

# Review

Questions from day 1?

# Setting up different case types

# Review of a case directory – Common files and commands

- Case documentation
  - README.case – detailed documentation of the case
  - CaseStatus – history of operations (configuration, builds, runs)
  - CaseDocs – model run-time settings files
- Commands
  - **xmlquery** – Find and display case information and settings
  - **xmlchange** – Change case settings
  - **pelayout** – Display how the model will use the machine processing nodes
  - **preview\_namelist** – Compute all the runtime model settings files
  - **preview\_run** – Display details of how the run will be submitted to the machine

## Review of a case directory – Configuration file groups

- **Component namelist (run-time) config files – (*Set runtime options here*)**
  - user\_nl\_biom user\_nl\_cam user\_nl\_cice user\_nl\_cism
  - user\_nl\_clm user\_nl\_cpl user\_nl\_mosart
  - These files are used to set component-model namelists at runtime.
- **Case configuration files (*FYI – you should not have to look at these*)**
  - env\_archive.xml – details of how to run the short-term archiver
  - env\_batch.xml – details of how the HPC machine's batch system works
  - env\_build.xml – details of how the executable will be built
  - env\_case.xml – details of this particular case
  - env\_mach\_pes.xml – details of how the run will use machine processors
  - env\_mach\_specific.xml – details of runtime environment settings
  - env\_run.xml – details of how the case will be run
  - env\_test.xml – used by the CIME testing system to run tests

# Setting up different case types

# Types of NorESM model run

- **Initial run**
  - Start NorESM with a set of initial-data files for each component
  - Used to begin a new experiment
- **Restart run**
  - Start NorESM with the exact state of the restart files
  - Used to continue a run in a case where the state was last saved
- **Branch run**
  - Similar to a restart run but using a consistent set of restart files from a previous run
- **Hybrid run**
  - Similar to an initial run but using initialization datasets *from a previous case*

## NorESM model **initial** run (aka **startup**)

- This is the default run type where the model is started from a set of initial value files or settings.
- Each component will have default initial value files for supported combinations of compset and resolution. In some unsupported settings, you may have to provide an initial value file (expert level).
- You can always find out what type of run you will be doing:  
    `./xmlquery RUN_TYPE`



## NorESM model **restart** run

- Restart runs are used to continue a run with minimal changes.
- When submitting a restart run, only changes to history (diagnostic) output fields and frequency are allowed.
- When a case is running and creates a set of restart files, these files can be used to restart the model such that the model state is bit identical with a normal model run.
- To create restart files, use the **REST\_OPTION** and **REST\_N** XML variables.
- Make sure **CONTINUE\_RUN** is **TRUE** before submitting (normally automatic after a successful run)

## NorESM model **branch** run

- In a branch run, all components are initialized using a consistent set of restart files from a previous run.
- The set of restart files is determined by the **RUN\_REFCASE** and **RUN\_REFDATE** XML variables
- Several reference cases are provided in /cluster/shared/noresm/inputdata/ccsm4\_init but you can change the location with the **RUN\_REFDIR** XML variable.
- Set **GET\_REFCASE** to **TRUE** so that the reference files are copied into your run directory.
- A branch run is a restart run so only limited changes to the namelists are allowed (see restart run).

## NorESM model **hybrid** run

- A hybrid run is initialized like an initial run but it uses initialization data sets from a previous case as in a branch run.
- Because a hybrid run is similar to an initial run, there is a lot more flexibility in the set of initial-data files and namelist settings.
- In a hybrid run, CAM requires an initial value file, not a restart file.
- The ocean component does not start until what would normally be the second ocean coupling interval.

Other (hopefully) useful tidbits

## More about xmlchange and xmlquery

- Search for an XML variable  
**./xmlquery --partial <STRING>**
- Find out more about an XML variable  
**./xmlquery --description <VARIABLE>**
- What about duplicate variables in different groups?  
**./xmlquery JOB\_WALLCLOCK\_TIME**  
Results in group case.st\_archive  
JOB\_WALLCLOCK\_TIME: 0:59:00  
Results in group case.run  
JOB\_WALLCLOCK\_TIME: 01:00:00
- Change just the run wallclock  
**./xmlchange --subgroup case.run**  
**JOB\_WALLCLOCK\_TIME=02:00:00**

## Another way to modify a case

- `create_newcase` has an argument, `--user-mods-dir`, that allows you to bundle a set of `user_nl_XXX` files along with `xmlchange` commands.
- NorESM and several component models provide common case customization sets in their `cime_config/usermods_dirs` directory.

# Really useful case control reference

<https://esmci.github.io/cime/versions/master/html/index.html>



Questions?