Using the Linux Cluster Lab

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1 Cluster Configuration

The Linux Cluster Lab in ET213-214 has one master node, named node00, and 32 compute nodes, named node01, . . . , node32, which are connected with a private Gigabit Ethernet switch.

The machine onyx.boisestate.edu is the master node and the only one that is on the public Internet. The master node onyx.boisestate.edu is connected to the private switch with the name node00. The figure below shows the layout of the Linux Cluster Lab.

To use the cluster you must login to onyx.boisestate.edu. Your home directory resides on the master node, which is the file server for the cluster. Thus you have the same home directory on all machines in the cluster.

2 Setup

Your home directory is shared among all nodes of the cluster. The master node acts as a network file server for the home directories.

When you log in to the master node, your account is already setup to use MPICH2 as the default MPI implementation.
3 Working on the Cluster

3.1 Accessing the cluster from a remote system

The following are some common commands for accessing the master node of the cluster from a remote site.

- **ssh onyx.boisestate.edu** starts a secure remote login session on the master node of the cluster.
- **ssh onyx.boisestate.edu <command>** runs the command remotely on the cluster and redirects the output to your terminal.
- **scp** lets you copy files to remote machines.

Note that the letter s stands for secure. All these commands encrypt your data on the network to protect against eavesdroppers. These commands replace the insecure equivalents (rlogin, rsh, rcp) that have been used commonly for accessing remote machines. Note that rsh commands work internally on the cluster (to support some software that needs it) but it is operating behind a firewall on a private network.

The command scp also has an option -r that lets you copy directories. In addition, you can specify a directory to copy to on the remote machine. Remember that the pathnames are relative to your home directory on the remote machine. For example to copy recursively a directory named prog1 on your local system to the master node onyx.boisestate.edu under the directory cs430 or tt cs530 in your home directory, use the following command:

```
scp -r prog1 onyx.boisestate.edu:cs430/prog1
```

3.2 Useful cluster commands

3.2.1 Cluster Monitoring

The following is a set of useful cluster commands that are available on the cluster lab. They can be run from the master node in the cluster.

- **cchk** Check the machines on the cluster by pinging the machines. Quick way to check if any machines are down.
- **cdate** Check the current time and date on each node of the cluster. The master node is a NTP time server (and also a client to another trusted source of accurate time). All the other nodes in the cluster are NTP time service clients of the master node.
- **cmips** Check the total computing power of the cluster. The computing power is reported in MIPS (using the bogoMIPS value reported by the kernel).
- **cfree** Reports memory usage across the cluster.
• **cdisks** Reports /tmp scratch disk space usage across the cluster.

• **ctemp** Reports CPU temperatures across the cluster. May not be available on all clusters.

### 3.2.2 Parallel Shell

The cluster comes with a simple parallel shell named **pdsh**. The **pdsh** shell is handy for running commands across the cluster. See the man page, which describes the capabilities of **pdsh** in detail.

First you need to setup ssh certificate with the **ssh-setup** to allow password less access among the cluster nodes with the command:

One of the useful features is the capability of specifying all or a subset of the cluster. For example:

- **pdsh -a <command>** targets the `<command>` to all nodes of the cluster, including the master.
- **pdsh -a -x node00 <command>** targets the `<command>` to all nodes of the cluster except the master.
- **pdsh -w node[01-08] <command>** targets the `<command>` to the 8 nodes of the cluster named node01, node02, ..., node08.

Another utility that is useful for formatting the output of **pdsh** is **dshbak**. Here we will show some handy uses of **pdsh**.

- Show the current date and time on all nodes of the cluster.
  ```bash
  pdsh -a date
  ```

- Show the current load and system uptime for all nodes of the cluster.
  ```bash
  pdsh -a uptime
  ```

- The utility **dshbak** formats the output from **pdsh** by consolidating the output from each node. The option `-c` shows identical output from different nodes just once.
  ```bash
  pdsh -a ls -l /etc/ntp | dshbak -c
  ```

Here is a sample output:

```
[amit@onyx amit]$ pdsh -a ls -l /etc/ntp | dshbak -c
-----------------
ws[01-16]-----------------
total 16
  -rw-r--r-- 1 root root 8 Jun 4 11:53 drift
  -rw------- 1 root root 266 Jun 4 11:53 keys
  -rw-r--r-- 1 root root 13 Jun 4 11:53 ntpservers
  -rw-r--r-- 1 root root 13 Jun 4 11:53 step-tickers
-----------------
ws00
-----------------
total 16
```
4 Using Swarm to run multiple sequential programs

Using the swatm utility, you can run multiple sequential programs on multiple nodes of the cluster. This helps you improve the throughput but doesn’t speed up any particular instance of your program.

List the sequential programs that you want to run in one file (one per line). For example, we have the following commands in the file named myjobs

```
program1; program1
program2
program3
```

Then submit your jobs with the following command

```
swarm -f myjobs
```

which creates PBS jobs and submits them. By default, swarm will run two processes per node (assuming each node is a dual-processor) and create one PBS job per node. So the output from the two commands will show up in the output file corresponding to the PBS job generated by swarm.

If you like to get email when your jobs are done, then use:

```
swarm -f myjobs -m e
```

You may want to set up a .forward file on beowulf so that the email notification from PBS gets routed to your favorite mail address.

If your program is I/O or memory intensive, then you may want to tell swarm to only run one process per node. This can be done as follows:

```
swarm -f myjobs -n 1 -m e
```

Check the status of your jobs with the following command.

```
qstat -a
```
To kill a job, use
qdel @job#\.beowulf
To kill all your jobs, use
qdelall
You should use the above command with caution! For more information about swarm, see the man page:
man swarm

5 Compiling Parallel Programs on the Cluster

5.1 Compiling MPI programs on the cluster

First setup the appropriate MPI system for your account. The default is MPICH2, which should be fine for most users. Compiling MPI programs is pretty simple: just use mpicc instead of the usual C compiler.

Here is a sample Makefile.

http://cs.boisestate.edu/~amit/teaching/430/lab/Makefile.mpich2.sample

6 Running parallel programs interactively

To run a parallel program interactively on the cluster requires two steps.

- Acquire desired number of compute nodes via the Portable Batch System that controls compute node allocation to users.
- Running a MPICH2 MPI program. Use the mpiexec command.

6.1 Acquiring nodes from the PBS system

The cluster uses the Portable Batch Scheduling system to manage the resources effectively. To run a parallel program, the user needs to request nodes from the PBS system. The master node is a shared resource and is always allocated to the user. The compute nodes are allocated in an exclusive mode. Currently there is a time limit of one hour for the use of compute nodes at a time.

To check the status of the nodes in the PBS system, there are two choices:

- qstat -a gives a list of all jobs running on the cluster.
- qstat -n gives a list of all jobs running on the cluster along with the nodes allocated to the jobs.

To request n nodes, use the command pbsget on the master node. Here is a sample session.
[amit@onyx ~]$ pbsget -4

Allocate cluster nodes via PBS for running interactive parallel jobs.

Trying for 4 nodes

Scheduling an interactive cluster session with PBS.
Please end session by typing in exit.

Use qstat -n to see nodes allocated by PBS.

You may now run MPI, pvm, xpvm or pvmrun. They will automatically use only the nodes allocated by PBS.
If you are using pvm or xpvm, then please always halt the pvm system before exiting the PBS cluster session.
If you are using MPI, then please always halt the MPI daemons system before exiting the PBS cluster session using mpdalleexit command.

For running LAM MPI programs use the following command:
   mpirun i-np <#copies> [options] <program> [<prog args>]
For running MPICH2 MPI programs use the following command:
   mpiexec -n <#copies> [options] <program> [<prog args>]
For running PVM programs use the following command:
Usage: pvmrun -np <#copies> <executable> {<args>,...}

qsub: waiting for job 3613.onyx.boisestate.edu to start
qsub: job 3613.onyx.boisestate.edu ready

[amit@onyx PBS ~]:qstat -n

onx.boisestate.edu:

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3613.onyx.boise amit interact STDIN 17742 5 -- -- 00:30 R
--
   node00/0+node20/0+node19/0+node18/0+node17/0
The command `pbsget` attempts to allocate the requested number of nodes from the PBS system. If it succeeds, it starts a new shell with the prompt modified to have PBS in the prompt. Note that the environment variable `PBS_NODEFILE` contains the name of a file that contains the list of nodes allocated by PBS to the user. Now the user can run either PVM or MPI parallel programs. When the user is done they would type `exit` to end the interactive PBS session.

If the required number of nodes are not available, then `pbsget` will wait. A user can cancel the request by typing in `Ctrl-c` and try again later. Remember to use `qstat -n` to check the status of the nodes.

### 6.2 Running PVM programs

#### 6.2.1 Using the PVM console

To run a PVM program the PVM daemon must be running. To start the daemon, startup the PVM console with the following command.

```
pvm
```

This invokes the PVM control program. The `pvm` console program is set to automatically add the nodes allocated to you by PBS. You can check what machines are in the PVM system with the `conf` command.

```
[amit@onyx amit]$ pbsget -4
******************************************************************************
Allocate cluster nodes via PBS for running interactive parallel jobs.
******************************************************************************
Trying for 4 nodes
...

qsub: waiting for job 806.onyx.boisestate.edu to start
qsub: job 806.onyx.boisestate.edu ready
```
pvm: Using list of machines from PBS.

```
HOST   DTID ARCH SPEED   DSIG
ws00   40000 LINUXI386 1000 0x00408841
ws04   80000  LINUXI386 1000 0x00408841
ws03   c0000  LINUXI386 1000 0x00408841
ws02   100000 LINUXI386 1000 0x00408841
ws01   140000 LINUXI386 1000 0x00408841
```

In the above example, we allocated 4 nodes using the PBS system. Note that the PVM console program adds the four compute nodes plus the master node to the PVM system. PBS always allocates the master node since that is a shared resource. Normally you would not run your programs on the master nodes but use it for monitoring purposes.

If you have already started the daemon, then the `pvm` console program would inform you. The simplest thing to do is to halt the already running daemon and start a fresh one, as shown below:

```
[pvm]
pvm: Using list of machines from PBS.
pvmd already running.
pvm> halt
Terminated
[pvm]
pvm: Using list of machines from PBS.
pvm>
```

You can type `help` in the `pvm` console program to get a list of all commands. You can run your application using the `spawn` command from the PVM console. Here is the help on the `spawn` command.

```
pvm> help spawn
spawn  Spawn task
Syntax:  spawn [ options ] file [ arg ... ]
Options:  - (count)  number of tasks, default is 1
          - (host)  spawn on host, default is any
          - (host):(wd) spawn on host, in directory 'wd'
          --host=(IP)  spawn on host with given IP addr
          --host=(IP):(wd)  spawn on IP, in directory 'wd'
          -(ARCH)  spawn on hosts of ARCH
          -(ARCH):(wd)  spawn on hosts of ARCH, in 'wd'
          -:(wd)  spawn in working directory 'wd'
          -?  enable debugging
          ->  redirect job output to console
          ->(file)  redirect output of job to file
          ->>(file)  append output of job to file
```

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The following figure continues the example by running a parallel program using the spawn command. The program is a SPMD style program. Thus we need to spawn a number of copies together. The output from the various nodes is captured together with the -> option to the spawn command. Note that each output is tagged with the task id of the task creating that output.

```
pvm> spawn -4 -> spmd_sum 10000
[1]
 4 successful
t80001
tc0001
t100001
t140001
pvm> [1:tc0001] EOF
[1:t80001] I got 2500 from 0
[1:t80001] I got 2500 from 1
[1:t80001] I got 2500 from 3
[1:t80001] I got 2500 from 2
[1:t80001] The total is 10000
[1:t80001] EOF
[1:t140001] EOF
[1:t100001] EOF
[1] finished
```

There is no limitation in PVM as to the number of tasks versus the number of nodes. So we could have invoked 12 tasks in the above example even though we acquired only 4 compute nodes. By default the PVM system distributes the tasks on the available nodes in a round robin fashion.

Finally you must remember to halt the PVM system and exit out of the PBS shell to release all resources.

```
pvm>
pvm> halt
Terminated
[amit@onyx PBS ~]:exit
logout
```

Note that MPMD (or master/slave) style programs can be invoked directly from the PBS shell if you have started the pvm daemon before. For example:

```
[amit@onyx amit]$ pbsget -4
```
Allocate cluster nodes via PBS for running interactive parallel jobs.

Trying for 4 nodes...
qsub: waiting for job 807.onyx.boisestate.edu to start
qsub: job 807.onyx.boisestate.edu ready

[amit@onyx PBS ~]:pvm
pvm: Using list of machines from PBS.
pvm> quit

Console: exit handler called
pvmd still running.

[amit@onyx PBS ~]:psum 10000 4
Starting 4 copies of spsum
I got 2500 from 2
I got 2500 from 0
I got 2500 from 1
I got 2500 from 3
The total is 10000

[amit@onyx PBS ~]:exit
logout

qsub: job 807.onyx.boisestate.edu completed

The above technique does not work for SPMD programs. It also will not show you the output from other nodes (unless you capture it using library calls in PVM). Finally, note that it is possible to embed the start up of PVM daemon and adding of nodes inside a program, making PVM transparent.

For more information on the pvm console program, see the man page for it.

6.2.2 Using the XPVM GUI

The xpvm graphical user interface to PVM is a nice visualization tool for developing, testing and debugging PVM programs. The xpvm program has built in help and is intuitive and simple to use. Similar to the PVM console program, pvm, the xpvm program starts up the PVM system with the list of nodes allocated by the PBS system. Then the user can use it to run parallel programs, visualize the execution, look at the output from all nodes, debug multiple nodes simultaneously, check utilization of the cluster during execution of the parallel program, check message queue length during execution and other features. A powerful feature of xpvm is its ability to graphically playback the sequence of events during an execution. This allows the user to investigate the behavior of a parallel program without having to spend a lot of time rerunning the parallel program. The following figure shows a screenshot of xpvm in action.
As with the PVM console program, remember to halt the PVM system from the xpvm GUI when you are done and then exit from the PBS session to release all resources.

To start xpvm, acquire nodes using `pbsget` and then start `xpvm`.

```
[amit@onyx amit]$ pbsget -4
...
qsub: waiting for job 820.onyx.boisestate.edu to start
qsub: job 820.onyx.boisestate.edu ready

[amit@onyx PBS ~]:xpvm
xpvm: Using list of machines from PBS.
New PVMD started... XPVM 1.2.5 connected as TID=0x40001.
[globs.tcl][procs.tcl][util.tcl]
Initializing XPVM.............................................. done.
```
The `xpvm` will start up in a new window. Initially it will ask you want to overwrite the trace file. You can answer “Yes” here. The main `xpvm` window will have two panels: *Network View* and *Space Time View*.

**Running a PVM program.** Go to *Tasks...* menu, then click on *Spawn* option to get the spawn window. In the *Command:* field you can type in the name of your program along with the command line arguments. Next select the *PvmHostCompl* button. Then select the *Host* button. A new field labeled *Host* will show up. Type the internal name of the master node, `node00`, here. The purpose of the last two steps to ensure that the PVM processes are not scheduled on the master node. Next type in the number of SPMD tasks you want to spawn in the *NTasks* field. Finally click on the *Start* button to start the parallel program.
Checking output from your program. As the parallel program runs, the Network View and the Space Time View are updated. You can also see the combined output from all the nodes by selecting the Views... menu and selecting the Task Output choice. A separate task output window will appear.

Other useful features. You can also go to the Views... menu and select the Utilization choice. This will show the node utilization in a separate window. Similarly you can look at message queue length with Views... and then Message Queue. The following screenshot shows some of the above examples.

As with the PVM console program, remember to halt the PVM system from the xpvm GUI when you are done and then exit from the PBS session to release all resources. To halt PVM, go to the File menu and then select Halt PVM choice.
6.2.3 Using the pvmrun application

With PVM you need to create the PVM system with either the pvm console or the xpvm application. Once the PVM system is operational, then you run your program. This is undesirable for several reasons. For example, if you forget to start the PVM system, your applications fails. If you left the PVM system running, then you will connect to what is already running. In order to avoid these problems, we have created an application named pvmrun. Running pvmrun with the help option shows its usage.

```
[amit@onyx amit]$ pvmrun -h
pvmrun: $Id: pvmrun.c,v 1.8 2004/09/02 00:00:28 amit Exp $
Usage: pvmrun -np <#tasks> <executable> {<args>,...}
************************************************************************
For a pvm master/slave program: pvmrun -np 1 <executable>
{<args>,...}
For a pvm n-way spmd program: pvmrun -np n <executable> {<args>,...}
************************************************************************
```

The list of machines to use for PVM is determined as follows:
if PBS_NODEFILE is set, then use the file specified by PBS
else: exit and ask user to run after getting nodes
using pbsget.

The pvmrun program starts the pvm system for you, runs your application (either master/slave MPMD style or SPMD style) and then cleanly shuts down the PVM system. If the PVM system is already running, it shuts it down and restarts it with the list of nodes supplied by the PBS system. **You must use pvmrun in your PBS batch jobs.**

The following shows the usage of pvmrun to run a parallel program. Note that we show the program being run twice. The second time we have redirected the standard error output from pvmrun to a file so only the output from the application shows up.

```
[amit@onyx parallel_sum]$ pbsget -4
************************************************************************
Allocate cluster nodes via PBS for running interactive parallel jobs.
************************************************************************
Trying for 4 nodes
...
qsub: waiting for job 850.onyx.boisestate.edu to start
qsub: job 850.onyx.boisestate.edu ready

[amit@onyx PBS ~/cs430/lab/PVM/parallel_sum]:pvmrun -np 4 spmd_sum 10000
pvmrun: Using host file: /var/spool/pbs/aux/850.onyx.boisestate.edu
pvmrun: Probing for pre-existing PVM.
************************************************************************
libpvm [pid10012] /tmp/pvmd.999: No such file or directory
libpvm [pid10012] /tmp/pvmd.999: No such file or directory
libpvm [pid10012] /tmp/pvmd.999: No such file or directory
libpvm [pid10012]: pvm_mytid(): Can’t contact local daemon
```
pvmrun: PVM not running. Ignore error messages above from pvm_mytid().
pvmrun: Succeeded in starting new PVM daemon!
Selected nodes from file: /var/spool/pbs/aux/850.onyx.boisestate.edu
ws00
ws04
ws03
ws02
ws01
pvmrun: Succeeded in adding 5 hosts.
pvmrun: Parallel Virtual Machine configuration
-----------------------------------------------
ws00
ws04
ws03
ws02
ws01
-----------------------------------------------
[tc0001] BEGIN
[t100001] BEGIN
[t140001] BEGIN
[t80001] BEGIN
[t80001] I got 2500 from 0
[t80001] I got 2500 from 1
[t80001] I got 2500 from 2
[t80001] I got 2500 from 3
[t80001] The total is 10000
[t100001] EOF
[tc0001] EOF
Received PvmTaskExit notification from task id 100001
Received PvmTaskExit notification from task id c0001
[t140001] EOF
[t80001] EOF
Received PvmTaskExit notification from task id 140001
Received PvmTaskExit notification from task id 80001

pvmrun: 4 tasks finished!
pvmrun: Halting PVM.
pvmrun: End of PVM application.
[amit@onyx PBS ~/cs430/lab/PVM/parallel_sum]:pvmrun -np 4 spmd_sum 10000 2> errorlog
[tc0001] BEGIN
[t140001] BEGIN
[t100001] BEGIN
[t80001] BEGIN
[tc0001] EOF
I got 2500 from 0
I got 2500 from 1
I got 2500 from 3
I got 2500 from 2
The total is 10000
EOF
EOF
EOF
EOF

6.3 Running MPI programs

- Running a MPI program
  - Log in to the head node onyx. Acquire nodes from the scheduler: `pbsget -4`
  - Next run your mpi program with the `mpiexec` command.
    `mpiexec -n 4 hello_world`
  - Exit and release all allocated nodes with the command: `exit`

Here is a sample session.

```
[amit@onyx hello-world]$ pbsget -4

******************************************************************************
Allocate cluster nodes via PBS for running interactive parallel jobs.
******************************************************************************

Trying for 4 nodes

******************************************************************************
Scheduling an interactive cluster session with PBS. 
Please end session by typing in exit.

Use qstat -n to see nodes allocated by PBS. 
or use xpbsmon to graphically see nodes allocated by PBS.

You may now run MPI, pvm, xpvm or pvmrun. They will automatically use only the nodes allocated by PBS.
If you are using pvm or xpvm, then please always halt the pvm system before exiting the PBS cluster session.
If you are using MPI, then please always halt the MPI daemons system before exiting the PBS cluster session using mpdallexit command.

For running LAM MPI programs use the following command:
  mpirun i-np <#copies> [options] <program> [<prog args>]
```
For running MPICH2 MPI programs use the following command:
  mpiexec -n <#copies> [options] <program> [<prog args>]
For running PVM programs use the following command:
Usage: pvmrun -np <#copies> <executable> {<args>,...}

qsub: waiting for job 3614.onyx.boisestate.edu to start
qsub: job 3614.onyx.boisestate.edu ready

[amit@onyx PBS ~/cs430/lab/MPI/hello-world]:mpdboot
Starting 5 mpds
[amit@onyx PBS ~/cs430/lab/MPI/hello-world]:mpiexec -n 4 spmd_hello_world
Hello! I am 0 of 4 running on node17.
Hello! I am 1 of 4 running on node18.
Hello! I am 3 of 4 running on node20.
Hello! I am 2 of 4 running on node19.

[amit@onyx PBS ~/cs430/lab/MPI/hello-world]:mpdallexit
[amit@onyx PBS ~/cs430/lab/MPI/hello-world]:exit
logout

qsub: job 3614.onyx.boisestate.edu completed
[amit@onyx hello-world]$

7 Running parallel programs in batch mode

7.1 Getting ready to run PBS batch jobs

To be able to run PBS jobs, you must check your .bash_profile and .bashrc files carefully. Any command that manipulates the terminal must be put in a conditional statement of the following form.

if test "$PBS_ENVIRONMENT" = "PBS_INTERACTIVE" -o -z "$PBS_ENVIRONMENT"
then
  # set up the prompt to the hostname
  #
  PS1="[\u@\h \W]":
fi

The reason is that there is no terminal when you are running in batch mode. If you try to manipulate the terminal in batch mode, your login will fail and your batch job will not run. The environment variable PBS_ENVIRONMENT is set by PBS to be either PBS_INTERACTIVE or PBS_BATCH. When you are not running under PBS, then the variable is unset.
## 7.2 Preparing a PBS batch job script

Any parallel program that takes more than a few minutes should normally be run as a PBS batch job. In order to run it as a PBS batch job, you will need to prepare a PBS batch script (which is just a shell script with some additional features). Here is a sample PBS batch job (~/amit/cs430/lab/PVM/tools/psort.pbs):

```bash
#!/bin/sh
#PBS -l nodes=16:node
# This is a PBS job submission script. It asks for the master node
# and 16 nodes in the PBS cluster to run the PVM application on.
#
# IMPORTANT NOTE: Be sure to modify the "cd" command below to switch
# to the directory in which you are currently working!
#
#---------------------------------------------------------------
cd /home/faculty/amit/cs430/lab/PVM/tools
pvmrun -np 16 psort 20000000 16
```

The line starting with `#PBS` is a PBS directive. There are many PBS directives but the one we will use is mainly the one that lists the nodes that we need to run our program. The following list shows some common options that can be used in the PBS directives:

<table>
<thead>
<tr>
<th>PBS option</th>
<th>Description</th>
</tr>
</thead>
<tbody>
<tr>
<td>-N jobname</td>
<td>name the job jobname</td>
</tr>
<tr>
<td>-l cput=N</td>
<td>request N seconds of CPU time; N can also be in hh:mm:ss form</td>
</tr>
<tr>
<td>-l mem=N[KMG][BW]</td>
<td>request N kilo— mega— gigabytes— words of memory</td>
</tr>
<tr>
<td>-l nodes=N:ppn=M</td>
<td>request N nodes with M processors per node</td>
</tr>
<tr>
<td>-m e</td>
<td>mail the user when the job completes</td>
</tr>
<tr>
<td>-m a</td>
<td>mail the user if the job aborts</td>
</tr>
<tr>
<td>-a 1800</td>
<td>Start job after 6pm</td>
</tr>
<tr>
<td>-o outfile</td>
<td>redirect standard output to outfile</td>
</tr>
<tr>
<td>-e errfile</td>
<td>redirect standard error to errfile</td>
</tr>
<tr>
<td>-j oe</td>
<td>combine standard output and standard error</td>
</tr>
</tbody>
</table>

For a full list, see the man page for `pbs_resources` on the cluster.

Here is another sample PBS batch job. Here the `psum` program is assumed to spawn processes to the 16 nodes.

```bash
#!/bin/sh
#PBS -l nodes=16:node
# This is a PBS job submission script. It runs a master/slave PVM program
# Note that even though we are specifying only one process to pvmrun, we
# need to reserve the appropriate number of nodes to match what the parallel
# program requires.
#
# IMPORTANT NOTE: Be sure to modify the "cd" command below to switch
# to the directory in which you are currently working!
```
# cd /home/faculty/amit/cs430/lab/PVM/parallel_sum
pvmrun -np 1 psum 10000 16

Here is a sample PBS batch script for a MPICH2 program.

```bash
#!/bin/sh
#PBS -l nodes=4:node
cd ${HOME}/MPI/hello_world
mpiexec -n 4 spmd_hello_world
```

### 7.3 Submitting a PBS batch job script

The command `qsub` can be used to submit a PBS job. Please see the man page for `qsub` for a variety of options that can be used with it. Continuing with the example script `psort.pbs` from the previous subsection, we can submit it for execution as follows.

```
cd ~amit/cs430/lab/PVM/tools/
qusub psort.pbs
```

The status of a job can be checked with the `qstat` command. Using `qstat -n` also shows the nodes that were allocated to your job.

```
[amit@onyx tools]:qsub psort.pbs
[amit@onyx tools]:qstat

116.onyx.boisestate.edu
```

```
Job id Name User Time Use S Queue
---------------- ------------------ -------------- -------- - ----- 
116.onyx.psort.pbs amit 0 Q default
```

You can delete jobs with the `qdel` command.

The standard output and standard error streams are redirected into the files `psort.pbs.oxxx` and `psort.pbs.exxx`, where `xxx` is the job number assigned by PBS.

In case of an error in running the job after it has been accepted in the queue, PBS sends an email to the user.
The following shows another example of submitting a job. This time we are specifying that the job should start anytime after the time specified in the `qsub` command, that we should be emailed at the start and end of the job, and the job is to be named `test1`.

```
[amit@onyx parallel_sum]$ qsub -m be -a 0400 -N test1 psum.pbs
853.onyx.boisestate.edu
[amit@onyx parallel_sum]$ qstat -n

onyx.boisestate.edu:

<table>
<thead>
<tr>
<th>Job ID</th>
<th>Username</th>
<th>Queue</th>
<th>Jobname</th>
<th>SessID</th>
<th>NDS TSK</th>
<th>Memory</th>
<th>Time</th>
<th>S Time</th>
</tr>
</thead>
<tbody>
<tr>
<td>853.onyx.boises amit default test1</td>
<td>17</td>
<td>--</td>
<td>00:30 W</td>
<td>--</td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
</tbody>
</table>

[amit@onyx parallel_sum]$ date
Thu Sep 9 03:57:33 MDT 2004
[amit@onyx parallel_sum]$ date
Thu Sep 9 04:00:23 MDT 2004
[amit@onyx parallel_sum]$ qstat -n
[amit@onyx parallel_sum]$ ls -l test1.*853
-rw------- 1 amit faculty 2047 Sep 9 04:00 test1.e853
-rw------- 1 amit faculty 1850 Sep 9 04:00 test1.o853

----------------------Mail message-----------------------------------
Date: Thu, 9 Sep 2004 04:00:00 -0600
From: adm <adm@onyx.boisestate.edu>
To: amit@onyx.boisestate.edu
Subject: PBS JOB 853.onyx.boisestate.edu

PBS Job Id: 853.onyx.boisestate.edu
Job Name: test1
Begun execution

----------------------Mail message-----------------------------------
Date: Thu, 9 Sep 2004 04:00:02 -0600
From: adm <adm@onyx.boisestate.edu>
To: amit@onyx.boisestate.edu
Subject: PBS JOB 853.onyx.boisestate.edu

PBS Job Id: 853.onyx.boisestate.edu
Job Name: test1
Execution terminated
Exit_status=0
resources_used.cput=00:00:00
resources_used.mem=312kb
resources_used.vmem=1440kb
resources_used.walltime=00:00:02

-------------------------------
```
8 Debugging PVM programs

8.1 Using print statements

Any print statements in the process running on the local machine will show up on the console. However, prints from processes running on remote machines will not show up on your console. To be able to see the output from all processes, you have three options.

- **pvm control program**: Run the application from the `pvm` control program using the `spawn` command. Giving the appropriate options will redirect all output from all processes in your application to the console. See the following example, which starts a master process that further spawns 3 worker processes.

```
pvm> spawn -1 -> psum 100 3
spawn -1 -> psum 100 3
[1]
1 successful
t40002
pvm> [1:t40002] Starting 3 copies of spsum
[1:t40003] Sending result back to master
[1:t40004] Sending result back to master
[1:t40002] I got 33 from 0
[1:t40002] I got 33 from 1
[1:t40003] EOF
[1:t40005] Sending result back to master
[1:t40002] I got 33 from 2
[1:t40002] The total is 99
[1:t40002] EOF
[1:t40004] EOF
[1:t40005] EOF
[1] finished
```

- **XPVM**: If you are running the PVM application under XPVM, then the output from all processes can be seen by going to the menu option `Views --> Task Output`, which shows the output from the processes in a separate window.

- **PVM daemon log**: The PVM daemon maintains a log in the file `/tmp/pvml.xxx`, where `xxx` is your numerical user id (find out your user id with the command `id`). This log file contains the output from all the remote processes.

8.2 Using a debugger

Using the debugger is recommended when running a small number of processes. If you supply the `PvmTaskDebug` flag in the function `pvm_spawn(...)`, it will attempt to start each task in a separate window under the control of a debugger. The same effect can be achieved by clicking on
the \texttt{PvmTaskDebug} button in the \textit{Spawn} subwindow in \texttt{xpv}m. Under LINUX the default debugger is \texttt{gdb}. The default debugger can be changed by setting the \texttt{PVM\_DEBUGGER} environment variable.

\section{How to Setup and use MPICH2 on your Linux PC or notebook}

\subsection{How to download and install MPICH2}

\begin{itemize}
\item Download the tarball of the software from \url{http://www-unix.mcs.anl.gov/mpi/mpich2/}. Unpack it somewhere, say in \texttt{/usr/local/src} with the command:

\begin{verbatim}
cd /usr/local/src
 tar xzvf mpich2-xyz.tar.gz
\end{verbatim}

\item Assuming that you have Sun Java installed in \texttt{/usr/local/java}, I recommend the following steps to build MPICH2.

\begin{verbatim}
mkdir /usr/local/src/mpich2
./configure --prefix=/usr/local/mpich2 --enable-mpe --enable-cxx --enable-romio --with-java-home=/usr/local/java/jre 2>&1 | tee configure.log
make 2>&1 | tee make.log
make install 2>&1 | tee install.log
\end{verbatim}

\item For more details on installation, please read the instructions in the README file in the MPICH2 source.

\item Add \texttt{/usr/local/mpich2/share/man} in the \texttt{/etc/man.config} file to enable viewing man pages in KDE Konqueror.
\end{itemize}

\subsection{Using MPICH2 on your computer}

\begin{itemize}
\item \textbf{Create MPI config file.} Create the file \texttt{~/.mpd.conf} and enter one line that contains a secret code word. The commands to do this are shown below. Remember to put your own secret word in your file.

\begin{verbatim}
touch ~/.mpd.conf
 echo "MPD_secretword=abracadabara" > ~/.mpd.conf
 chmod 600 ~/.mpd.conf
\end{verbatim}

\item \textbf{Setting up MPICH2.} Assume that you installed MPICH2 in the folder \texttt{/usr/local/mpich2} on your computer. Then add the following at the end of your \texttt{~/.bashrc} file and source it with the command \texttt{source ~/.bashrc} (or log in again).
\end{itemize}
# MPICH2 setup
export PATH=/usr/local/mpich2/bin:$PATH
export MANPATH=/usr/local/mpich2/man:$MANPATH

• Running a MPI program
  – Run your mpi program with the mpiexec command.
    mpiexec -n 4 hello_world

10 Documentation

Here are some sources of documentation:

• There are man pages for all MPI calls. For example, to read the man page for MPI_Send(), type:
  
  man MPI_Send

  In the KDE desktop, you can use Alt+F2 to get a command window, where you can type
  man:MPI_Send to get the man page in hypertext.

• The MPI reference book is also available online at