



Using Gaussian16 software On Paramshakti supercomputer

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Outline

- About Gaussian Software
- Gaussian Input File Preparation
- Access of PARAMSHAKTI (PS)
 1. Create login at PS
 2. Connect with PS
- Submitting Gaussian jobs
 1. Generate batch script for SLURM workload manager
 2. Submitting job
- Monitor jobs in PS
- Gaussian16 performance on CPUs



About Gaussian 16

- Gaussian is a general purpose electronic structure package for use in computational chemistry.
- Most widely used computational chemistry package. The latest release is Gaussian16 Rev C.01
- Vendor's website: <http://www.gaussian.com>



About Gaussian 16

Pricing for Gaussian Products

USA	Outside the USA (excluding India, China & Hong Kong)	China & Hong Kong	India
Academic special bundle pricing	Academic special bundle pricing	Academic special bundle pricing	Academic special bundle pricing
Academic customers	Academic customers	Academic customers	Academic customers
Commercial customers	Commercial customers	Commercial customers	Commercial customers
US Government customers			

Due to recommendations by several international groups that monitor intellectual property protection laws and the enforcement of these laws, there are a number of countries to which Gaussian, Inc. does not license source code.

Maintenance pricing for Gaussian 16, GaussView 6, and Linda can be found by clicking this link.

Last updated on: 27 January 2017.



About Gaussian 16

LICENSING: A signed license is required prior to shipping

Shrink wrap license

Gaussian 16 (New orders include *Exploring Chemistry*)

UNIX, Linux, Mac OS X			New License	Upgrade From G09
Site license	All machine types	Binary code	\$6,950	\$6,250
	Single Machine Type		\$3,475	\$2,775
			Add \$1,400 for each addl. machine type	
Single computer	Mac OS 32-bit single core		\$1,275	\$975
Additional Charge for TCP Linda License			New License	Upgrade From Linda 8
Linda license	Each machine type		\$2,775	\$2,250



Gaussian 16 Functionality

□ Types of calculation:

- Gradients/Geometry optimizations
- Frequencies (IR/Raman, NMR, etc.)
- Transition state search
- IRC for reaction path
- Excited state calculation
- Spin-orbit coupling
- Several solvation models (PCM, SMD, etc), NonEquilibrium solvent model
- Two and three layer ONIOM model
- B-O molecular dynamics

□ Methods:

- Semiempirical: MNDO, AM1, PM3 etc.
- HF: restricted/unrestricted
- DFT: many pure/hybrid functionals
- MP: 2nd-5th order
- CI: single and double
- CC: single, double, triples contribution
- High accuracy methods: Gn, CBS, etc.
- MCSCF: including CASSCF
- MM: AMBER, Dreiding, UFF force field
- EOM-CCSD with analytic gradient
- DFTB and DFTBA methods

[illegible]



Molecular modeling and visualizer

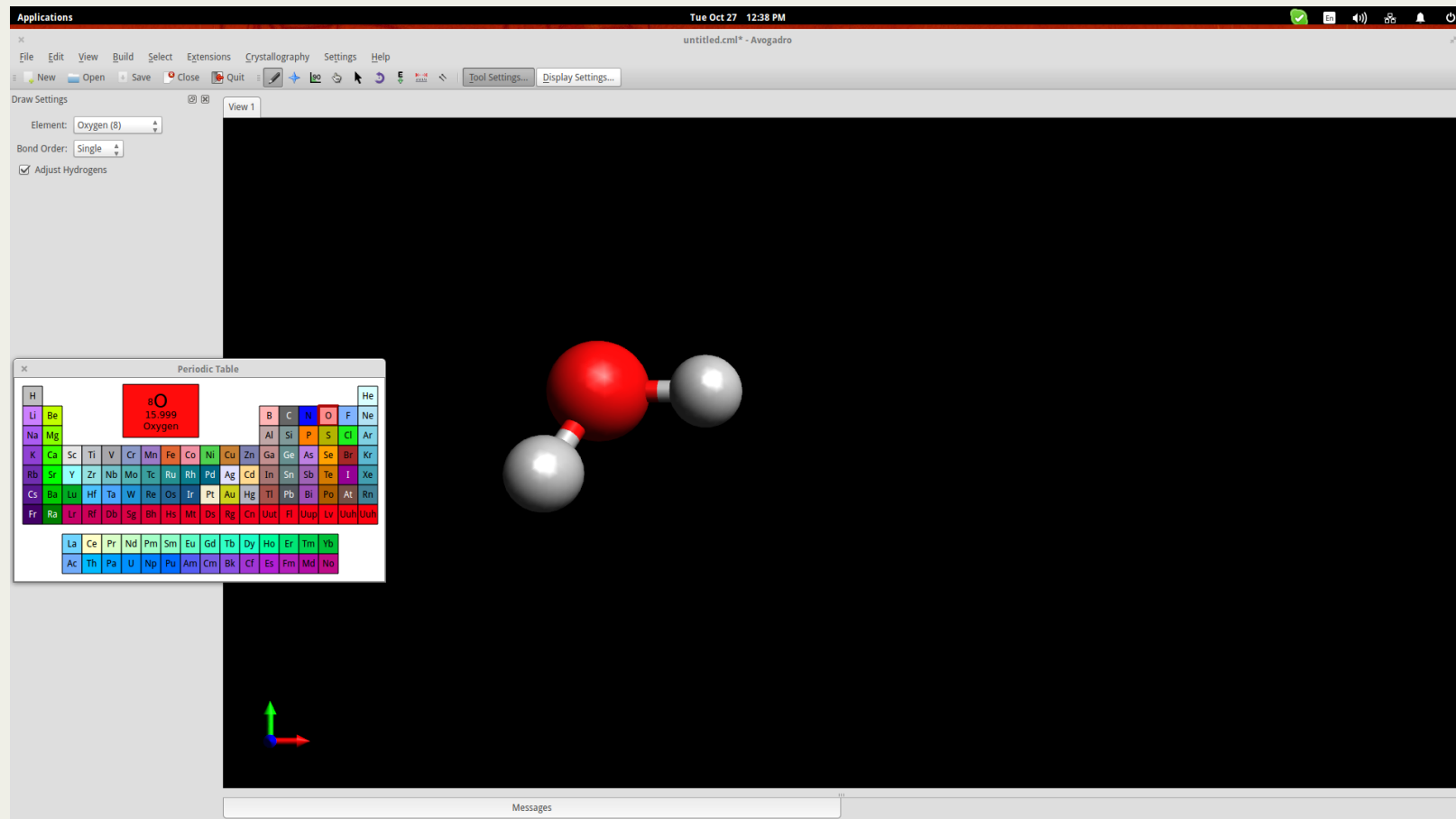
Generate Cartesian Coordinate

□ For Small molecules:

- Avogadro
- Gaussview
- Jmol
- Molekel
- Molden

□ For Biomolecules:

- VMD
- PyMOL
- Chimera





Input File - Keyword Specification in the Route Section

- Keyword line(s) - specify calculation type and other job options
- Start with # symbol in the route section
- Can be multiple lines
- Terminate with a blank line
- Format
 - keyword=option
 - keyword(option)
 - keyword(option1,option2,...)
 - keyword=(option1,option2,...)
- User's guide provides list of keywords, options, and basis set notion
<https://gaussian.com/keywords/>

Quick Links

Basis Sets

Density Functional (DFT) Methods

Solvents List SCRF

References

Users Ref Index

Keyword

[Keyword=]Option

in Follow

Follow @gaussian

YouTube

YOUKU

List of Gaussian Keywords

- #
- ADMP
- BD
- BOMD
- CacheSize
- CASSCF
- CBS Methods
- CBSExtrapolate
- CCD and CCSD
- Charge
- ChkBasis
- CID and CISD
- CIS
- CNDO
- Complex
- Constants
- Counterpoise
- CPHF
- Density
- DensityFit and NoDensityFit
- DFT Methods
- DFTB and DFTBA
- EET
- EOMCCSD
- EPT
- External
- ExtraBasis & ExtraDensityBasis
- Field
- FMM
- Force
- Freq
- Gen and GenECP
- GenChk
- Geom
- GFIInput
- GFPrint
- Gn Methods
- Guess
- GVB
- HF
- Huckel
- INDO
- Integral
- IOP
- IRC
- IRCmax
- Link0 Commands
- LSDA
- MaxDisk
- MINDO3
- MNDO
- MM Methods
- MP Methods
- Name
- NMR
- ONIOM
- Optimization
- Output
- PBC
- Polar
- Population
- Pressure
- Prop
- Pseudo
- Punch
- QCI
- Restart
- SAC-CI
- Scale
- Scan
- SCF
- SCRF
- Semi-Empirical Methods
- SP
- Sparse
- Stable
- Symmetry
- TD
- Temperature
- Test
- TestMO
- TrackIO
- Transformation
- Units
- Volume
- W1 Methods
- Window Keyword and Frozen Core Options
- ZIndo

Create login in PARAMSHAKTI



High Performance Computing Facility Indian Institute of Technology Kharagpur

HPCNC User Support

Home

System Overview

Quick Start Guide

Using the System

Tutorials

User Guidelines

Best Practices for HPC

Publications

HPC Advisory Committee

Virtual Workshop

User Support

User Forms

FAQs

Quick Start Guide

Quick Start Guide

1. All users are requested to go through the [User policy](#) and [Best Practices](#) to know the detailed guidelines approved by the Advisory Committee before requesting access to Paramshakti HPC systems
2. All the First Time Users will get access to the HPC systems through their respective Faculty/Project Investigator/Guide, who will be the account coordinator. The Faculty/Project Investigator/Guide must have submitted the [User Forms](#) as a pre-requisite.
3. Any User other than Faculty Adviser needs to follow the 2-step procedure for user account creation. For a particular user if the Faculty Adviser account is already registered then he can skip the Step-1.

Step-1

Faculty Adviser can submit a request using [Faculty Form](#). The signed hard copy needs to be submitted at Room-F7, Lab1, First Floor JC Bose Annex building. Account creation of the Faculty Adviser may take upto 48 hours after the submission of the duly signed hard copy of the form. The Faculty Adviser will be notified via email after account creation is completed.

Step-2

After Faculty Adviser account creation is complete, the students/users other than Faculty in the respective group of the Faculty Adviser can start filling the [User Forms](#) and submit the duly signed hard copy at Room-F7, Lab1, First Floor JC Bose Annex building. The user will be notified via email after account creation is completed.

** Due to current pandemic situation users may send the scanned copy of the signed form duly endorsed by the respective Head of the Department/Centre/School by e-mail to:shaktisupport@iitkgp.ac.in.



Indian Institute of Technology Kharagpur

High-Performance Computing Facility

www.hpc.iitkgp.ac.in

PARAM SHAKTI ACCOUNT REQUEST FORM
for

Internal User other than faculty

IITKGP Faculty

Create login in PARAMSHAKTI



Indian Institute of Technology Kharagpur
High-Performance Computing Facility
www.hpc.iitkgp.ac.in

PARAM SHAKTI ACCOUNT REQUEST FORM

(internal users other than IIT Kharagpur faculty)

1. User Details

Name(Full)

Name of the Project
Guide/Advisor

Please type few characters of the first name of Faculty Advisor and select from available list. Manual Entry is invalid *

Department/School/Center

Select Department/School/Center



Degree
Registered/Designation

Roll No. / Institute ID

Date of joining the

mm/dd/yyyy





Connect with PARAMSHAKTI

Remote System Access

✓ Connect PS through ssh via port 22 (Inside Campus) or 4422 (Outside Campus)

Linux OS:

- ☐ ssh command to login/access from terminal
- ☐ sftp/scp command to copy file and directory
- ☐ vi command for editing file

```
ssh -p 4422 user1@paramshakti.iitkgp.ac.in
```

```
scp -P 4422 user2@hostname:[address2] user1@paramshakti.iitkgp.ac.in:[address1]
```

Windows OS:

- ☐ login via ssh from Putty, MobaXterm, Tera Term etc.
- ☐ copy file between servers using FileZilla, WinSCP etc.

Category:

- ☒ Session
 - Logging
- ☒ Terminal
 - Keyboard
 - Bell
 - Features
- ☒ Window
 - Appearance
 - Behaviour
 - Translation
 - ☒ Selection
 - Colours
- ☒ Connection
 - Data
 - Proxy
 - Telnet
 - Rlogin
 - ☒ SSH
 - Serial

Basic options for your PuTTY session

Specify the destination you want to connect to

Host Name (or IP address) Port

paramshakti.iitkgp.ac.in 22

Connection type:

☐ Raw
 ☐ Telnet
 ☐ Rlogin
 ☒ SSH
 ☐ Serial

Load, save or delete a stored session

Saved Sessions

paramshakti.iitkgp.ac.in

Default Settings

paramshakti.iitkgp.ac.in

Load

Save

Delete

Close window on exit:

☐ Always
 ☐ Never
 ☒ Only on clean exit

About

Help

Open

Cancel

login as:

Keyboard-interactive authentication prompts from server:

If you truly desire access to this host, then you must indulge me in a simple challenge.

Observe the picture below and answer the question listed afterwards:

```

  /\  /\  /\  /\  /\  /\  /\  /\
 ( J | k | q | v | y | v | x | T )
  \/  \/  \/  \/  \/  \/  \/  \/
    
```

Type the string above: JkqVvxxT

Password:

End of keyboard-interactive prompts from server

Last login: Tue Sep 21 23:31:35 2021 from 42.110.131.111

on Mon Nov 22 00:44:23 IST 2021

File System utilization for user : 18tspd202

Filesystem	used	quota	limit	grace	files	quota	limit	grace
/home	36.78G	40G	50G	-	382783	0	0	-
/scratch	677.6G	2T	0k	-	16090	0	0	-

Cluster utilization

(Usage reported in Percentage of Total)

Cluster	CPU	GPU
Paramshakti	76%	90%

For any issues and queries, open a ticket at:

<https://paramshakti.iitkgp.ac.in/support/open.php>

@login01 ~]\$





Job-file/batch-script

```
[user@hostname ~]$ module avail          # lists all the available modules
[user@hostname ~]$ module load apps/gaussian/16/gnu  # load modules the gaussian with gnu compilers into
                                                    your environment
[user@hostname ~]$ module unload apps/gaussian/16/gnu # remove all environment related to
                                                    apps/gaussian/16/gnu
```

#Serial/Minimal batch script for Slurm Workload manager:

```
#!/bin/bash
#SBATCH -J job_name          # name of the job
#SBATCH -p standard-low     # name of the partition: available options "standard,standard-low,gpu,gpu-low,hm"
#SBATCH -t 01:00:00         # walltime in HH:MM:SS, Max value 72:00:00
#SBATCH --nodes=1           # number of nodes (-N) should be 1 for serial
#SBATCH -n 1                # no of processes

#list of modules you want to use, for example
module load apps/gaussian/16/gnu

#name of the executable
exe=g16                      # name of the executable

#run the application
$exe input.com
```

Batch script for parallel gaussian job:

```
[user@hostname ~]$ vi job_name.sh
```

```
#!/bin/bash
#SBATCH -J job_name           #Job name(--job-name)
#SBATCH -o %j-out-myjob       #Name of stdout output file(--output)
#SBATCH -e %j-err-myjob       #Name of stderr error file(--error)
#SBATCH -p standard-low       #Queue (--partition) name
#SBATCH -N 1                  #number of nodes should be 1 for g16 without linda)
#SBATCH -n 10                 # Number of procs
#SBATCH -t 72:00:00           # specifies walltime(--time maximum duration)of run
#SBATCH --mem=10000           # Memory per node specification is in MB. It is optional.
#SBATCH --mail-user=user@email.com  # user email ID where job status info will be sent
#SBATCH --mail-type=ALL       # Send Mail for all type of event regarding the job

# list of modules you want to use, for example
# module load apps/gaussian/16/gnu
# name of the executable
g16root="/home/apps"
GAUSS_SCRDIR="/scratch/$USER/gaussian"
export g16root GAUSS_SCRDIR
. $g16root/g16/bsd/g16.profile
exe=g16
# Executables along with right Parameters and right input files
#mpirun -bootstrap slurm -n $SLURM_NTASKS $exe < input.com > output.log
$exe input.com
```

```
[user@hostname ~]$ sbatch job_name.sh
```

```
[user@hostname ~]$ Submitted batch job 546965
```



Monitor Jobs Status

by
SLURM WORKLOAD MANAGER

[user@hostname ~]\$ squeue

lists all the submitted jobs

```
[user@login08 ~]$ squeue
JOBID PARTITION NAME USER ST TIME NODES NODELIST(REASON)
545616 standard- pdi-thio 19 j0 R 1-04:28:58 1 cn308
545472 standard- pdi_q4.c 19 j0 R 1-13:30:03 1 hm027
545584 standard- pdi_q4.c 19 j0 R 1-13:21:42 1 hm025
545587 standard- pdi_q4.c 19 j0 R 1-13:18:35 1 hm033
545583 standard- pdi-tri- 19 j0 R 2-10:59:23 1 cn076
545588 standard- pdi-tri- 19 j0 R 2-10:59:23 1 cn276
545615 standard- pdi-thio 19 j0 R 2-10:59:23 1 hm008
545572 standard- trimer_1 19 j0 R 2-13:49:09 1 cn031
545271 standard- Fe-2b-W4 18 j2f1 R 2-23:44:06 12 cn[010,032-033,037,039-040,084,093,138,273-275]
```

[user@hostname ~]\$ scontrol show jobid <job id>

Show the details of the job

[user@hostname ~]\$ scancel <job id>

Kill a job

<http://www.hpc.iitkgp.ac.in/HPCF/jobControl>

<https://slurm.schedmd.com/tutorials.html>

<https://slurm.schedmd.com/documentation.html>



Monitor Jobs Status

`[user@hostname ~]$ less job-name.log`

```
Any fool can criticize, condemn, and complain -- and most do.
```

```
-- Dale Carnegie
```

```
Job cpu time:      0 days  0 hours  0 minutes 25.3 seconds.
```

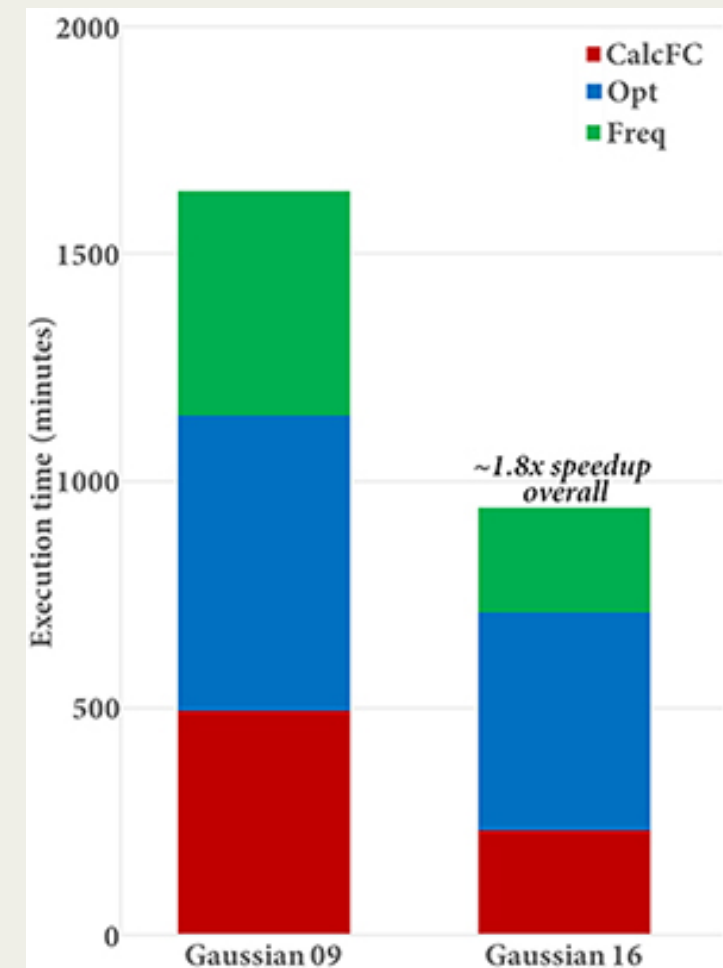
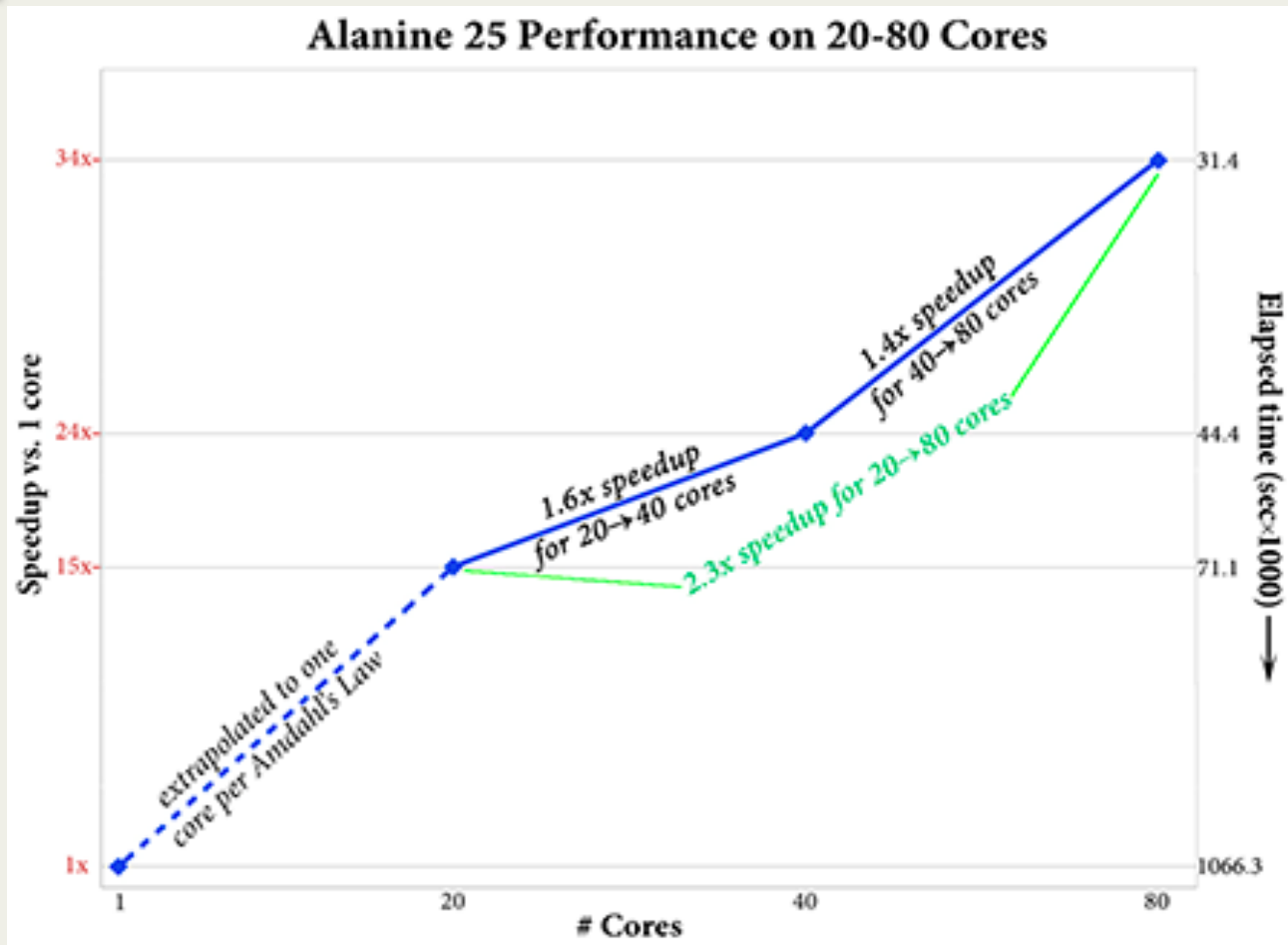
```
Elapsed time:      0 days  0 hours  0 minutes  7.5 seconds.
```

```
File lengths (MBytes):  RWF=      6 Int=      0 D2E=      0 Chk=      93 Scr=      1
```

```
Normal termination of Gaussian 16 at Thu Nov 25 10:19:45 2021.
```



Gaussian 16 on CPUs



THANK YOU