Computing cluster of the Institute of Botany, Czech Academy of Sciences

Overview, features, usage, discussion

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n Login

New research tool

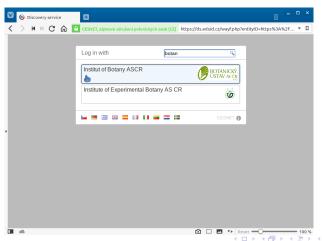
- Purchased in 2019 from grant of the Czech Academy of Sciences (almost 3 500 000 CZK)
- Part of MetaCentrum national grid infrastructure connecting computing clusters
 - Maintained by MetaCentrum admins
 - All software and other resources available (as for any MetaCentrum computing node)
 - Can be used by any MetaCentrum user
- Connected together by 100 GB network cables
- Connected to the Internet (and rest of MetaCentrum) by optical fibre





Login to all MetaCentrum/CESNET services

- Login to all MetaCentrum/CESNET services, e.g. MetaCentrum computing grid, ownCloud or FileSender is done via EduID
- Logins are redirected to EduID login page



ction Overview How to use Limits Discussion Login

Login for employees of the Institute of Botany, CAS

 Our employees are from EduID login page redirected to custom login page (different for every institution)

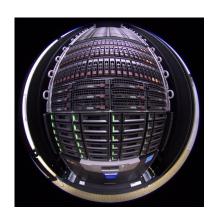


 User without password must first go to https://praha. verso.eis.cas.cz/ and obtain new login name and password



Hardware overview

- Data storage and file server tilia-nfs
- HPC nodes for massive parallelization draba1-3
- 'Standard' computing nodes carex1-6
- Frontend tilia
- Database server sorbus



Overview Computing nodes Frontend Database server Spark

Disk system

- File server tilia-nfs.ibot.cas.cz (alias storage-pruhonice1-ibot.metacentrum.cz)
- Disk array with capacity 179 TB



HPC nodes for massive parallelization

- draba1, draba2 and draba3, each equipped with 4 CPU Intel Xeon Gold 6230 (4x 20 cores (4x 40 threads), 2.1 GHz, turbo 3.9 GHz), 1536 GB RAM and 1920 GB NVMe disk (RAID 0)
- Primarily intended for running computations requiring massive parallelization such as Apache Hadoop and Spark



Standard computing nodes

Each of the six nodes
 (carex1 to carex6) is
 equipped with 1 CPU AMD
 EPYC Naples 7261 (8 cores
 (16 threads), 2.5 GHz, turbo
 2.9 GHz), 512 GB RAM and
 1920 GB NVMe disk
 (RAID 0)

 Intended for any general purpose computations, running of any application





Frontend



• 4 vCPU, 2.4 GHz, 8 GB RAM

- Standard MetaCentrum frontend — mainly advantageous to log there in order to process data stored on the Průhonice data array and to send requests to the Průhonice cluster
- Home directories are at Průhonice storage
- Can be reached by SSH at tilia.ibot.cas.cz (alias tilia.metacentrum.cz)



Database server

- 8 vCPU, 2.4 GHz, 32 GB RAM and 60 GB SSD disk
- sorbus.ibot.cas.cz accessible only for dedicated users
- Mainly intended for running NoSQL databases such as Neo4j and MongoDB that are used during Apache Hadoop and Spark computations on the HPC nodes
- Can be used for any special supportive task



Apache Spark

- A computing framework for large-scale data processing in parallel
- The primary interface language is Scala but there are wrappers for java, Python and R
- Spark performs in memory computing which speeds up computation by at least x1000 compared to regular parallel implementations...it is the FUTURE
- See also https://spark.apache.org/ and https://hadoop.apache.org/



When to use Spark?

What is big-data? Data larger than 0.5 GB

- Tabular data (genetic variants, species distribution, geographic location of features...)
- Genomic data (several genomes)
- Metagenomes comparisons
- Multiple sequence alignments (more than 1000) for gene expression, phylogenetics...

Large-scale spatial data with GeoSpark, SpatialSpark, Spark-GIS

- Co-Location Pattern Mining (e.g. in Africa do lions co-locate with zebras?)
- Multivariate Spatial Patterns (e.g. species density correlations, biomass correlations...)
- Classification and clustering (soil types, types of vegetation, response to perturbations)

Bioinformatics

- Read processing (quality cleaning, trimming) (QMSpy)
- Alignment (SparkSW, SparkBWA)
- Mapping (MetaSpark)
- Metagenomics (MetaSpark, SpaRC)
- Assembly (Spaler)
- Blasting (SparkBLAST)
- General sequence analysis (variants discovery, gene expression...) (GATK-Spark)
- Phylogenetics (CloudPhylo)
- Population genetics studies (SEQSpark)



How to use Limits Discussion Connection Priority access Storage Tasks

Usage basics

- As any MetaCentrum nodes
- Knowledge of Linux command line is required...
- To copy data using graphical application like FileZilla connect with SFTP (and MetaCentrum credentials) on port 22 to tilia-nfs.ibot.cas.cz (alias storage-pruhonice1-ibot.metacentrum.cz)

```
# Connect to frontend

2 ssh USER@tilia.ibot.cas.cz # or

3 ssh USER@tilia.metacentrum.cz

4 # Copy data to/from data storage

5 scp -r local/data USER@tilia-nfs.ibot.cas.cz:~/

6 scp -r USER@tilia-nfs.ibot.cas.cz:~/remote/data ldir/

7 # To connect directly to any computing node, e.g.

8 ssh USER@carex1.ibot.cas.cz

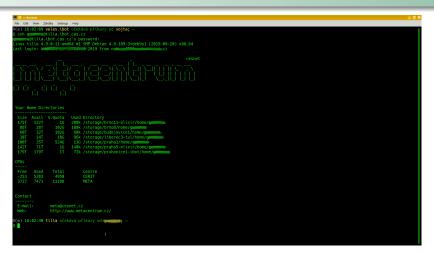
9 ssh USER@carex4.metacentrum.cz

10 ssh USER@draba2.ibot.cas.cz

11 ssh USER@draba3.metacentrum.cz
```

How to use Limits Discussion Connection Priority access Storage Tasks

SSH access



 Check e.g. PDF https://github.com/V-Z/course-linuxcommand-line-bash-scripting-metacentrum/releases/tag/v2019



o use Limits Discussion Connection Priority access Storage Tasks

Priority access to storage and computing resources

- Resources are available to all MetaCentrum users
 - Any user may send task of the length up to 24 hours (-1 walltime=24:0:0)
 - Default user quota for data storage is 10 GB
- Members of group **ibot** can submit tasks of any length requesting any resources (no limits) — tasks are running immediately as the resources are available
- Members of group ibot have default quota on the storage 2 TB (can be anything higher — mail cluster@ibot.cas.cz)
- Group ibot is available for anyone having affiliation with the Institute of Botany, or for collaborators
- To became member of the ibot group mail cluster@ibot.cas.cz
- It is possible to establish space for data shared by group of users on the storage — such requests must be send to cluster@ibot.cas.cz

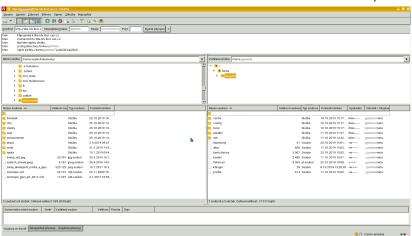


How to use Limits Discussion Connection Priority access Storage Task

Graphical connection to the storage

Use any favourite SFTP/SCP client...

 MetaCentrum USER connecting on port 22 to tilia-nfs.ibot.cas.cz via SFTP/SCP/SSHFS/rsync/...



use Limits Discussion Connection Priority access Storage Tasks

Access to data from any MetaCentrum node and from scripts

- Home of fontend tilia is on the storage tilia-nfs
- Storage is available from any MetaCentrum node in /storage/pruhonice1-ibot/home/USER/
- As any other MetaCentrum storage, just note particular path...

```
# In scripts use something like
cp -a /storage/pruhonice1-ibot/home/USER/data \
"$SCRATCHDIR"/
# to copy data to computing node, and
cp -a "$SCRATCHDIR"/results \
/storage/pruhonice1-ibot/home/USER/
# to copy results back to the storage
```

use Limits Discussion Connection Priority access Storage Tasks

Sending tasks to Průhonice cluster

- Submit task script as usually using qsub command, just add parameter -q ibot
- Try more options in https://metavo.metacentrum. cz/pbsmon2/qsub_pbspro

```
# Submit task by something like

2 qsub -l walltime=1:0:0 -q ibot \
3    -l select=1:ncpus=4:mem=4gb:scratch_local=1gb \
4    -m abe script.sh

5 # Or specifically select carex node
6 qsub -l walltime=1:0:0 -q ibot -l cluster=carex \
7    -l select=1:ncpus=4:mem=4gb:scratch_local=1gb \
8    -m abe script.sh

9 # Or specifically select draba node
10 qsub -l walltime=1:0:0 -q ibot -l cluster=draba \
11    -l select=1:ncpus=4:mem=4gb:scratch_local=1gb \
12    -m abe script.sh
```



use Limits Discussion Connection Priority access Storage Tasks

Requesting whole node and using hyperthreading

- Queuing system reserves only physical CPU cores and does not work well with hyperthreading
- To use hyperthreading (advantageous for some applications, like Java and some parallelization), user must request whole node using -1 place=exclhost
- MetaCentrum admins are working on better reservation of whole nodes and handling of hyperthreading

```
# To reserve whole carex node use something like

2qsub -l walltime=1:0:0 -q ibot \

-l select=1:ncpus=8:mem=500gb:scratch_local= \

1600gb:hyperthreading=True:cluster=carex \

-l place=exclhost -m abe script.sh

# To reserve whole draba HPC node use something like

qqsub -l walltime=1:0:0 -q ibot \

-l select=1:ncpus=80:mem=1500gb:scratch_local= \

1600gb:hyperthreading=True:cluster=draba \

-l place=exclhost -m abe script.sh
```



Default limits

- Members of group ibot
 - No limits for computing resources
 - Default quota for data storage is 2 TB requests to change to be send to cluster@ibot.cas.cz
 - We can discuss and set different default limits for our users
- Any other MetaCentrum users
 - Any number of CPU, RAM, disk space, etc., but the task must end up within 24 hours (no longer walltime than 24:0:0)
 - Default quota for data storage is 10 GB
 - We can discuss and set different default limits for other users
- Content of shared directories is counted to quotas of respective users — their storage quotas must be set accordingly



- The cluster is managed by administrators of the MetaCentrum

 regarding general problems related to computation,
 application etc. please use the MetaCentrum help support
- Specific questions related to the Průhonice cluster, e.g. issues connected with the usage of the database server, should be addressed to cluster administrators Vojtěch Zeisek and Yann JK Bertrand
- If you desire to join the **ibot** group, create or modify shared folder and change your allowed disk space on the disk storage, please contact Vojtěch Zeisek
- Questions related to Apache Hadoop, Spark, NoSQL databases (Neo4j and MongoDB) and other issues specific computing on the HPC nodes should be directed to Yann JK Bertrand
- Contact Vojtěch Zeisek and/or Yann JK Bertrand on cluster@ibot.cas.cz



Thank you for your attention...

Let's discuss...

- Which default limits do we wish to set?
- How do we wish to use shared data folders? Any use case? Any requests?
- Any requests to higher quota on the storage?
- Any requests to run something special on the database server?
- Any question from anyone? :-)