Instructions for using NEMO on SISU

Yongmei Gong 8 March, 2018 The following instruction consist text instructions and a number of well commented bash files on how to launch basic nemo simulations. They are created for the environment of the super computer Sisu in CSC-IT center for science.

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1 Get the code

- 1. Sign up for the wiki (http://forge.ipsl.jussieu.fr/nemo/wiki/Users) by sending an email containing your username (5 characters minimum length) to nemo@forge.ipsl.jussieu.fr;
- 2. Get the conformation email then reset your password.
- 3. Now download the source code from the distribution

```
#!/bin/bash
   set -ex
   # 2018-02-20, juha.lento@csc.fi'
   # modified 2018-03-08, yongmei.gong@helsinki.fi
   # NEMO is "research code", which for NEMO means that:
   # - "install" in the NEMO documentation actually is what
      is usually referred as "build"
   # The following build instructions are based on:
   # - https://forge.ipsl.jussieu.fr/nemo/wiki/Users/
      ModelInstall#ExtracttheNEMOcode
   # - https://forge.ipsl.jussieu.fr/nemo/wiki/Users/
      ModelInterfacing/InputsOutputs#
      ExtractingandinstallingXIOS
   #Your user name used for registering to nemo user wiki
   USER = "YOUR_USER_NAME"
   # Load system I/O libraries (that is how we use an
       existing software on Sisu)
   # now use xios2.0 by default
   module load cray-hdf5-parallel cray-netcdf-hdf5parallel
      xios/
   # Create the main NEMO directory (a folder) where you want
       to store the source code
   # Here I put in in the user application directory in Sisu,
       $USERAPPL, where you can build (install) your own
       software
   mkdir -p $USERAPPL/nemo_test
   cd $USERAPPL/nemo_test
30
   # Checkout (download) the source code
```

```
# Type your password on the screen and type yes when asked
.
svn --username $USER co http://forge.ipsl.jussieu.fr/nemo/
    svn/branches/2015/nemo_v3_6_STABLE/NEMOGCM
```

2 Declare the compilers

All compiler options in NEMO are controlled using files in NEMOGCM/ARCH/arch. Now we create a file to declare the compilers we use to build nemo accroding to what we have in Sisu.

```
#!/bin/bash
set -ex
# NEMO 3.6 STABLE + XIOS-2 build instructions for sisu.csc
# 2018-02-20, juha.lento@csc.fi
# modified 2018-03-08, yongmei.gong@helsinki.fi
# NEMO is "research code", which for NEMO means that:
# - "install" in the NEMO documentation actually is what
   is usually referred as "build"
# The following build instructions are based on:
# - https://forge.ipsl.jussieu.fr/nemo/wiki/Users/
   ModelInstall#ExtracttheNEMOcode
# - https://forge.ipsl.jussieu.fr/nemo/wiki/Users/
   ModelInterfacing/InputsOutputs#
   ExtractingandinstallingXIOS
# NEMO build
# All compiler options in NEMO are controlled using files
   in NEMOGCM/ARCH/arch-'my_arch'.fcm where 'my_arch' is
   the name of the computing architecture.
# Now we create a file to declare the compilers we use to
   build nemo accroding to what we have in Sisu
cd $USERAPPL/nemo_test/NEMOGCM
cat > ARCH/arch-XC40-SISU.fcm <<EOF
%NCDF_HOME
                     $NETCDF_DIR
%HDF5_HOME
                     $HDF5_DIR
                     $(pkg-config --variable=CRAY_prefix
%XIOS_HOME
   xios)
```

```
%NCDF_INC
                      -I%NCDF_HOME/include -I%HDF5_HOME/
    include
%NCDF_LIB
                     -L%HDF5_HOME/lib -L%NCDF_HOME/lib -
    lnetcdff -lnetcdf -lhdf5_hl -lhdf5 -lz
%XIOS_INC
                      -I%XIOS_HOME/inc
%XIOS_LIB
                      -L%XIOS_HOME/lib -lxios
%CPP
                      cpp
%FC
                      ftn
                                                    -02 -
%FCFLAGS
                      -emf -s real64 -s integer32
    hflex_mp=intolerant -Rb
%FFLAGS
                      -emf -s real64 -s integer32 -00 -
    hflex_mp=strict -Rb
                      -P -E -traditional-cpp
%FPPFLAGS
%LDFLAGS
                      -hbyteswapio
%AR
                      ar
%ARFLAGS
                     -r
%MK
                      gmake
%USER_INC
                      %XIOS_INC
%USER_LIB
                     %XIOS_LIB
%CC
%CFLAGS
                      -00
EOF
```

3 Build NEMO for experiment - GYRE_XIOS

Now we build an executable for the experiment GYRE_XIOS. We use the up-to-date version of XIOS - xios2.0.990 instead the oldder xios1.0. This requires declaring the active cpp keys in the cpp_*_fcm files.

```
#!/bin/bash
set -ex

# NEMO 3.6 STABLE + XIOS-2 build instructions for sisu.csc
    .fi

#
# 2018-02-20, juha.lento@csc.fi
# modified 2018-03-08, yongmei.gong@helsinki.fi

# NEMO is "research code", which for NEMO means that:
# - "install" in the NEMO documentation actually is what
    is usually referred as "build"

# The following build instructions are based on:
# - https://forge.ipsl.jussieu.fr/nemo/wiki/Users/
    ModelInstall#ExtracttheNEMOcode
```

```
# - https://forge.ipsl.jussieu.fr/nemo/wiki/Users/
      ModelInterfacing/InputsOutputs#
      ExtractingandinstallingXIOS
   # Load system I/O libraries
   # might change to xios2.0
   module load cray-hdf5-parallel cray-netcdf-hdf5parallel
      xios/2.0.990
   # Declare your NEMO code directory
   NEMOBUILD="$USERAPPL/nemo_test3"
   # NEMO build
   # Declare your configuration for the simulation, e.g. the
      GYRE_XIOS experiment
   cd $NEMOBUILD/NEMOGCM/CONFIG
30
   #you need to add new keys in .fcm file
   sed -i 's/$/ key_nosignedzero/' GYRE_XIOS/cpp_GYRE_XIOS.
   # Here you compile a executable for the experiment
      GYRE_XIOS in either $TMPDIR or $WRKDIR/DONOTREMOVE
   ./makenemo -t $TMPDIR -m XC40-SISU -r GYRE_XIOS -n
      MY_GYRE_XIOS
   # NEMO test
   # For a quick test, only!
   # cd MY_GYRE_XIOS/EXP00
   # cp $TMPDIR/MY_GYRE_XIOS/BLD/bin/nemo.exe .
   \# aprun -n 4 nemo.exe
   # For actual experiments:
   # Copy the experiments in $WRKDIR/DONOTREMOVE
   cp GYRE_XIOS/ $WRKDIR/DONOTREMOVE/MY_GYRE_XIOS/
   cd $WRKDIR/DONOTREMOVE/MY_GYRE_XIOS/EXPOO
   # Copy the executable to the EXP00 directory
   cp $TMPDIR/MY_GYRE_XIOS/BLD/bin/nemo.exe .
  # Creat a script for Using SLURM commands to execute batch
   jobs
```

```
in Sisu queue
   # More about the SLURM commands can be found in
   # - https://research.csc.fi/sisu-using-slurm-commands-to-
       execute-batch-jobs
   cat > batch_job.sh <<EOF
   #!/bin/bash -1
   #SBATCH -t 00:29:00
   \#SBATCH - J \ gyre\_xios
   \#SBATCH - p test
   #SBATCH -o gyre_xios.%j
   #SBATCH -e gyre_xios_err.%j
   \#SBATCH - N 4
   aprun -n 4 nemo.exe
   EOF
70
   # Then submit the job in the queue
   sbatch batch_job.sh
```

4 Build NEMO for experiment - ORCA2_LIM3

Now we build an executable for the experiment ORCA2_LIM3.

ORCA is the generic name given to global ocean configurations. Its specificity lies on the horizontal curvilinear mesh used to overcome the North Pole singularity found for geographical meshes. LIM (Louvain-la-Neuve sea-ice model, multi-category model LIM3 is used) is a thermodynamic-dynamic sea ice model specifically designed for climate studies.

Similarly this requires declaring the active cpp keys in the cpp_*_.fcm files. And this time the experiment needs input data.

```
# The following build instructions are based on:
   # - https://forge.ipsl.jussieu.fr/nemo/wiki/Users/
      ModelInstall#ExtracttheNEMOcode
  # - https://forge.ipsl.jussieu.fr/nemo/wiki/Users/
      ModelInterfacing/InputsOutputs#
      ExtractingandinstallingXIOS
   # Load system I/O libraries
   # might change to xios2.0
   module load cray-hdf5-parallel cray-netcdf-hdf5parallel
      xios/2.0.990
   # Declare your NEMO code directory
  NEMOBUILD="$USERAPPL/nemo_test3"
   # NEMO build
   # Declare your configuration for the simulation, e.g. the
      ORCA2_LM3 experiment
   cd $NEMOBUILD/NEMOGCM/CONFIG
   #you need to add new keys in .fcm file
   sed -i 's/$/ key_nosignedzero key_xios2/' ORCA2_LIM3/
      cpp_ORCA2_LIM3.fcm
   # Here you compile a executable for the experiment
      ORCA2_LIM3 in $TMPDIR
   ./makenemo -t $TMPDIR -m XC40-SISU -r ORCA2_LIM3 -n
      MY_ORCA2_LIM3
   # NEMO test
40
   cd ORCA2_LIM3/EXPOO
   # If you use XIOS2.0 you need to copy all the .xml file
      from GYRE_XIOS
   cp $NEMOBUILD/NEMOGCM/CONFIG/GYRE_XIOS/EXPOO/*xml .
   # For actual experiments:
   # Copy the experiments in $WRKDIR/DONOTREMOVE
   cd ../..
   cp ORCA2_LIM3/ $WRKDIR/DONOTREMOVE/MY_ORCA2_LIM3/
  cd $WRKDIR/DONOTREMOVE/MY_ORCA2_LIM3/
   cp $TMPDIR/MY_ORCA2_LIM3/BLD/bin/nemo.exe .
```

```
# now you need to download input data ORCA2_LIM_nemo_v3.6.
   tar from http://forge.ipsl.jussieu.fr/nemo/wiki/Users/
   ReferenceConfigurations/ORCA2_LIM3_PISCES
# Then extract them in EXP00
tar xvf ORCA2_LIM_nemo_v3.6.tar
gzip -d *gz
# Creat a script for Using SLURM commands to execute batch
    jobs
in Sisu queue
# More about the SLURM commands can be found in
# - https://research.csc.fi/sisu-using-slurm-commands-to-
   execute-batch-jobs
cat > batch_job.sh <<EOF
#!/bin/bash -1
#SBATCH -t 00:29:00
#SBATCH -J ORCA2_LIM3
#SBATCH -p test
#SBATCH -o ORCA2_LIM3.%j
#SBATCH -e ORCA2_LIM3_err.%j
#SBATCH -N 4
aprun -n 4 nemo.exe
EOF
# Then submit the job in the queue
sbatch batch_job.sh
```

5 Build NEMO for experiment - ORCA1_LIM3

The difference between ORCA1 and ORCA2 is that the spacial resolution of the former is 1 degree and the latter is 2 degree. So ORCA1 has higher resolution and also needs more input data.

You will need files and data in two folders: ORCA1_cfg and ORCA1_data to create ORCA1 experiment configuration

```
#!/bin/bash
set -ex

# NEMO 3.6 STABLE + XIOS-2 build instructions for sisu.csc
    .fi
#
# 2018-02-20, juha.lento@csc.fi
# modified 2018-03-06, yongmei.gong@helsinki.fi
```

```
# NEMO is "research code", which for NEMO means that:
# - "install" in the NEMO documentation actually is what
    is usually referred as "build"
# The following build instructions are based on:
# - https://forge.ipsl.jussieu.fr/nemo/wiki/Users/
    ModelInstall#ExtracttheNEMOcode
# - https://forge.ipsl.jussieu.fr/nemo/wiki/Users/
    ModelInterfacing/InputsOutputs#
    ExtractingandinstallingXIOS
# Load system I/O libraries
# might change to xios2.0
module load cray-hdf5-parallel cray-netcdf-hdf5parallel
    xios/2.0.990
# Declare your NEMO code directory
NEMOBUILD="$USERAPPL/nemo_test3"
# NEMO build
# There is no ORCAl experiment, which means in the
    standard distribution therefore we need to creat it
    ourselves
cd $NEMOBUILD/NEMOGCM/CONFIG
mkdir ORCA1
cd ORCA1
# put the cpp file here
cp /homeappl/home/ygong/appl_sisu/NEMO/NEMO_local/
    ORCA1_cfg/cpp_ORCA1_LIM3.fcm .
# Creat a experiment dir EXP00 and put all the xml and
    namelist files and data there
# Make sure that you have all the file_* and field_* xml
    files listed in context_nemo.xml (you don't need *
    pisces.xml files if you don't want to couple with
    biochemistry).
# you can put the line 'nn_msh
    (=1) a mesh file or not (=0)' under &namdom in the file
     namelist_cfg.
# This prevents nemo to creat the mesh files, which will
   cause trouble when you re-launch the experiment in the
```

```
same directory as they cannot be over-written
   mkdir EXP00
   cp /homeappl/home/ygong/appl_sisu/NEMO/NEMO_local/
      ORCA1_cfg/*xml .
   cp /homeappl/home/ygong/appl_sisu/NEMO/NEMO_local/
      ORCA1_cfg/namelist* .
   # Here you compile a executable for the experiment ORCA1
      in $TMPDIR
   ./makenemo -t $TMPDIR -m XC40-SISU -r ORCA1 -n MY_ORCA1
   # NEMO test
  # now the experiment gets bigger so we do it in $WRKDIR
   cd $WRKDIR/DONOTREMOVE/
   mkdir MY_ORCA1
   cd ORCA1
   cp $NEMOBUILD/NEMOGCM/CONFIG/ORCA1/EXPOO .
  cp $TMPDIR/MY_ORCA2_LIM3/BLD/bin/nemo.exe .
   # Data includes all the data describing the boundray and
      initial conditions and the climatology in NetCDF format
       . And a samll *.dat file describing the humidity
   cp /homeappl/home/ygong/appl_sisu/NEMO/NEMO_local/
      ORCA1_data/* .
   # Creat a script for Using SLURM commands to execute batch
       jobs
   in Sisu queue
   # More about the SLURM commands can be found in
   # - https://research.csc.fi/sisu-using-slurm-commands-to-
      execute-batch-jobs
   cat > batch_job.sh <<EOF
   #!/bin/bash -1
   #SBATCH -t 12:00:00
   \#SBATCH - J ORCA1
   \#SBATCH - p \ small
  #SBATCH - O ORCA1.%j
   #SBATCH -e ORCA1_err.%j
   \#SBATCH - N 4
   aprun -n 24 nemo.exe
  EOF
80
   # submit the job
```

sbatch batch_job.sh

6 Clean nemo build

If something has gone wrong with or has been changed for the build of nemo (nemo.exe)

- Clean a bad configuration

./makenemo -C YOUR_CONFIG clean_config

- Uninstalling (Clean up the whole thing)

./makenemo -n YOUR_BUILD clean

(e.g. ./makenemo ./makenemo -t \$TMPDIR -m XC40-SISU -n MY_GYRE_XIOS clean)

7 Check the run status and outputs

- If run crushes, try to find 'E R R O R' section in ocean.output - Use neview (now it is only in Petteris directory) to check the results *nc files cd /homeappl/home/puotila/bin

export HDF5_DISABLE_VERSION_CHECK=1

./ncview /YOUR_OUTPUT_DIR/_5d_00010101_00011230_grid_T.nc &