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خوش آمدید



Ministry of Electronics and
Information Technology
Government of India



PARAM GANGA @ IIT Roorkee

23rd Aug 2021

Agenda

PARAM-Ganga Architecture

Technical Specification of PARAM-Ganga

PARAM Ganga Software Stack

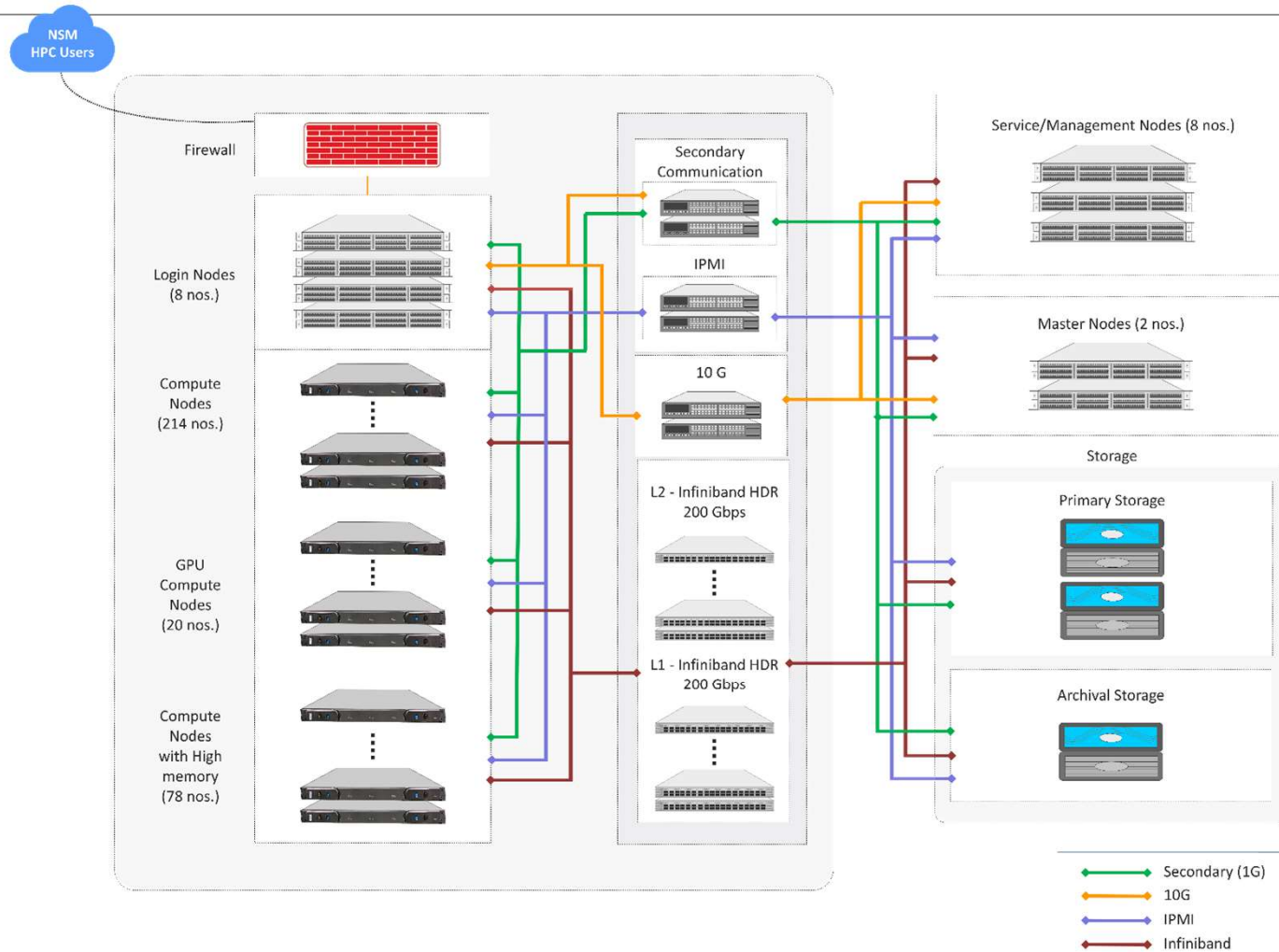
How to access Param-Ganga.

Ticketing Tool and Helpdesk

PARAM GANGA



PARAM Ganga Architecture Diagram



One Vision. One Goal... Advanced Computing for Human Advancement...

PARAM Ganga - System Configuration



System Specifications

Theoretical Peak Floating-point Performance Total (Rpeak)	1.67 PFLOPS
Base Specifications (Compute Nodes)	2 X Intel Xeon Cascadelake 8268, 24 Cores, 2.9 GHz, Processors per node, 192 GB Memory, 480 GB SSD
Master/Service/Login Nodes	20 nos.
CPU only Compute Nodes (Memory)	214 nos. (192GB)
GPU Compute Nodes (Memory)	20 (192 GB)
High Memory Compute Nodes	78 nos. (768GB)
Total Memory	104.832 TB
Interconnect	Primary: 100Gbps Mellanox Infiniband Interconnect network 100% non blocking, fat tree topology Secondary: 10G/1G Ethernet Network Management network: 1G Ethernet

CPU Only Compute Nodes

- ✦ 214 Nodes
- ✦ 10272 Cores
- ✦ Compute power of Rpeak 953.2 TFLOPS
- ✦ Each Node with
 - + 2 X Intel Xeon Cascadelake 8268, 24 cores, 2.9 GHz, processors
 - + 192 GB memory
 - + 480 GB SSD

GPU Compute Nodes

- ✦ 20 Nodes
- ✦ 800 CPU Cores
- ✦ 204800 CUDA Cores
- ✦ Rpeak CPU 64 TFLOPS + GPU 312 TF
- ✦ Each Node with
 - + 2 X Intel Xeon Cascadelake 6248, 20 cores, 2.5 GHz, processors
 - + 192 GB Memory
 - + 2 x NVIDIA V100 SXM2 GPU Cards
 - + 480 GB SSD

High Memory Compute Nodes

- ✦ 78 Nodes
- ✦ 3744 Cores
- ✦ Compute power of Rpeak 347.33 TFLOPS
- ✦ Each Node with
 - + 2 X Intel Xeon Cascadelake 8268, 24 cores, 2.9 GHz, processors
 - + 768 GB Memory
 - + 480 GB SSD

Technical Specifications



CPU only Compute Nodes

- 214 Nodes
- 10272 Cores
- Compute power of Rpeak 953.156 TFLOPS
- Each Node with
 - 2* Intel Xeon Platinum 8268, 24 cores, 2.9GHz, processors
 - 192 GB memory
 - 480 GB SSD

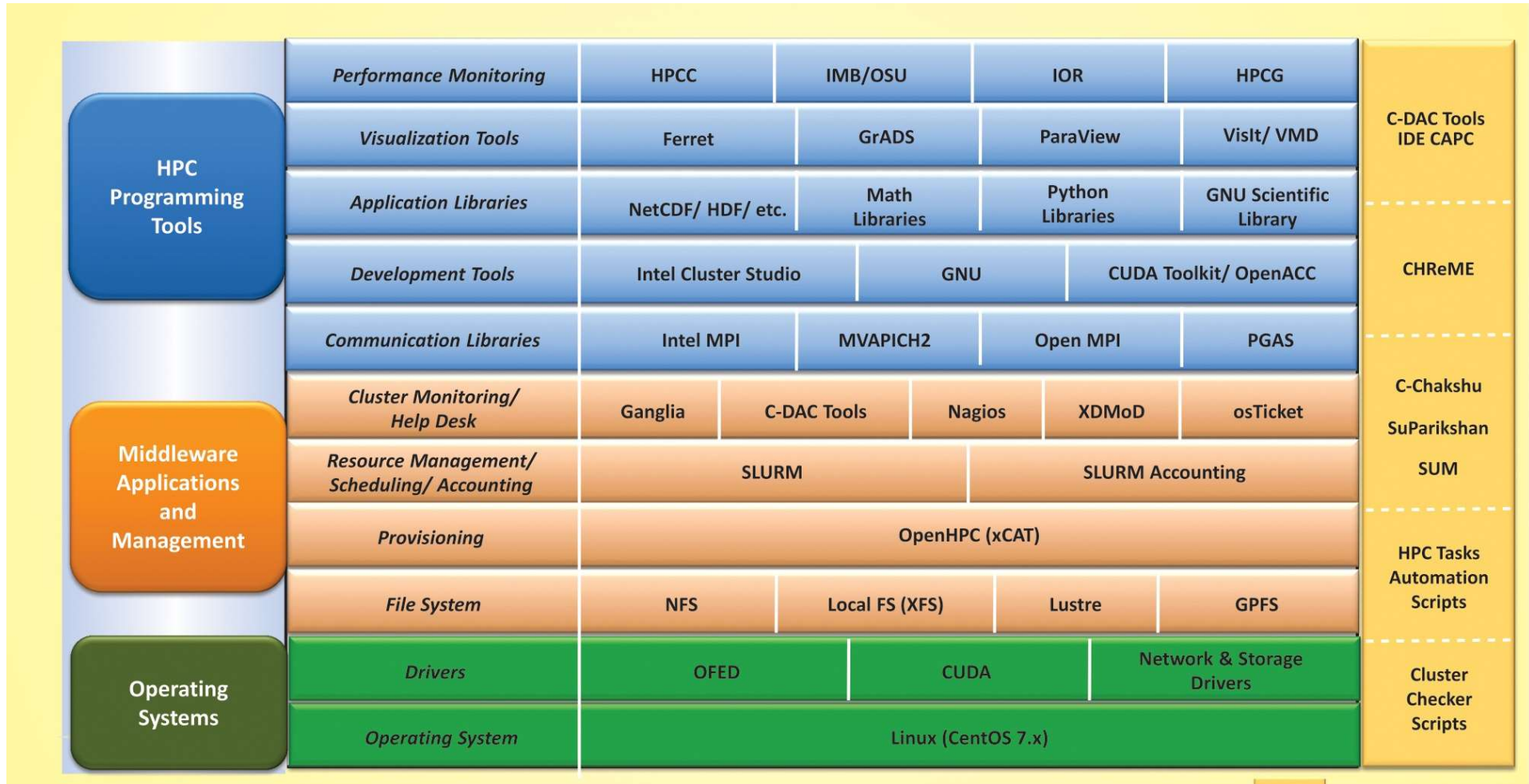
High Memory Compute Nodes

- 78 Nodes
- 3744 Cores
- Compute power of Rpeak 347.412 TFLOPS
- Each Node with
 - 2* Intel Xeon Platinum 8268, 24 cores, 2.9GHz, processors
 - 768 GB memory
 - 480 GB SSD

GPU Compute Nodes

- 20 Nodes
- 800 CPU cores
- 204800 GPU Cores
- Compute power of Rpeak 64 TFLOPS + 280 TF = 344
- Each Node with
 - 2* Intel Xeon Gold G-6248, 20 cores, 2.5 GHz, processors
 - 192 GB Memory
 - 480 GB SSD
 - 2xNvidia V100 SXM2 GPU cards each with 5120 CUDA cores

Software Stack



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Software Components

Operating System –

- HPC clusters generally are build with Linux operating system as a base OS (Centos7.6)
- It includes all the device drivers for the H/W connected to each node.

Cluster Manager/Orchestrator

- Tools in this category builds a centralized architecture where a controller node builds and manages the cluster.
- xCAT – Is an open-source cluster Manager, Maintained by community, is the widely used tool for HPC as well as cloud clusters.
- It provides flexibility to handle objects within the cluster with its easy manageable methods
- It provide methods to deploy nodes with a very light weight stateless images.



Software Components

Resource Manager (SLURM)

- As there are a lot of resource within a cluster like : CPU-Cores, Memory banks, GPU accelerator cards managing which becomes a tedious task for a user and a system administrator.
- Resource manager with in “slurm” tool helps to manage and represent resources to the users in a simplest way.

Job Scheduler (SLURM)

- A scheduler checks the available resources within a cluster and manages which jobs run where and when.
- Allocating resources to each users for optimal utilization of system resources.
- Provides multiple algorithm, which provides different ways to initiate jobs on the resources.
- BACKFILL scheduling is the widely used and the most efficient algorithm.
- Provider batch jobs as well as Interactive jobs submission methods.

NSM Clusters – Applications, Tools, Programming Models



HPC Applications	Bio-informatics	MUMmer, HMMER, MEME, PHYLIP, mpiBLAST, ClustalW	Visualization Programs	GrADS, ParaView, VisIt, VMD
	Molecular Dynamics	NAMD (CPU & GPU), LAMMPS(CPU & GPU), GROMACS	Dependency Libraries	NetCDF, PNETCDF, Jasper, HDF5, Tcl, Boost, FFTW
	Material Modeling, Quantum Chemistry	Quantum-Espresso, Abinit, CP2K, NWChem,	Programming Models	MPI, OpenMP, OpenACC, CUDA, PGAS, Pthreads
	CFD	OpenFOAM, FDS, SU2	Additional applications, libraries, tools on different NSM systems as per requirements from users of respective systems	
	Weather, Ocean, Climate	WRF, RegCM, MOM, ROMS		
	Disaster Management	ANUGA Hydro		
AI/ ML/ DL Tools/ Technologies		DL Frame work: TensorFlow , keras, theano, pytorch, scikit-learn,scipy, cuDNN Data Science: Numpy , RAPIDS Distributed DL Framework: TensorFlow with Horovod Container Technology: enroot JupyterHub: DL application development platforms and web based IDE		

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Accessing the cluster

Login Environment

- The cluster can be accessed through 8 general login nodes.
- The login nodes are the primary gateway to the rest of the cluster.
- All libraries, compilers, preinstalled applications, user installed applications are available over login nodes.

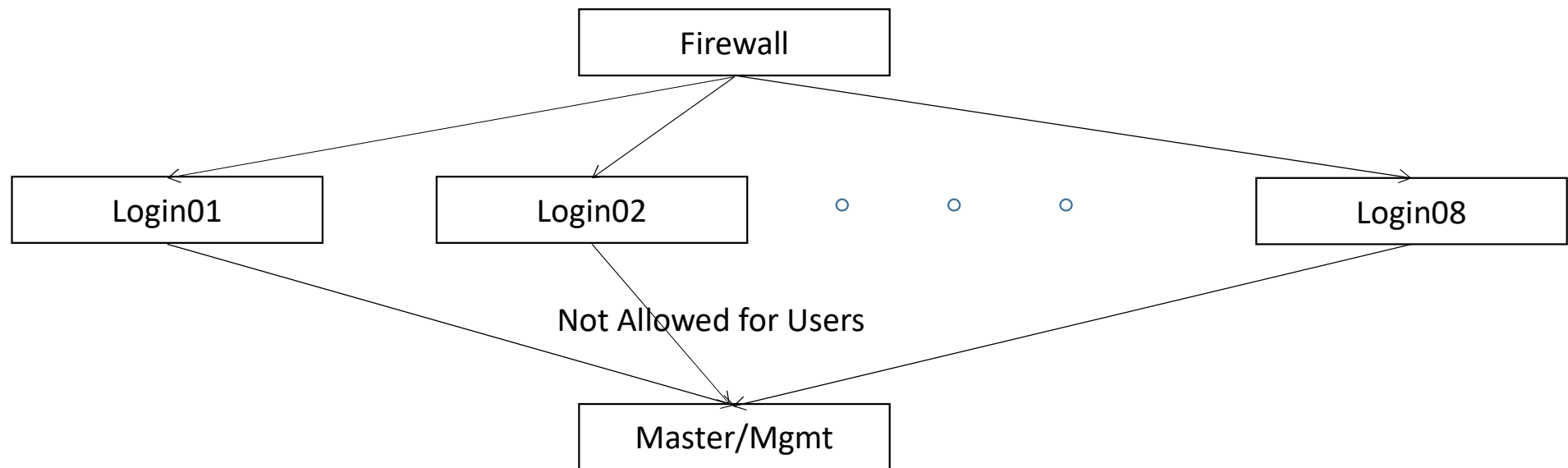
Remote Login

- You may access login nodes through ssh.
- Using SSH in Windows (Putty, Moab-xterm, etc).
- Using SSH in Linux via terminal.
- For example, to connect to the PARAM Ganga Login Node, with the username.
- **Ex:** `ssh -p 4422 testuser@paramganga.iitr.ac.in`

Access Policy



- Access to Login nodes are in Round-Robin Mode.
- Users are not allowed to access Master/Management Nodes



How to Access PARAM Ganga ?



- If you are using windows you can access via(SSH Clients):

- MobaXterm
- Putty, etc

- Within IIT Roorkee Campus:

ssh [username@paramganga.iitr.ac.in](ssh://username@paramganga.iitr.ac.in)

- Outside IIT Roorkee Campus:

ssh [username@paramganga.iitr.ac.in](ssh://username@paramganga.iitr.ac.in) -p 4422

Ticketing Tool



- A Support Portal will be created for Assisting the Users.

<https://paramganga.iitr.ac.in/support>



Frequently Asked Questions

[General FAQ](#)

[Environment](#)

[Job Submission](#)

[Applications](#)

[ML / DL](#)

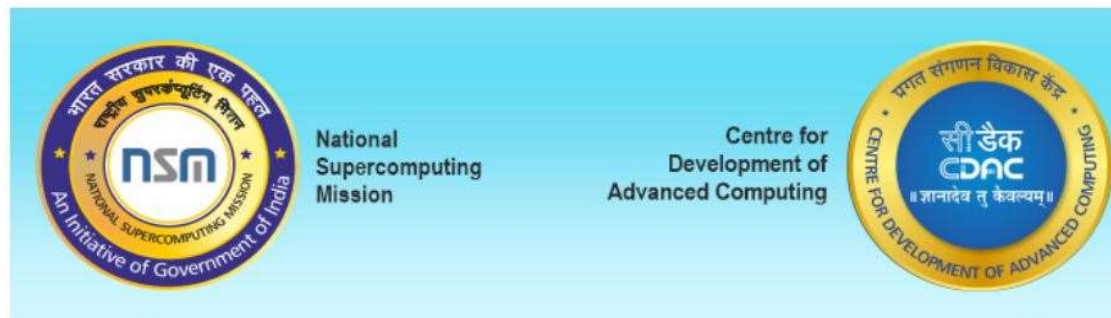
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General FAQ

How to get account on HPC cluster ?

- Get 'User Account Creation Form'
- Fill the relevant details.
- Get the signatures of your Head of the Department and the 'approving authority'.
- You will receive an Email in your official Email ID intimating the creation of your account

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[Can I run MS Windows applications on HPC?](#)

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[How much of the file/space quota I have used ?](#)

[My account expired ! What should I do? Is my data gone forever?](#)



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if your application is hybrid :

```
export I_MPI_FALLBACK="0" [ Do not switch to other available network ]
```

```
export I_MPI_FABRICS="shm:ofa"
```

```
export I_MPI_FABRICS="shm:dapl" (if using DAPL)
```

For OpenMP :

```
export I_MPI_FABRICS="shm:shm"
```

To check which fabric is currently used, you can set the I_MPI_DEBUG environment variable to 2:

```
mpirun -np n -genv I_MPI_DEBUG=2 your_command/command_path ; where "n" => number of
```

processes.

For Ex. : `mpirun -np 48 -genv I_MPI_DEBUG=2 myprog`

You can also specify above variables in your mpirun command :

```
mpirun -np n -genv I_MPI_FALLBACK=0 -genv I_MPI_FABRICS="shm:ofa"
```

`your_command/command_path`

For Ex. : `mpirun -np 48 -genv I_MPI_FALLBACK=0 -genv I_MPI_FABRICS="shm:ofa" myprog`

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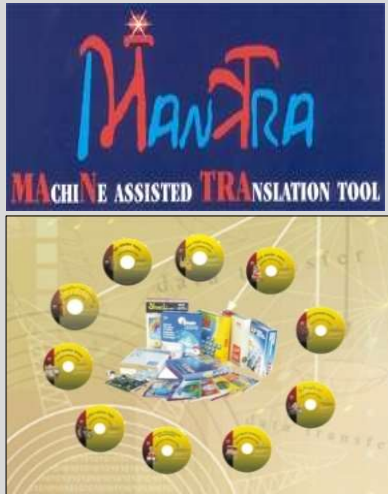
[My account expired ! What](#)

[Cannot find your queries ? Click here to create a ticket.](#)

Next

[Environment](#)





One Vision.

One Goal..

Advanced Computing

for Human Advancement...

Thank You !



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धन्यवाद आभार आभार यन्त्राद धन्यवाद नुंन्नि നന്മ ഭനന്തന