HPC User Group Meeting

September 21st, 2022

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Ask SCDA

SCDA Answers

Storage Request System

We now have a process!

Saion Update

New

system version!

New

module command!

New

GPU job settings!

The Bioinformatics Group Does it Again!



Refreshed Saion

Changes:

- OS updated to CentOS 7.9
- module system now same as Deigo ('ml' command now works)
- X11-related issues on the nodes are fixed
- Newer CUDA versions are available

Major Change:

We now limit GPUs per user

- 4/8 GPUs per user
- 36 CPU cores per user
- 7/2 Days per job



Refreshed Saion

- The lack of GPU limits was a problem with old Slurm versions
- Very few users have ever used more than 8 GPUs at once
- Most GPU jobs also need CPU
 - they will speed up with more CPU cores available than before

Major Change:

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- 36 CPU cores per user
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Ask SCDA

- IT section and SCDA have both grown
- We work differently, have different requirements
 - → We need our own issue tracker



ask-scda@oist.jp

https://ask-scda.oist.jp



SCDA Answers

- Work in progress
- Right now: Desktop scientific computing
- In the future there will be more

https://scda-answers.oist.jp



Storage Requests

- Bucket is our main storage
 - Initial allocation is ~50TB
 - Can be increased if needed
- Specialized storage solutions sometimes possible
- Other storage is fixed
 - can not be increased



- Today requests are ad-hoc:
 - Requests are informal
 - very short timeframe
 - No concern about data management, data sources etc.

This is a problem



Belief:

We have spare storage

adding storage is easy

keeping data is free



Belief:

We have spare storage

adding storage is easy

keeping data is free

Reality:

- We can't buy "spare" storage
 - everything needs to have concrete justification
 - future projected need
 - new unit allocations
 - Based on what you tell us



Belief:

We have spare storage

adding storage is easy

keeping data is free

Reality:

- New storage is a large capital investment
 - Needs scientific justification
 - Acquisition follows OIST procurement rules and calendar
 - → takes ~months to a year



Belief:

We have spare storage

adding storage is easy

keeping data is free

Reality:

- Storage is a consumable
 - Hardware has ~5 year lifetime
 - Needs maintenance
 - Must be replaced periodically
- Storage is difficult, critical
 - specialists, maintenance contracts are necessary



Belief:

Reality:

We have spare storage

→ •

- all capacity is accounted for
 - existing and future units

adding storage is easy



- adding capacity takes months
 - need scientific justification

keeping data is free



- keeping data is expensive
 - maintenance, replacement



Data Management

Data at scale is a research resource

Small data:

- Desktop computer
- Bench loupe
- Rotation student
- → simple to get little planning simple management

Big data:

- Cluster
- Electron microscope
- Project research group
- → complex, expensive lots of planning heavy management burden



A new Workflow

1.Fill in the storage request form:

https://groups.oist.jp/scs/request-storage

- 2. Wait for us to respond:
 - Your request is easy to accommodate:
 - → you get the allocation
 - Your request needs system changes:
 - → Discuss with us, wait for implementation
 - Your request can't be fulfilled
 - → Talk with us to find a different solution

Fill in the Form

- You need to be the unit leader to ask for storage
- You can't ask for more storage on the in-cluster systems Flash or Work
- We ask you three questions; tell us as much as you can, and as much you think is needed
- Also please clean up and archive data in your existing allocation



#1: Storage Amount

- How much do you need?
- When do you need it?
- Are there any specific conditions?

What storage system is appropriate? How can we provide it to you?

- Be concrete
- Include future need if you know
- be as early as possible
 - → but we only increase when you're actually running out...

Specific Conditions:

- could be anything
- no guarantee that we can fulfill it
 - → don't ask unless important



#2: Data Plan

- What kind of data, and how much?
- It comes from where? How will you use it?
- How long do you need to store it?
- What will happen with it at the end?

Determine data flow and needed connectivity. Storage retention estimations.

- What other systems are you going to connect with
- What kind of bandwidth, intermediate storage needed

- Storage is not free over time
 need to know the time frame
- Do you know what to do with the data when project ends and people leave?



#3: Research Project

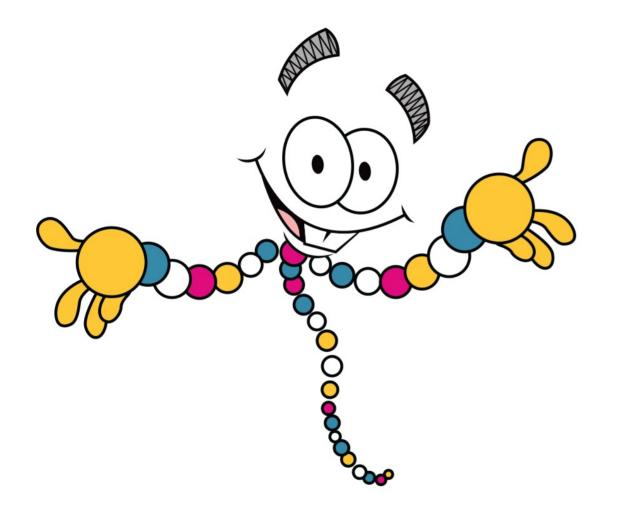
- Project title or description
- Ignore this for small or recurring request
- If you do have more info, you're very welcome to give it to us!

Justify our budget requests. Show the scientific impact of HPC investment.

Goal: Show how this helps OIST mission

- Be brief
- not compulsory use your own best judgement





OIST Bioinformatics User Group



OIST's Bioinformatics User Group



Why?

- Targeted (bio)informatic tools (<u>DebianMed</u> collection, <u>Other</u>, AlphaFold, nanopore basecall)
- Access to wide range of curated bioinformatic pipelines: OIST's profile for <u>Nextflow</u>
- Overcome unit's storage limits for installation of databases (NCBI's nt, nr, etc.)

ssh deigo

load modules
ml bioinfo-ugrp-modules
load DebianMed
ml bioinfo-ugrp-modules DebianMed

list Other softwares (= out of DebianMed distribution)
ml av Other
list databases
ml av DB

ssh saion

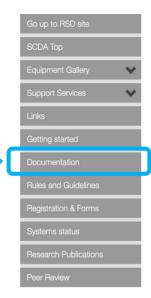
load modulesmodule bioinfo-ugrp-modules
module available

(AlphaFold, Guppy, Rerio,...)

Where to find us?

On SCDA's Documentation:

https://groups.oist.jp/scs/documentation



Scientific Computing & Data Analysis Section

Documentation

Your Own Live Information Page:

HighSci

The Deigo Cluster Getting Started

Open Hours

15:30-17:30 every day

The User Survey results are in

Software Resources

- Running Comsol jobs
- Running Matlab jobs
 - GPU with Matlab
- Running Mathematica jobs
- Software on Sango
 - Software usage notes
- Desktop HPC Software

Repository GitHub:

https://github.com/oist/BioinfoUgrp

The OIST Bioinformatics User Group

Local version control systems (Obsolete)

How to contact us?



Via Microsoft <u>Teams</u> (looking at alternatives..)



OIST's bioinformatics user group

Communication channel

Prioritized communication channel is on Microsoft Teams: BioinfoUgrp Do not hesitate to use the ping function (putting @ and then the name, like in other chat systems), because the discussions on the Team app are a bit easy to miss otherwise. Please "Google" the issues prior to contacting us. Very often, the main issues will already be reported and the solution available on the reference webpage of the program: in the Issues tab of GitHub for some, in GoogleGroups for others (e.g. for IQ-TREE). Other great platforms are StackOverflow, or Biostars.



Bimonthly meetings of the BioinfoUgrp

What: afternoon of Q&A

When: November-December

To be announced on Teams (?)

This and That

Comspace

- Share data between units at OIST
- Restricted data for a subgroup of lab members
- 5TB/unit
- GUI management

New Software

Geneious Prime now available

Copyright Infringement

A researcher was caught sharing movies using the OIST systems

Storage is *only* for research data!

- no copyrighted data
- no PC backups
- no personal files
- no unit administrative documents



Software Week!

https://tida.oist.jp/announcements/software-week-back-oct-4th-oct-7th

Learn about your software!

One-on-one help and advice directly from the vendor!

Amira Tue 4th, Wed 5th

Matlab Wed 5th

COMSOL Fri 7th

Introduction to HPC and Scientific Computing

Tuesday 4th and Thursday 6th 13-15, C700 or Zoom

Advanced Blender!

Wednesday 5th 13-15, C700 or Zoom

