

# Sample PBS Script for Pleiades

## Category: PBS on Pleiades

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#PBS -S /bin/csh
#PBS -N cfd
# This example uses the Harpertown nodes
# User job can access ~7.6 GB of memory per Harpertown node.
# A memory intensive job that needs more than ~0.9 GB
# per process should use less than 8 cores per node
# to allow more memory per MPI process. This example
# asks for 64 nodes and 4 MPI processes per node.
# This request implies 64x4 = 256 MPI processes for the job.
#PBS -l select=64:ncpus=8:mpiprocs=4:model=har
#PBS -l walltime=4:00:00
#PBS -j oe
#PBS -W group_list=a0801
#PBS -m e

# Currently, there is no default compiler and MPI library set.
# You should load in the version you want.
# Currently, MVAPICH or SGI's MPT are available in 64-bit only,
# you should use a 64-bit version of the compiler.

module load comp-intel/11.1.072
module load mpi-sgi/mpt.2.04.10789

# By default, PBS executes your job from your home directory.
# However, you can use the environment variable
# PBS_O_WORKDIR to change to the directory where
# you submitted your job.

cd $PBS_O_WORKDIR

# use of dplace to pin processes to processors may improve performance
# Here you request to pin processes to processors 2, 3, 6, 7 of each node.
# This helps for using the Harpertown nodes, but not for Nehalem-EP or
# Westmere-EP nodes

# The resource request of select=64 and mpiprocs=4 implies
# that you want to have 256 MPI processes in total.
# If this is correct, you can omit the -np 256 for mpiexec
# that you might have used before.

mpiexec dplace -s1 -c2,3,6,7 ./grinder < run_input > output

# It is a good practice to write stderr and stdout to a file (ex: output)
# Otherwise, they will be written to the PBS stderr and stdout in /PBS/spool,
# which has limited amount of space. When /PBS/spool is filled up, any job
# that tries to write to /PBS/spool will die.

# -end of script-
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<http://www.nas.nasa.gov/hecc/support/kb/entry/190/?ajax=1>