

# Modern HPC Tools

**PEARC 2023**  
*July 2023*

**PRESENTED BY:**

***Robert McLay***  
***TACC HPC***

# Please Note:

- These modern tools are portable and effective on almost all supercomputers around the world.
- Almost all tools covered in this presentation are open-source. Most of them can be installed without system administrator privileges.
- Please feel free to ask for access or installation assistance when necessary.
- Demos and labs are available to all attendees.

# Tools Make a Difference on HPC

- Great tools make a profound difference
- Require less effort to achieve some desired goals
- Save a lot of time and energy for both new and experienced users
- Enhance the user experience particularly on large-scale systems
- Under active development by experienced TACC members
- (Most of them) Available under open source licenses to public

# Outline

- User Environment  
Lmod, SanityTool
- Workflow Assistance  
ibrun, Launcher, Launcher-gpu, Pylauncher
- Job Monitoring  
core\_usage, show\_affinity, amask,
- Runtime resource monitoring  
Remora

# **Lmod**

## **Manage your environment on a Supercomputer**

# User Environment (1)

There are **environment variables** for defining values used by the shell (e.g., bash, tcsh) and programs executed on command line.

An **environment management package** provides a command-line interface to manage the collection of environment variables associated with various software packages, and to automatically modify environment variables as needed.

# User Environment

- Environment Variables (mostly)
  - PATH (where to find command)
  - MANPATH (where to find help)
  - LD\_LIBRARY\_PATH (where compilers find libs, like MKL, etc.)
  - Package environment variables (TAU\_METRICS, etc.)
  - Site environment variables for package (TACC\_NETCDF\_LIB)
- Functions and aliases
- Other possibilities: anything “unixy”

# Lm d



# Lmod

- A Lua based module system
- A convenient way to dynamically change the users' environment through modulefiles.
- Add or remove environment variable easily
- Handle MODULEPATH hierarchical problem for complicated user environment
  - Only have one version active
  - Only load one compiler or MPI stack at a time

# Basic Module Commands (1)

# List the modules already loaded

\$ **module list**

# Show what modules are available to be loaded

\$ **module avail**

# Load a package

\$ **module load matlab**

# Unload a package

\$ **module unload matlab**

# Basic Module Commands (2)

# Change from impi to mvapich2

\$ **module sw impi mvapich2**

# Go back to an initial set of modules

\$ **module reset**

# Access a modulefile's help

\$ **module help lammps**

# Show the description section of a module

\$ **module whatis petsc**

# ml: A Convenient Tool

# This means module list

\$ ml

# Module load and unload

\$ ml matlab

\$ ml -matlab

# Do it in one single line

\$ ml netcdf hdf5 -gsl

# Save/load Your Own Collection (1)

# Save the designed collection of modules

\$ module save

# Restore the designed collection

\$ module restore

# List the collections

\$ module savelist

# Save/load Your Own Collection (2)

Users can have as many collections as they like.

# Save to a named collection

\$ module save my\_collection

# List the contents of a collection (default)

# module describe

# Restore that named collection with

\$ module restore my\_collection

# Define and Use Your Own Modulefiles

Define your own module files

- Start with an existing modulefile
- Easy to share with your colleagues

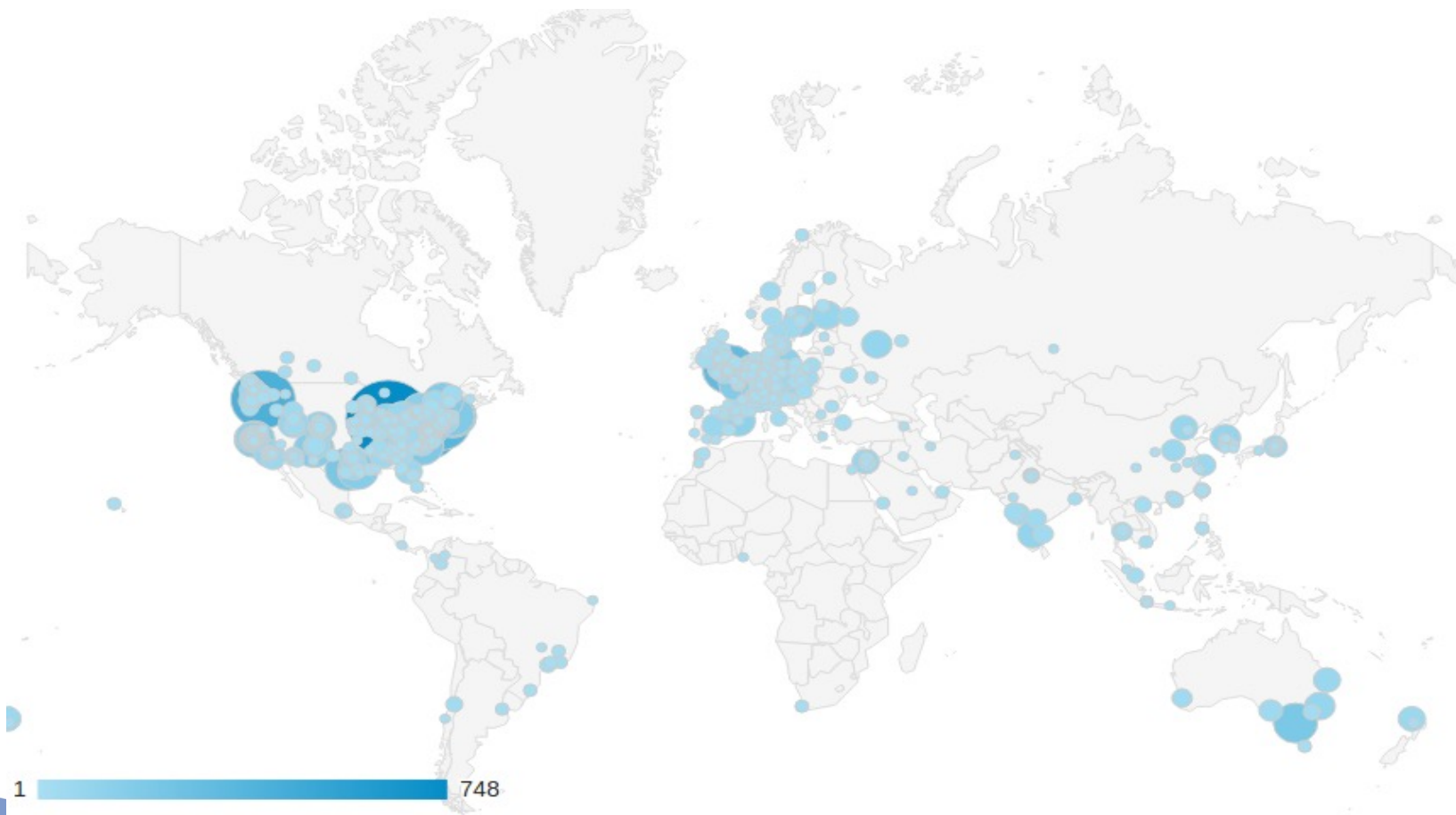
```
$ module use /scratch1/01255/siliu/mvapich2/modulefiles/
```

```
$ module load mvapich2-test/2.x-intel19
```

# Create Your Own Modulefile

- Start with existing one built by a module expert
- An introduction of writing Modulefiles:  
[https://lmod.readthedocs.io/en/latest/015\\_writing\\_modules.html](https://lmod.readthedocs.io/en/latest/015_writing_modules.html)
- Mkmod: A tool automatically creates a modulefile  
<https://github.com/milfeld/mkmod>





# Lmod References

- Lmod Documentation  
<https://lmod.readthedocs.io>
- Monthly discussion via Zoom: See  
<https://github.com/TACC/Lmod/wiki/home>
- TACC/Lmod on github  
<https://github.com/TACC/Lmod>

# **SanityTool**

## **Make my user environment valid**

# Why SanityTool

- Improper or incorrect user account configurations slow down/impede work progress
- These problems could be difficult to detect (or remember), but not difficult to fix most of the time
- There are so many tools and scripts at each site, each focusing on a few tests

A lightweight integrated tool to diagnose and resolve these problems is necessary

# SanityTool

- A lightweight generic and integrated tool
- Free and open-source software
- Created in a relatively standardized format
- Contains many useful and practical tests
- Can be conveniently used whenever necessary

SanityTool

# Running SanityTool

*\$ module load sanitytool*

*\$ sanitycheck --help*

*Sanity Tool Version: 2.0*

*Texas Advanced Computing Center*

*High Performance Computing Group*

*[-h, --help]            Help information*

*[-s, --silent]        Silent mode*

*[-v, --verbose]    Verbose mode (default)*

1: Check SSH permissions:

**Failed**

**Error: group permission on \$HOME will cause RSA to fail!**

**Error: other permission on \$HOME will cause RSA to fail!**

Make sure you have a .ssh directory under your \$HOME directory.

You can use the following commands to set the proper permissions:

\$ chmod 700 \$HOME # (750 and 755 are also acceptable)

\$ chmod 700 \$HOME/.ssh

\$ chmod 600 \$HOME/.ssh/authorized\_keys

\$ chmod 600 \$HOME/.ssh/id\_rsa

\$ chmod 644 \$HOME/.ssh/id\_rsa.pub

2: Check SSH keys:

**Passed**

3: Check environment variables (e.g. HOME, WORK, SCRATCH) and file system access:

**Passed**

4: Check user's queue accessibility (Stampede2 Only):

**Passed**

5: Check allocation balance:

**Warning: One of your projects 'ABC-123' has negative balance -1511.194.**

**Passed**

6: Check quota for \$HOME and \$WORK spaces:

**Passed**

7: Check module environment:

**Passed**

8: Check compilers:

**Failed**

**Error: Compiler icc is not available at this time!**

**Error: Compiler icpc is not available at this time!**

**Error: Compiler ifort is not available at this time!**

Please check your \$PATH again, compilers are missing.

If you unload the compilers on purpose, please ignore this test.

9: Check scheduler commands:

**Passed**

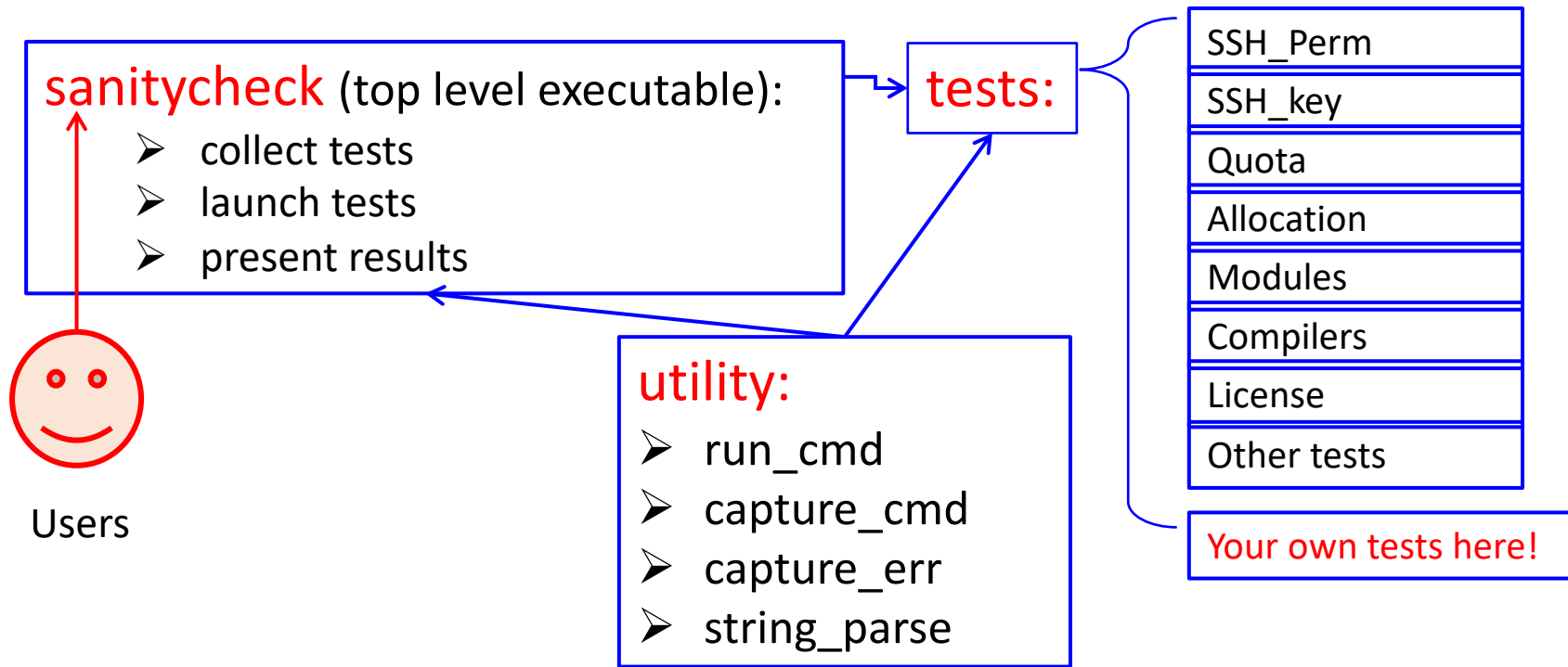
-----  
**2(out of 9) failure in sanitycheck.**  
-----

# SanityTool Features

- Applicable to almost all supercomputer systems  
(and personal computers)
- Independent of system configurations or settings
- Work for different kinds of shell (bash, csh, zsh, etc.)
- Easy for users to remember and run
- Flexible to be run almost any time
- Full of practical tests (and still extending)



# Overall Design



# Currently Supported Tests

## Generic Tests:

- Valid ssh configurations
- File system accessibility
- Proper permission of file systems
- Usage and quota of file systems
- Necessary software licenses
- Current module environments
- ... ..

## Customized Tests:

- Necessary preloaded modules
- Necessary compiler commands
- Necessary scheduler commands
- Whether the user is blocked
- Users' allocations and balance
- Permission to access to protected data
- ... ..

# Customized Testset

The new sanitytool version 2.0 allows users to create/use their own tests.

```
$ sanitycheck -t mytestdir
```

Create your own test case as simple as:

```
def execute(self):  
    Flag=True  
    output=capture("type h5copy")  
    if "not found" in output:  
        Flag = False  
        self.error_message+="      ERROR: h5copy is not available!"  
    return Flag
```

# Obtain SanityTool

- Obtain the source code of Sanity Tool  
<https://github.com/siliu-tacc/sanitytool>
- Make sure “python” and “sanitycheck” are accessible
- Go through the tests directory and choose proper tests
- Add more tests modules when necessary
- Run the “*sanitycheck*” command

# **1st hands-on/homework session: LMOD and SanityTool**

# LMOD Lab (A):

- Display all available modules on the Frontera system
- View the help information for any specific module if necessary
- Load a few modules you will need for your research
- Make the new collection as the default

# LMOD Lab (B):

- Learn more about the Mvapich2 module
- Run “`echo $MPICH_HOME`”
- Switch to mvapich2 from impi
- Run “`echo $MPICH_HOME`” again

# Sanitytool Lab:

- Load the “sanitytool” module
- Run “sanitycheck” in your account
- Run “whyblockme” in your account
- Load the “sanitytool” module
- “unset SCRATCH” and run “sanitycheck” again