

Assignment04

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1.

modifications are marked with ####

hmd.c :

```
/*-----
Program pmd.c performs parallel molecular-dynamics for Lennard-Jones
systems using the Message Passing Interface (MPI) standard.
-----*/
#include "hmd.h" // ####

/*-----*/
int main(int argc, char **argv) {
/*-----*/
    double cpu1;

    MPI_Init(&argc,&argv); /* Initialize the MPI environment */
    MPI_Comm_rank(MPI_COMM_WORLD, &sid); /* My processor ID */
    /* Vector index of this processor */
    vid[0] = sid/(vproc[1]*vproc[2]);
    vid[1] = (sid/vproc[2])%vproc[1];
    vid[2] = sid%vproc[2];

    omp_set_num_threads(nthrd); // ####

    init_params();
    set_topology();
    init_conf();
    atom_copy();
    compute_accel(); /* Computes initial accelerations */

    cpu1 = MPI_Wtime();
    for (stepCount=1; stepCount<=StepLimit; stepCount++) {
        single_step();
        if (stepCount%StepAvg == 0) eval_props();
    }
    cpu = MPI_Wtime() - cpu1;
    if (sid == 0) printf("CPU & COMT = %le %le\n",cpu,comt);

    MPI_Finalize(); /* Clean up the MPI environment */
    return 0;
}

/*-----*/
void init_params() {
/*-----
Initializes parameters.
-----*/

    int a;
    double rr,ri2,ri6,r1; // ####
    FILE *fp;
```

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/* Read control parameters */
fp = fopen("pmd.in","r");
fscanf(fp,"%d%d%d",&InitUcell[0],&InitUcell[1],&InitUcell[2]);
fscanf(fp,"%le",&Density);
fscanf(fp,"%le",&InitTemp);
fscanf(fp,"%le",&DeltaT);
fscanf(fp,"%d",&StepLimit);
fscanf(fp,"%d",&StepAvg);
fclose(fp);

/* Compute basic parameters */
DeltaTH = 0.5*DeltaT;
for (a=0; a<3; a++) al[a] = InitUcell[a]/pow(Density/4.0,1.0/3.0);
if (sid == 0) printf("al = %e %e %e\n",al[0],al[1],al[2]);

/* Compute the # of cells for linked cell lists */
for (a=0; a<3; a++) {
    lc[a] = al[a]/RCUT;

    /* Size of cell block that each thread is assigned */
    thbk[a] = lc[a]/vthrd[a];
    /* # of cells = integer multiple of the # of threads */
    lc[a] = thbk[a]*vthrd[a]; /* Adjust # of cells/MPI process */

    rc[a] = al[a]/lc[a];
}
if (sid == 0) {
    printf("lc = %d %d %d\n",lc[0],lc[1],lc[2]);
    printf("rc = %e %e %e\n",rc[0],rc[1],rc[2]);
}

/* Constants for potential truncation */
rr = RCUT*RCUT; ri2 = 1.0/rr; ri6 = ri2*ri2*ri2; r1=sqrt(rr);
Uc = 4.0*ri6*(ri6 - 1.0);
Duc = -48.0*ri6*(ri6 - 0.5)/r1;
}

/*-----*/
void set_topology() {
/*-----*/
Defines a logical network topology. Prepares a neighbor-node ID table,
nn, & a shift-vector table, sv, for internode message passing. Also
prepares the node parity table, myparity.
/*-----*/

/* Integer vectors to specify the six neighbor nodes */
int iv[6][3] = {
    {-1,0,0}, {1,0,0}, {0,-1,0}, {0,1,0}, {0,0,-1}, {0,0,1}
};
int ku,a,k1[3];

/* Set up neighbor tables, nn & sv */
for (ku=0; ku<6; ku++) {
    /* Vector index of neighbor ku */
    for (a=0; a<3; a++)
        k1[a] = (vid[a]+iv[ku][a]+vproc[a])%vproc[a];
    /* Scalar neighbor ID, nn */
    nn[ku] = k1[0]*vproc[1]*vproc[2]+k1[1]*vproc[2]+k1[2];
    /* Shift vector, sv */
    for (a=0; a<3; a++) sv[ku][a] = al[a]*iv[ku][a];
}

/* Set up the node parity table, myparity */
for (a=0; a<3; a++) {

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        if (vproc[a] == 1)
            myparity[a] = 2;
        else if (vid[a]%2 == 0)
            myparity[a] = 0;
        else
            myparity[a] = 1;
    }
}

/*-----*/
void init_conf() {
/*-----
r are initialized to face-centered cubic (fcc) lattice positions.
rv are initialized with a random velocity corresponding to Temperature.
-----*/

    double c[3],gap[3],e[3],vSum[3],gvSum[3],vMag;
    int j,a,nX,nY,nZ;
    double seed;
    /* FCC atoms in the original unit cell */
    double origAtom[4][3] = {{0.0, 0.0, 0.0}, {0.0, 0.5, 0.5},
                             {0.5, 0.0, 0.5}, {0.5, 0.5, 0.0}};

    /* Set up a face-centered cubic (fcc) lattice */
    for (a=0; a<3; a++) gap[a] = al[a]/InitUcell[a];
    n = 0;
    for (nZ=0; nZ<InitUcell[2]; nZ++) {
        c[2] = nZ*gap[2];
        for (nY=0; nY<InitUcell[1]; nY++) {
            c[1] = nY*gap[1];
            for (nX=0; nX<InitUcell[0]; nX++) {
                c[0] = nX*gap[0];
                for (j=0; j<4; j++) {
                    for (a=0; a<3; a++)
                        r[n][a] = c[a] + gap[a]*origAtom[j][a];
                    ++n;
                }
            }
        }
    }

    /* Total # of atoms summed over processors */
    MPI_Allreduce(&n,&nglob,1,MPI_INT,MPI_SUM,MPI_COMM_WORLD);
    if (sid == 0) printf("nglob = %d\n",nglob);

    /* Generate random velocities */
    seed = 13597.0+sid;
    vMag = sqrt(3*InitTemp);
    for(a=0; a<3; a++) vSum[a] = 0.0;
    for(j=0; j<n; j++) {
        RandVec3(e,&seed);
        for (a=0; a<3; a++) {
            rv[j][a] = vMag*e[a];
            vSum[a] = vSum[a] + rv[j][a];
        }
    }
    MPI_Allreduce(vSum,gvSum,3,MPI_DOUBLE,MPI_SUM,MPI_COMM_WORLD);

    /* Make the total momentum zero */
    for (a=0; a<3; a++) gvSum[a] /= nglob;
    for (j=0; j<n; j++)
        for(a=0; a<3; a++) rv[j][a] -= gvSum[a];
}

/*-----*/

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void single_step() {
/*-----*/
r & rv are propagated by DeltaT using the velocity-Verlet scheme.
/*-----*/
    int i,a;

    half_kick(); /* First half kick to obtain v(t+Dt/2) */
    for (i=0; i<n; i++) /* Update atomic coordinates to r(t+Dt) */
        for (a=0; a<3; a++) r[i][a] = r[i][a] + DeltaT*rv[i][a];
    atom_move();
    atom_copy();
    compute_accel(); /* Computes new accelerations, a(t+Dt) */
    half_kick(); /* Second half kick to obtain v(t+Dt) */
}

/*-----*/
void half_kick() {
/*-----*/
Accelerates atomic velocities, rv, by half the time step.
/*-----*/
    int i,a;
    for (i=0; i<n; i++)
        for (a=0; a<3; a++) rv[i][a] = rv[i][a]+DeltaTH*ra[i][a];
}

/*-----*/
void atom_copy() {
/*-----*/
Exchanges boundary-atom coordinates among neighbor nodes: Makes
boundary-atom list, LSB, then sends & receives boundary atoms.
/*-----*/
    int kd,kdd,i,ku,inode,nsd,nrc,a;
    int nbnew = 0; /* # of "received" boundary atoms */
    double com1;

/* Main loop over x, y & z directions starts-----*/

    for (kd=0; kd<3; kd++) {

        /* Make a boundary-atom list, LSB-----*/

        /* Reset the # of to-be-copied atoms for lower&higher directions */
        for (kdd=0; kdd<2; kdd++) lsb[2*kd+kdd][0] = 0;

        /* Scan all the residents & copies to identify boundary atoms */
        for (i=0; i<n+nbnew; i++) {
            for (kdd=0; kdd<2; kdd++) {
                ku = 2*kd+kdd; /* Neighbor ID */
                /* Add an atom to the boundary-atom list, LSB, for neighbor ku
                according to bit-condition function, bbd */
                if (bbd(r[i],ku)) lsb[ku][++(lsb[ku][0])] = i;
            }
        }

        /* Message passing-----*/

        com1=MPI_Wtime(); /* To calculate the communication time */

        /* Loop over the lower & higher directions */
        for (kdd=0; kdd<2; kdd++) {

            inode = nn[ku=2*kd+kdd]; /* Neighbor node ID */

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/* Send & receive the # of boundary atoms-----*/

nsd = lsb[ku][0]; /* # of atoms to be sent */

/* Even node: send & recv */
if (myparity[kd] == 0) {
    MPI_Send(&nsd,1,MPI_INT,inode,10,MPI_COMM_WORLD);
    MPI_Recv(&nrc,1,MPI_INT,MPI_ANY_SOURCE,10,
             MPI_COMM_WORLD,&status);
}
/* Odd node: recv & send */
else if (myparity[kd] == 1) {
    MPI_Recv(&nrc,1,MPI_INT,MPI_ANY_SOURCE,10,
             MPI_COMM_WORLD,&status);
    MPI_Send(&nsd,1,MPI_INT,inode,10,MPI_COMM_WORLD);
}
/* Single layer: Exchange information with myself */
else
    nrc = nsd;
/* Now nrc is the # of atoms to be received */

/* Send & receive information on boundary atoms-----*/

/* Message buffering */
for (i=1; i<=nsd; i++)
    for (a=0; a<3; a++) /* Shift the coordinate origin */
        dbuf[3*(i-1)+a] = r[lsb[ku][i]][a]-sv[ku][a];

/* Even node: send & recv */
if (myparity[kd] == 0) {
    MPI_Send(dbuf,3*nsd,MPI_DOUBLE,inode,20,MPI_COMM_WORLD);
    MPI_Recv(dbuf,3*nrc,MPI_DOUBLE,MPI_ANY_SOURCE,20,
             MPI_COMM_WORLD,&status);
}
/* Odd node: recv & send */
else if (myparity[kd] == 1) {
    MPI_Recv(dbuf,3*nrc,MPI_DOUBLE,MPI_ANY_SOURCE,20,
             MPI_COMM_WORLD,&status);
    MPI_Send(dbuf,3*nsd,MPI_DOUBLE,inode,20,MPI_COMM_WORLD);
}
/* Single layer: Exchange information with myself */
else
    for (i=0; i<3*nrc; i++) dbufr[i] = dbuf[i];

/* Message storing */
for (i=0; i<nrc; i++)
    for (a=0; a<3; a++) r[n+nbnew+i][a] = dbufr[3*i+a];

/* Increment the # of received boundary atoms */
nbnew = nbnew+nrc;

/* Internode synchronization */
MPI_Barrier(MPI_COMM_WORLD);

} /* Endfor lower & higher directions, kdd */

comt += MPI_Wtime()-com1; /* Update communication time, COMT */

} /* Endfor x, y & z directions, kd */

/* Main loop over x, y & z directions ends-----*/

/* Update the # of received boundary atoms */

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    nb = nbnew;
}

/*-----*/
void compute_accel() {
/*-----
Given atomic coordinates, r[0:n+nb-1][], for the extended (i.e.,
resident & copied) system, computes the acceleration, ra[0:n-1][], for
the residents.
-----*/

    int i,j,a,lc2[3],lcyz2,lcxyz2,mc[3],c,mcl[3],cl; // #####
    //int bintra; // #####
    double rrCut,lpe;
    double lpe_td[nthrd]; // #####

    /* Reset the potential & forces */
    lpe = 0.0;
    for (i=0; i<nthrd;i++) lpe_td[i] = 0.0; // #####
    for (i=0; i<n; i++) for (a=0; a<3; a++) ra[i][a] = 0.0;

    /* Make a linked-cell list, lscl-----*/

    for (a=0; a<3; a++) lc2[a] = lc[a]+2;
    lcyz2 = lc2[1]*lc2[2];
    lcxyz2 = lc2[0]*lcyz2;

    /* Reset the headers, head */
    for (c=0; c<lcxyz2; c++) head[c] = EMPTY;

    /* Scan atoms to construct headers, head, & linked lists, lscl */
    for (i=0; i<n+nb; i++) {
        for (a=0; a<3; a++) mc[a] = (r[i][a]+rc[a])/rc[a];

        /* Translate the vector cell index, mc, to a scalar cell index */
        c = mc[0]*lcyz2+mc[1]*lc2[2]+mc[2];

        /* Link to the previous occupant (or EMPTY if you're the 1st) */
        lscl[i] = head[c];

        /* The last one goes to the header */
        head[c] = i;
    } /* Endfor atom i */

    /* Calculate pair interaction-----*/

    rrCut = RCUT*RCUT;

#pragma omp parallel private(a,mc,c,mcl,cl,i,j)
{

    double dr[3],rr,ri2,ri6,r1,rrCut,fcVal,lpe,f,vVal;
    int std, vtd[3],mofst[3];

    std = omp_get_thread_num();
    vtd[0] = std/(vthrd[1]*vthrd[2]);
    vtd[1] = (std/vthrd[2])%vthrd[1];
    vtd[2] = std%vthrd[2];
    for (a=0; a<3; a++)
        mofst[a] = vtd[a]*thbk[a];

    // #####
    /* Scan inner cells */

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//for (mc[0]=1; mc[0]<=lc[0]; (mc[0])++)
//for (mc[1]=1; mc[1]<=lc[1]; (mc[1])++)
//for (mc[2]=1; mc[2]<=lc[2]; (mc[2])++) {
for (mc[0]=mofst[0]+1; mc[0]<=mofst[0]+thbk[0]; (mc[0])++)
for (mc[1]=mofst[1]+1; mc[1]<=mofst[1]+thbk[1]; (mc[1])++)
for (mc[2]=mofst[2]+1; mc[2]<=mofst[2]+thbk[2]; (mc[2])++) {

/* Calculate a scalar cell index */
c = mc[0]*lcyz2+mc[1]*lc2[2]+mc[2];
/* Skip this cell if empty */
if (head[c] == EMPTY) continue;

/* Scan the neighbor cells (including itself) of cell c */
for (mc1[0]=mc[0]-1; mc1[0]<=mc[0]+1; (mc1[0])++)
for (mc1[1]=mc[1]-1; mc1[1]<=mc[1]+1; (mc1[1])++)
for (mc1[2]=mc[2]-1; mc1[2]<=mc[2]+1; (mc1[2])++) {

/* Calculate the scalar cell index of the neighbor cell */
c1 = mc1[0]*lcyz2+mc1[1]*lc2[2]+mc1[2];
/* Skip this neighbor cell if empty */
if (head[c1] == EMPTY) continue;

/* Scan atom i in cell c */
i = head[c];
while (i != EMPTY) {

/* Scan atom j in cell c1 */
j = head[c1];
while (j != EMPTY) {

/* No calculation with itself */
if (j != i) {
/* Logical flag: intra(true)- or inter(false)-pair atom */
//bintra = (j < n); // #####

/* Pair vector dr = r[i] - r[j] */ // #####
for (rr=0.0, a=0; a<3; a++) {
dr[a] = r[i][a]-r[j][a];
rr += dr[a]*dr[a];
}

/* Calculate potential & forces for intranode pairs (i < j)
& all the internode pairs if rij < RCUT; note that for
any copied atom, i < j */
//if (i<j && rr<rrCut) { // #####
if (rr<rrCut) {
ri2 = 1.0/rr; ri6 = ri2*ri2*ri2; r1 = sqrt(rr);
fcVal = 48.0*ri2*ri6*(ri6-0.5) + Duc/r1;
vVal = 4.0*ri6*(ri6-1.0) - Uc - Duc*(r1-RCUT);
//if (bintra) lpe += vVal; else lpe += 0.5*vVal; // #####
lpe_td[std] += 0.5*vVal;
for (a=0; a<3; a++) {
f = fcVal*dr[a];
ra[i][a] += f;
//if (bintra) ra[j][a] -= f; // #####
}
}
} /* Endif not self */

j = lscl[j];
} /* Endwhile j not empty */

i = lscl[i];

```

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        } /* Endwhile i not empty */

    } /* Endfor neighbor cells, c1 */

} /* Endfor central cell, c */

} /*End for parallel cell*/ // #####

    //Thresd reduction // #####
    for (i=0; i<nthrd; i++) lpe+= lpe_td[i];
    /* Global potential energy */
    MPI_Allreduce(&lpe,&potEnergy,1,MPI_DOUBLE,MPI_SUM,MPI_COMM_WORLD);
}

/*-----*/
void eval_props() {
/*-----
Evaluates physical properties: kinetic, potential & total energies.
-----*/

    double vv,lke;
    int i,a;

    /* Total kinetic energy */
    for (lke=0.0, i=0; i<n; i++) {
        for (vv=0.0, a=0; a<3; a++) vv += rv[i][a]*rv[i][a];
        lke += vv;
    }
    lke *= 0.5;
    MPI_Allreduce(&lke,&kinEnergy,1,MPI_DOUBLE,MPI_SUM,MPI_COMM_WORLD);

    /* Energy paer atom */
    kinEnergy /= nglob;
    potEnergy /= nglob;
    totEnergy = kinEnergy + potEnergy;
    temperature = kinEnergy*2.0/3.0;

    /* Print the computed properties */
    if (sid == 0) printf("%9.6f %9.6f %9.6f %9.6f\n",
        stepCount*DeltaT,temperature,potEnergy,totEnergy);
}

/*-----*/
void atom_move() {
/*-----
Sends moved-out atoms to neighbor nodes and receives moved-in atoms
from neighbor nodes. Called with n, r[0:n-1] & rv[0:n-1], atom_move
returns a new n' together with r[0:n'-1] & rv[0:n'-1].
-----*/

/* Local variables-----

mvque[6][NBMAX]: mvque[ku][0] is the # of to-be-moved atoms to neighbor
ku; MVQUE[ku][k>0] is the atom ID, used in r, of the k-th atom to be
moved.
-----*/

    int mvque[6][NBMAX];
    int newim = 0; /* # of new immigrants */
    int ku,kd,i,kdd,kul,kuh,inode,ipt,a,nsd,nrc;
    double com1;

    /* Reset the # of to-be-moved atoms, MVQUE[][0] */
    for (ku=0; ku<6; ku++) mvque[ku][0] = 0;

```



```

/* Main loop over x, y & z directions starts-----*/
for (kd=0; kd<3; kd++) {

    /* Make a moved-atom list, mvque-----*/

    /* Scan all the residents & immigrants to list moved-out atoms */
    for (i=0; i<n+newim; i++) {
        kul = 2*kd ; /* Neighbor ID */
        kuh = 2*kd+1;
        /* Register a to-be-copied atom in mvque[kul|kuh][] */
        if (r[i][0] > MOVED_OUT) { /* Don't scan moved-out atoms */
            /* Move to the lower direction */
            if (bmv(r[i],kul)) mvque[kul][++(mvque[kul][0])] = i;
            /* Move to the higher direction */
            else if (bmv(r[i],kuh)) mvque[kuh][++(mvque[kuh][0])] = i;
        }
    }

    /* Message passing with neighbor nodes-----*/

    com1 = MPI_Wtime();

    /* Loop over the lower & higher directions-----*/
    for (kdd=0; kdd<2; kdd++) {

        inode = nn[ku=2*kd+kdd]; /* Neighbor node ID */

        /* Send atom-number information-----*/

        nsd = mvque[ku][0]; /* # of atoms to-be-sent */

        /* Even node: send & recv */
        if (myparity[kd] == 0) {
            MPI_Send(&nsd,1,MPI_INT,inode,110,MPI_COMM_WORLD);
            MPI_Recv(&nrc,1,MPI_INT,MPI_ANY_SOURCE,110,
                    MPI_COMM_WORLD,&status);
        }
        /* Odd node: recv & send */
        else if (myparity[kd] == 1) {
            MPI_Recv(&nrc,1,MPI_INT,MPI_ANY_SOURCE,110,
                    MPI_COMM_WORLD,&status);
            MPI_Send(&nsd,1,MPI_INT,inode,110,MPI_COMM_WORLD);
        }
        /* Single layer: Exchange information with myself */
        else
            nrc = nsd;
        /* Now nrc is the # of atoms to be received */

        /* Send & receive information on boundary atoms-----*/

        /* Message buffering */
        for (i=1; i<=nsd; i++)
            for (a=0; a<3; a++) {
                /* Shift the coordinate origin */
                dbuf[6*(i-1) +a] = r[mvque[ku][i]][a]-sv[ku][a];
                dbuf[6*(i-1)+3+a] = rv[mvque[ku][i]][a];
                r[mvque[ku][i]][0] = MOVED_OUT; /* Mark the moved-out atom */
            }

        /* Even node: send & recv, if not empty */
        if (myparity[kd] == 0) {

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        MPI_Send(dbuf,6*nsd,MPI_DOUBLE,inode,120,MPI_COMM_WORLD);
        MPI_Recv(dbufr,6*nrc,MPI_DOUBLE,MPI_ANY_SOURCE,120,
                MPI_COMM_WORLD,&status);
    }
    /* Odd node: recv & send, if not empty */
    else if (myparity[kd] == 1) {
        MPI_Recv(dbufr,6*nrc,MPI_DOUBLE,MPI_ANY_SOURCE,120,
                MPI_COMM_WORLD,&status);
        MPI_Send(dbuf,6*nsd,MPI_DOUBLE,inode,120,MPI_COMM_WORLD);
    }
    /* Single layer: Exchange information with myself */
    else
        for (i=0; i<6*nrc; i++) dbufr[i] = dbuf[i];

    /* Message storing */
    for (i=0; i<nrc; i++)
        for (a=0; a<3; a++) {
            r [n+newim+i][a] = dbufr[6*i +a];
            rv[n+newim+i][a] = dbufr[6*i+3+a];
        }

    /* Increment the # of new immigrants */
    newim = newim+nrc;

    /* Internode synchronization */
    MPI_Barrier(MPI_COMM_WORLD);

} /* Endfor lower & higher directions, kdd */

comt=comt+MPI_Wtime()-com1;

} /* Endfor x, y & z directions, kd */

/* Main loop over x, y & z directions ends-----*/

/* Compress resident arrays including new immigrants */

ipt = 0;
for (i=0; i<n+newim; i++) {
    if (r[i][0] > MOVED_OUT) {
        for (a=0; a<3; a++) {
            r [ipt][a] = r [i][a];
            rv[ipt][a] = rv[i][a];
        }
        ++ipt;
    }
}

/* Update the compressed # of resident atoms */
n = ipt;
}

/*-----
Bit condition functions:

1. bbd(ri,ku) is .true. if coordinate ri[3] is in the boundary to
   neighbor ku.
2. bmv(ri,ku) is .true. if an atom with coordinate ri[3] has moved out
   to neighbor ku.
-----*/
int bbd(double* ri, int ku) {
    int kd,kdd;
    kd = ku/2; /* x(0)|y(1)|z(2) direction */

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kdd = ku%2; /* Lower(0)|higher(1) direction */
if (kdd == 0)
    return ri[kd] < RCUT;
else
    return al[kd]-RCUT < ri[kd];
}
int bmv(double* ri, int ku) {
    int kd,kdd;
    kd = ku/2; /* x(0)|y(1)|z(2) direction */
    kdd = ku%2; /* Lower(0)|higher(1) direction */
    if (kdd == 0)
        return ri[kd] < 0.0;
    else
        return al[kd] < ri[kd];
}

```

2.

hmd.out :

```

-----
Begin SLURM Prolog Wed 10 Oct 2018 09:12:09 PM PDT
Job ID:          2019069
Username:        liangsiq
Accountname:     lc_an2
Name:            hmd.sl
Partition:       quick
Nodes:           hpc[1118-1119]
TasksPerNode:    1(x2)
CPUSPerTask:     4
TMPDIR:          /tmp/2019069.quick
SCRATCHDIR:      /staging/scratch/2019069
Cluster:         uschpc
HSDA Account:    false
End SLURM Prolog
-----
al = 4.103942e+01 4.103942e+01 2.051971e+01
lc = 16 16 8
rc = 2.564964e+00 2.564964e+00 2.564964e+00
nglob = 55296
0.050000 0.877345 -5.137153 -3.821136
0.100000 0.462056 -4.513097 -3.820013
0.150000 0.510836 -4.587287 -3.821033
0.200000 0.527457 -4.611958 -3.820772
0.250000 0.518668 -4.598798 -3.820796
0.300000 0.529023 -4.614343 -3.820808
0.350000 0.532890 -4.620133 -3.820798
0.400000 0.536070 -4.624899 -3.820794
0.450000 0.539725 -4.630387 -3.820799
0.500000 0.538481 -4.628514 -3.820792
CPU & COMT = 7.283849e+00 5.137362e-01

```

3.

Plot:

