

Free Surface with LBM D3Q19

Group 6

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0.1 Compilation.

Requirements: UNIX based OS.

Use **make** command to compile:

```
>> make
```

There is debug mode available. This mode includes additional checks for velocity and density, computes total mass in the domain and checks that there is no FLUID cell that is neighboring with a GAS cell directly. Also it calculates a time spent on the following parts: streaming, collision, flag updating and boundary treatment.

```
>> make debug
```

0.2 Usage

To run the program use

```
>> ./lbsim <exampleName> <number of threads>
```

For each example it is required to have `<exampleName>.dat` and `<exampleName>.pgm` files. They have to be placed in the **examples** directory.

`<number of threads>` specifies how many threads will be used by OpenMP.

Output is vtk-files and they can be found in the **vtk-output** directory (if there is no such directory it will be created).

0.2.1 Description of input files

We added a few new fields to the `input.dat` file:

Parameter	Description
radius	Radius of a droplet
exchange_factor	Factor to increase speed of a mass exchange (mainly to get more beautiful videos)
forces_x	Gravity force in the x direction
forces_y	Gravity force in the y direction
forces_z	Gravity force in the z direction

0.3 What we have done

New cell types First of all we added 2 new cell types: FLUID and INTERFACE. Here is table representing cell types, which should be used in `<exampleName>.pgm` file:

Cell type	Value	Cell type	Value
FLUID	0	PRESSURE_IN	5
INTERFACE	1	OBSTACLE	6
MOVING_WALL	2	FREESLIP	7
INFLOW	3	NOSLIP	8
OUTFLOW	4	GAS	9

Test functions In the `checks.c` one can find three check functions:

1. `check_in_rank` checks that every FLUID cell has a density between 0.9 and 1.1 and norm of the velocity vector is less then `sqrt(3) * C_S`;
2. `check_flags` checks that no FLUID cell has a GAS neighbor;
3. `check_mass` compute total mass in the domain (FLUID and INTERFACE cells).

This checks will be activated in debug mode and be executed every time when we write output file.

External forces influence In collision step additional term for external forces was added.

Related function in `collision.c`: `computeExternal(int i, float density, float * extForces)`, where `extForces` is a force vector specified in input file.

Mass and fluid fraction fields For every cell we store 2 additional float values: mass and fluid fraction. We update mass field during streaming step. Fluid fraction field is updated during collide step.

To be able get more fast we introduced `exchange_factor` in an input file. By this factor we increase mass exchange between cells.

DF from gas cells In the streaming step was added reconstruction from GAS cells. Streaming works normally for all FLUID cells and for INTERFACE cells we are checking whether a neighbor is GAS. If yes, we are using outflow boundary formula in this direction.

Flag update From the mass fraction for each cell we determine whether that cell emptied or filled. We track these cells using two arrays, `emptiedCells` and `filledCells`. We perform the filling and emptying in separate phases, `performFill()` and `performEmpty()`. These phases are asymmetric in order

to handle the case of a cell filling adjacent to a cell emptying. If this happens, the ‘filling’ operation cancels the neighbor’s ‘emptying’ operation and the neighboring cell is stricken from `emptiedCells`.

Both phases maintain the loop invariant of a contiguous interface layer. When an `INTERFACE` cell is converted to `FLUID`, all `GAS` neighbors are converted to `INTERFACE`. Correspondingly, when an `INTERFACE` is converted to `GAS`, all `FLUID` cells are converted to `INTERFACE`.

Parallelization with OpenMP We have added `pragma omp parallel` for streaming, collision, treatment boundaries and update flags.

Other optimizations At the beginning our main bottleneck was `computeFeq` function which took approximately 42% percent of the whole time. We splitted the Q-loop in this function in a way that compiler was able to vectorize it, then we got rid of all divisions and replaced them by corresponding multiplications. Then we changed types of all double fields to float. It gave us approximately 2x speedup.