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#include <iostream>
#include <iomanip>
#include <math.h>
#include <stdlib.h>
#include <fstream>
#include <ctime>
#include <complex>
#include <cstdio>
#include <string>
#include "../MASTER-FILES/AtomParams.h"
using namespace std;

////////////////////////////////////
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void xAxisRotation(AtomParams **Atom, double rotationAngle, int nAtoms)
{
    struct vect tempPosition, position;

    for (int i=0; i<nAtoms; i++)
    {
        position = Atom[i]->getPosition();

        tempPosition.x = position.x;
        tempPosition.y = position.y*cos(rotationAngle) - position.z*sin(rotationAngle);
        tempPosition.z = position.y*sin(rotationAngle) + position.z*cos(rotationAngle);

        Atom[i]->setPosition( tempPosition );
    }
}

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void yAxisRotation(AtomParams **Atom, double rotationAngle, int nAtoms)
{
    struct vect tempPosition, position;

    for (int i=0; i<nAtoms; i++)
    {
        position = Atom[i]->getPosition();

        tempPosition.x = position.x*cos(rotationAngle) - position.z*sin(rotationAngle);
        tempPosition.y = position.y;
        tempPosition.z = position.x*sin(rotationAngle) + position.z*cos(rotationAngle);

        Atom[i]->setPosition( tempPosition );
    }
}

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void zAxisRotation (AtomParams **Atom, double rotationAngle, int nAtoms)
{
    struct vect tempPosition, position;

    for (int i=0; i<nAtoms; i++)
    {
        position = Atom[i]->getPosition();

        tempPosition.x = position.x*cos(rotationAngle) - position.y*sin(rotationAngle);
        tempPosition.y = position.x*sin(rotationAngle) + position.y*cos(rotationAngle);
        tempPosition.z = position.z;

        Atom[i]->setPosition( tempPosition );
    }
}

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void Translate (AtomParams **Atom, struct vect translationVector, int nAtoms)
{
    struct vect tempPosition, position;

    for (int i=0; i<nAtoms; i++)
    {
        position = Atom[i]->getPosition();
        tempPosition = position + translationVector;
        Atom[i]->setPosition( tempPosition );
    }
}

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int main(int argc, char *argv[])
{
    ////////////
    // USER INPUT //
    ////////////

    if (argc != 3) {
        cout << "Must have 2 command line arguments (nCarbon, nQuarter)!" << endl;
    }

    int nCarbon = atoi(argv[1]);

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int nQuarter = atoi(argv[2]);

float n1, n2;
double latticeSpacing, domainAngle;
if (nCarbon == 6) {

    n1 = 11.0 + float(nQuarter)*0.25;
    n2 = 3.0 + float(nQuarter)*0.25;

    if (nQuarter == 0) {
        latticeSpacing = 31.401;
        domainAngle = 11.742;
    } else if (nQuarter == 1) {
        latticeSpacing = 32.414;
        domainAngle = 12.331;
    } else if (nQuarter == 2) {
        latticeSpacing = 33.430;
        domainAngle = 12.885;
    } else if (nQuarter == 3) {
        latticeSpacing = 34.450;
        domainAngle = 13.407;
    } else if (nQuarter == 4) {
        latticeSpacing = 35.472;
        domainAngle = 13.898;
    } else {}

} else if (nCarbon == 10) {

    n1 = 15.0 + float(nQuarter)*0.25;
    n2 = 1.0 + float(nQuarter)*0.25;

    if (nQuarter == 0) {
        latticeSpacing = 38.182;
        domainAngle = 3.198;
    } else if (nQuarter == 1) {
        latticeSpacing = 39.135;
        domainAngle = 3.901;
    } else if (nQuarter == 2) {
        latticeSpacing = 40.095;
        domainAngle = 4.571;
    } else if (nQuarter == 3) {
        latticeSpacing = 41.059;
        domainAngle = 5.209;
    } else if (nQuarter == 4) {
        latticeSpacing = 42.028;
        domainAngle = 5.818;
    } else {}

} else {}

int nAtomsPerTSB = 54+6*(3*nCarbon+1);
int nAtomsTotal = nAtomsPerTSB;
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// Create Instance of the class AtomParams
AtomParams *Atom[nAtomsTotal];

//////////
// Read TSB input files (from Gaussian09) //
//////////
stringstream inStream1;
inStream1 << "../cpp-input/tsb35-c" << nCarbon << "_cpp-input.dat";
ifstream in1(inStream1.str().c_str());

cout << endl;
cout << "TSB input file to be read: " << inStream1.str() << endl;

string tempAtomName, tempAtomType, tempElementType;
struct vect tempPosition;
double tempCharge;

for (int i=0; i<nAtomsPerTSB; i++)
{
    in1 >> tempAtomName >> tempAtomType >> tempPosition.x >> tempPosition.y >> tempPosition.z >>
        tempElementType >> tempCharge;
    Atom[i] = new AtomParams(tempAtomName, tempAtomType, tempPosition, tempElementType,
        tempCharge);
}

cout << "TSB input file read and stored in class AtomParams/Atom." << endl;
cout << endl;

// Create TSB output PDB files
stringstream outStream1;
outStream1 << "tsb35-c" << nCarbon << "_0" << nQuarter << "-quarter_TSBA-PDB.pdb";
ofstream OUT1(outStream1.str().c_str());
OUT1 << setiosflags(ios::fixed) << setprecision(3);

stringstream outStream2;
outStream2 << "tsb35-c" << nCarbon << "_0" << nQuarter << "-quarter_TSBB-PDB.pdb";
ofstream OUT2(outStream2.str().c_str());
OUT2 << setiosflags(ios::fixed) << setprecision(3);

cout << "TSB output files created: " << outStream1.str() << endl;
cout << "and " << outStream2.str() << endl;
cout << endl;

//////////
// Build lattice vectors //
//////////
struct vect a1, a2, a3, b1, b2, b3;
double a = sqrt(3.0)*1.42;
double triAngle = 60.0*(PI/180.0);

// Define graphene lattice vectors
a1.x = a;
a1.y = 0.0;

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a1.z = 0.0;
a2.x = a*cos(triAngle);
a2.y = a*sin(triAngle);
a2.z = 0.0;
a3 = (1.0/3.0)*(a1+a2);

// Define TSB lattice vectors
b1 = n1*a1 + n2*a2;
b2 = (-1.0)*n2*a1 + (n1+n2)*a2;
b3 = (-1.0/3.0)*(b1+b2);

double Bmag = 4.0*sqrt(b1.x*b1.x + b1.y*b1.y + b1.z*b1.z);

// Calculate center of mass for the TSB molecule
double mass;
double totalMass=0.0;

struct vect position;
struct vect CoM;
CoM.x=0.0;
CoM.y=0.0;
CoM.z=0.0;

for (int i=0; i<nAtomsPerTSB; i++)
{
    mass = Atom[i]->getMass();

    position = Atom[i]->getPosition();
    Atom[i]->setPosition(position);

    CoM.x += mass * position.x;
    CoM.y += mass * position.y;
    CoM.z += mass * position.z;

    totalMass += mass;
}

CoM = (1.0/totalMass)*CoM;

Translate(Atom, (-1.0)*CoM, nAtomsPerTSB);
double rotationAngle = 60.0*(PI/180.0);
zAxisRotation(Atom, rotationAngle, nAtomsPerTSB);

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struct vect hexagonCenter;
hexagonCenter.x = 0.5*a;
hexagonCenter.y = 0.5*a/sqrt(3.0);
hexagonCenter.z = 3.366;

//go to a reasonable distance above graphene sheet, centered on the first graphene carbon

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Translate (Atom, hexagonCenter, nAtomsPerTSB);

//define number of TSB molecule "rows" in directions of b1 and b2
int nMolecules1 = 4;
int nMolecules2 = 4;

struct vect comTSBA, comTSBB;

int moleculeCount = 0;
// Write TSBA molecules
OUT1 << "CRYST1 " << Bmag << " " << Bmag << " 100.000 " << domainAngle << " " <<
domainAngle << " " << domainAngle << " P 1 1" << endl;
for (int i=0; i<nMolecules1; i++)
{
    for (int j=0; j<nMolecules2; j++)
    {
        for (int k=0; k<nAtomsPerTSB; k++)
        {
            position = Atom[k]->getPosition();
            position = position + i*b1 + j*b2;
            Atom[k]->setPosition(position);

            OUT1 << "ATOM " << setw(6) << k + moleculeCount*nAtomsPerTSB + 1;
            OUT1 << " " << setw(4) << Atom[k]->getAtomName() << " TSB " << setw(3) << moleculeCount
            +1 << " ";
            OUT1 << setw(7) << position.x << " " << setw(7) << position.y << " " << setw(7) <<
            position.z << " ";
            OUT1 << "0.00 0.00 " << setw(4) << "TSBA" << endl;

            if (i==0 && j==0)
            {
                mass = Atom[k]->getMass();
                position = Atom[k]->getPosition();
                Atom[k]->setPosition(position);

                comTSBA.x += mass * position.x;
                comTSBA.y += mass * position.y;
                comTSBA.z += mass * position.z;

                totalMass += mass;
            }

            position = Atom[k]->getPosition();
            position = position - i*b1 - j*b2;
            Atom[k]->setPosition(position);
        }
        moleculeCount++;
    }
}

comTSBA = (1.0/totalMass)*comTSBA;

// Place molecule on origin by subtracting: (r_i - r_com)

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Translate(Atom, (-1.0)*CoM, nAtomsPerTSB);
// Rotate by 60 degrees, interdigitating the rest of the TSB molecules
zAxisRotation(Atom, rotationAngle, nAtomsPerTSB);
// Replace molecule at (r_i + r_com)
Translate(Atom, CoM, nAtomsPerTSB);

moleculeCount = 0;
// Write rotated molecules, using the TSB basis vector b3
OUT2 << "CRYST1 " << Bmag << " " << Bmag << " 100.000 " << domainAngle << " " <<
domainAngle << " " << domainAngle << " P 1 1" << endl;
for (int i=0; i<nMolecules1; i++)
{
    for (int j=0; j<nMolecules2; j++)
    {
        for (int k=0; k<nAtomsPerTSB; k++)
        {
            position = Atom[k]->getPosition();
            position = position + i*b1 + j*b2 + b3;
            Atom[k]->setPosition(position);

            OUT2 << "ATOM " << setw(6) << k + moleculeCount*nAtomsPerTSB + 1;
            OUT2 << " " << setw(4) << Atom[k]->getAtomName() << " TSB " << setw(3) << moleculeCount
            +1 << " ";
            OUT2 << setw(7) << position.x << " " << setw(7) << position.y << " " << setw(7) <<
            position.z << " ";
            OUT2 << "0.00 0.00 " << setw(4) << "TSBB" << endl;

            if (i==0 && j==0)
            {
                mass = Atom[k]->getMass();
                position = Atom[k]->getPosition();
                Atom[k]->setPosition(position);

                comTSBB.x += mass * position.x;
                comTSBB.y += mass * position.y;
                comTSBB.z += mass * position.z;

                totalMass += mass;
            }

            position = Atom[k]->getPosition();
            position = position - i*b1 - j*b2 - b3;
            Atom[k]->setPosition(position);
        }
        moleculeCount++;
    }
}

comTSBB = (1.0/totalMass)*comTSBB;

return 0;
}

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