

NEMATIC ORDERING, ORDERING DEPENDENT ANISOTROPIC ELASTICITY, AND MORPHOLOGY OF FLUID MEMBRANES

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BIBLIOGRAPHY

This code may be used to reproduce most of the results reported in the following articles

1. N. Ramakrishnan, P. B. Sunil. Kumar, John. H. Ipsen, *Phys. Rev. E* 81, 41922 (2010)
 2. N. Ramakrishnan, P. B. Sunil. Kumar, John. H. Ipsen, *Macromolecular Theory and Simulations* 20, 446–450 (2011)
 3. N. Ramakrishnan, John. H. Ipsen, P. B. Sunil. Kumar, P. B. Sunil. Kumar, *Soft Matter* 8, 3058–3061 (2012)
 4. N. Ramakrishnan, P. B. Sunil. Kumar, John. H. Ipsen, *Biophysical Journal* 104, 1–11 (2013)
 5. N. Ramakrishnan, P. B. Sunil. Kumar, Ravi Radhakrishnan, *Physics Reports* 543(1), 1–60, (2014)
 6. N. Ramakrishnan, Ravi Radhakrishnan, *Advances in Planar Lipid Bilayers and Liposomes*, (2015)
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OVERVIEW

Here, I give a brief overview of the physics underlying this code. It aims to evolve the morphology of an elastic surface, of a specified topology, in its corresponding configurational space. **In this version, the code only handles surfaces without boundaries.**

SYSTEM

There are three two components in the model:

1. **A membrane surface** whose configurations is governed by the Canham-Helfrich Hamiltonian

$$H_{el} = \frac{\kappa}{2} \int (2H - C_0)^2 dS$$

In the implementation here, we have set $C_0 = 0$.

2. **A unit in-plane nematic field** \hat{m} , whose lateral organization and texture are governed by Lebwhol-Lasher interactions.

$$H_{NN} = \frac{\epsilon_{LL}}{2} \int (\nabla \theta)^2 dS$$

This is an one constant approximation of the Frank's Free energy. *It should be remembered that the angle between the neighboring nematic field vectors in the calculations in the codes is computed using a parallel transport technique.*

3. **An anisotropic elastic term that couples the orientation of the nematic field to the curvature of the membrane**

$$H_{NC} = \frac{\kappa_{\parallel}}{2} \int (\hat{m} K \hat{m} - C_0^{\parallel})^2 dS + \frac{\kappa_{\perp}}{2} \int (\hat{m}^{\perp} K \hat{m}^{\perp} - C_0^{\perp})^2 dS$$

Terms 1 and 2 corresponding to anisotropic stiffness and curvature along directions parallel and perpendicular to \hat{m} .

Total energy of the system is given by:

$$H_{tot} = H_{el} + H_{NN} + H_{NC}$$

MODEL

The two dimensional surface is discretized into a triangulated surface with N vertices, T triangles and L links. These are related to the surface topology through the relation $\chi = N + T - L$, where χ is the Euler characteristic.

The morphology of the surface is evolved through a set of four Monte Carlo moves, which are aimed to change each degree of freedom independent of the other. The moves are as follows:

- **Vertex move:** A randomly chosen vertex is displaced by an arbitrary displacement. This move is designed to simulate thermal fluctuations in a membrane surface.
- **Link flip:** A randomly chosen link, which shares two triangles, is reconnected to the two previously unconnected vertices. This move is to simulate the diffusion of lipids in the membrane.
- **Rotation of the in-plane nematic field:** The nematic vector \hat{m} at a randomly chosen vertex is rotated in the tangent plane of the vertex to a new orientation \hat{m}' . This move aims to simulate the thermal noise in the orientation of the in-plane field (*breathing modes of a protein*).
- **Kawasaki exchange of the in-plane nematic field:** This move is relevant for partly decorated nematic membranes. Here, the nematic field \hat{m} at a randomly chosen vertex is exchanged with that at one of the randomly chosen neighboring vertices.

All the above are accepted using the Metropolis criterion:

$$P_{acc} = \min(1, \exp(-\Delta H_{tot}))$$

where ΔH_{tot} is the change in the total energy after a move. *You need to follow a similar approach if you want to augment this code with additional degrees of freedom.*

TECHNICAL DESCRIPTION

- The entire code has been written in `FORTRAN`, in an `objective oriented` fashion, and adheres to specifications laid out for `2003` or later standards.
- `MPI` based parallelization has also been implemented and the current implementation only handles `SINGLE INSTRUCTION MULTIPLE DATA (SIMD)` mode. This means, for a given set of parameters, you can specify the number of processors `N` and the code will automatically generate data for `N` independent ensembles.
- The code uses `FORTRAN TYPE` structures which are the equivalent of a `typedef` in `C` or `C++`. The type structures used are as follows:

DATASTRUCTURE FOR THE VERTICES ON THE SURFACE

Type vertex

```
REAL(KIND=8),DIMENSION(3,1)::splo,spgl,vnor,vcoord,t1,t2
INTEGER,DIMENSION(10)::vneipt,vneitr
REAL(KIND=8),DIMENSION(3,3)::L2G
INTEGER :: nonei,phase,clno,neigh
REAL(KIND=8)::mcur,cur1,cur2,totarea,spen,op,WN
REAL(KIND=8)::kap,kpar,kper,cpar,cper,Nema1,Isa1
REAL(KIND=8)::rkap,rkpar,rkper,rcpar,rcper,rNema1,rIsa1
REAL(KIND=8)::ukpar,ukper
INTEGER :: linkcell,next_vert,prev_vert
```

End Type vertex

DATASTRUCTURE FOR THE TRIANGLES ON THE SURFACE

Type triangle

```
REAL(KIND=8)::ar,vol
INTEGER,DIMENSION(3) ::li,vert
REAL(KIND=8),DIMENSION(3,1)::fnor
```

End Type triangle

DATASTRUCTURE FOR THE LINKS ON THE SURFACE

Type link

```
INTEGER ::tr
INTEGER,DIMENSION(2) ::sep
```

End Type link

DATASTRUCTURE FOR VARIOUS PROPERTIES OF THE MEMBRANE SURFACE

Type membrane_prop

```
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```

END Type membrane_prop

DATASTRUCTURE FOR THE LINKLIST

Type Linklist

```
Real(Kind=8) :: start_coord(3,1),end_coord(3,1)
Integer :: first_vert,last_vert,num_vert,buffer
Integer :: neigh_cell(27)
```

End Type Linklist

ORGANIZATION

The code is divided into the following modules:

```
module_datastruct.f -- definition of the datastructure
module_rerun.f -- modules to start from and dump restart files
module_dataformat.f -- vtu and jvx file writers for visualization
module_initialize_system.f -- modules to initialize the datastructure
module_linklist_calc.f -- linklist definition and calculations
module_curvcalc.f -- compute the curvature at each vertex
module_energy_calculations.f -- energy calculations
module_mcsmoves.f -- definition of the various Monte Carlo moves
module_compute_analytic_measures.f -- compute various measures
maincode.f -- definition of the MAIN function
```

SYSTEM PARAMETERS

- All the required inputs are defined in a file named `parameters.in`
 - **RESTART FILES**
If `restart_mode` is defined as `MEMBRANE` then the code will look for pre-equilibrated configuration file named `startdet.in`, which is dumped from `module_rerun.f`.
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COMPILATION & EXECUTION

- **REQUIRED COMPILER:** `mpif90`
- There are three makefiles supplied along with the code:
`Makefile` -- generic make file that makes call to
`Makefile.gnu` -- for `gnu` based compiler
`Makefile.intel` -- for `intel` based compiler
- Compile as `make gnu` or `make intel`, depending on your system, which would produce an executable `nematic-membrane`.
- The executable may be executed as:
`mpirun -np xx ./nematic-membrane`, where `xx` is the required number of

processors.

So far, I have tested this code using `mpif90` provided with `MPICH`, `OPENMPI`, and `MVAPICH2`, compiled with both `GNU` and `INTEL` compilers. *In general, any stock compiler should work without a problem.*

VISUALIZATION

Upon execution, the code dumps a number of files which includes `conf-*.vtu` and `conf-*.jvx` which are VTK/XML based formats for the morphology of the nematic membrane. These files may be visualized either using:

`paraview` (www.paraview.org) for the `.vtu` files

`javaview` (www.javaview.de) for the `.jvx` files

In the case of `paraview`, in order to view the nematic field, you should load the `glyph` object with `line` representation (from -0.5 to 0.5) and choose `scale as vector`, with a scaling factor of 1.0.
