

Introduction to MSI



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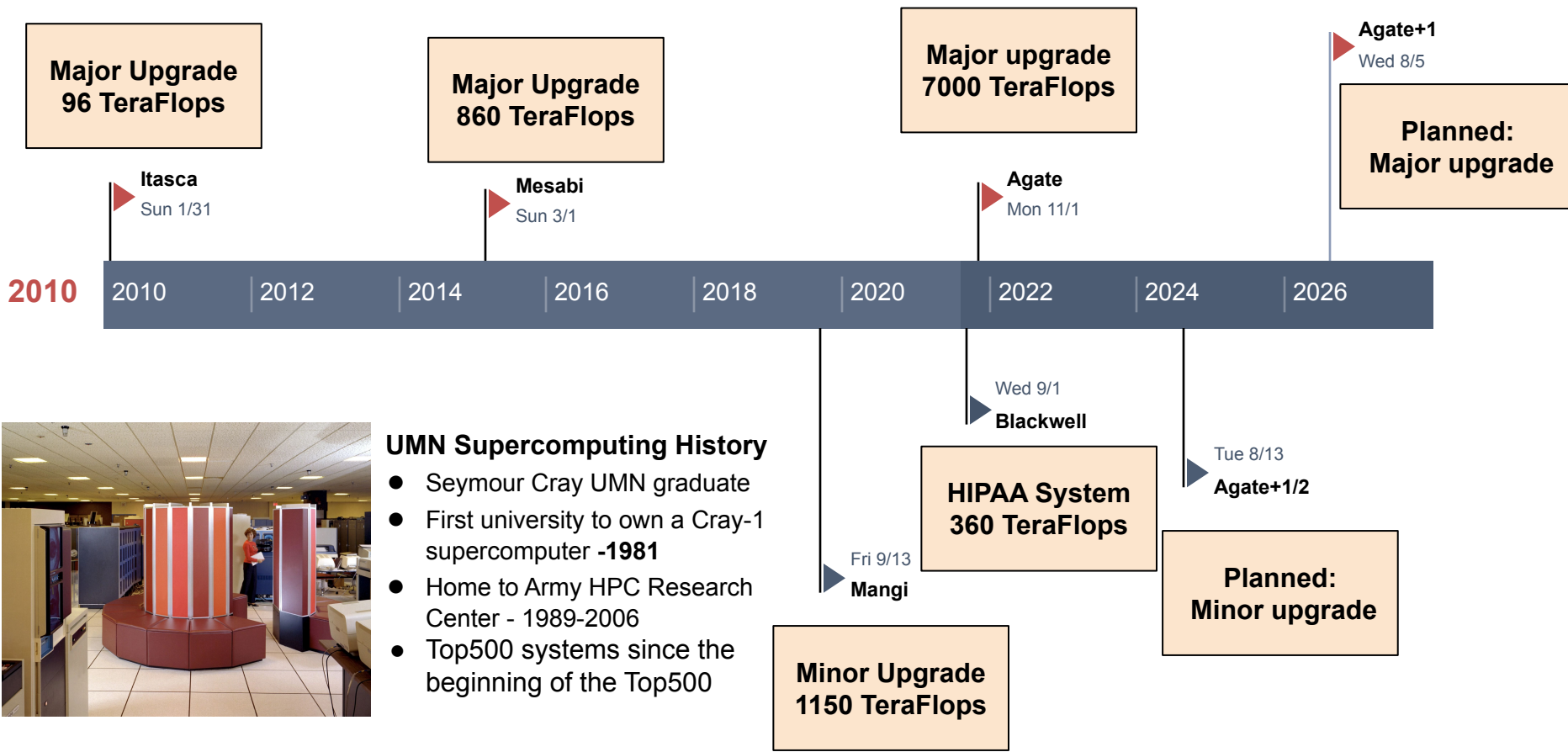
Overview

- Mission
- Organization
- Resources and Services
- Eligibility, Access, and Accounts

Minnesota Supercomputing Institute

- To provide advanced **research computing infrastructure** and **expertise** to the University of Minnesota research and scholarly community and the State of Minnesota in order to advance and accelerate research and foster innovation and discoveries through **advanced computing technologies**, **scientific computing and informatics**, **application development**, and **services**.
- Academic unit under the Office of the Vice President of Research
- 50 full time employees and ~6 students
 - Six working teams
- 900+ active Principle Investigator groups using MSI resources
- 4,800+ active users
- Significant resources and expertise provided for FREE to UMN PIs.
- UMN have been leaders in academic supercomputing since the 1980s.

Minnesota Supercomputing Institute: Strong HPC History & Regular Upgrade Cycle



UMN Supercomputing History

- Seymour Cray UMN graduate
- First university to own a Cray-1 supercomputer -1981
- Home to Army HPC Research Center - 1989-2006
- Top500 systems since the beginning of the Top500

Office of the Vice President for Research

Research Computing

**Minnesota
Supercomputing
Institute**

**UM
Informatics
Institute**

U-Spatial

**User Gateway
Group**

**Scientific
Computing
Solutions**

**Research
Informatics
Solutions**

**Application
Development
Solutions**

**Advanced
Systems
Operations**

- First Line User Support
- User Training
- On Boarding
- Communications
- Outreach

- Optimization
- Benchmarking
- HPC Research
- Workflow & pipeline Development

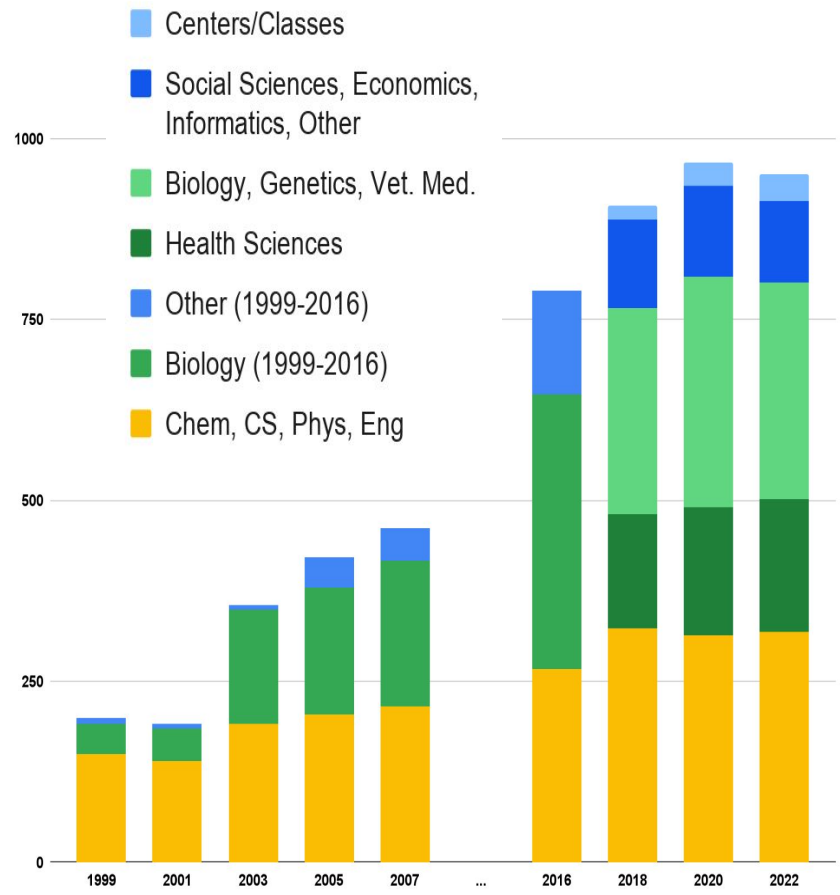
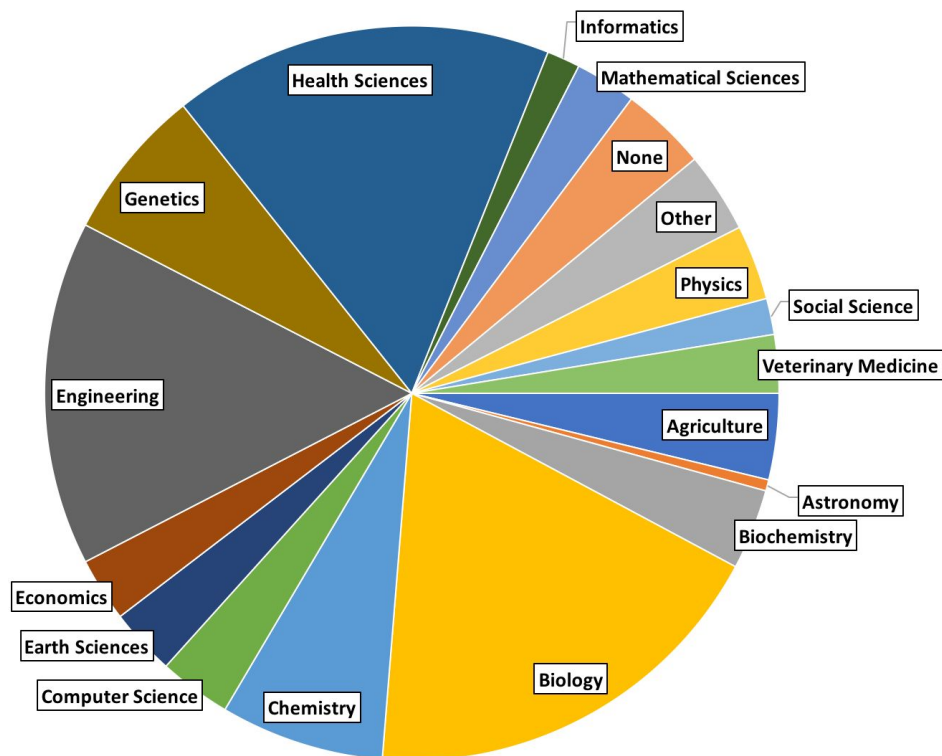
- Informatics education
- Informatics research
- Informatics services
- Life Science Computing

- Custom App Dev
- System Programming

- Common Services
- HPC Systems
- Storage Systems
- Hosted Services

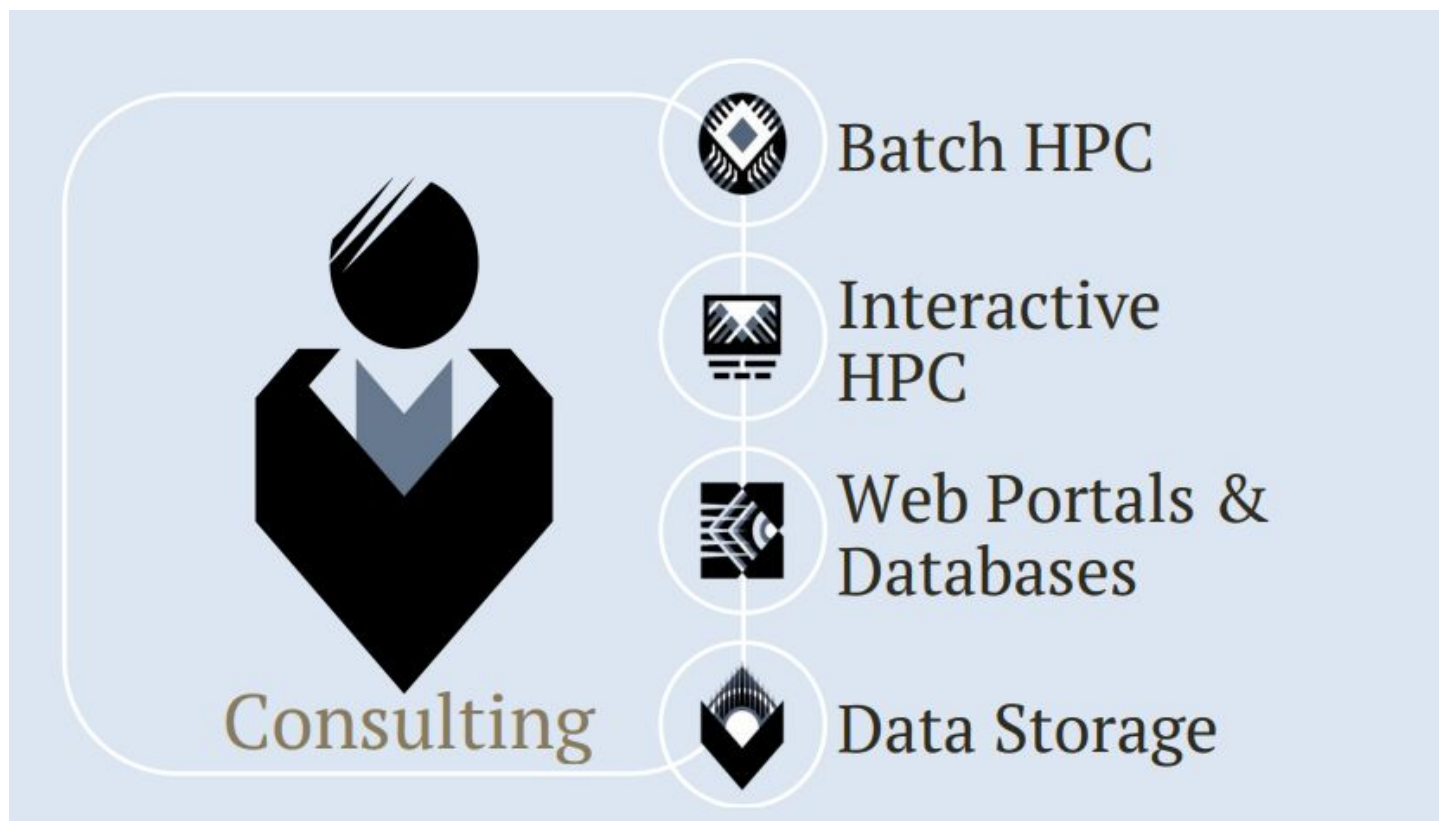
The Long Tail of Research Computing

Groups in 2020: 900 User Groups
4,555 Active users



Biggest increasing in Life Sciences

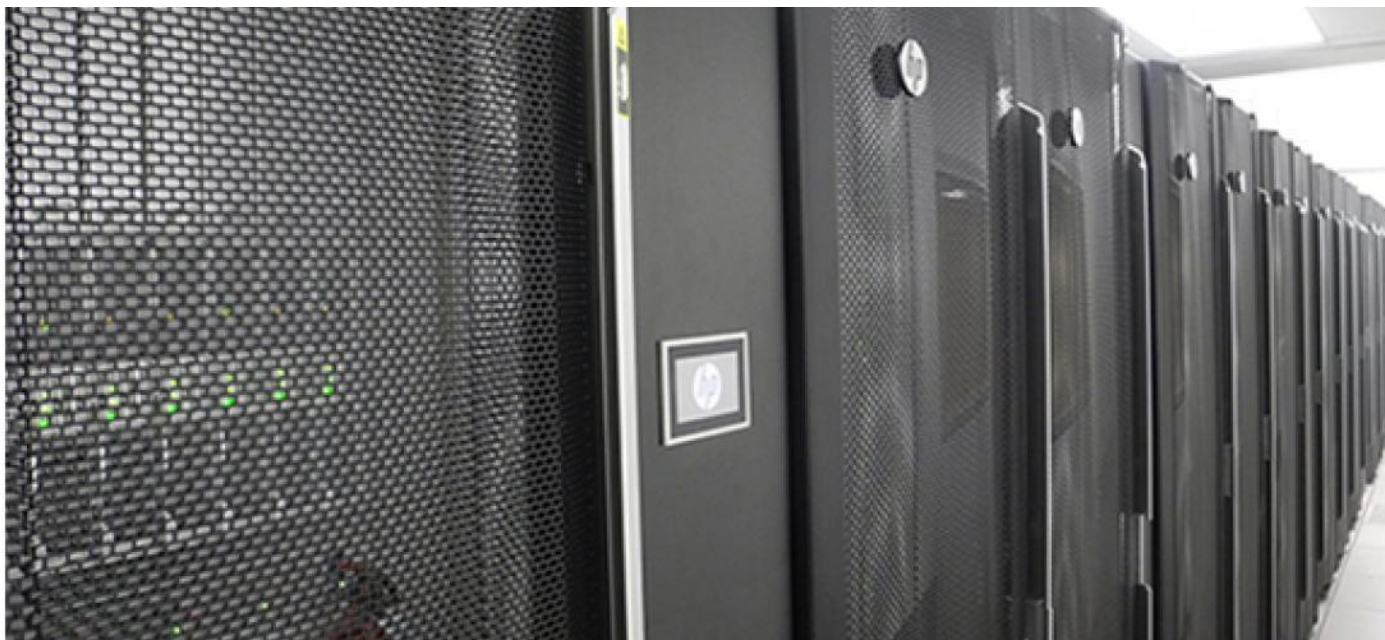
Resources and Services



High Performance Computing: Batch & Interactive

High Performance Computing Services

- Highly parallel computing environment with high-speed network connections, graphics processing units, high memory, and high-performance storage
- Offer both **batch** HPC and **interactive** HPC
- HPC access controlled by the “Fair share” SLURM scheduler



Mesabi/Mangi HPC System

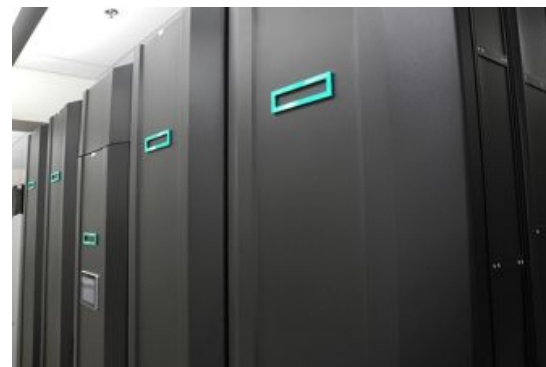
Mesabi

- 741 nodes
- 17,784 cores
- 40 nodes with 2x NVidia Tesla GPUs
- 32 nodes with 480GB SSDs



Mangi

- 164 AMD ROME Nodes
- 20,992 cores
- Amd2tb Nodes
- v100 GPU nodes
- v100-4 GPU nodes
- v100-8 GPU nodes



Agate HPC System

- 264 NVIDIA A100 GPUs are available in two configurations.
 - 50 nodes have 4 A100 GPUs and 512 GB of memory
 - 8 nodes have 8 A100 GPUs and 1 TB of memory
 - Multi-GPU nodes are connected via NVLink
- 344 CPU-only compute nodes are available in two configurations
 - 244 with 512 GB of memory
 - 100 with 2TB of memory
- 10 GPU nodes equipped with 8 A40 GPUs, 128 cores, and 512 GB of memory are made available for interactive work through Jupyter or command line sessions.

Batch HPC

- Using batch scheduling is the primary way to make use of MSI's shared supercomputing resources.
- You will '**submit**' jobs to the cluster scheduler, and your analysis eventually runs on the computer without any further intervention.
 - Requires jobs to be written as **shell scripts** and submitted via the command line.
- Jobs get **queued** and run based on a combination of the size of the request, the current load of the systems, how many jobs your group has run recently.
 - This is called "FairShare" - details can be found at this link:
<https://www.msi.umn.edu/content/hpc>
- This will probably be where you spend most of your time interacting with MSI, unless you have a specialized interactive application.

Batch HPC: SLURM

Job Submission and Scheduling (SLURM Scripts)

```
#!/bin/bash -l
#SBATCH --time=8:00:00
#SBATCH --ntasks=8
#SBATCH --mem=10g
#SBATCH --tmp=10g
#SBATCH --mail-type=ALL
#SBATCH --mail-user=sample_email@umn.edu
#
cd ~/program_directory
module load intel
module load omp/intel
mpirun -np 8 program_name < inputfile > outputfile
```


Batch partitions

SLURM scripts generally have a line to designate a partition (or partitions) where your batch job can execute. It looks like this:

```
$SBATCH -p <NameOfPartition>
```

A table of the current partitions on all MSI HPC clusters can be found at

<https://www.msi.umn.edu/partitions>

Research groups who lease private nodes will have private partition names they can use to access their purchased nodes.

Interactive HPC

- MSI also offers interactive HPC access
 - Very useful for real-time data visualization and exploration
 - Allows use of specialized hardware and software without labs having to purchase it
- Types of interactive HPC offered:
 - srun: SLURM batch method to get an interactive shell on HPC queue
 - NICE: web-based desktop
 - CITRIX: remote Windows desktop on HPC hardware
 - NX NoMachine: remote Linux graphical desktop
 - Jupyter Notebooks (beta): interactive Python/R on MSI
 - *Coming Soon: OpenOnDemand*
- See <https://www.msi.umn.edu/content/interactive-hpc>

Interactive HPC

Interactive queue use with srun, using an applications that needs X windows:

```
ssh -X mesabi.msi.umn.edu
```

```
your_user_id@ln0004 [~] % srun -N 1 --ntasks-per-node=4  
--mem-per-cpu=1gb -t 1:00:00 -p interactive --x11 --pty bash
```

```
your_user_id@cn0001 [~] % module load a-module-you-need
```

```
your_user_id@cn0001 [~] % ./my_program
```

Software Resources

- Scientific and numerical software packages are built into “modules” and available to be loaded as you need them.
- Common software tools used in many fields are available on our systems
 - Sequence analysis
 - Genetics
 - Proteomics
 - Image processing
 - Computational chemistry
 - Etc...
- 500+ software applications
 - [MSI Software Search](#)
- We can work with your research groups to install software that you need
 - Contact the help desk!

Service Units

- Service Units is a term MSI uses to reference the total compute resources consumed by a group or individual.
- Commercial and Industrial partners still purchase SUs, and their compute, storage, and memory usage all are accounted for.
- Formerly, MSI used SUs to allocate different resources for academic research groups.
- Allocation of resources is now done among groups via the SLURM “Fair share” algorithm.
- **Full deployment of SLURM commenced in January 2021, and the share of resources among groups is being monitored by an advisory group of MSI principal investigators.**

Data Storage at MSI

Primary Storage

- Primary storage is where you will do most of your work
 - GNU/Linux filesystem
- Watch your disk usage:
 - If you are up against file number or disk usage quota, **your entire group** cannot make new files
- Important directories in your primary storage space:

[/home/groupname/YOUR_X.500](#)

Your home directory. Store scripts and small files here

[/home/groupname/shared](#)

PI group shared directory. Store shared software, common datasets, reference databases, etc, here.

Primary Storage Scratch

- Quotas are large, so there should be leeway to make very large files here
 - Use it for intermediate/temporary files in analytical workflows
- Files are kept for 30 days, so be sure to copy important files to your group directory
 - Scratch space is also not backed up by snapshot
 - **We know the tricks that exist to keep data in scratch for longer than 30 days. Consider data on scratch as having no warranty.**
- Location:

[/scratch.global/](#)

Second Tier Storage

- Sometimes referred to as “Ceph”, the name of the software defined storage utility in use in this storage layer.
- Second tier storage is separate from primary storage, it is an “object store” - you must use special tools to access it.
- From the command line, you can use the “s3cmd” software package to interact with your storage:
 - See: <https://www.msi.umn.edu/support/faq/how-do-i-use-second-tier-storage-command-line>
- For a graphical interface, you can use Globus
 - See: <https://www.msi.umn.edu/support/faq/how-do-i-use-globus-transfer-data-msi-0>
- **Not** backed up, so if you delete a file, it will be gone for good
 - Quotas are much more relaxed on second tier storage
 - **Resilient system to store data**

Data Storage Allocation

- Default Primary storage allocation: 150GB for each group
- **Requests of up to 1TB:**
 - No justification or review required
- **For requests of up to 5TB:**
 - Justification
 - Review by MSI staff
- **For requests of more than 5TB:**
 - Justification
 - Estimate of amount needed
 - Estimate of amount actively used
 - Estimate of duration of storage
 - Review by MSI staff
- ANNUAL renewals for all non-default storage allocations

PI Groups, Accounts and Support

Service Costs

- **Free** for UMN faculty and their associates
- **Free** for other MN academic institutions
- **Free** for expert consultation

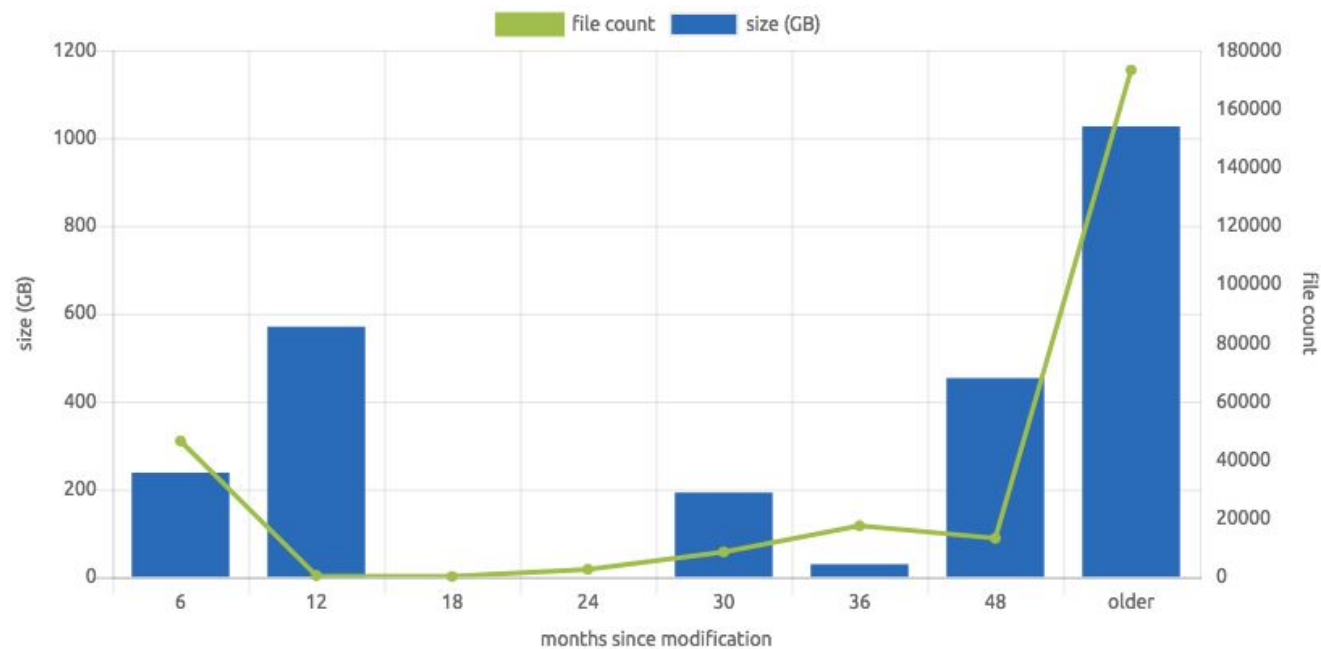
- Fee for resources over allocations limits
- Fee for expert collaboration
- Fee for service for external organizations

PI Group Structure

- **PI:** Controls everything
- **Group Administrators:** Users within the group designated to do everything a PI can do except add other Group Administrators
- **Users:** Added by PI, usually lab members (postdocs, techs, undergrads, grad students)
 - Users may be part of multiple accounts if they perform analysis for multiple PIs
 - The PI is a user, too, and their SU and disk usage counts the same as their users'

Checking Group Usage

- Check disk usage:
<https://www.msi.umn.edu/group/GROUPNAME/storage>



Accounts and Renewals

- UMN faculty members, department staff, and researchers at other MN state institutions are eligible for MSI accounts
- **PI accounts are renewed annually**
 - Renewal period starts in October
 - Extra storage allocations all must be checked and justified

Research Support Services

- MSI has ~25 consulting staff, most with an advanced degree.
- We provide 20-30 workshops and tutorials annually
- 1-2 hour meetings on computational or informatics methodologies
- Informational sessions for proposal development
- Expertise including:
 - Research Computing; Informatics
 - Big Data Analysis
 - Parallel Algorithm Optimization and Development
 - Custom Analysis Pipeline Development
 - Genomics
 - Custom Application Development

Training and Tutorials

- We host tutorials on a range of topics
<https://www.msi.umn.edu/tutorials>
- **Introductory tutorials**
 - Intro to MSI
 - Intro to Linux
 - MSI Batch Job Submission
- **Bioinformatics tutorials**
 - Proteomic and mass spec data analysis
 - RNAseq data analysis
- **Advanced tutorials**
 - Parallelization
 - Python for scientific computing
- **And many more!**
- **We also collaborate with The Carpentries and LATIS at UMN, have a look at our Events page:** <https://www.msi.umn.edu/events>

Links to Resources

- MSI Homepage
 - <https://www.msi.umn.edu>
- PI eligibility and access policies
 - <https://www.msi.umn.edu/content/eligibility-getting-access>
- SLURM batch partition limits page
 - <https://www.msi.umn.edu/partitions>
- Tutorials page
 - <https://www.msi.umn.edu/tutorials>
- Check Primary storage
 - <https://www.msi.umn.edu/group/GROUPNAME/storage>
- Software tools
 - <https://www.msi.umn.edu/software>

Questions?

- MSI Help Desk
- Every Day help available **using Email or Video**
- **Walk-Ins**
 - Monday, Wednesday, Friday (587 Walter Library, UM campus)
9:00 AM – 4:00 PM
 - Tuesday, Thursday (Zoom virtual walk-in)
9:00 - 11:00 AM, and 1:00 - 3:00 PM**Zoom Meeting Room Link:** <https://umn.zoom.us/my/msi.helpdesk>
- Call x6-0802 (1-612-626-0802)
- Email **help@msi.umn.edu**
- Schedule a Zoom Consultation
<https://www.msi.umn.edu/content/helpdesk-video-consultations>