell, the number of neighboring cells and the label of each one of them. The number of neighboring cells is stored in the diagonal of the matrix. Let c be the number of a cell containing fluid and n_c the number of neighbors for the cell n , the diagonal elements are defined as follows:

$$A_{n,n} = -n_c$$

Let i be the set of fluid cells neighboring n , we then have:

$$A_{n,i} = 1$$

There are at most 4 neighbors for each fluid cell.

6.2 Velocity field

Knowing the potential in all cells of the domain, the velocity can be computed by calculating the gradient of the potential. The method used is the finite difference:

$$\begin{aligned} v_x(i, j) &= -\frac{\phi_{i+1,j} - \phi_{i-1,j}}{2h} \\ v_y(i, j) &= -\frac{\phi_{i,j+1} - \phi_{i,j-1}}{2h} \end{aligned}$$

However, this method only works if the cell (i, j) is surrounded by fluid cells. For the cells at the edge of the domain, the method used is slightly different. As mentionned in the scope statement[1], if the cell $(i + 1, j)$ is not a fluid cell, we consider it as a fictitious cell with the value of the cell (i, j) . The same rule applies for each of the neighboring cells that are not fluid cells.

6.3 Streamlines

In the alpha version, the streamlines are computed using matplotlib.

In the beta version, the method of Euler is used.

In the gold version, the planned method is Runge-Kutta 4, however I had not the time to complete the gold version.

6.4 Pressure field

In order to compute the pressure, we need to know several things. Firtly, the velocity field must be know at each cell of the domain. Then we need the initial pressure of the domain. Moreover, there are certain conditions to meet regarding the system:

- the fluid is ideal
- the fluid is incompressible
- the fluid is irrotationnal

Knowing all that, we can use BERNOULLI's Theorem:

$$\mathcal{H} = P + \frac{1}{2}\rho v^2 \quad [2]$$

\mathcal{H} stays the same on each point of the domain. Using this constant, we can then compute the pressure at each point of the domain.

7 Constitutive elements

7.1 `main.py`

This file is executable (without parameters input in the command).

After the imports, it will first call the file `check_data.py` to check the type of the input data.

Then it will call the file `matrices.py` to initiate the values of `G`, `M`, `cell_coords`, `A` and `b`.

It then checks for the `recompute` value, if `True` or if there are no existing **dat** files for the current parameters, it will call the function `domain_check()` to display a warning if the domain is large. It then calls the subroutine `make_data()` to compute `phi`, `grad_y`, `grad_norm`, `pressure` and `pressure` and saves them in **dat** files.

Else, it will use the existing `dat` files to generate the plots.

Then it calls the function `load_data()` to read the **dat** files and stores them in a dictionary and it initiate the file `plot.py` which will be used to plot the different values.

The last lines call the subroutine `plot_graphs` with different arguments to plot and save all the wanted graphs.

7.2 `data_check.py`

This file has several subroutines and functions to rule the execution of the program.

7.2.1 `data_check()`

The subroutine `data_check()` reads the value of each parameter and checks that the type and the value are correct and will not cause a crash of the program. If the value/type is incorrect, it will raise an error and stop the program.

7.2.2 `existing_data()`

The function `existing_data()` will check for specific files in the **dat/** subfolder. If at least one files is missing, it returns `True` and the program will recompute all the data. Else it does nothing.

7.2.3 `domain_check()`

The subroutine `domain_check()` is just a warning. It read the max value of the matrix `M` plus one which is the number of fluid cells to compute. If this number is higher than 5000, it will display a message warning the user that the program can take some time to run. Then it asks if the user wants to keep going. `"Yes"` will continue, `"No"` will stop the program and anything other than that will stop the program aswell.

7.3 `matrices.py`

This class is where everything is computed, from start to end.

7.3.1 `__init__()`

`__init__()` stores the value of `G`, `M`, `cell_coords`, `b` and `A` which will be used several times in the class.

7.3.2 `make_data()`

The subroutine `make_data{}` will call the functions that compute `phi`, `grax_x`, `grad_norm`, `pressure` and `pressure`. Each function will write the data in a **dat** file.

7.3.3 `load_data()`

The function `load_data()` reads the **dat** files and return the values in a dictionary.

7.3.4 `build_g()`

The function `build_g()` takes 4 input arguments.

- `Nx: int`
- `Ny: int`
- `geometry: str`
- `angle: int` of size 8 bits, in radians

It acts as a selector. Given a geometry, it will create a matrix `G` filled with zeros and call the function `build_geometry()` with different input values. There is no real point in this function in the alpha version since the straight, widening and shrinkage geometries are generated from the same function, with only the angle changing. However, it will prove useful in the beta and gold version.

7.3.5 `build_geometry()`

The function `build_geometry()` takes 2 input arguments.

- `G: np.array`
- `angle: int` of size 8 bits, in radians

The function returns the matrix `G`.

The function starts by computing `alpha` as the tangent of the angle.

It then uses it in a `for` loop to compute an offset which will be used to build the geometry.

In case of a shrinkage geometry, the angle inputed is negative, the function will flip horizontally the matrix. It then sets the inlet and outlet of the domain.

Since the program will always compute `G` no matter what and to avoid some problem, it will save the matrix only if `recompte == True`.

7.3.6 `build_index_matrices()`

The function `build_index_matrices()` takes 2 input arguments.

- `Nx: int`
- `Ny: int`

The function returns the matrix `M` and the array `cell_coords`.

It starts by creating the matrix `M` the shape of `G`, filled with zeros and initiate a counter `count`. It then parses `M` and for each fluid cell (`G[c, r] != 0`), it sets the value of `M[r, c]` to the value of the counter and increment the counter by 1. All the other values are set to -1.

Then it create an array `cell_coords` of size $(M.max() + 1, 2)$ and sets the value of the counter back to zero. Then it parses the matrix `M` and for each cell which value is not -1, it will store the coordinates of that cell in the index count of the array `cell_coords` and increment the counter.

7.3.7 `build_a()`

The function `build_a()` takes no input, it uses the imported data from `parameters.py`.

The function returns the matrix `A`.

It starts by creating an array `A` of size $(N_{\text{fluid}}, N_{\text{fluid}})$, filled with zeros. It then parses the array `cell_coords` and for each inlet and fluid cell (the outlet is not counted there), it counts the neighbors, stores their coordinates, set the value of the diagonal depending on the neighbors and sets each neighbor associated cell to 1. The diagonal cells corresponding to the outlet are all set to 1.

7.3.8 `build_b()`

The function `build_b()` takes no input, it uses the imported data from `parameters.py`.

It returns the array `b`.

It starts by creating an array `b` of size `(cell_coords.shape[0], 1)`, filled with zeros.

It then parses the array. For each cell corresponding to the inlet, it sets the value to $-v_x * h$ and for each value corresponding to the outlet, it sets the value to `phi_ref`. The rest of the array is not modified.

7.3.9 `build_phi()`

The function `build_phi()` takes no input, it uses the imported data from `parameters.py`.

This function is the slowest one since it compute the solution of $Ax = b$. For that it uses the function `linalg.solve()` from `numpy`. Then it creates an empty matrix `phi` and fills it with `numpy.nan`. It then parses `x` and sets the value of each `phi` cell corresponding to the `x` value. The other cells are not modified.

Finally, it saves the values in a **dat** file.

7.3.10 `build_gradient()`

The function `build_gradient()` takes no input, it uses the imported data from `parameters.py`.

This function is rather big, but it is only a variation of the `gradient` function from `numpy`. The function `numpy.gradient` does not work the way I want with the `numpy.nan` values so I made my own.

It starts by loading the **dat** file containing the values of `phi` and creates 2 empty matrices `grad_y` and `grad_norm` the same shape as `G` and fills them with `numpy.nan`. It then parse the `cell_coords` array and sets the values of `grad_y` and `grad_norm` following the centered difference with step $2h$ mentioned in the scope statement. I then found that the function `numpy.gradient` uses the same method. However, since the borders of the matrix `phi` is filled with `numpy.nan` values, the gradient is not computed at the outer edge of the domain.

To compute there values, I applied the impermeable wall condition specified in the section 5.1 of the scope statement. We consider the `numpy.nan` cell with coordinates `[i - 1, j]` as a fictious cell with the value of the cell `[i, j]`. This rule applies to the cells to the left and to the right for `grad_y` and to the top and to the right for `grad_norm`.

It then computes the norm of the gradient (used later to plot the normalized gradient).

Finally, it saves the 3 matrices in **dat** files.

7.3.11 `build_pressure()`

The function `build_pressure()` takes no input, it uses the imported data from `parameters.py`.

The function starts by loading the **dat** files containing the values of pressure.

It then creates an empty matrix `pressure` and fills it with `numpy.nan`. Then, it computes the Bernoulli's constant \mathcal{H} using the initial values.

It then parses the array `cell_coords`, sets the inlet cells of the pressure matrix to `pressure_init`, computes and sets the values for each cell.

Finally, it saves the matrix in a **dat** file.

7.4 `plot.py`

The class is used to plot every piece of data.

7.4.1 `plot_graphs()`

The function `plot_graphs()` takes 2 input arguments.

- `display`: `str`
- `data`: `dict`

This subroutine creates the figure, shows the matrix G (without interpolation) and disables the axes as they are not needed in that kind of plot. Then, depending on the `display` value, it will call other functions to plot the graphs and save them as **pdf** files.

7.4.2 Data specific plotting functions

Each one of the 4 next functions take 1 input argument.

- data: `dict`

They return the plotted graph.

The 4 functions work the same way, they set the name of the graph and plot the desired data.

Part III

Optimization

In order to reduce the memory usage of the program, I tried to limit as much as possible the size allocated to each array by setting the datatype at declaration.

At runtime, the memory usage stays relatively low since the program does not have to store anything big. However, if you set a domain ludicrously big, the program will crash telling you there is not enough memory to store the data since the matrix A increases exponentially with the size of the domain.

Concerning the CPU, I tried running the program on a really big domain to see how it reacts. The CPU usage rises to the maximum on one core (since python natively uses only one core) until the end of the `linalg.solve` computation. Aside from this function, the CPU usage stays low at runtime.

Part IV

Outcome examples

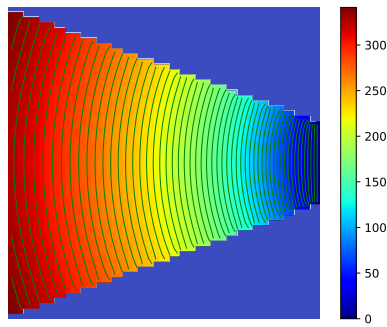
8 Input data

The program needs some data to run. For the example, I found that a box of size 60×60 is good enough. Obviously we can use a bigger box, but the computing time increases exponentially as we increase the box size. A smaller box can also produce acceptable results. I also chose to use cells of size 1 ($h = 1$). The geometry used is the shrinkage geometry.

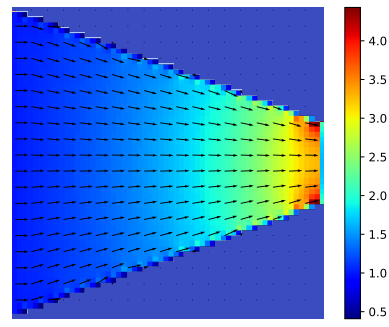
Concerning the fluid, here are the data I choosed :

- $\rho = 1$, relative density of water
- $v_x = 1 \text{ m s}^{-1}$, typical speed in a watering pipe
- $\phi_{\text{ref}} = 0$
- $p_{\text{init}} = 5 \text{ bar}$, typical pressure in a watering pipe

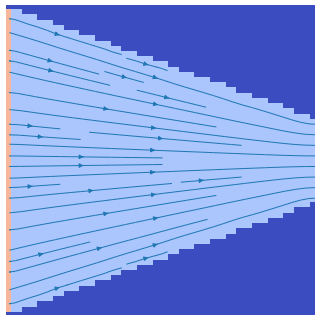
9 Output



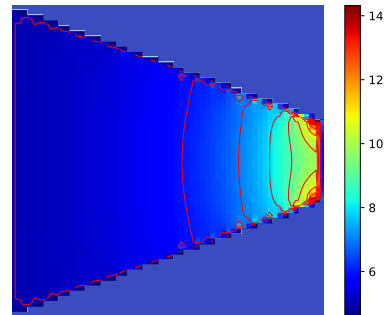
(a) Velocity potential field



(b) Velocity field



(c) Streamlines



(d) Pressure field

Figure 1: Results for $N_x = N_y = 60$ in a shrinkage geometry

10 Interpretation

10.1 Expectations

The expected behavior for this kind of geometry (bottleneck) is that the fluid (water in this case) will bump into the boundaries of the domain and converge toward the center as it crosses the domain. Regarding the pressure of the fluid, since the number of fluid particles is constant throughout the simulation, we should observe a rise of pressure as the domain narrows.

10.2 Observations

Firstly, looking at [Figure 1b](#) and [Figure 1c](#), the water crosses the domain correctly and converges toward the center of the domain. Despite some accuracy problems (the velocity in the edges seems off), I can say that the simulation meets the expected behavior. It is mandatory to normalize these vectors, else the output graph is not readable. However, the bigger the domain, the bigger the number of vectors. As the size of the domain is 60×60 , there are quite a lot of vectors to show. With the default display, there are too much arrows on the graph and they are too small to analyze anything, I added a small algorithm using interpolation to reduce the number of arrows while keeping the data as accurate as possible. Since the problem does not appear without the interpolation, I think it comes from there. I will try to correct it in the beta. In the alpha version, the streamlines are computed using the velocity, there should not be any problem with them, they show the path of the fluid and it seems to be the right one.

On the [Figure 1d](#), it is clear that the pressure increases as the fluid travels the domain. It is once again the expected behavior.

10.3 Pressure verifications

I ran some tests regarding the pressure field when changing the relative density of the fluid and the results are satisfying.

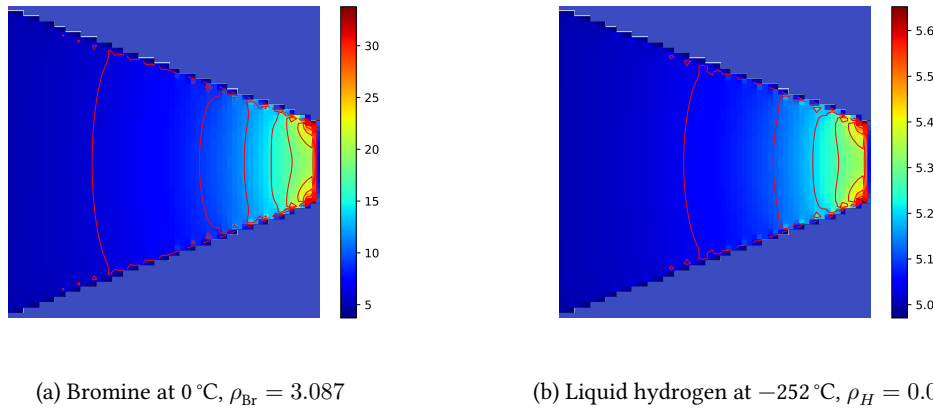


Figure 2: Comparison of the pressure field for bromine and liquid hydrogen

According to the BERNOULLI's Theorem, when you have 2 different fluids in the same conditions, the heavier the fluid, the higher the pressure. In this example, the bromine is heavier than the liquid hydrogen, and the graphs are showing that property really well. With the same initial pressure and the same conditions, the pressure of the bromine rises higher than the one of the liquid hydrogen. The isobaric lines are also slightly different.

References

- [1] Vincent Ballenegger. *Planar potential flows*. 2020. [2] Wikipedia. *Bernoulli's Principle*. Apr. 11, 2020. URL: https://en.wikipedia.org/wiki/Bernoulli%27s_principle.