

# Improving NeSI researchers' productivity with a consultancy service

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# Outline

- ① NeSI Consultancy service
- ② Some recent projects
- ③ Summary



# What is NeSI Consultancy?

## Introduction

- Goals of NeSI's Consultancy service
  - Enable increased scale and complexity of research
  - Help researchers use NeSI's computing capabilities to increase research outputs and impacts
- How we achieve this
  - Working on projects with researchers; typically 1 day per week for 3-4 months
  - The scope of projects we work on is very broad and covers topics such as:
    - Code optimisation and/or parallelisation
    - Workflows
    - Improving software sustainability
    - Porting code to NeSI's platforms
    - Custom code development
    - Visualisation

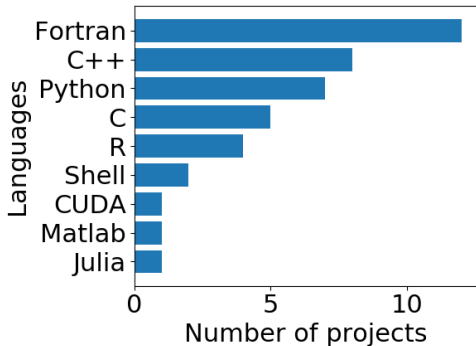
# Consultancy projects

## Applying for a project

- Usually researchers who meet the following criteria can access consultancy at no cost to themselves, based on their institution's or MBIE's investment into NeSI:
  - Researchers with Merit or postgrad projects
  - Researchers based at collaborator institutions
    - University of Auckland
    - NIWA
    - University of Otago
    - Manaaki Whenua - Landcare Research
- Others may be able to pay to access the service
- Application form
  - Usually filled out after an initial meeting
  - Defines the current state of the problem
  - Defines the scope of the work

# Consultancy projects

## Languages



- The programming languages that we have worked with
- Some projects had more than one language
- Fortran, C++/C still dominate
- Python and R are popular too
- We started work on our first project with Julia recently



# Getting closer to more accurate climate predictions for New Zealand

## Researchers:

**Erik Behrens & Jonny Williams**

*National Institute of Water & Atmospheric Research (NIWA)*

**“The NZESM is a community project. With this upgrade we have an exciting tool with which to study the climate.”**

## NeSI delivered:

- High performance computing resources
- Computational science expertise





# Code development

NZESM project

- Multiple models that need to be coupled together (atmosphere, ocean, . . .)
- Using a tool called OASIS for coupling these models together
- We added a new feature to OASIS to enable an ocean coupling scheme for coupling a high resolution regional ocean component
- Important step to setting up customised earth, ocean and weather system modelling for New Zealand





# Cascade trajectories project

- ^ Infrastructure
  - ^ Added a CMake build system
    - ^ Easy to port to different systems
    - ^ Easy to try different compilers Cray compiler 40% faster
  - ^ Adding tests
- ^ Optimisation
  - ^ Optimised performance of serial Fortran code
    - ^ 22 % boost
- ^ Work flows
  - ^ Instead of running one long calculation they can run many shorter calculations
  - ^ Created Python/Slurm scripts to enable this parallel workflow and give a big boost in performance



# Fractal analysis project

## Code optimisation

- ^ Matlab code
- ^ Identified two functions with big loops that were performance bottlenecks
- ^ Converted these functions to C++ and call from Matlab mex'ing
- ^ Total speedup of 8x from converting these functions
- ^ Other outcomes
  - ^ Taught the researcher how to use git version control
  - ^ Importance of testing adding a checksum to easily check result

# Trinity project overview

## Work ows and runtime tuning

- ^ Genomics code - de novo assembly
- ^ Complex work ow consisting of two distinct computational phases
  - ^ Initial phase is multithreaded with high memory requirements (100s GB)
  - ^ Second phase has many millions of short, serial embarrassingly parallel jobs with low memory requirements ( 5 GB)
- ^ By default Trinity runs everything at once within a single high memory node not optimal

# Trinity project solution

Work ows and runtime tuning

- ^ HPCGridRunner launch embarrassingly parallel jobs on a compute grid (cluster)
  - ^ Accumulated I/O bandwidth of many nodes gives much better performance

Type	Num cores	Elapsed time (hh:mm:ss)	Core hours
Single node	16	24:09:36	387
Grid	20	07:59:58	168
Grid	40	04:10:45	171
Grid	60	02:36:58	160

The NeSI consultancy service has drastically improved my work ow by reducing RNA de novo assembly time from 7 days to 24 hours  
Alexis Marshall, University of Waikato





