

Dipoles Implementation in NEMOH Deployment Guide

Revision History

| Author | Revision Number | Date |
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| yedtoss | 1.0 | 10/01/2015 |
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Deployment Instructions

1. Organization of Submission

Nemoh/ Contains the modified source of Nemoh Fortran software

docs/ Contains this deployment guide

docs/GreenFunctionDerivativeODE.pdf Contains ODE for the derivative of the Green's function docs/DipolesImplementationDetails.pdf Contains further details about the dipoles implementation

NemohPython/ Contains the modified Nemoh python code

README.txt note about testing previous old Fortran code

2. Application Setup

Linux

- GCC with GFortran >= 4.8
- BLAS
- LAPACK
- OpenMP provided by GCC
- HDF5 >=1.8.11 http://www.hdfgroup.org/HDF5/ Optionally provided by Anaconda
- HDFView http://www.hdfgroup.org/products/java/release/download.html
- Python 2.7 Optionally provided by Anaconda
- H5py >= 2.3.1 Optionally provided by Anaconda
- Numpy Optionally provided by Anaconda
- CMake >= 2.8 http://www.cmake.org/cmake/resources/software.html
- Anaconda (with Python 2.7) >= 2.1.0 http://continuum.io/downloads

Windows

- MinGW 4.8.1 http://sourceforge.net/projects/MinGWbuilds/files/host-windows/releases/4.8.1/
- BLAS http://icl.cs.utk.edu/lapack-for-windows/lapack/#libraries
- LAPACK http://icl.cs.utk.edu/lapack-for-windows/lapack/#libraries
- OpenMP provided by MinGW
- HDF5 >=1.8.11 http://www.hdfgroup.org/HDF5/ Optionally provided by Anaconda
- HDFView http://www.hdfgroup.org/products/java/release/download.html
- Python 2.7 Optionally provided by Anaconda
- H5py >= 2.3.1 Optionally provided by Anaconda
- Numpy Optionally provided by Anaconda
- CMake >= 2.8 http://www.cmake.org/cmake/resources/software.html
- Anaconda (with Python 2.7) >= 2.1.0 http://continuum.io/downloads

MAC OS X

- Brew http://brew.sh/
- GCC >= 4.8 installed by brew
- XCode Command Line Tools installed by brew



- BLAS provided by XCode commands
- LAPACK Provided by XCode commands
- OpenMP provided by GCC
- HDF5 >=1.8.11 http://www.hdfgroup.org/HDF5/ Optionally provided by Anaconda
- HDFView http://www.hdfgroup.org/products/java/release/download.html
- Python 2.7 Optionally provided by Anaconda
- H5py >= 2.3.1 Optionally provided by Anaconda
- Numpy Optionally provided by Anaconda
- CMake >= 2.8 installed by brew
- Anaconda (with Python 2.7) >= 2.1.0 http://continuum.io/downloads

Variables

Let's call:

\$NEMOH_FORTRAN the directory Nemoh/ in the root of the submission directory **\$NEMOH_PYTHON** the directory NemohPython/ **\$FORTRAN_BUILD** the build directory for the FORTRAN version of Nemoh **\$MINGW_ROOT** the directory where MINGW will be installed

3. Configuration

The python code can be configured using the file **\$NEMOH_PYTHON/nemoh/settings.py Newly added properties are in red**

| Property | Definition | Example |
|-----------------------------|------------------------------------------------------------------------------------------------------------------------------------------------------------------|---------|
| USE_DIPOLES_IMPLEMENTAT ION | Whether or not to use the dipoles implementation | True |
| THIN_PANELS | A list containing the index of the panel which are thin dipoles. The index should be 0-based. Set the list to [-1] to indicate that all panels are thin dipoles. | [1, 2] |
| USE_HIGHER_ORDER | Whether or not to use higher order panel method. | True |
| NUM_PANEL_HIGHER_ORDER | The number of panel per patch in the higher order method. Read section 4 and 7 before attempting to increase this | 1 |
| B_SPLINE_ORDER | The order of the B-Spline for the potential in the higher order panel method. Read section 4 and 7 before attempting to increase this | 1 |
| USE_ODE_INFLUENCE_COEFF | Indicate whether or not to use the | True |



| Property | Definition | Example |
|--------------------------------------|-----------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|-------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| ICIENTS | ode method to compute the influence coefficients | |
| GREEN_TABULATION_NUMX | Number of points in x direction of tabulated data | 500 |
| GREEN_TABULATION_NUMZ | Number of points in z direction of tabulated data | 60 |
| GREEN_TABULATION_SIMPSO N_NPOINTS | Number of sub intervals used to approximate the Green's function integral using simpson rule | 551 |
| HDF5_FILE | The path to the HDF5 file where to save and load the results and input. Required | 'db.hdf5' No need to change |
| NEMOH_CALCULATIONS_FILE | The old nemoh calculation file. Not required but it is needed to automatically convert the old Nemoh.cal file to HDF5 storage | 'Nemoh.cal' No need to change but then make sure you have the nemoh calculation file Nemoh.cal in your working directory. |
| | | Also make sure the path to the mesh file references in Nemoh.cal exists. For example if in the Nemoh.cal file you have 'Cylinder.dat', then you should have the Cylinder.dat file in your current directory |
| NEMOH_INPUT_FILE | Same as above but applied to the nemoh input file | 'input.txt' No need to change but then make sure you have the nemoh input file input.txt in your working directory |
| NEMOH_INT | Represents the integer type to use when performing computations. It should be a valid numpy integer type. See http://docs.scipy.org/doc/numpy/reference/arrays.scalars.html#arrays-scalars-built-in for possible values | 'i' |
| NEMOH_FLOAT | Represents the float type to use when performing computations. It should be a valid numpy float type. See http://docs.scipy.org/doc/numpy/reference/arrays.scalars.html#arrays-scalars-built-in | 'f' |



| Property | Definition | Example |
|-------------------------------------|-----------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|-----------------------------------------------|
| | for possible values | |
| NEMOH_COMPLEX | Represents the complex type to use when performing computations. It should be a valid numpy complex type. See http://docs.scipy.org/doc/numpy/reference/arrays.scalars.html#arrays-scalars-built-in for possible values | 'F' |
| MESH_TEC_FILE | The path to the file where to save the mesh tec file | No need to change. The default value is fine. |
| FK_FORCE_TEC_FILE | The path to the froudkrylov force data in tec format | No need to change. The default value is fine. |
| RADIATION_COEFFICIENTS_T EC_FILE | The path to the file where to save the added mass and damping forces for the radiation problems in tec format. | No need to change. The default value is fine. |
| DIFFRACTION_FORCE_TEC_FI LE | The path to the file where to save the diffraction force for the diffraction problems in tec format. | No need to change. The default value is fine. |
| EXCITATION_FORCE_TEC_FIL E | The path to the file where to save the excitation force for the diffraction problems in tec format. | No need to change. The default value is fine. |
| IRF_TEC_FILE | The path to the file where to save the IRF tec file | No need to change. The default value is fine. |
| WAVE_FIELD_TEC_FILE | The path to the file where to save the wave field tec file | No need to change. The default value is fine. |

4. Implementation details

The references are listed in section 9.

The main code for the dipoles implementation is written in "\$NEMOH_FORTRAN/Solver/Core/SOLVE_BEM_DIPOLES_THIN.f90".

In the dipoles implementation, the body is assumed to be composed of two parts which can be empty. One part consists of conventional element S_s, the other is a surface consisting of thin 'dipoles' elements. To avoid singularity of the original potential (or source) formulation of the problem when the thickness of part of the body tends to zero, this method used an alternative integral equation. The alternative form is equation 15.46 and 15.47 of reference R2 (or equation 6.2 and 6.3 of reference R1). It is a pair of two simultaneous integral equations which is solved with a technique similar to the one describe in equation 15.3 of reference R2 or to Chapter 6 of reference R1. For further information please read Chapter 6 of reference R1 or section 15.7 of reference R2



5. Deployment Instructions

5.1. Generic Instructions

- 5.1.1.Install a Fortran and C/C++ compiler which support OpenMP (Currently gcc and ifort are supported)
- 5.1.2.Install Python 2.7
- 5.1.3.Install pip
- 5.1.4. Install CMake version greater or equal to 2.8
- 5.1.5. Install BLAS and LAPACK and make sure there are in the library search paths
- 5.1.6.Install HDF5 libraries version greater or equal to 1.8.11 and make sure they are in the library search path
- 5.1.7. Compile the Fortran version of Nemoh Solver:
 - 5.1.7.1. Create a build directory \$FORTRAN_BUILD different from \$NEMOH_FORTRAN
 - 5.1.7.2. Go to \$FORTRAN_BUILD folder.
 - 5.1.7.3. Delete the file CMakeCache.txt and the directory CMakeFiles if it exists
 - 5.1.7.4. Run the command cmake \$NEMOH_FORTRAN
 - 5.1.7.5. Run make. The Nemoh library will be created
- 5.1.8. Compile the Nemoh python against the Nemoh Fortran
 - 5.1.8.1. Go to \$NEMOH PYTHON
 - 5.1.8.2. Install the python module prerequisites pip install -r requirements.txt
 - 5.1.8.3. Make sure the \$FORTRAN_BUILD is in the library search paths for compilation and for linking
 - 5.1.8.4. Run python setup.py build_ext --inplace

5.2. Linux Instructions with Ubuntu commands

(If you are not using Ubuntu, you should be able to use your distribution package manager to install equivalent commands)

- 5.2.1.Install GFortran and gcc by running: sudo apt-get install build-essential gfortran gcc
- 5.2.2.Install CMake by running sudo apt-get install cmake
- 5.2.3.Install BLAS and LAPACK and make sure there are in the library search paths by running sudo apt-get install liblapack-dev libblas-dev
- 5.2.4.Install python 2.7, HDF5 libraries, and the nemoh python module requirements. To do so, we just need to install Anaconda:
 - 5.2.4.1. Download Anaconda >= 2.1.0 from http://continuum.io/downloads.

 For linux 64 bits the direct link is (with no space) http://o9c8d0b2229f813c1b93-c95ac804525aac4b6dba79b00b39d1d3.r79.cf1.rackcdn.com/Anaconda-2.1.0-Linux-x86_64.sh
 - 5.2.4.2. Install Anaconda by running bash Anaconda-2.1.0-Linux-x86_64.sh
 - 5.2.4.3. When prompted, accept to add it's path to your ~/.bashrc.
 - 5.2.4.4. Make sure the python version you are using is the one from Anaconda by logout then login or by running source ~/.bashrc



- 5.2.4.5. If successful, when you run python --version you should see something like Python 2.7.8 :: Anaconda 2.1.0 (64-bit)
- 5.2.5. Compile the Nemoh Fortran
 - 5.2.5.1. Create a build directory \$FORTRAN_BUILD different from \$NEMOH_FORTRAN
 - 5.2.5.2. Go to \$FORTRAN BUILD folder by running cd \$FORTRAN BUILD.
 - 5.2.5.3. Delete the file CMakeCache.txt and the directory CMakeFiles if it exists by running rm -rf CMakeCache.txt CMakeFiles/
 - 5.2.5.4. Run cmake -DCMAKE_Fortran_COMPILER="gfortran" \$NEMOH_FORTRAN to generate the Makefiles
 - 5.2.5.5. Run make to build the library
 - 5.2.5.6. The library libnemoh.so will be created
- 5.2.6. Compile the nemoh python against the nemoh Fortran
 - 5.2.6.1. Go to \$NEMOH_PYTHON/nemoh
 - 5.2.6.2. Make sure the \$FORTRAN_BUILD is in the library search paths for compilation and for linking by running:

export LD_LIBRARY_PATH=\$FORTRAN_BUILD export LDFLAGS="-L\$FORTRAN_BUILD"

5.2.6.3. Run python setup.py build_ext --inplace to build the python module in place

5.3. Windows Instructions

5.3.1.Install MinGW 4.8.1

5.3.1.1. You should install posix threads MinGW from http://sourceforge.net/projects/MinGWbuilds/files/host-windows/releases

The 64 bits version is located at

http://sourceforge.net/projects/MinGWbuilds/files/host-windows/releases/4.8.1/64-bit/threads-posix/sjlj/x64-4.8.1-release-posix-sjlj-rev5.7z/download

The 32 bits version is located at

http://sourceforge.net/projects/MinGWbuilds/files/host-windows/releases/4.8.1/32-bit/threads-posix/sjli/x32-4.8.1-release-posix-sjli-rev5.7z/download

- 5.3.1.2. Download the binaries for your platform and extract it somewhere. By default the 64 bits get extracted in a directory named "MinGW64". Let's call this directory \$MINGW_ROOT
- 5.3.1.3. Now add the full path to the folder \$MINGW_ROOT\bin and \$MINGW_ROOT\lib to your Path. Make sure those directories are at the leftmost (the beginning) of the Path. See Setting Path on Windows sub section for more information
- 5.3.2.Install CMake 2.8
 - 5.3.2.1. Download and install CMake from http://www.cmake.org/files/v2.8/cmake-2.8.12.2-win32-x86.exe
 - 5.3.2.2. Choose to add CMake to the path for all users. By default CMake doesn't put the path at the beginning.

So, you need to make sure the CMake path is at the beginning of your path. It is by default "C:\Program Files (x86)\CMake 2.8\bin". See **Setting Path on Windows** sub section for more information



5.3.3.Install LAPACK and BLAS

Download and install LAPACK and BLAS from http://icl.cs.utk.edu/lapack-for-windows/lapack/#libraries Choose the dll libraries for MinGW

The 64 bits BLAS is http://icl.cs.utk.edu/lapack-for-windows/libraries/VisualStudio/3.5.0/Dynamic-MINGW/Win64/liblapack.dll

5.3.3.1. Copy them to **\$MINGW_ROOT\lib** (the lib directory inside MinGW installation root)

5.3.4.Install Anaconda >= 2.1.0

- 5.3.4.1. Download and install Anaconda Windows version greater than or equal to 2.1.0 from http://continuum.io/downloads. The 64 bits Windows version is located at http://o9c8d0b2229f813c1b93-c95ac804525aac4b6dba79b00b39d1d3.r79.cf1.rackcdn.com/Anaconda-2.1.0-Windows-x86 64.exe
- 5.3.4.2. When installing accept adding anaconda to the path and using it's python version as the default python.

By default Anaconda would not put it's path to the beginning of the Windows Path.

You need to move the anaconda paths to the beginning of the Path list. See **Setting Path on Windows** sub section for more information

For me the paths were C:\Users\yedtoss\Anaconda (the most important) and C:\Users\yedtoss\Anaconda\Scripts. You should replace C:\Users\yedtoss\Anaconda by the location where you install Anaconda

5.3.5. Compile the Nemoh Fortran

- 5.3.5.1. Start Powershell (Or windows cmd if you prefer it)
 Make sure that all paths were correctly set. If not then you should close
 Powershell/Cmd, set the path and reopen it. Basically run python --version, cmake --version, gfortran --version and verify that they come from the one you just installed
- 5.3.5.2. Create a build directory \$FORTRAN_BUILD different from \$NEMOH_FORTRAN
- 5.3.5.3. Go to \$FORTRAN BUILD folder.
- 5.3.5.4. Delete the file CMakeCache.txt and the directory CMakeFiles if it exists or better make sure \$FORTRAN BUILD is empty
- 5.3.5.5. Run cmake -DCMAKE_Fortran_COMPILER="gfortran" "\$NEMOH_FORTRAN" -G "MinGW Makefiles"

Please note the "" surrounding \$NEMOH_FORTRAN. You need it even if the directory does not contain space. Make sure "\$NEMOH_FORTRAN" is a full path to avoid any CMake bug

- 5.3.5.6. Run mingw32-make and you will get libnemoh.dll and libnemoh.dll.a
- 5.3.5.7. Copy both generated files to \$MINGW ROOT\lib



5.3.6. Compile the nemoh python against the nemoh Fortran

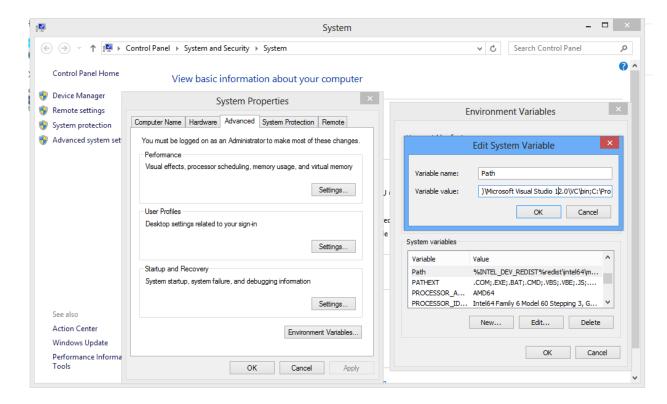
5.3.6.1. Go to \$NEMOH PYTHON\nemoh

5.3.6.2. Run python setup.py build_ext --inplace

5.3.7. Setting Path on Windows

You need to go to Computer → Right click and choose Properties → Advanced System Settings → Advanced Tabs → Environment Variables → Look for path.

A screenshot



Note that when setting the path, you need to separate the different directories by J Also make sure your new directory is not at the end of the list but at the beginning because Windows reads the environment variables from left to right.

5.4. MAC OS X Instructions

We will use homebrew to install most software

5.4.1.Install brew: ruby -e "\$(curl -fsSL https://raw.githubusercontent.com/Homebrew/install/master/install)" (If you did not have XCode Command Line Tools, it will request it, install it and when done press enter on the terminal)



- 5.4.2.Run brew doctor
- 5.4.3.Install GCC and GFortran brew install gcc You can ignore any warning about multilib. It won't affect us
- 5.4.4. Install CMake brew install cmake
- 5.4.5. Download and Install anaconda from http://continuum.io/downloads

You can get the 64 bits version from http://09c8d0b2229f813c1b93-c95ac804525aac4b6dba79b00b39d1d3.r79.cf1.rackcdn.com/Anaconda-2.1.0-MacOSX-x86 64.pkg

We will export anaconda bin to the Path before compiling and using the nemoh python module

- 5.4.6. Compile the Nemoh Fortran
 - 5.4.6.1. Create a build directory \$FORTRAN_BUILD different from \$NEMOH_FORTRAN
 - 5.4.6.2. Go to \$FORTRAN_BUILD folder cd \$FORTRAN_BUILD.
 - 5.4.6.3. Delete the file CMakeCache.txt and the directory CMakeFiles if it exists rm -rf CMakeCache.txt CMakeFiles/
 - 5.4.6.4. Run cmake -DCMAKE Fortran COMPILER="gfortran" \$NEMOH FORTRAN
 - 5.4.6.5. Generate nemoh Fortran library by running make The library libnemoh.dylib will be created (If you get a warning about CMake policy ignore it)
- 5.4.7. Compile the nemoh python against the nemoh Fortran
 - 5.4.7.1. Go to \$NEMOH_PYTHON/nemoh
 - 5.4.7.2. Make sure you are using python from Anaconda

 By running export PATH=/Users/tcs/anaconda/bin:\$PATH

 Replace /Users/tcs/anaconda/bin according to the location where anaconda was installed
 - 5.4.7.3. Make sure the \$FORTRAN_BUILD is in the library search paths for compilation and for linking by running export LD_LIBRARY_PATH=\$FORTRAN_BUILD
 - 5.4.7.4. Run python setup.py build ext --inplace

export LDFLAGS="-L\$FORTRAN BUILD"

Note that in the above process we did not explicitly install LAPACK or BLAS libraries. This is because it is implicitly installed with brew (XCode Command Line Tools). If for some reason you receive an error when linking against LAPACK or BLAS you can install a custom version by running brew install https://raw.githubusercontent.com/Homebrew/homebrew-dupes/master/lapack.rb

Also note that the Intel Fortran Compiler (ifort) is fully supported on all three platforms. It comes bundled with LAPACK and BLAS so if you choose it, you won't need them.

Installing VMTK (Optional, you can skip)

VMTK is used by \$NEMOH_PYTHON/nemoh/export_tec.py to export the generated .tec files to other formats. To install it for Mac, Windows or Linux follow instructions at http://www.vmtk.org/documentation/installation.html

Note that unless you are using the .egg for Anaconda (Windows only), you should not use it with the python from Anaconda. More specifically, on Linux and Mac you have to use it with the built-in python



when installing or using it.

6. Starting

You need to setup all environment variables as described in the deployment instructions.

You should also configure the application as described in the configuration section

Enter the directory \$NEMOH_PYTHON/nemoh/

Run python preprocessor.py to run the preprocessor

Then run the solver with *python solver.py*

Finally run the post processor with *python postprocessor.py*

7. Verification

Setup your environment using the deployment instructions.

Then run the tools by following the Starting section

By default, the cylinder example has been configured. You can run other example by modifying the configuration. You can find additional cases files in \$NEMOH_FORTRAN/Verification folder The HDF5 file db.hdf5 will be generated.

You can visualize it with HDFView from http://www.hdfgroup.org/products/java/release/download.html It has a version for Windows, Mac and Linux

For example, to install the Ubuntu 64 bits:

- Download http://www.hdfgroup.org/ftp/HDF5/hdf-java/current/bin/HDFView-2.10.1-centos5-static64.tar.gz
- Extract it to /tmp so that you have /tmp/HDFView-2.10.1-Linux, then locate the file hdfview.sh inside it. It should be located at /tmp/HDFView-2.10.1-Linux/HDF_Group/HDFView/2.10.1/bin/hdfview.sh
- Open the file hdfview.sh and set INSTALLDIR to ■■
- Enter the bin directory (/tmp/HDFView-2.10.1-Linux/HDF_Group/HDFView/2.10.1/bin)
- Make sure you have java from Sun, JRE is enough. You can use JRE 6 or JRE 7
- Run bash hdfview.sh

Then open the HDF5 file (Click File → Open)

Enter the directory \$NEMOH PYTHON/nemoh/

Run python preprocessor.py to run the preprocessor

Then run the solver with python solver.py

Finally run the post processor with python postprocessor.py

Ignore the warning of the type "Warning: normal vector of panel 14 points towards the x axis". It is caused by the input mesh used.

The default example for test is \$NEMOH PYTHON/test files/thin body dipoles

To run the dipoles implementation, set USE_DIPOLES_IMPLEMENTATION settings to True (the default) and set THIN_PANELS to the list of 'dipoles' panels or to [-1] to include all panels. To run previous lower order panel method (source formulation) set USE_DIPOLES_IMPLEMENTATION to False.



The lower order results are located in \$NEMOH_PYTHON/test_files/thin_body_dipoles/db_old.hdf5 and the dipoles implementation results in \$NEMOH_PYTHON/test_files/thin_body_dipoles/db.hdf5

ACCURACY VERIFICATION

To test the accuracy of the dipoles implementation we can perform either one of the following tests:

We use a body containing thin mesh elements with a zero thickness. We run both the previous source formulation and the dipoles implementation against this test case. It is expected that the previous source formulation method will fail. Indeed the linear system to solve will be singular and the GAUSS solver will lead to singularities error. The dipoles implementation will successfully run.

We could also use a body containing thin mesh elements with thickness near zero (but not zero). The body will be modeled with N panels

The previous could fail as before or could pass. If it passes, then we will take the same body and model it with 2N smaller panels. We will observe that the forces and potential will not converge and will be different.

Doing the same test with the dipoles implementation will converge

Finally it will be useful to use a body which has been already solved analytically (in the literature for example.) We could then compare the dipoles implementation and the previous source formulation against the analytical result.

We could not perform any of these tests because of the issue described here http://apps.topcoder.com/forums/?module=Thread&threadID=844161&start=0

Instead we used a conventional body and we assumed that all its panel are thin dipoles ones. Although this assumption is not correct, it shows that the dipoles implementation can be run successfully. The results (dynamic forces) are different from the previous method because the integral equations we are solving is different in both cases. But the Froude Krylov Forces match between the dipoles implementation and the previous source formulation. This can be checked manually in the HDF5 files.

8. Limitations of the current dipoles implementation

- The current implementation does not compute the Kochin function and thus does not compute the drift forces, yaw moment and wave elevation (in postprocessing module). The Kochin function is not computed because the previous method to compute it uses the source strength which is not available in the implementation of the dipoles implementation. Other technique to compute the Kochin function should be performed.
- The dipoles implementation does not support the technique for symmetry in the body. You should describe the whole geometry of the body to use it. It should be possible to extend the dipoles implementation to support fast computation when there is symmetry.



To run your problem containing a symmetry in the mesh, disable the symmetry switch by setting the second number in the mesh file to 0 before running the preprocessor. Alternatively you can also set the second number in input/bodies/body1/mesh of the input HDF5 file to 0. You should do this for all bodies of your problem.

An error message will be given and program will stop if one attempts to use the new method with symmetry.

 The current dipoles implementation does not support computing the Green's function using an ODE as supported in the lower order panel method. If such an attempt is made, a warning will be shown and the method to compute the Green's function will fall back to the default (tabulation, series approximation and interpolation).

9. References

- R1. A Rankine panel method as a tool for the hydrodynamic design of complex marine vehicles. http://dspace.mit.edu/bitstream/handle/1721.1/50364/39696248.pdf?sequence=1
- R2. Wamit User Manual http://www.wamit.com/manualupdate/V70_manual.pdf

10. Resource Contact List

| Name | Resource Email |
|---------|----------------|
| yedtoss | |