



# How to install the integrated code: TASK

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1. Preparation for macOS
2. Preparation for Ubuntu
3. Install PETSc
4. Introduction to git
5. Install task and related libraries
6. Introduction to the task code

# Preparation for macOS (1)

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- **Install Xcode**

- Xcode: development environment on macOS
- Use App Store
- Category: Development
- Choose and install Xcode

- **Install Command\_Line\_Tools**

- Command\_Line\_Tools: various Unix commands for development
- Input command on terminal
- `xcode-select --install`

- **Install XQartz**

- Download and install XQartz from <https://www.xquartz.org>

- **Install java**

- Download and install java from [https://www.java.com/en/download/mac\\_download.jsp](https://www.java.com/en/download/mac_download.jsp)

# Preparation for macOS (2)

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- **Install Macports**

- Download the latest macports binary from <https://www.macports.org>
- Select tab: Installing MacPorts
- Quickstart: download and install MacPorts installer for appropriate macOS version
- MacPorts is mostly installed at /opt/local
- Update to the latest Macports
  - `sudo port selfupdate`
  - `sudo port upgrade outdated`

- **Install compiler and related modules**

- gfortran: `sudo port install gcc11`
- mpich: `sudo port install mpich`
- others: `sudo port install gmake cmake imake`

# Preparation for Ubuntu

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## 1. Install required modules

```
sudo apt-get install gfortran-11
```

```
sudo apt-get install gcc-11
```

```
sudo apt-get install g++-11
```

```
sudo apt-get install git
```

```
sudo apt-get install xorg-dev
```

```
sudo apt-get install valgrind
```

```
sudo apt-get install cmake
```

```
sudo apt-get install python
```

```
sudo apt-get install mpich
```

# Install PETSc (1)

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- **PETSc**: Parallel matrix solver library
  - blas,lapack: matrix solver tolls
  - scalapack, metis, parmetis, blacs, superlu: parallel solver tools
  - MUMPS: Direct matrix solver for real and complex
  - PETSc: Iterative matrix solver for real or complex
- **Make PETSc directory and change its owner**
  - `sudo mkdir /opt/PETSc`
  - `sudo chown /opt/petsc $USERNAME`
  - `cd /opt/PETSc`
- **Download latest PETSc library package by git**
  - **First download of PETSc source**
    - `git clone -b release https://gitlab.com/petsc/petsc.git petsc`
  - **In order to update PETSc source**
    - `git pull`

# Install PETSc (2)

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- **Provide environment variables for PETSC in ~/.profile or .zprofile**
  - `export PETSC_DIR=/opt/PETSc/petsc`
  - `export PETSC_ARCH=default`
- **Configure script in python**
  - **Copy** `default.py` **to** `/opt/PETSc/petsc`
  - **Provide exec attribute to** `default.py`
    - `chmod 755 default.py`
  - **Execute configuration script** (It may take half an hour.)
    - `./default.py`
  - **Additional libraries are created in** `default/externalpackages`
- **Make and check PETSc library**
  - `make` (It may take half an hour.)
  - `make check`

# How to use git (1)

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- **git**: version and remote repository control facility
- **Repositories**
  - **local**: in your machine
  - **remotes**: in remote servers
  - **remotes/origin**: in default server: bpsi.nucleng.kyoto-u.ac.jp
- **Branches**
  - **There are several branches for code development**
    - **master**: default, stable version, often rather old
    - **develop**: latest version, where I am working
    - **others**: branches for working specific modules
  - **cd task**
  - **git branch** : list branch names, local only
  - **git branch -a** : list branch names, local and remote

# How to use git (2)

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- **To use develop branch**
  - **Create local branch develop and associate it with remote develop**
  - `git checkout -t -b develop origin/develop`
  - `git branch`
- **Change working branch**
  - `git checkout master`
  - `git checkout develop`
- **Update working branch:** download from remote repository
  - `git pull`
    - Your modification is kept, if committed.
    - If uncommitted modification remains, no overwrite.
    - use `git stash` to keep away your modification.
    - If there are conflicts with your committed modification, the conflicts are indicated in the file. Correct them and `git pull` again.



# How to use git (3)

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- **To check your modification**
  - `git status`
- **To commit your modification with message:** only local depository is updated. message is required.
  - `git commit -a -m'message'`
- **To list all modification**
  - `git log`
- **To show difference from committed repository**
  - `git diff [filename]`
- **For more detail, visit**
  - <https://git-scm.com/documentation>

# Install TASK (1)

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- **Check availability of git:** just command input “git”
- **Set your identity:** To record who changed the code?
  - `git config --global user.name “[your-full-name]”`
  - `git config --global user.email [your-mail-address]`
  - For example,
    - `git config --global user.name “Atsushi Fukuyama”`
    - `git config --global user.email fukuyama@nucleng.kyoto-u.ac.jp`
  - Data is saved in `$HOME/.gitconfig`
- **Create a working directory:** any directory name is OK
  - `mkdir git`
  - `cd git`

# Install TASK (2)

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- **Download TASK and necessary libraries** for download only
  - `git clone https://bpsl.nucleng.kyoto-u.ac.jp/pub/git/gsaf.git`
  - `git clone https://bpsl.nucleng.kyoto-u.ac.jp/pub/git/bpsd.git`
  - `git clone https://bpsl.nucleng.kyoto-u.ac.jp/pub/git/task.git`
- **Download TASK and necessary libraries** for download and upload
  - `git clone ssh://username@bpsl.nucleng.kyoto-u.ac.jp/pub/git/gsaf.git`
  - `git clone ssh://username@bpsl.nucleng.kyoto-u.ac.jp/pub/git/bpsd.git`
  - `git clone ssh://username@bpsl.nucleng.kyoto-u.ac.jp/pub/git/task.git`
  - `username@` can be omitted if the usernames at remote and local are same.
- **Three directories are created**
  - **gsaf**: graphic library
  - **bpsd**: data interface library
  - **task**: main TASK directory

# Install TASK (3)

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- **Install graphic library GSAF**

- `cd git/gsaf/src`
- Copy Makefile.arch appropriate for your environment
  - for macOS: `cp ../arch/macos-gfortran/Makefile.arch .`
  - for Ubuntu: `cp ../arch/ubuntu-gfortran64-static/Makefile.arch .`
- **Edit Makefile.arch**: adjust BINPATH and LIBPATH to available ones
  - BINPATH: graphic commands are located, should be included in \$PATH in ~/.profile or ./zprofile
  - LIBPATH: graphic libraries are located, should be included in library path for compiling applications using the graphic libs.
- `make`
- `make install`
  - if BINPATH is protected, use “`sudo make install`”

# Install TASK (4)

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- **Check the availability of GSAF library**
  - `cd test`
  - `make`
  - Applications using GSAF library must be started from X11 window such as xterm, not from Terminal on macOS.
  - `./bsctest`
  - `5` : Choose the size of window
  - `c` : Continue the run
  - `m`
  - New graphic window opens and marks and lines are drawn.
  - To go back to the original window, enter CR.
  - If focus does not change, click the original window and check XQartz preferences.
  - `e` : Close the graphic window
  - `cd ../../..`

# Install TASK (5)

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- **Use the latest develop branch and setup make.header file**
  - `cd task`
  - `git checkout -t -b develop origin/develop`
  - `cp make.header.org make.header`
  - **Edit make.header** to remove comments for target OS and compiler
- **Compile data exchange library BPSD**
  - `cd ../bpsd`
  - `make`
  - `cd ../task`
- **Compile TASK**: eq for example
  - `cd eq`
  - `make`
  - `./eq`

# Install TASK (6)

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- **Type of parallel matrix solver configuration**
  - `.nompi`: only single- processing solver without MPI environment
  - `.mpi`: only single-processing solver with MPI environment
  - `.petsc`: various parallel iterative solvers in with MPI
  - `.petsc+mumps`: various parallel iterative and direct solvers with MPI
- **Setup matrix solver library**
  - `cd mtxp`
  - `cp make.mtxp.XXX make.mtxp`
  - `make`
- **Compile modules:**
  - **Edit the beginning of Makefile:** Select matrix solver
    - **Real** matrix equation (fp,ti): any mtxp library
    - **Complex** matrix equation (wm,wf2d,wf3d): band matrix or MUMPS

# How to use GSAF

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- **At the beginning of the program**
  - **Set graphic resolution** (0: metafile output only, no graphics)
  - **commands**
    - **c**: continue
    - **f**: set metafile name and start saving
- **At the end of one page drawing**
  - **commands**
    - **c** or **CR**: change focus to original window and continue
    - **f**: set metafile name and start saving
    - **s**: start saving and save this page
    - **y**: save this page and continue
    - **n**: continue without saving
    - **d**: dump this page as a bitmap file “gsdumpn”
    - **b**: switch on/off bell sound
    - **q**: quit program after confirmation



# Graphic Utilities

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- **Utility program**

- **gsview**: View metafile
- **gsprint**: Print metafile on a postscript printer
- **gstoeeps**: Convert metafile to eps files of each page
- **gstops**: Convert metafile to a postscript file of all pages
- **gstotgif**: Convert metafile to a tgif file for graphic editor tgif
- **gstotsvg**: Convert metafile to a svg file for web browser

- **Options**

- **-a**: output all pages, otherwise interactive mode
- **-s ps**: output from page ps
- **-e pe**: output until page pe
- **-p np**: output contiguous *np* pages on a sheet
- **-b**: output without title
- **-r**: rotate page
- **-z**: gray output

# Typical File Name of TASK

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- **XXcomm.f90**: Definition of global variables, allocation of arrays
- **XXmain.f90**: Main program for standalone use, read XXparm file
- **XXmenu.f90**: Command input
- **XXinit.f90**: Default values
- **XXparm.f90**: Read input parameters
- **XXview.f90**: Show input parameters
- **XXprep.f90**: Initialization of run, initial profile
- **XXexec.f90**: Execution of run
- **XXgout.f90**: Graphic output
- **XXfout.f90**: Text file output
- **XXsave.f90**: Binary file output
- **XXload.f90**: Binary file input

# Typical input command

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- When input line includes **=**, interpreted as a namelist input (e.g., **rr=6.5**)
- When the first character is not an alphabet, interpreted as line input
- **r**: Initialize profiles and execute
- **c**: Continue run
- **p**: Namelist input of input parameters
- **v**: Display of input parameters
- **s**: Save results into a file
- **l**: Load results from a file
- **q**: End of the program
- **Order of input parameter setting**
  - Setting at the subroutine **XX\_init** in **XXinit.f90**
  - Read a namelist file **XXparm** at the beginning of the program
  - Setting by the input line