

Documentation

This package contains several python codes to run simulations of rigid bodies made out of rigidly connected *blobs* (aka rigid multiblobs), confined above a single wall (floor). These codes can compute the mobility of complex shape objects, solve mobility or resistance problems for suspensions of many bodies or run deterministic or stochastic dynamic simulations.

We explain in the next sections how to use the package. For the theory consult the references:

1. **Brownian Dynamics of Confined Rigid Bodies**, S. Delong, F. Balboa Usabiaga and A. Donev. The Journal of Chemical Physics, **143**, 144107 (2015). [DOI](#) [arXiv](#)
2. **Hydrodynamics of suspensions of passive and active rigid particles: a rigid multiblob approach** F. Balboa Usabiaga, B. Kallemov, B. Delmotte, A. Pal Singh Bhalla, B. E. Griffith, and A. Donev, Communications in Applied Mathematics and Computational Science, **11**, 217 (2016). [DOI](#) [arXiv](#)
3. **Brownian dynamics of condined suspensions of active micro-rollers**, F. Balboa Usabiaga, B. Delmotte and A. Donev, The Journal of Chemical Physics, **146**, 134104 (2017). [DOI](#) [arXiv](#)
4. **Large Scale Brownian Dynamics of Confined Suspensions of Rigid Particles**, B. Sprinkle, F. Balboa Usabiaga, N. Patankar and A. Donev. *In preparation*, 2017.

Note: The key blob-blob translational mobility in the presence of a wall is computed using the Rotne-Prager-Blake tensor in the appendices B and C of: **Simulation of hydrodynamically interacting particles near a no-slip boundary**, James Swan and John Brady, Phys. Fluids **19**, 113306 (2007)[DOI](#). We modify the mobility to allow overlaps between blobs and between blobs and the wall, see Ref. [3].

1. Prepare the package

The codes are implemented in python (version 2.x) and it is not necessary to compile the package to use it.

However, we provide alternative implementations in *C++* (through the Boost Python library) and *pycuda* for some of the most computationally expensive functions. You can skip to section 2 but come back if you want to take fully advantage of this package.

1.1 Prepare the mobility functions

The codes use functions to compute the blob mobility matrix \mathbf{M} and the matrix vector product $\mathbf{M}\mathbf{f}$. For some functions we provide a *C++* implementation which can be around five times faster than the python version. We also provide *pycuda* implementations which, for large systems, can be orders of magnitude faster. To use the *C++* implementation move to the directory `mobility/` and compile `mobility_ext.cc` to a `.so` file using the Makefile provided (which you will need to modify slightly to reflect your Python version, etc.).

To use the *pycuda* implementation all you need is *pycuda* and a GPU compatible with CUDA; you don't need to compile any additional file in this package.

1.2 Blob-blob forces

In dynamical simulations it is possible to include blob-blob interactions to, for example, simulate a colloid suspension with a given steric repulsion. Again, we provide versions in *python*, *C++* and *pycuda*. To use the *C++* version move to the directory `multi_bodies/` and compile `forces_ext.cc` to a `.so` file using the Makefile provided (which you will need to modify slightly to reflect your Python version, etc.).

To use the *pycuda* implementation all you need is *pycuda* and a GPU compatible with CUDA; you don't need to compile any additional file in this package.

2. Rigid bodies configuration

We use a vector (3 numbers) and a quaternion (4 numbers) to represent the location and orientation of each body, see Ref. 1 for details. This information is saved by the code in the `*.clones` files, with format:

```
number_of_rigid_bodies
vector_location_body_0 quaternion_body_0
vector_location_body_1 quaternion_body_1
.
.
.
```

For example, the file `multi_bodies/Structures/boomerang_N_15.clones` represents the configuration of a single body with location (0, 0, 10) and orientation given by the quaternion (0.5, 0.5, 0.5, 0.5).

The coordinates of the blobs forming a rigid body in the default configuration (location (0, 0, 0) and quaternion (1, 0, 0, 0)) are given to the codes through `*.vertex` files. The format of these files is:

```

number_of_blobs_in_rigid_body
vector_location_blob_0
vector_location_blob_1
.
.
.

```

For example, the file `multi_bodies/Structures/boomerang_N_15.vertex` gives the structure of a boomerang-like particle formed by 15 blobs.

3. Active slip

The blobs can have an active slip as described in the Ref. 2, therefore, we can simulate the dynamics of active bodies like bacteria or self-propelled colloids.

For each structure the user can include the name of a slip file in the input file (in the line `strucutre` see below). In this case the code reads the slip at the beginning of the simulation. This active slip is assumed to be time independent and fixed in the reference body frame and will be converted to the lab frame as the body moves. You can see an example in `multi_bodies/examples/squirmer/`. If the slip file is not given the slip is set to zero.

Note that in practice the slip probably depends on the position of the particles relative to the boundary and each other, and the user can provide a function to compute the slip (this time in the lab frame) at runtime, see Section 5.3 for details.

4. Run static simulations

We start explaining how to compute the mobility of a rigid body close to a wall. First, move to the directory `multi_bodies/` and inspect the input file `inputfile_body_mobility.dat`:

```

# Select problem to solve
scheme                                body_mobility

# Select implementation to compute the blobs mobility
mobility_blobs_implementation        python

# Set fluid viscosity (eta) and blob radius
eta                                  1.0
blob_radius                          0.25

```

```

# Set output name
output_name                                data/run.body_mobility

# Load rigid bodies configuration, provide
# *.vertex and *.clones files
structure  Structures/boomerang_N_15.vertex Structures/boomerang_N_15.clones

```

Now, to run the code, use

```
python multi_bodies_utilities.py --input-file inputfile_body_mobility.dat
```

The code writes the results in files with the prefix `data/run.body_mobility`. You can inspect the file `*.body_mobility.dat` to see the 6x6 rigid body mobility.

List of input file options:

- **scheme:** (string) Options: `mobility`, `resistance` and `body_mobility`. Select the problem to solve. `mobility` computes the velocities of a suspension of rigid bodies subject to external forces and torques (see below). `resistance` computes the forces and torques on rigid bodies moving with given velocities (see below). `body_mobility` computes the mobility **matrix** of one rigid body as in the above example.
- **mobility_blobs_implementation:** Options: `python`, `C++`, `python_no_wall` and `C++_no_wall`. It selects which implementation is used to compute the blob mobility matrix **M**. See section 1 to use the C++ versions. The options ended with `_no_wall` use the Rotne-Prager tensor, the others include wall corrections (Rotner-Prager-Blake tensor) as explained in the introduction.
- **mobility_vector_prod_implementation:** Options: `python`, `C++`, `pycuda`, `python_no_wall`, `C++_no_wall` and `pycuda_no_wall`. It selects the implementation to compute the matrix vector product **Mf**. See section 1 to use the C++ or pycuda implementations. The options ended with `_no_wall` use the Rotne-Prager tensor, the others include wall corrections (Rotner-Prager-Blake tensor) as explained in the introduction.
- **eta:** (float) the fluid viscosity.
- **blob_radius:** (float) the hydrodynamic radius of the blobs.
- **solver_tolerance:** (float) the tolerance for the iterative mobility solver.
- **output_name:** (string) the prefix used to save the output files.

- **velocity_file**: (string) name of a file with the velocities of the rigid bodies used in the **resistance** problem. The format of the file is one line per body and six floats per line corresponding to linear (first three) and angular velocities (last three).
- **force_file**: (string) name of a file with the forces and torques used in the **mobility** problem. The format of the file is one line per body and six floats per line corresponding to force (first three) and torque (last three). If no file is given the code compute the forces on the bodies as explained in the section 5.2.
- **structure**: (two or three strings) name of the vertex, clones and optionally slip files with the rigid bodies configuration and the active slip, see sections 2 and 3. To simulate bodies with different shapes add to the input file one **structure** option per each kind of body and give their **vertex** and **clones** files, see multibodies/inputfile.dat for an example. If the slip file is not given the active slip is set to zero for that structure (i.e. it will be a passive particle).
- **plot_velocity_field**: (x_0 x_1 N_x y_0 y_1 N_y z_0 z_1 N_z) if the code is run with this options and the schemes **mobility** or **resistance** the code plots the velocity field of the fluid to a **vtk** file. The velocity field is plotted in a rectangular box with the lower corner located at (x_0 , y_0 , z_0), the upper corner located at (x_1 , y_1 , z_1) and using a grid of dimensions (N_x , N_y , N_z). The **vtk** file can be postprocessed with external software like **VisIt** from the Lawrence Livermore National Laboratory or **ParaView** from Sandia National Laboratories to generate an image of the velocity field. See figure 1 as an example.
- **tracer_radius**: (float (default 0)) effective radius of the nodes where the fluid velocity field is computed with the option **plot_velocity_field**. The default value, zero, computes the pointwise fluid velocity, a larger value average the fluid velocity over a region of radius **tracer_radius**.

The output files are:

- **.inputfile**: It is a copy of the input file to better keep track of input options.
- **.body_mobility.dat**: the 6x6 rigid body mobility computed with the option **scheme body_mobility**. When this mobility is apply to the vector (force, torque) generates the vector (velocity, angular_velocity).
- **.force.dat**: forces and torques on the bodies computed with the option **scheme resistance**. The format of the file is one line per body and six floats per line corresponding to force (first three) and torque (last three).

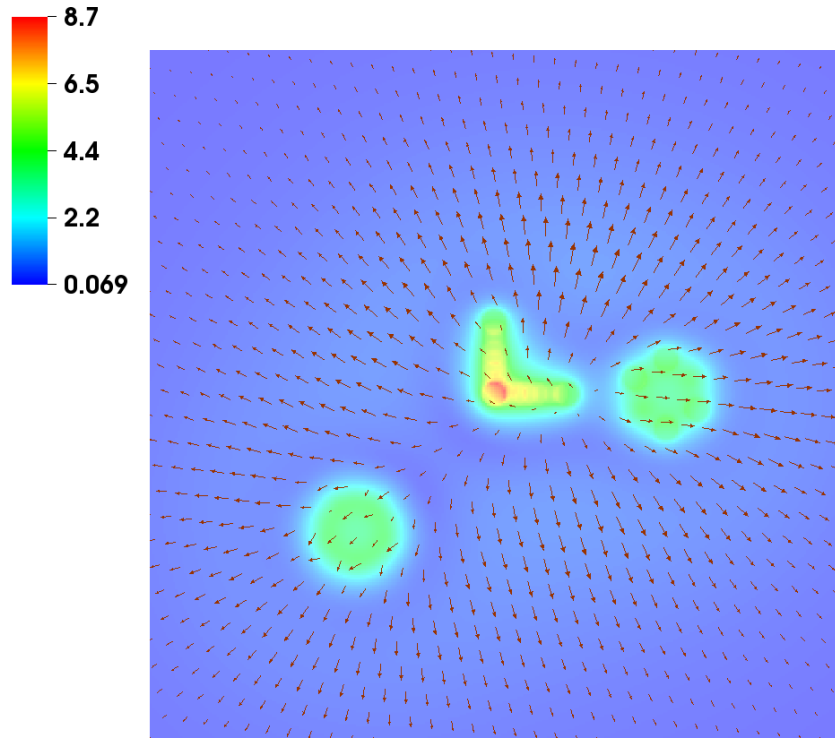


Figure 1: Velocity field around two spheres and a boomerang shaped colloid falling towards a bottom wall. Velocity field computed with the option `plot_velocity_field` and image generated with the software VisIt.

- `.velocities.dat`: velocities of the rigid bodies computed with the option `scheme mobility`. The format of the file is one line per body and six floats per line corresponding to linear (first three) and angular velocities (last three).

5. Run dynamic simulations

5.1 Rigid multiblob simulations

We have two python codes to run dynamic simulations. The first, in the directory `boomerang/`, allows to run stochastic Brownian simulations for a single body. See the instruction in `boomerang/README.md`. Here, we explain how to use the other code which allows to run deterministic and stochastic simulations for many bodies.

First, move to the directory `multi_bodies/` and inspect the input file `inputfile_dynamic.dat`:

```
# Select integrator
scheme                                deterministic_adams_bashforth

# Select implementation to compute M and M*f
mobility_blobs_implementation        python
mobility_vector_prod_implementation  python

# Select implementation to compute the blob-blob interactions
blob_blob_force_implementation       python

# Set time step, number of steps and save frequency
dt                                    0.1
n_steps                               10
n_save                                1

# Set fluid viscosity (eta), gravity (g) and blob radius
eta                                    1.0
g                                      1.0
blob_radius                           0.25

# Set parameters for the blob-blob interaction
repulsion_strength                    1.0
debye_length                          1.0

# Set interaction with the wall
```

```

repulsion_strength_wall          1.0
debye_length_wall                1.0

# Set output name
output_name                      data/run

# Load rigid bodies configuration, provide
# *.vertex and *.clones files
structure Structures/boomerang_N_15.vertex Structures/boomerang_N_15.clones
structure Structures/shell_N_12_Rg_1.vertex Structures/shell_N_12_Rg_1.clones

```

With this input we can run a simulation with three rigid bodies, one with a boomerang shape and two with a spherical shape; see structures given to the options `structure`. To run the simulation use

```
python multi_bodies.py --input-file inputfile_dynamic.dat
```

Now, you can inspect the output, `ls data/run.*`. The output files are:

- `.bodies_info`: it contains the number of bodies and blobs in the simulation. Also how many types of bodies (i.e. different shapes) and how many bodies of each type.
- `.clones`: For each time step saved and each structure type the code saves a file with the location and orientation of the rigid bodies. The name format is (output_name + structure_name + time_step + .clones) The format of the files is the same that in the input .clones files.
- `.info`: It records the number of invalid configurations generated during the simulation (number of times a blob cross the wall), and the total number of iterations to solve the mobility problems (GMRES iterations) and to generate the Brownian noise (Lanczos iterations).
- `.inputfile`: a copy of the input file.
- `.random_state`: it saves the state of the random generator at the start of the simulation. It can be used to run a simulation with the same random numbers.
- `.time`: the wall-clock time elapsed during the simulation (in seconds).

List of options for the input file:

- `scheme` (string). There are several schemes implemented to run deterministic and stochastic simulations. See reference [4] for a discussion. The input file should select one of the following ones

Name	Solver type	Notes
deterministic_forward_euler	Iterative	first order accuracy
deterministic_forward_euler_ dense_algebra	Direct solve	cost $O(\text{number_of_blobs}^3)$
deterministic_adams_bashforth	Iterative	second order accuracy
stochastic_first_order_RFD	Iterative	it uses three mobility solves and one Lanczos call per step
stochastic_adams_bashforth	Iterative	primarily used for microrollers
stochastic_first_order_RFD_ dense_algebra	Direct solve	cost $O(\text{number_of_blobs}^3)$ but faster for small systems
stochastic_traction_EM	Iterative	first order. two mobility solves and one Lanczos call per step.
Fixman	Direct solve	cost $O(\text{number_of_blobs}^3)$
stochastic_Slip_Trapz	Iterative	second order accurate deterministically and weakly first order accurate stochastically. It uses three mobility solves and one Lanczos call per step
stochastic_traction_AB	Iterative	(Not tested) Cost and accuracy like stochastic_traction_AB
stochastic_Slip_Mid	Iterative	(Not tested) Accuracy like in stochastic_traction_AB. It uses three mobility solves and two Lanczos calls per step
stochastic_Slip_Mid_DLA	Direct solve	cost $O(\text{number_of_blobs}^3)$ (dense LA version of Slip_Mid)

We recommend `deterministic_adams_bashforth` for deterministic simulations since it costs no more than forward Euler but is more accurate. For Brownian simulation with many rigid multiblobs we recommend the trapezoidal slip scheme `stochastic_Slip_Trapz` [4]. If time step is restricted by stability rather than accuracy you can try the traction

Euler-Maruyama scheme `stochastic_traction_EM` (cheaper but less accurate) [4]. For minimally-resolved BD as with micro-rollers (see [3]), we recommend `stochastic_adams_bashforth`, which adds Brownian motion to `deterministic_adams_bashforth` using Lanczos+RFD [3] (and is more expensive). For small systems using dense linear algebra may be faster and then the Fixman scheme can be used.

With schemes that use iterative methods you can print the residual of GMRES and the Lanczos algorithm to the standard output using the flag `--print-residual`.

- `mobility_blobs_implementation`: Options: `python`, `C++`, `python_no_wall` and `C++_no_wall`. This option indicates which implementation is used to compute the blob mobility matrix \mathbf{M} . See section 1 to use the C++ version. The options ended with `_no_wall` use the Rotne-Prager tensor, the others include wall corrections (Rotner-Prager-Blake tensor) as explained in the introduction.
- `mobility_vector_prod_implementation`: Options: `python`, `C++`, `pycuda`, `python_no_wall`, `C++_no_wall` and `pycuda_no_wall`. This option select the implementation to compute the matrix vector product $\mathbf{M}\mathbf{f}$. See section 1 to use the C++ or pycuda implementation. The options ended with `_no_wall` use the Rotne-Prager tensor, the others include wall corrections (Rotner-Prager-Blake tensor) as explained in the introduction.
- `blob_blob_force_implementation`: Options: `None`, `python`, `C++` and `pycuda`. Select the implementation to compute the blob-blob interactions between all pairs of blobs. If `None` is selected the code does not compute blob-blob interactions. The cost of this function scales like $(\text{number_of_blobs})^2$, just like the product $\mathbf{M}\mathbf{f}$. The default soft repulsion is described under `repulsion_strength` below. See Section 5.3 for details on how to change this.
- `body_body_force_torque_implementation`: Options: `None` and `python`. Select the implementation to compute the body-body interactions between all pairs of bodies. This function provides an alternative way to compute force between bodies without iterating over all the blobs in the system. Note that this force will be added to the forces coming from other terms like blob-blob interactions. If `None` is selected the code does not compute body-body interactions directly but it can compute blob-blob interactions which lead to effective body-body interactions. The cost of this function scales like $(\text{number_of_bodies})^2$. The default soft repulsion is described under `repulsion_strength` below. See Section 5.3 for more details on how to implement your own force law in python.
- `eta`: (float) the fluid viscosity.
- `blob_radius`: (float) the hydrodynamic radius of the blobs.

- **solver_tolerance**: (float) the relative tolerance for the iterative mobility solver.
- **output_name**: (string) the prefix used to save the output files.
- **dt**: (float) time step length to advance the simulation.
- **n_steps**: (int) number of time steps.
- **initial_step**: (int (default 0)) Use this option to restart a simulation. If **initial_step** > 0 the code will run from time step **initial_step** to **n_steps**. Also, the code will try to load **.clones** files with the name (output_name + structure_name + initial_step + .clones).
- **n_save**: (int) save the bodies configuration every **n_save** steps.
- **repulsion_strength**: (float) the blobs interact through a soft potential of the form $(U = \text{eps} + \text{eps} * (d-r)/b)$ if $r < d$ and $U = \text{eps} * \exp(-(r-d)/b)$ if $r \geq d$ where r is the distance between blobs, b is the characteristic length, **eps** is the strength and $d=2*a$ is twice the blob radius. This is the strength of the potential, **eps** in the above expression (see section 5.3 to modify blobs interactions).
- **debye_length**: (float) the characteristic length of the blob-blob soft potential, b in the expression given above under **repulsion_strength**. (see section 5.3 to modify blobs interactions).
- **repulsion_strength_wall**: (float) the blobs interact with the wall with a soft potential. The potential is $(U = \text{eps} + \text{eps} * (d-r)/b)$ if $r < d$ and $U = \text{eps} * \exp(-(r-d)/b)$ if $r \geq d$ where h is the distance between the wall and the particle, $d=a$ is the blob radius, b is the characteristic potential length and **eps** is the strength. This is the strength of the Yukawa potential, **eps** in the above formula (see section 5.3 to modify blobs interactions).
- **debye_length_wall**: (float) the characteristic length of the Yukawa blob-wall potential, b in the expression given above under **repulsion_strength_wall** (see section 5.3 to modify blobs interactions).
- **random_state**: (string) name of a file with the state of the random generator from a previous simulation. It can be used to generate the same random numbers in different simulations.
- **seed**: (unsigned int) seed for the random number generator. It is not used if a **random_state** is defined in the input file. If neither **seed** nor **random_state** are defined the code will automatically initialized to a pseudorandom state (see documentation for `numpy.random.RandomState`).
- **structure**: (two or three strings) name of the vertex, clones and optionally slip files with the rigid bodies configuration and the active slip, see sections 2 and 3. To simulate bodies with different shapes add to the input file

one **structure** option per each kind of body and give their **vertex** and **clones** files, see `multibodies/inputfile.dat` for an example. If the slip file is not given the active slip is set to zero for that structure (i.e. it will be a passive particle).

- **save_clones:** (string (default `one_file_per_step`)) options `_one_file_per_step_` and `onefile_`. With the option `_one_file_per_step_` the clones configuration are saved in one file per kind of structure and per time step as explained above. With the option `_one_file_` the code saves one file per kind of structure with the configurations of all the time steps; configurations of different time steps are separated by a line with the number of rigid bodies.
- **periodic_length:** (three floats (default 0 0 0)) length of the unit cell along the x, y and z directions. If the length of the unit cell along the x or y directions is larger than zero the code uses Pseudo Periodic Boundary Conditions (PPBC) along that axis, otherwise the system is considered infinite in that direction. With PPBC, particles forces are computed using the minimum image convention as with standard periodic boundary conditions. Hydrodynamic interactions are computed between particles in the unit cell and the first neighbor cells along the pseudo-periodic axis. PPBC along the z axis are not supported.

5.2 Rollers simulations

We can also use the code `multi_bodies.py` to run simulations of bodies discretized with a single blob interacting hydrodynamically with a grand-mobility matrix that includes couplings between the linear and angular velocities, see Ref. [3] for a detailed description. To run a simulation use:

```
python multi_bodies_utilities.py --input-file inputfile_body_mobility.dat
```

The input file options are the same than for a rigid multiblob simulation (see section 5.1 and the file `inputfile_body_mobility.dat`) except for the following differences:

- **scheme:** (string) Options: `deterministic_forward_euler_rollers`, `stochastic_first_order_rollers`, `deterministic_adams_bashforth_rollers`, `stochastic_adams_bashforth_rollers`, `stochastic_mid_point_rollers`, `stochastic_trapezoidal_rollers`. We provide several schemes for deterministic and stochastic simulations, but we recommend using `'stochastic_adams_bashforth_rollers'`.
- **structure:** (two strings) name of the vertex and clones files with the rigid bodies configuration, see section 2. However, this code only accepts bodies discretized with a single blob so the vertex file is trivial, see file `multi_bodies/Structures/blob.vertex`.

- **free_kinematics**: (string (default True)) if **free_kinematics** is True the angular velocity of the blobs is not fixed and each blob is subject to a torque $T=8\pi\eta a^3\omega_{\text{one_roller}}$ (see below). If **free_kinematics** is False all the blobs rotate with a prescribed angular velocity given with the option **omega_one_roller** but they are free to translate. The torque acting on the blobs is a Lagrangian multiplier that enforces the prescribed angular velocity.
- **omega_one_roller**: (three floats (default 0 0 0)) prescribed angular velocity of the blobs if the option **free_kinematics** is set to False. If **free_kinematics** is set to True the blobs are subject to a constant torque $T=8\pi\eta a^3\omega_{\text{one_roller}}$.
- **domain**: (string) Options: **single_wall** and **no_wall**. With the option **single_wall** (default) the mobilities include wall corrections, i.e. the code uses the Rotne-Prager-Blake tensor as explained in the introduction. With the option **no_wall** the mobilities do not include wall corrections, the code uses the Rotne-Prager mobilities.

5.3 Modify the codes

We provide default implementations to calculate the slip on the rigid bodies and the interactions between blobs and between bodies as explained above. However, the user can override the default implementation with their own functions. We explain here how to do it. First, we provide two alternatives to compute the interactions between bodies. A direct method that uses the locations and orientations of the bodies and a indirect form that compute the forces between all the blobs forming the rigid bodies and then it uses those forces to compute the forces and torques on the bodies as explained in the Refs 1 and 2. The second approach can be more expensive, since in general `number_of_blobs >> number_of_bodies` but it can be used with bodies with arbitrary shapes, while the first method it is hard to generalize to non-spherical bodies. Note that the code can use both methods at the same time. You can override the following functions:

- blob-blob interactions: to override the *python* implementation create your own function `blob_blob_force` in the file `user_defined_functions.py` as we show in the example in `multi_bodies/examples/boomerang_suspension/`. To override the *pycuda* version create your own function `blob_blob_force` in `forces_pycuda_user_defined.py` as we show also in `multi_bodies/examples/boomerang_suspension/`. `blob_blob_force` in the file `multi_bodies/forces_pycuda.py` However, to modify the *C++* implementation you need to edit the function `blobBlobForce` in the file `multi_bodies/forces_ext.cc` and recompile the *C++* code, note that this not override the default implementation but it modifies it.

- blob-wall interaction: to override the *python* implementation create your own function `blob_external_force` in the file `user_defined_functions.py` as we show in the example in `multi_bodies/examples/boomerang_suspension/`. There are not *C++* or *pycuda* versions of this function since it is not an expensive operation.
- body-body interactions: to override the *python* implementation create your own function `body_body_force_torque` as we show in the example in `multi_bodies/examples/boomerang_suspension/`.
- body external forces: to override the one-body forces, for example gravity or interactions with the wall, create your own function `bodies_external_force_torque` in the file `user_defined_functions.py`.
- active slip: The code assigns a constant slip to each body if a slip file is passed in the inputfile (see section 3). However, you can use more general slips like functions that depend on time, distance to the wall etc. To override the default python implementation create your own function `set_slip_by_ID` in the file `user_defined_functions.py`. For a simple example see `multi_bodies/examples/pair_active_rods/`.

6. Run Monte Carlo simulations

We have a Markov Chain Monte Carlo code to generate equilibrium configurations. To use this code move to the directory `many_bodyMCMC` and inspect the input file `inputMCMC.dat`. The options are similar to the ones to run dynamic simulations, however, some options like the time step size are not necessary. We have only implemented the subroutines to compute the body-body interactions in *pycuda*, therefore, it is necessary to have a GPU with CUDA capabilities to use this code. To run a simulation use

```
python many_body_MCMC.py inputMCMC.dat
```

The output files are similar to the ones generated with dynamic simulations.

The user can override the default interactions by creating its own functions in the file `potential_pycuda_user_defined.py`. In the folder `many_bodyMCMC/examples/boomerang_suspension/` we show how to override the potentials to simulate a boomerang suspension as in Ref. [4].

7. Software organization

- `doc/`: documentation.
- `body/`: it contains a class to handle a single rigid body.
- `boomerang/`: older stochastic example from [1], see documentation `doc/boomerang.txt`.

- **sphere/**: the folder contains an example to simulate a sphere whose center of mass is displaced from the geometric center (i.e., gravity generates a torque), sedimented near a no-slip wall in the presence of gravity, as described in Section IV.C in [1]. Unlike the boomerang example this code does not use a rigid multiblob model of the sphere but rather uses the best known (semi)analytical approximations to the sphere mobility. See documentation `doc/boomerang.txt`.
- **many_bodyMCMC/**: Markov Chain Monte Carlo code for rigid bodies.
- **mobility/**: it has functions to compute the blob mobility matrix \mathbf{M} and the product $\mathbf{M}\mathbf{f}$ using CPUs or GPUs, see [2].
- **multi_bodies/**: codes to run many-body simulations, based on [3] (minimally-resolved active rollers) and primarily on [4] (general many-particle case).
- **quaternion_integrator/**: it has a small class to handle quaternions and the schemes to integrate the equations of motion, see [1] and [4].
- **stochastic_forcing/**: it contains functions to compute the product $\mathbf{M}^{1/2}\mathbf{z}$ necessary to perform Brownian simulations, see [3] and [4].
- **utils.py**: this file has some general functions that would be useful for general rigid bodies (mostly for analyzing and reading trajectory data and for logging).