

THE MITGCM USER'S GUIDE TO ARCHER2

*Mike Mineter¹, Emma Boland², Dan(i) Jones²,
Kaitlin Naughten², Dan Goldberg¹*

1: University of Edinburgh, 2: British Antarctic Survey

POLAR SCIENCE
FOR PLANET EARTH



BACKGROUND

- Based on work by Mike Mineter, University of Edinburgh, under eCSE02-06 funded by EPCC
- ARCHER2 “embedded CSE” (eCSE) projects are funded through regular calls to develop sustainable software and improve research on ARCHER2
- Provide funds for a research software engineer - can be provided by the CSE team or from your/other institute.
- eCSE02-06 was titled “Optimising MITgcm on ARCHER2” and provided funding to install and optimised MITgcm
- Find out more at www.archer2.ac.uk/ecse/



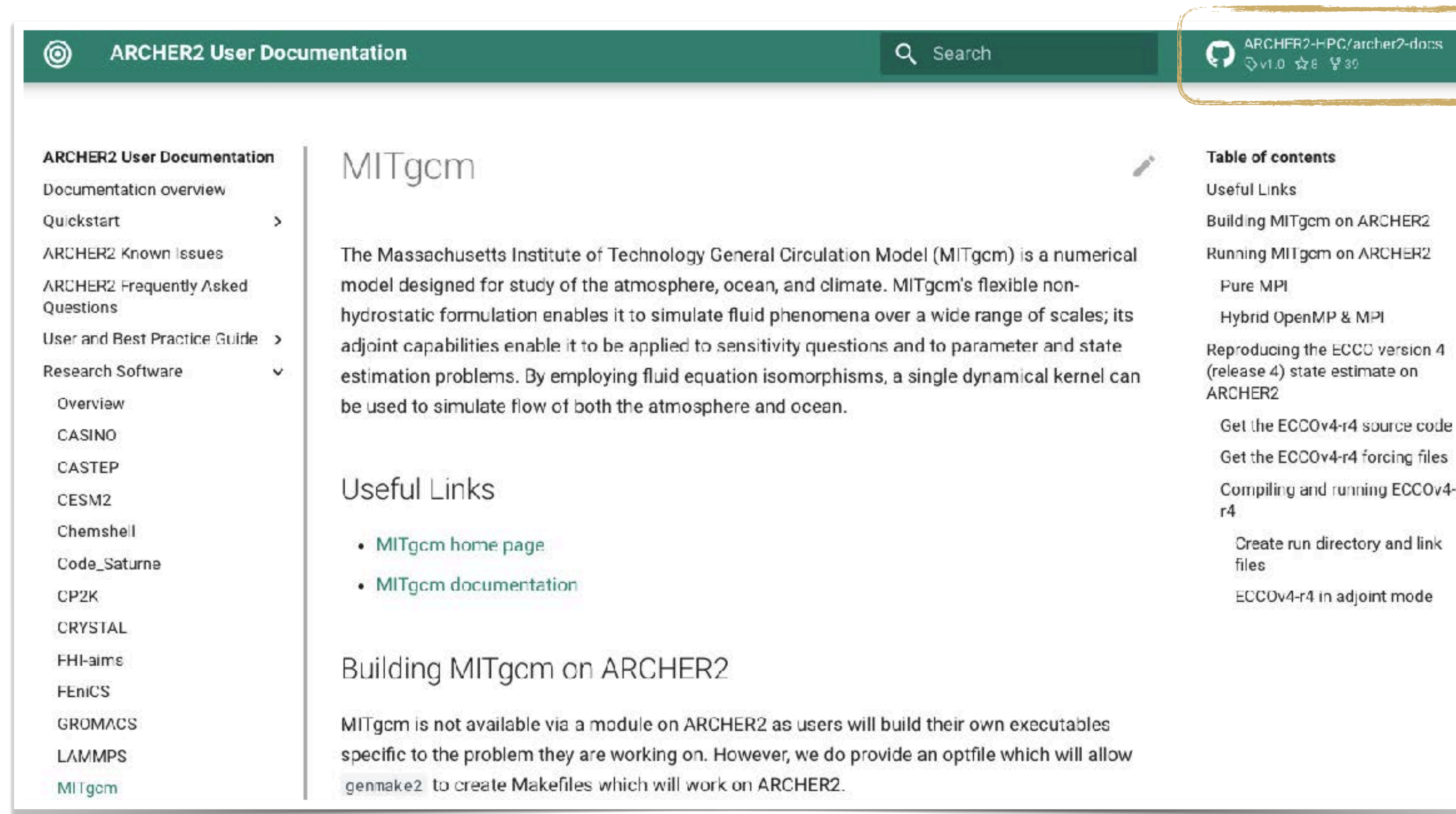
OUTLINE

- Getting started with MITgcm on ARCHER2
 - Installing
 - Running
- Tips and tricks for optimal running
 - Cheap vs fast runs
- Advanced usage:
 - Job chaining for long runs
 - Adjoint models
 - Containerisation for bundling extra code
- OptClim - parameter optimisation for climate models



GETTING STARTED WITH MITGCM ON ARCHER2

- All this and more is covered in the ARCHER2 documentation:
docs.archer2.ac.uk/research-software/mitgcm/



The screenshot shows the ARCHER2 User Documentation website. The header is green with the title 'ARCHER2 User Documentation' and a search bar. A sidebar on the left lists various documentation sections, with 'Research Software' expanded to show a list of software packages including MITgcm. The main content area is titled 'MITgcm' and contains a description of the model, a 'Useful Links' section with links to the MITgcm home page and documentation, and a 'Building MITgcm on ARCHER2' section. A right sidebar contains a 'Table of contents' and 'Useful Links' for the MITgcm page. A GitHub repository link for 'ARCHER2-HPC/archer2-docs' is highlighted in the top right corner of the screenshot.

ARCHER2 User Documentation

Search

ARCHER2-HPC/archer2-docs v1.0 ⭐ 8 🗨 39

ARCHER2 User Documentation

- Documentation overview
- Quickstart >
- ARCHER2 Known Issues
- ARCHER2 Frequently Asked Questions
- User and Best Practice Guide >
- Research Software ▾
 - Overview
 - CASINO
 - CASTEP
 - CESM2
 - Chemshell
 - Code_Saturne
 - CP2K
 - CRYSTAL
 - FHI-aims
 - FEniCS
 - GROMACS
 - LAMMPS
 - MITgcm

MITgcm

The Massachusetts Institute of Technology General Circulation Model (MITgcm) is a numerical model designed for study of the atmosphere, ocean, and climate. MITgcm's flexible non-hydrostatic formulation enables it to simulate fluid phenomena over a wide range of scales; its adjoint capabilities enable it to be applied to sensitivity questions and to parameter and state estimation problems. By employing fluid equation isomorphisms, a single dynamical kernel can be used to simulate flow of both the atmosphere and ocean.

Useful Links

- [MITgcm home page](#)
- [MITgcm documentation](#)

Building MITgcm on ARCHER2

MITgcm is not available via a module on ARCHER2 as users will build their own executables specific to the problem they are working on. However, we do provide an optfile which will allow `genmake2` to create Makefiles which will work on ARCHER2.

Table of contents

Useful Links

- Building MITgcm on ARCHER2
- Running MITgcm on ARCHER2
 - Pure MPI
 - Hybrid OpenMP & MPI
- Reproducing the ECCO version 4 (release 4) state estimate on ARCHER2
 - Get the ECCOv4-r4 source code
 - Get the ECCOv4-r4 forcing files
 - Compiling and running ECCOv4-r4
 - Create run directory and link files
 - ECCOv4-r4 in adjoint mode

ARCHER2 documentation is open source – fork it from GitHub and add your own useful info!



GETTING STARTED WITH MITGCM ON ARCHER2

- All this and more is covered in the ARCHER2 documentation:
docs.archer2.ac.uk/research-software/mitgcm/
- Install MITgcm in your **home** directory, e.g. /home/n01/n01/emmomp
- Best practice is to clone from git:
“git clone <https://github.com/MITgcm/MITgcm.git>”
- Compile using our opt file “dev_linux_amd64_cray_archer2”

```
[emmomp@ln03:~> ls MITgcm/tools/build_options/  
bgl_gnu_ncar          darwin_ppc_xlf_panther_baylor    linux_amd64_gfortran_greenplanet  linux_amd64_pathf90  
bgl_ncar              darwin_ppc_xlf_panther+wienders  linux_amd64_ifort                  linux_amd64_pathf90+redhatlam  
cygwin_ia32_g77       darwin_ppc_xlf_tiger_baylor      linux_amd64_ifort11                linux_amd64_pgf77  
darwin_absoft_f77     dev_linux_amd64_cray_archer2     linux_amd64_ifort_beagle           linux_amd64_pgf77+mpi_ncar  
darwin_amd64_gfortran dev_linux_amd64_gfortran_archer2  linux_amd64_ifort_discover         linux_amd64_pgf77+mpi_xd1
```



GETTING STARTED WITH MITGCM ON ARCHER2

- All this and more is covered in the ARCHER2 documentation:
docs.archer2.ac.uk/research-software/mitgcm/
- Install MITgcm in your **home** directory, e.g. /home/n01/n01/emmomp
- Best practice is to clone from git:
“git clone <https://github.com/MITgcm/MITgcm.git>”
- Compile using our opt file “dev_linux_amd64_cray_archer2”
- Remember to specify the number of processors (nP_x/y) in SIZE.h **before** compiling!



GETTING STARTED WITH MITGCM ON ARCHER2

- Set up a folder in your **work** directory with required namelists, inputs etc.
- If you're using large input files, place them in a shared directory like */work/n01/shared* so they don't end up downloaded multiple times.
- Copy over your executable (normally *mitgcmuv*) from your home directory.
- Create a submission script specifying options like names, nodes, time, queues etc. See examples in the ARCHER2 documentation:

```
#!/bin/bash

# Slurm job options (job-name, compute nodes, job time)
#SBATCH --job-name=MITgcm-simulation
#SBATCH --time=1:0:0
#SBATCH --nodes=2
#SBATCH --tasks-per-node=128
#SBATCH --cpus-per-task=1

# Replace [budget code] below with your project code (e.g. t01)
#SBATCH --account=[budget code]
#SBATCH --partition=standard
#SBATCH --qos=standard

# Set the number of threads to 1
# This prevents any threaded system libraries from automatically
# using threading.
```



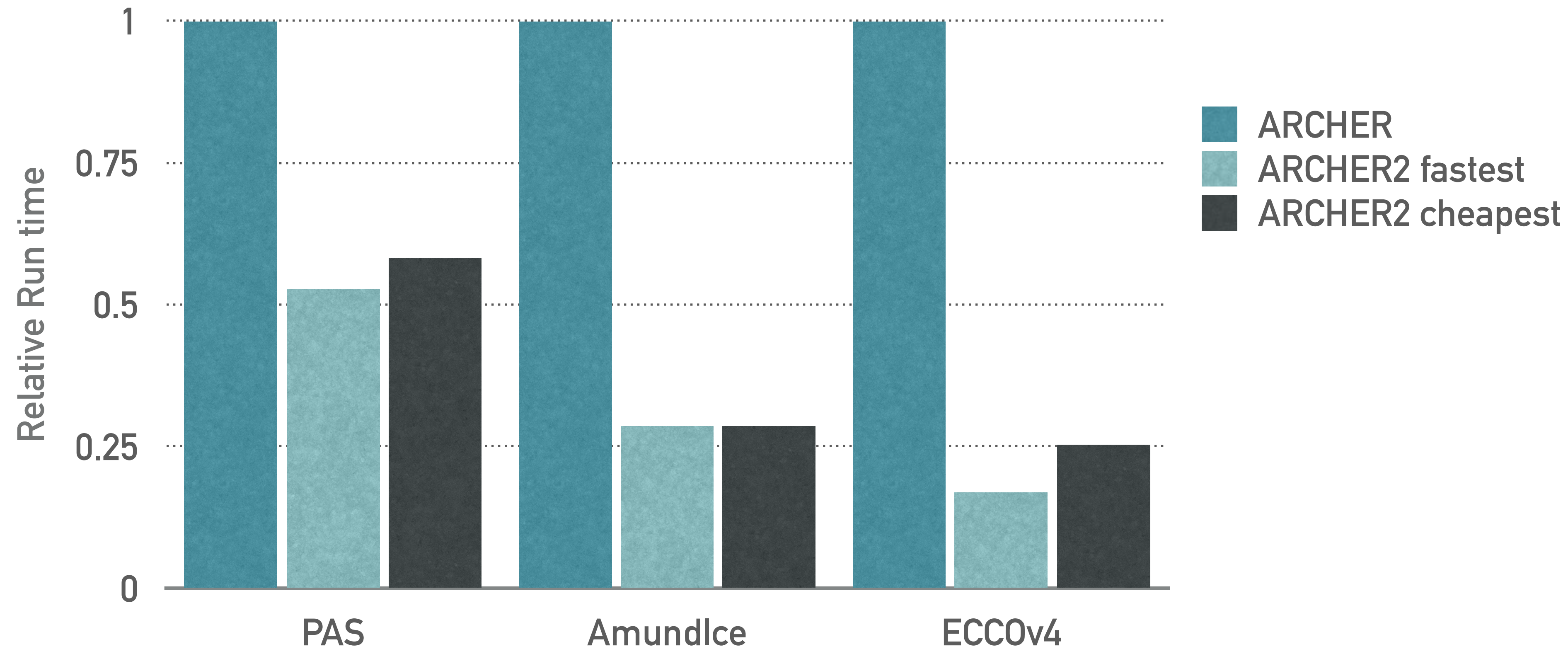
GETTING STARTED WITH MITGCM ON ARCHER2

- Set up a folder in your **work** directory with required namelists, inputs etc.
- If you're using large input files, place them in a shared directory like */work/n01/shared* so they don't end up downloaded multiple times.
- Copy over your executable (normally *mitgcmuv*) from your home directory.
- Create a submission script specifying options like names, nodes, time, queues etc. See examples in the ARCHER2 documentation.
- Submit using a command like "*sbatch run_script.slurm*"
- Aaaand Relax(!)



TIPS AND TRICKS FOR RUNNING

➤ ARCHER2 is fast!



Data courtesy of M Mineter from eCSE02-06



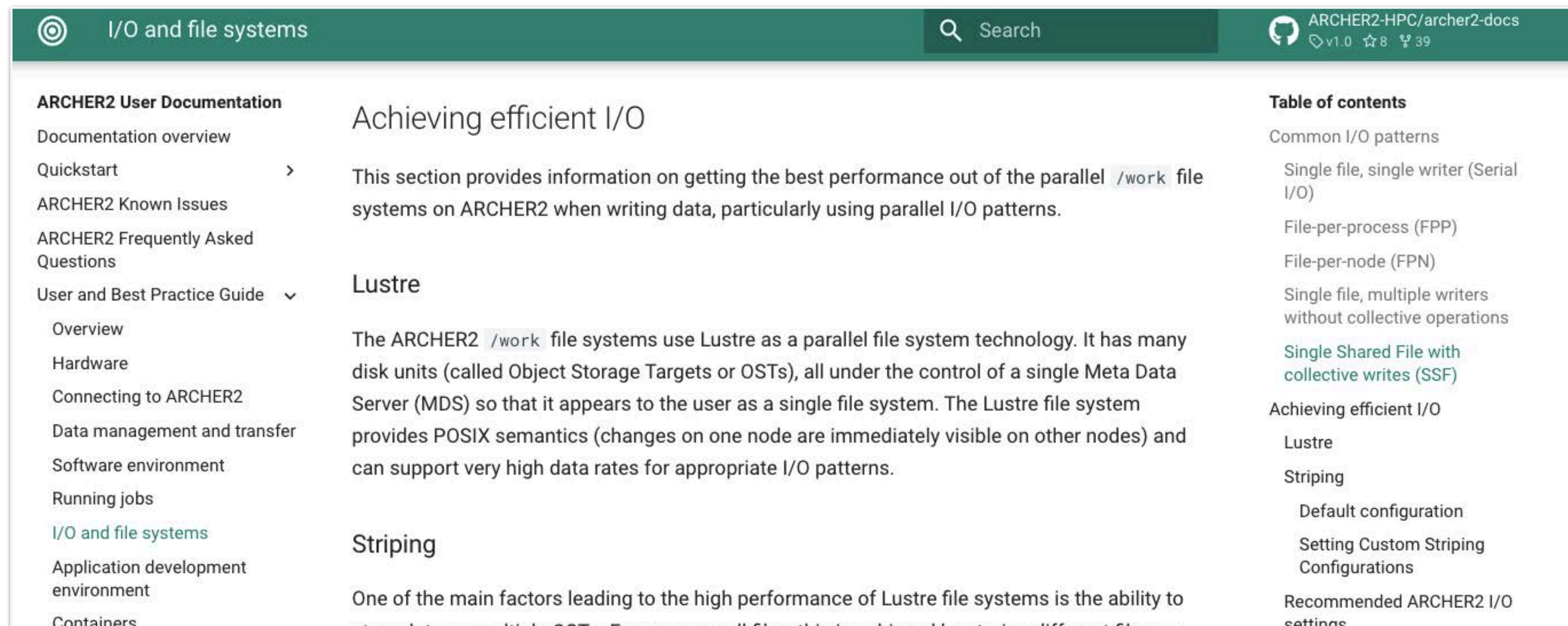
TIPS AND TRICKS FOR RUNNING

- ARCHER2 is fast!
- How fast depends on particular cases - we saw speed ups of 2-5 x over ARCHER
- Fastest runs were with **more** nodes...
- ...but beware of diminishing returns
- Cheapest runs were with **fewest** nodes
- Our options file is optimised for the test cases here, if you have a particularly complicated setup it may be worth experimenting...
- We found the Cray compiler most reliable but we also have a Gnu options file, just ask...



OPEN QUESTIONS

- Our study focused on runtime efficiency.
- I/O options remain unexplored - see the ARCHER2 documentation for some guidance.
- Please share with the community if you have useful tips!



The screenshot shows the ARCHER2 User Documentation page for 'I/O and file systems'. The page has a green header with a search bar and the repository name 'ARCHER2-HPC/archer2-docs'. The left sidebar contains a table of contents for the documentation, with 'I/O and file systems' highlighted. The main content area is titled 'Achieving efficient I/O' and contains two sections: 'Lustre' and 'Striping'. The 'Lustre' section explains that the ARCHER2 /work file systems use Lustre as a parallel file system technology. The 'Striping' section mentions that one of the main factors leading to the high performance of Lustre file systems is the ability to spread data across multiple OSTs. The right sidebar contains a 'Table of contents' for the current page, listing sections like 'Common I/O patterns', 'Single Shared File with collective writes (SSF)', 'Achieving efficient I/O', and 'Recommended ARCHER2 I/O settings'.

I/O and file systems ARCHER2-HPC/archer2-docs

ARCHER2 User Documentation

- Documentation overview
- Quickstart
- ARCHER2 Known Issues
- ARCHER2 Frequently Asked Questions
- User and Best Practice Guide
 - Overview
 - Hardware
 - Connecting to ARCHER2
 - Data management and transfer
 - Software environment
 - Running jobs
 - I/O and file systems**
 - Application development environment
 - Containers

Achieving efficient I/O

This section provides information on getting the best performance out of the parallel `/work` file systems on ARCHER2 when writing data, particularly using parallel I/O patterns.

Lustre

The ARCHER2 `/work` file systems use Lustre as a parallel file system technology. It has many disk units (called Object Storage Targets or OSTs), all under the control of a single Meta Data Server (MDS) so that it appears to the user as a single file system. The Lustre file system provides POSIX semantics (changes on one node are immediately visible on other nodes) and can support very high data rates for appropriate I/O patterns.

Striping

One of the main factors leading to the high performance of Lustre file systems is the ability to spread data across multiple OSTs. For example, all files in a directory are striped across different OSTs.

Table of contents

- Common I/O patterns
 - Single file, single writer (Serial I/O)
 - File-per-process (FPP)
 - File-per-node (FPN)
 - Single file, multiple writers without collective operations
 - Single Shared File with collective writes (SSF)
- Achieving efficient I/O
 - Lustre
 - Striping
 - Default configuration
 - Setting Custom Striping Configurations
- Recommended ARCHER2 I/O settings



ADVANCED USAGE



Job chaining: motivation

What if your simulation needs more than the maximum walltime? (24h)

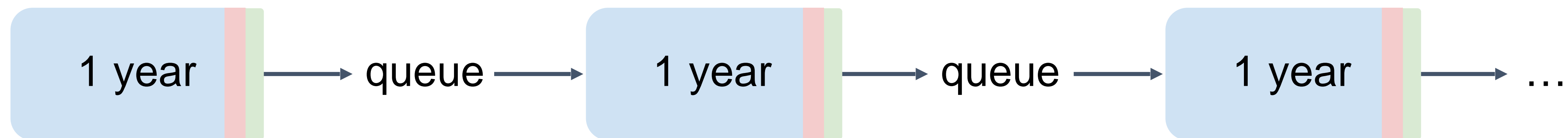


When do you decide to stop and restart?



Job chaining: old approach

Run a set length of simulation (eg 1 year) for each segment



Disadvantages:

- Have to overestimate walltime for each segment
- Run risk of not finishing a segment in time
- More time spent on model initialisation
- Queue more often, so more opportunity for bottlenecks



Job chaining: new approach

Request maximum queue length and run MITgcm for as long as possible

Stop a few minutes before the end (**timeout** utility), identify latest restart point, edit namelist, resubmit

As many years as possible in 24 hours

queue

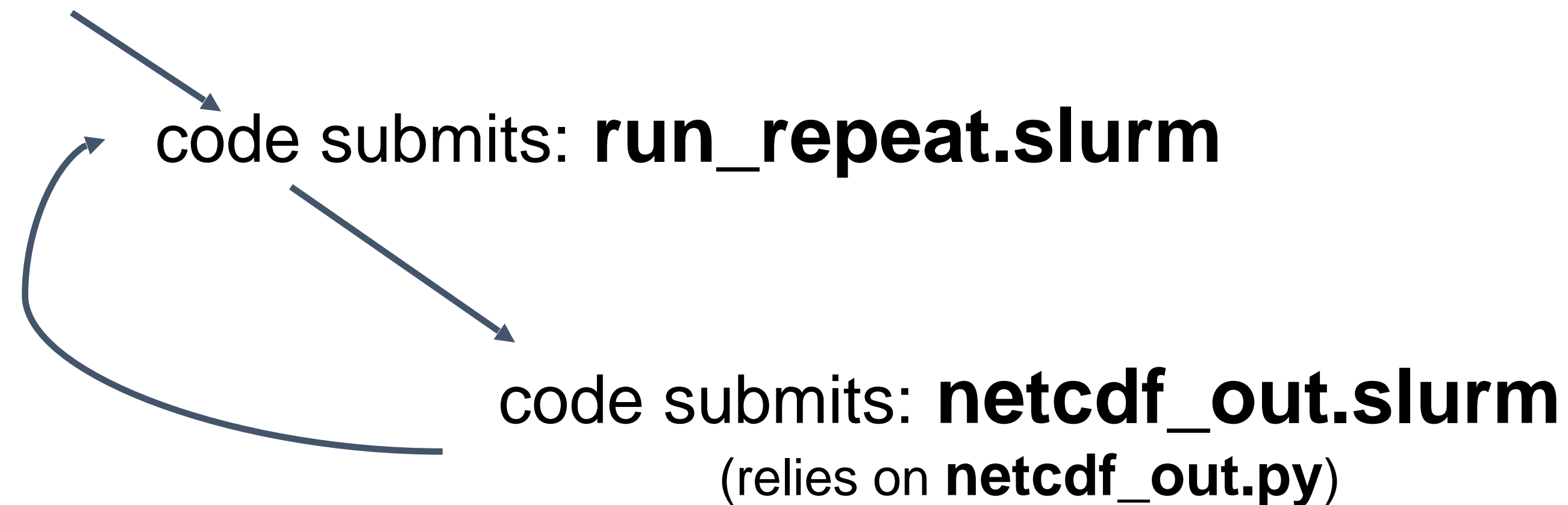
Makes most efficient use of the queue, and reduces guesswork for user



Job chaining: code structure

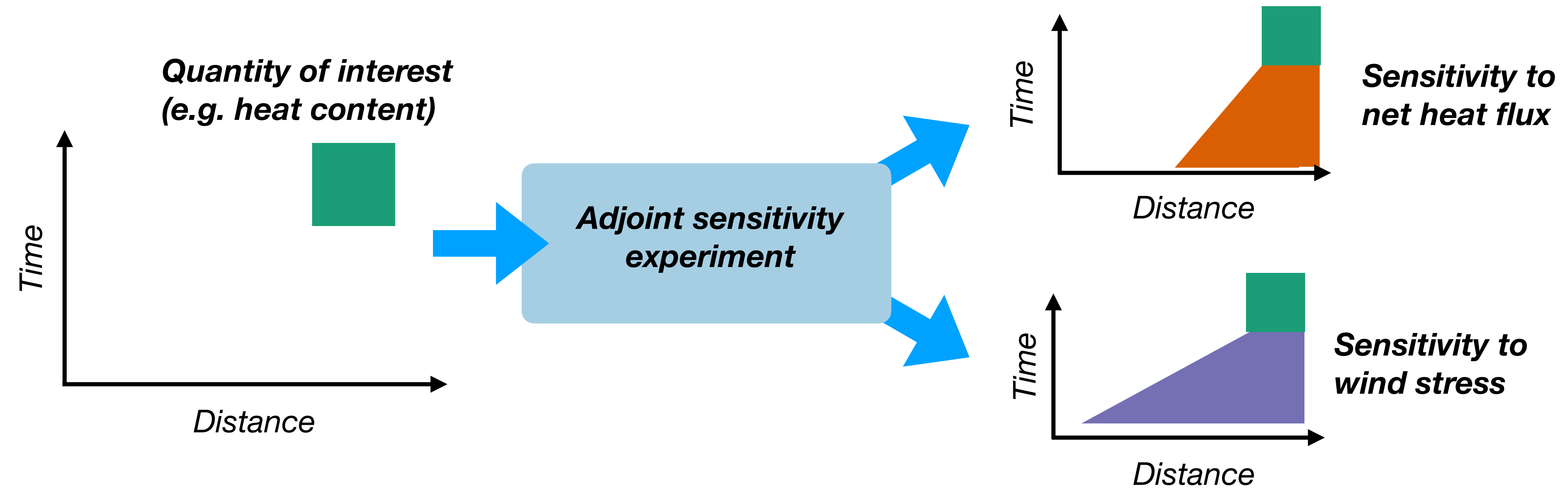
https://github.com/knaughten/UaMITgcm/tree/archer2/example/PAS_999/mitgcm_run/scripts/standalone_mit

user runs: **sub_run.sh**



ADVANCED USAGE – ADJOINT MODELS

- What is an adjoint model?
- Unlike a traditional sensitivity experiment (perturb then run fwd to see what happens), adjoints run in reverse:



ADVANCED USAGE – ADJOINT MODELS


Software to auto-differentiate your model:

- OpenAD:
 - Open Source
 - Harder to use off the shelf
- TAF:
 - Requires a license
 - Works reliably
 - Used for the ECCOv4 state estimate



ADVANCED USAGE – ADJOINT MODELS

- Follow the ARCHER2 documentation for generating the ECCOV4 adjoint, a state estimate which uses the adjoint sensitivities to minimise model-observation differences.



The screenshot shows the MITgcm website interface. The top navigation bar is dark green with the MITgcm logo and a search bar. The left sidebar, titled 'ARCHER2 User Documentation', lists various links: Documentation overview, Quickstart, ARCHER2 Known Issues, ARCHER2 Frequently Asked Questions, User and Best Practice Guide, and Research Software (which is expanded to show Overview, CASINO, CASTEP, and CESM2). The main content area is titled 'ECCOV4-r4 in adjoint mode' and contains a paragraph explaining that if you have access to the commercial TAF software from <http://FastOpt.de>, you can compile and run the ECCOV4-r4 instance of MITgcm in adjoint mode. It then instructs users to create a new code directory and a new build directory from the MITgcm/ECCOV4/release4 directory. Below this text is a code block with the following commands:

```
mkdir code_ad
cd code_ad
ln -s ../code/* .
cd ..
mkdir build_ad
cd build_ad
```



ADVANCED USAGE – ADJOINT MODELS

- Instead of “make all” -> “make adtaf” then “make adall”
- Creates *mitgcmuv_ad* instead of *mitgcmuv*
- Turn on the ECCO package for calculating your cost function/QoI
- Turn on the CTRL package to generate sensitivities
- Define your QoI in *data.ecco* following the MITgcm documentation:

Table 10.5 Implemented *gencost_barfile* options (as of checkpoint 67x) that can be used via *cost_gencost_boxmean.F* (Section 10.1.2).

variable name	description	remarks
<code>m_boxmean_theta</code>	mean of theta over box	specify box
<code>m_boxmean_salt</code>	mean of salt over box	specify box
<code>m_boxmean_eta</code>	mean of SSH over box	specify box
<code>m_boxmean_shifwf</code>	total shelfice freshwater flux over box	specify box
<code>m_boxmean_shihf</code>	total shelfice heat flux over box	specify box
<code>m_horflux_vol</code>	volume transport through section	specify transect



ADVANCED USAGE – ADJOINT MODELS

- Everything is documented in section 10 of the MITgcm documentation
- There is a dedicated ECCO mailing list that can help with problems

See ecco-group.org



Introduction to Containers: an MITgcm user perspective

Dan Goldberg
School of GeoSciences
University of Edinburgh



Overview

- ▶ Singularity containers - from a User perspective
- ▶ Use Cases:
 - ▶ Open-Source algorithmic differentiation tool
 - ▶ Visualisation on ARCHER2
 - ▶ MITgcm/PETSc on rack server

What are containers?

- ▶ Virtual Machines - Virtualise the *HARDWARE*
- ▶ Containers - Virtualise the *OPERATING SYSTEM* but interact with physical file server
- ▶ *Why?*
 - ▶ You might not have root privileges on the machine you are using
 - ▶ You might not know how to install software in the target environment, even if you can do so on your local computer
 - ▶ The target environment may not have libraries that you need
 - ▶ Reproducibility!

Docker versus Singularity

- ▶ Singularity images are *files* which can be transferred, moved, etc - while Docker images must be in a central location
- ▶ Singularity is “closer to the hardware” so may yield better performance
- ▶ Singularity is *more secure* than Docker

But

- ▶ Docker is more mature, with better documentation/support (Singularity cannot be easily installed on Mac/Windows)

Use Case -- OpenAD



- ▶ OpenAD is an *open-source, source-to-source* AD tool from Argonne National Laboratory
 - ▶ Has been applied to MITgcm - not used as extensively
 - ▶ Adjoint performance is not great - *except with STREAMICE*

Problem: OpenAD is no longer updated, and binaries cannot be compiled with gcc 5.X or later!

Note - this is distinct from using it to generate compilable source code

ARCHER2 administrators were asked to install - recommended singularity

Use Case -- OpenAD



► Strategy:

- Compile OpenAD executables in a Singularity container (*.sif file*). Requires Singularity installation on “home” computer, along with *.def file*, and “user group” permissions
- Build MITgcm-OpenAD adjoint on ARCHER2, calling OpenAD in container when necessary. Requires singularity installation and *.sif file*.
- MITgcm executable is NOT compiled in container - and hence is not run in a container.

Use Case -- OpenAD



► Resources:

- <https://www.archer2.ac.uk/training/courses/220119-containers/>
- https://docs.sylabs.io/guides/3.5/user-guide/definition_files.html
- <https://mitgcm.readthedocs.io/en/latest/autodiff/autodiff.html#adjoint-code-generation-using-openad>
- Dr Magnus Hagdorn, School of GeoSciences

```
Bootstrap: library
From: ubuntu:18.04
Stage: build

%setup
  touch /file1
  touch ${SINGULARITY_ROOTFS}/file2

%files
  /file1
  /file1 /opt

%environment
  export LISTEN_PORT=12345
  export LC_ALL=C

%post
  apt-get update && apt-get install -y netcat
  NOW=`date`
  echo "export NOW=${NOW}" >> $SINGULARITY_ENVIRONMENT

%runscript
  echo "Container was created $NOW"
  echo "Arguments received: $*"
  exec echo "$@"

%startscript
  nc -lp $LISTEN_PORT
```

```
3670 # canonicalizer
3671 ad_input_code_sf.pre.f90 : \$(CB2M_AD_FILES)
3672   \${SINGULARITYCMD} \${OPENADFORTTK_BASE}/tools/SourceProcessing/preProcess.py --timing --r8 -H -S -o \${@} \${^}
3673
3674 # replace stop statements (to avoid the implied unstructured control flow) with print statements
3675 ad_input_code_sf.pre.s2p.f90 : ad_input_code_sf.pre.f90
3676   cat \${< | sed -f \$(OADTOOLS)/stop2print.sed > ad_input_code_sf.pre.s2p.f90
3677
3678 # F -> WHIRL
3679 ad_input_code_sf.pre.s2p.B: ad_input_code_sf.pre.s2p.f90
3680   \${SINGULARITYCMD} \${OPEN64ROOT}/crayf90/sgi/mfef90 -r8 -z -F ad_input_code_sf.pre.s2p.f90
3681
3682 # WHIRL -> XAIF
3683 ad_input_code_sf.pre.s2p.xaif : ad_input_code_sf.pre.s2p.B
3684   \${SINGULARITYCMD} \${OPENADFORTTK}/bin/whirl2xaif -s -n --debug 1 -o \${@} \${<
3685
3686 # XAIF -> XAIF'
3687 ad_input_code_sf.pre.s2p.xb.xaif : ad_input_code_sf.pre.s2p.xaif xaif.xsd xaif_base.xsd xaif_inlinable_intrinsics.xsd xaif_deriv
3688   \${SINGULARITYCMD} \${XAIFBOOSTERROOT}/xaifBooster/algorithms/BasicBlockPreaccumulationReverse/driver/oatDriver -f -t fo
```

SINGULARITYCMD = singularity exec \${SINGULARITYFILE}

Use Case -- OpenAD

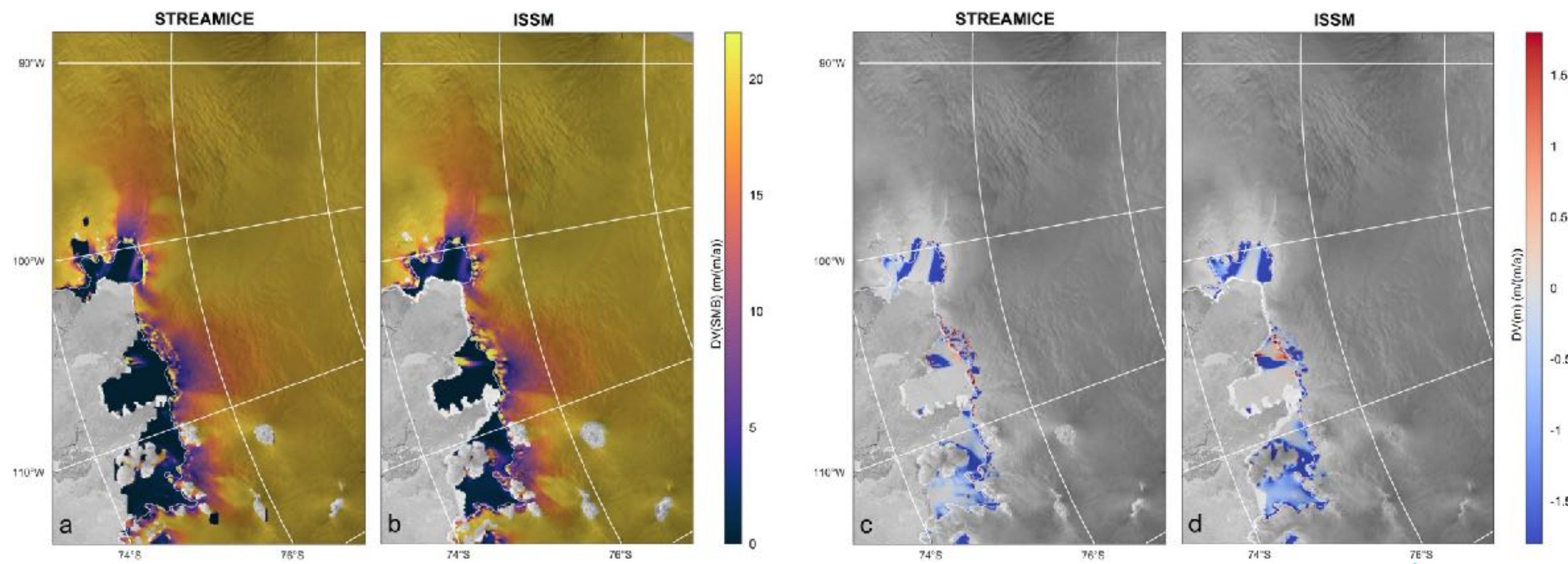


.... And then science!

Geophysical Research Letters*

Research Letter | Full Access

Mapping the Sensitivity of the Amundsen Sea Embayment to Changes in External Forcings Using Automatic Differentiation



Other use cases: `display`

- ▶ I like to check progress of ARCHER2 batch by plotting diagnostics
- ▶ Problem - was unable to generate interactive figures with IPython on ARCHER2 login nodes (unable to set correct "backend" in matplotlib)
- ▶ Can save figures to file - but where is `display`???
- ▶ Correct solution: email helpdesk. My solution: install `display` in a container, move image to ARCHER2

```
dispfunc() {  
    singularity exec -B $PWD display.sif display $1  
}
```


Other use cases: MITgcm on rack server

- ▶ The catch - you need a native installation of MPI, and the image must be "consistent" with the MPI installation (i.e. mpich→mpich)

```
singularity exec -B [file share] mitgcm.sif MITgcm/tools/genmake2 -mods=../code  
-mpi=/opt/mpi
```

```
singularity exec -B [file share] mitgcm.sif make depend
```

```
singularity exec -B [file share] mitgcm.sif make -j 8
```

```
mpirun -n 24 singularity exec -B [file share] mitgcm.sif ./mitgcmuv
```

OR

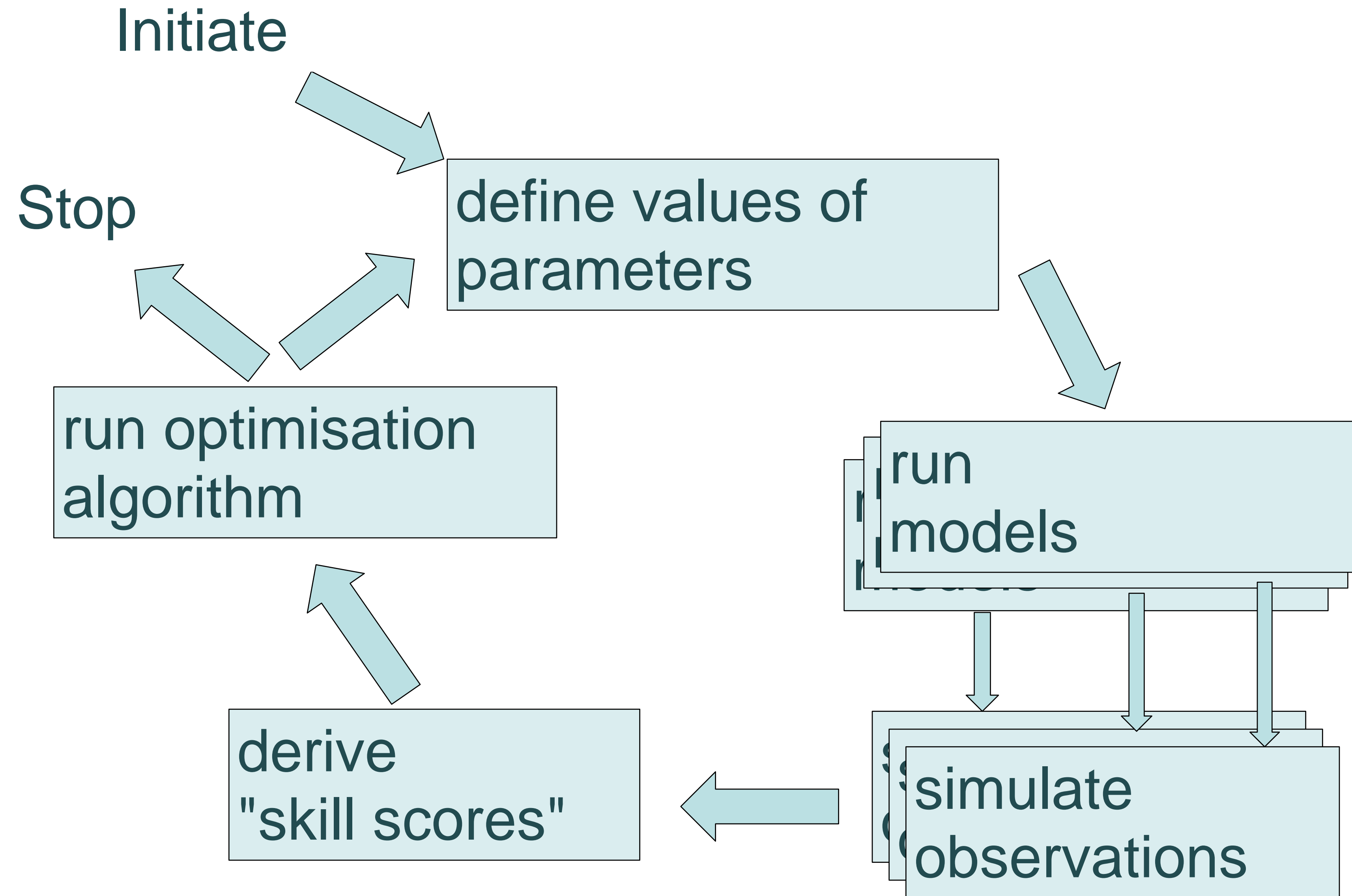
```
singularity exec -B [file share] mitgcm.sif mpirun -n 24 ./mitgcmuv
```

OptClim: software to optimise models

Mike Mineter, Simon Tett
GeoSciences, University of Edinburgh

Coralia Cartis
Mathematics, University of Oxford

m.mineter@ed.ac.uk



OptClim

- Ported to ARCHER2
- Explores parameter values to optimise a model against observation
- Works with MITgcm, CESM, UKESM
- Ask for more information
m.mineter@ed.ac.uk