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Solvation Gibbs Energy Calculation

To run, **PyGBe** needs two input files: a **config** file (`*.config`) and a **parameters** file (`*.param`), which need to be located in the `./bem_pycuda/input_files` directory. The following shows an example of a run script:

```
python main_asymmetric.py input_files/problem.param input_files/problem.config --asymmetric --chargeForm
```

1. The first argument (`main_asymmetric.py`) specifies the python script the carries out solvation free energy calculations
2. The second argument (`input_files/problem.param`) specifies the relative path to the **param** file location
3. The third argument (`input_files/problem.config`) specifies the relative path to the **config** file location
4. The `--asymmetric` flag activates the asymmetric SLIC boundary condition
5. The `--chargeForm` denotes the type of charge distribution

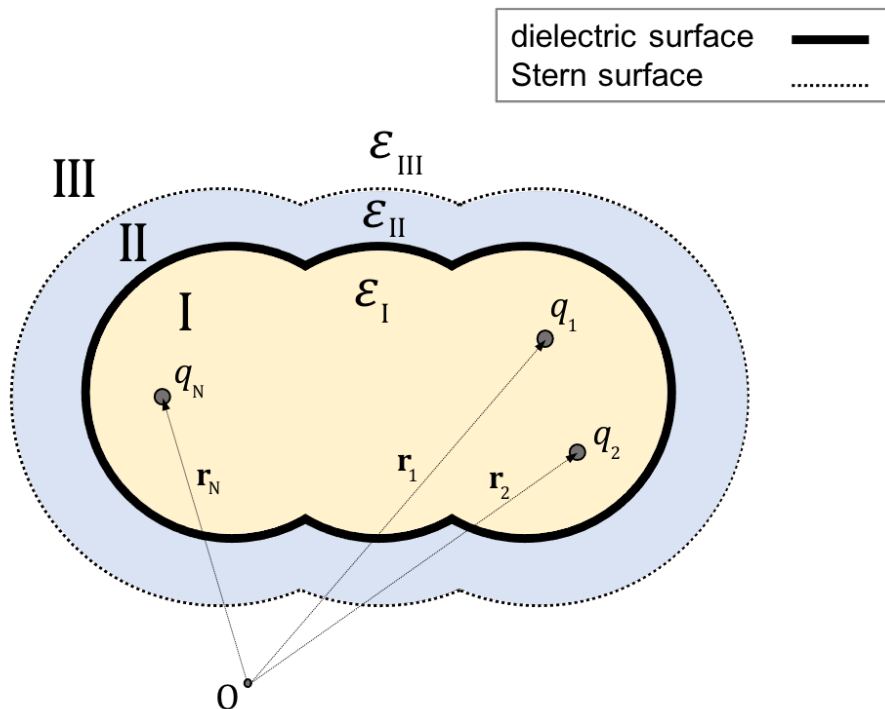
config file

config file specifies the geometry and physical properties of the system. The following is an example of a config file.

FILE	path/to/stern_mesh	stern_layer									
FILE	path/to/diel_mesh	dielectric_interface									

PARAM	LorY	E?	Dielec	kappa	charges?	coulomb?	charge_file	Nparent	parent	Nchild	children
FIELD	1	0	78.36	1e-12	0	0	NA	0	NA	1	0
FIELD	1	0	78.36	1e-12	0	0	NA	1	0	1	1
FIELD	1	1	1	1e-12	1	1	path/to/mol.pqr	1	1	0	NA

which corresponds to the system shown below:



config file consists of two parts, separated with dashes, as seen above.

FILE section

Lines starting with **FILE** indicate that the line specifies the geometry and type of one interface. The first argument after **FILE** determines the relative path to the interface mesh files, `filename.vert` and `filename.face`. `filename.vert` contains the location of the vertices of the discretized triangular elements and `filename.vert` is the connectivity table for the vertices. The `*.face` and `*.vert` files can be generated using Michael Sanner's Molecular Surface program **MSMS**. This link shows an example on how to use **MSMS**. To account for more than one surface (ie. for Stern layers, solvent filled cavities, several proteins), more than one **FILE** line is needed. For instance, the sample **config** file accounts for two surface, described in lines 1 and 2. After `filename`, the user must specify what kind of surface the `filename` describes. It can be:

- `stern_layer`: surface separates Stern layer (region shown in light blue) from solvent
- `dielectric_interface`: surface separates low dielectric (inside protein shown in yellow) and high dielectric (outside protein) regions.
- `internal_cavity`: surface is an internal cavity (a result of `'-all_components'` flag in **MSMS**). This is important to specify

because by default `MSMS` changes the vertex ordering for internal cavities.

- `dirichlet_surface` : surface of specified potential. The value of this potential is read from a text file which has to be specified next to 'dirichlet_surface'.
- `neumann_surface` : surface of specified potential. The value of this potential is read from a text file which has to be specified next to 'neumann_surface'.

The first line in the sample `config` file specifies

1. the relative path to the Stern surface (shown with dots in the schematic) geometry where `path/to` is a folder containing `stern_mesh.face` and `stern_mesh.vert`.
2. the kind of surface which is `stern_layer`.

Similarly, the second line in the sample `config` file specifies this information for the dielectric interface (shown with thick black line in the schematic).



FIELD section

Lines starting with `FIELD` indicate that the line specifies physical parameters of one region. These parameters include:

- `LorY` indicates that the electrostatic potential in the region is
 - 1: Laplace
 - 2: Yukawa

- `E?`
 - 0: don't calculate the energy in this region
 - 1: calculate the energy in this region

Note: if region is surrounded by a *dirichlet* or *neumann surface*, surface energy will be calculated.

- `Dielec` : dielectric constant of the region
- `kappa` : inverse Debye-length
- `charges?`
 - 0: no charges inside the region
 - 1: there are charges in the region
- `coulomb?`
 - 0: don't calculate Coulomb energy in this region
 - 1: calculate Coulomb energy in this region
- `charge_file`
 - `NA` if the region does not contain charges
 - relative path to the `*.pqr` file if the region contains charges. For example, the dielectric region in the schematic above (region `I` shown in ) has a corresponding charge distribution file `mol.pqr` located at `path/to/` directory relative to the config file location.
- `Nparent` : number of 'parent' surfaces (surface containing this region).
 - 0: if the region corresponds to an infinite region, e.g., the solvent region `III` in the schematic
 - 1: if the region is bounded by a surface, e.g., the Stern layer `II` bounded by the Stern surface in the schematic
- `parent` : the region's parent surface mesh index (starting from 0), according to their position in the `FILE` section. Takes `NA` if the region is infinite, otherwise an `int`. For instance in the sample `config` file, the first line in the `FIELD` section describes the properties of solvent as an infinite region (region `III` in the schematic). Because the solvent region is infinite and not bounded by another surface, the `parent` parameter takes the value `NA`. Similarly, the second line in the `FIELD` section corresponds to the Stern layer (shown in ) in the schematic). This region is bounded by the Stern surface, therefore, the `parent` parameter takes the value `0` because the Stern surface is the first surface (hence the `0` index) defined in the `FILE` section.
- `Nchild` : number of child surfaces, i.e., surfaces completely contained in this region. The Stern surface in the schematic is completely contained in the solvent region `III` so the `Nchild` parameter for the solvent region is `1`. Similarly, the dielectric surface is contained in the Stern layer (region `II`) so the `Nchild` parameter for the Stern layer is `1`.
- `children` : mesh file index (indices if the region has more than one child) for the children surfaces completely contained in this region, using the same convention as the `parent` parameter.

Param file

`parameters` file controls the computational parameters of the job. The following shows an example of a `*.param` file

Precision	double
K	7
Nk	13
K_fine	19
threshold	0.5
BSZ	128
restart	500
tolerance	1e-5
max_iter	1000
P	6
eps	1e-12
NCRIT	500
theta	0.5
GPU	1

- **Precision** : double or float. (float not supported yet!).
- **K** : number of Gauss quadrature points per element (1, 3, 4, and 7 are supported).
- **Nk** : number of Gauss quadrature points per triangle edge for semi-analytical integration.
- **threshold** : defines region near singularity where semi-analytical technique is used. if $\sqrt{2 \cdot \text{Area}} / r > \text{threshold}$, integration is done semi-analytically.
- **BSZ** : CUDA block size.
- **restart** : number of iterations for GMRES to do restart.
- **tolerance** : GMRES tolerance.
- **max_iter** : maximum number of GMRES iterations.
- **P** : order of expansion in treecode.
- **eps** : epsilon machine.
- **NCRIT** : maximum number of boundary elements per twig box of tree structure.
- **theta** : multipole acceptance criterion of treecode.
- **GPU**
 - 0: don't use GPU.
 - 1: use GPU.

More details regarding the parameter definitions can be found in this [paper](#).