

Lemina

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Chapter 1

Lamina: A Molecular Dynamics Package

Welcome to the **Lamina** documentation!

1.1 Overview

Lamina is a modular 2D molecular dynamics (MD) simulation package designed for simulating hybrid soft solids, including spring networks and finite-size discs. Written in C, it models a wide variety of soft and condensed matter systems. It supports robust time evolution integrators and a range of thermostats, providing accurate force evaluations for bonded and non-bonded interactions.

Originally developed for 2D bonded systems, **Lamina** now supports broader research goals including active matter, granular solids, and complex fluids.

1.2 Why "Lamina"?

The word **Lamina** comes from Latin, meaning "a thin layer", "a plate", or "a sheet". In nature and science, laminae often refer to flat, two-dimensional structural elements such as leaves, thin metal sheets, or tissue membranes. This name reflects both the **two-dimensional (2D)** nature of the simulations and the types of materials **Lamina** is built to study: **liquids**, **soft solids**, and **networked structures** confined to thin sheets or layers. Just as natural laminae exhibit rich structural and dynamic behaviors in simple geometry, this code explores the complexity of emergent phenomena in 2D soft matter systems.

1.3 Key Features

1.3.1 Interaction Potentials

- · Yukawa potential (screened Coulomb interactions)
- Lennard-Jones potential (standard 12-6)
- · Harmonic bond potential (elastic network models)
- · Hookean granular contact potential (for soft granular matter)

1.3.2 Thermostats and Temperature Control

- · Gaussian thermostat
- · Nose-Hoover thermostat
- · Langevin thermostat
- · Configurational temperature evaluation and control

1.3.3 Time Integration

- · Leap-Frog integrator
- · Velocity-Verlet integrator
- · Langevin (stochastic) integrator
- · Brownian (overdamped) integrator

1.3.4 Physical Observables

- · Radial Distribution Function (RDF)
- Velocity Autocorrelation Function (VACF)
- Root-Mean-Square Velocity (VRMS)
- · Stress tensor and momentum
- · Center-of-mass motion
- · Space-time correlation functions

1.3.5 Output and Utilities

- Output files saved to ../output folder ensure this directory exists relative to where you run the code
- · Run the simulation with:

```
./main prefix
```

- Structured output files: .xyz, .bond, .pair, .com, .result
- Restart and resume capability: .restart and .state files
- · Clear separation of source code, unit tests, and output
- Support for Lees–Edwards boundary conditions (sheared systems)
- · Configurable halting conditions based on VRMS or custom metrics
- · Modular design for easy extension of potentials and features

1.4 Project Structure

```
Lamina/
  |-- source/
         ource/  # C source files; avoid

|-- main.c  # Main driver

|-- *.c, *.h  # Modular source files

nittest/  # Unit test suite (plans

|-- test_*.c  # Individual test cases

utput/  # Runtime output files

repros/  # Preprocessing scripts.
                                        # C source files; avoid placing README.md here to prevent extra pages
        |-- main.c
|-- *.c, *.h
  |-- unittest/
                                        # Unit test suite (planned or implemented)
  |-- output/
                                        # Preprocessing scripts/tools (.sh, .py, etc.)
   |-- prepros/
                               # Postprocessing scripes, ...
# Doxygen configuration and au
# Doxygen config file
# Custom LaTeX header for docs
" Optional CSS for HTML styl.
                                        # Postprocessing scripts/tools (.sh, .py, etc.)
   |-- postpros/
                                        # Doxygen configuration and auxiliary files
   -- doxygen/
         |-- Doxyfile
|-- header.tex
 \-- extra_stylesheet.css # Optional CSS for HTML styling
gures/ # Figures, logos, icons used in docs/code
          \-- workflows/  # GitHub Actions workflows for CI/CD  # CI workflow file
```

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1.5 Installation Instructions

1.5.1 Prerequisites

1. GCC Compiler

Install gcc to compile C code:

· Ubuntu/Debian:

```
sudo apt-get install build-essential
```

Fedora/CentOS:

```
sudo dnf install gcc
```

• macOS (via Homebrew):

bash brew install gcc

2. MPICH (MPI Library)

Required for parallel computations:

· Ubuntu/Debian:

```
sudo apt-get install libmpich-dev
```

Fedora/CentOS:

```
sudo dnf install mpich
```

· macOS:

bash brew install mpich

3. CMake (Recommended for modern builds)

Install cmake to build with the CMake system:

· Ubuntu/Debian:

```
sudo apt-get install cmake
```

Fedora/CentOS:

```
sudo dnf install cmake
```

· macOS:

bash brew install cmake

1.6 Building Lamina

You can build Lamina either using the traditional Makefile or the CMake build system.

1.6.1 Using Makefile

cd Lamina/source make clean

To build and run unit tests:

cd ../unittest
make clean
make all
make run

1.6.2 Using CMake (Recommended)

This builds the project in a clean isolated directory and manages dependencies automatically.

cd Lamina
mkdir -p build
cd build
cmake ..
make -j\$(nproc)

To run unit tests (assuming they are built in unittest and mpirun is used):

cd ../unittest
make clean
make all
make run

1.7 Continuous Integration (CI) with GitHub Actions

The project includes a GitHub Actions workflow (.github/workflows/ci.yml) that automates building and testing on Ubuntu runners with MPI installed.

The workflow performs the following:

- Checks out the latest code on push or pull requests to the main branch.
- Installs build dependencies including build-essential, cmake, mpich, and MPI development libraries.
- Configures and builds Lamina using CMake in the build directory.
- · Runs the unit tests in parallel using MPI.

You can view the build status and logs on the Actions tab of the GitHub repository.

1.8 Documentation

- Browse full HTML documentation
- Download Source code PDF manual
- User manual Physics PDF manual
- Documentation generated with Doxygen 1.10.0

Thank you for your interest in Lamina! Contributions and feedback are welcome.

Please check the repository for the latest updates and contact information.

1.9 Adding a Build Status Badge

Chapter 2

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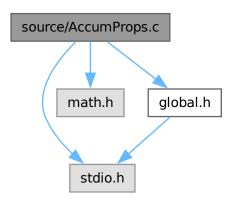
Chapter 3

File Documentation

3.1 README.md File Reference

3.2 source/AccumProps.c File Reference

```
#include <stdio.h>
#include <math.h>
#include "global.h"
Include dependency graph for AccumProps.c:
```



Functions

• void AccumProps (int icode)

3.2.1 Function Documentation

3.2.1.1 AccumProps()

```
00030 sTotEnergy = ssTotEnergy = 0.;
00031 svirSum = 0.;
00032 }else if(icode == 1){
00033 sPotEnergy += potEnergy;
00034    ssPotEnergy += Sqr(potEnergy);
00035    sKinEnergy += kinEnergy;
00036    ssKinEnergy += Sqr(kinEnergy);
00037 sTotEnergy += totEnergy;
00038 ssTotEnergy += Sqr(totEnergy);
00039 sPressure += pressure;
00040 ssPressure += Sqr(pressure);
00041 svirSum += virSum;
00042 }else if(icode == 2){
00043 sPotEnergy /= stepAvg;
00044 ssPotEnergy /= sqrt(ssPotEnergy/stepAvg - Sqr(sPotEnergy));
00045 sTotEnergy /= stepAvg;
00046 ssTotEnergy = sqrt(ssTotEnergy/stepAvg - Sqr(sTotEnergy));
00047 sKinEnergy /= stepAvg;
00048 ssKinEnergy = sqrt(ssKinEnergy/stepAvg - Sqr(sKinEnergy));
00049 sPressure /= stepAvg;
00050 ssPressure = sqrt(ssPressure/stepAvg - Sqr(sPressure));
00051
        svirSum /= stepAvg;
00052 } }
```

References kinEnergy, potEnergy, pressure, sKinEnergy, sPotEnergy, sPressure, Sqr, ssKinEnergy, ssPotEnergy, ssPressure, ssTotEnergy, stepAvg, sTotEnergy, svirSum, totEnergy, and virSum.

Referenced by SetupJob().

Here is the caller graph for this function:



3.3 AccumProps.c

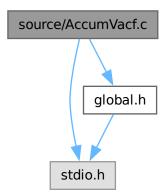
Go to the documentation of this file.

```
00001 /*
00002
       * This file is part of Lamina.
00003
00004
        * Lamina is free software: you can redistribute it and/or modify
00005
          it under the terms of the GNU General Public License as published by
00006
        * the Free Software Foundation, either version 3 of the License, or
00007
        \star (at your option) any later version.
80000
00009 * Lamina is distributed in the hope that it will be useful,
          but WITHOUT ANY WARRANTY; without even the implied warranty of
00011
        * MERCHANTABILITY or FITNESS FOR A PARTICULAR PURPOSE. See the
00012
        \star GNU General Public License for more details.
00013
* You should have received a copy of the GNU General Public License 00015 * along with Lamina. If not, see <a href="https://www.gnu.org/licenses/">https://www.gnu.org/licenses/</a>.
00017 Copyright (C) 2025 Harish Charan, University of Durham, UK
00018
00019
00020
00021 #include<stdio.h>
00022 #include<math.h>
00023 #include"global.h"
00024
00025 void AccumProps(int icode){
00026 if(icode == 0) {
00027 sPotEnergy = ssPotEnergy = 0.;
00028 sKinEnergy = ssKinEnergy = 0.;
00029 sPressure = ssPressure = 0.;
00030 sTotEnergy = ssTotEnergy = 0.;
00031 svirSum = 0.;
00032 }else if(icode == 1){
00033 sPotEnergy += potEnergy;
00034 ssPotEnergy += Sqr(potEnergy);
00035 sKinEnergy += kinEnergy;
```

3.4 source/AccumVacf.c File Reference

```
#include <stdio.h>
#include "global.h"
```

Include dependency graph for AccumVacf.c:



Functions

- double Integrate (double *, int)
- void PrintVacf ()
- void ZeroVacf ()
- void AccumVacf ()

3.4.1 Function Documentation

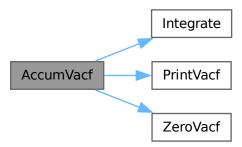
3.4.1.1 AccumVacf()

```
00037    if(countAcfAv == limitAcfAv) {
00038       fac = 1./(kinEnergy*region[1]*region[2]*limitAcfAv);
00039       viscAcfInt = fac*stepAcf*deltaT*Integrate(viscAcfAv, nValAcf);
00040       PrintVacf();
00041       ZeroVacf();
00042    } } }
```

References countAcfAv, deltaT, indexAcf, Integrate(), kinEnergy, limitAcfAv, nBuffAcf, nValAcf, PrintVacf(), region, stepAcf, viscAcf, viscAcfAv, viscAcfInt, and ZeroVacf().

Referenced by EvalVacf().

Here is the call graph for this function:



Here is the caller graph for this function:



3.4.1.2 Integrate()

Referenced by AccumVacf().

Here is the caller graph for this function:



3.4.1.3 PrintVacf()

References deltaT, fpvisc, nValAcf, stepAcf, viscAcfAv, and viscAcfInt.

Referenced by AccumVacf().

Here is the caller graph for this function:

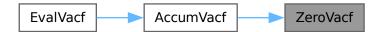


3.4.1.4 ZeroVacf()

References countAcfAv, nValAcf, and viscAcfAv.

Referenced by AccumVacf().

Here is the caller graph for this function:



3.5 AccumVacf.c

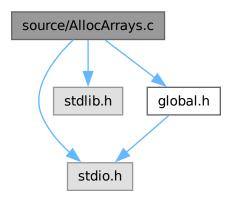
Go to the documentation of this file.

```
00002
        * This file is part of Lamina.
00003
        \star Lamina is free software: you can redistribute it and/or modify \star it under the terms of the GNU General Public License as published by
00004
00005
         * the Free Software Foundation, either version 3 of the License, or
        * (at your option) any later version.
80000
00009 \star Lamina is distributed in the hope that it will be useful,
00010 * but WITHOUT ANY WARRANTY; without even the implied warranty of 00011 * MERCHANTABILITY or FITNESS FOR A PARTICULAR PURPOSE. See the
       * GNU General Public License for more details.
00012
00013
00014 * You should have received a copy of the GNU General Public License 00015 * along with Lamina. If not, see <a href="https://www.gnu.org/licenses/">https://www.gnu.org/licenses/</a>.
00016
00017 Copyright (C) 2025 Harish Charan, University of Durham, UK
00018
00019
00020
00021 #include<stdio.h>
00022 #include"global.h"
00023
00024 double Integrate(double *, int);
00025 void PrintVacf();
00026 void ZeroVacf();
00027
         void AccumVacf() {
00028 double fac;
        int j, nb;
for(nb = 1 ; nb <= nBuffAcf ; nb ++) {
  if(indexAcf[nb] == nValAcf) {</pre>
00029
00030
00031
          for(j = 1; j <= nValAcf; j ++) {
  viscAcfAv[j] += viscAcf[nb][j];</pre>
00032
00033
00034
00035
          indexAcf[nb] = 0;
00036
          countAcfAv ++:
          if (countAcfAv == limitAcfAv) {
00037
          fac = 1./(kinEnergy*region[1]*region[2]*limitAcfAv);
00039
           viscAcfInt = fac*stepAcf*deltaT*Integrate(viscAcfAv, nValAcf);
00040
           PrintVacf();
00041
           ZeroVacf();
00042 } } }
00043
```

3.6 source/AllocArrays.c File Reference

```
#include <stdio.h>
#include <stdlib.h>
#include "global.h"
```

Include dependency graph for AllocArrays.c:



Functions

· void AllocArrays ()

3.6.1 Function Documentation

3.6.1.1 AllocArrays()

```
void AllocArrays ( )
Definition at line 25 of file AllocArrays.c.
00025
00026
        // SPACETIME CORRELATIONS
00028 cfOrg = (double **) malloc ((nBuffCorr+1)*sizeof(double *));
00029 for (n = 0; n <= nBuffCorr; n++)</pre>
00030
        cfOrg[n] = (double *) malloc ((2*nFunCorr+1)*sizeof(double));
00031
00032 cfVal = (double *) malloc ((2*nFunCorr+1)*sizeof(double));
00033 indexCorr = (int *) malloc ((nBuffCorr+1)*sizeof(int));
00034
00035 spacetimeCorr = (double **) malloc ((nBuffCorr+1)*sizeof(double));
00036 for (n = 0; n <= nBuffCorr; n++)
00037 spacetimeCorr[n] = (double *) malloc ((nFunCorr*nValCorr+1)*sizeof(double));
00038
00039 spacetimeCorrAv = (double *) malloc ((nFunCorr*nValCorr+1)*sizeof(double));
00040 // VISCOSITY
00041 indexAcf = (double *)malloc((nBuffAcf+1)*sizeof(double));
00042 viscAcf = (double **)malloc((nBuffAcf+1)*sizeof(double *));
00043 for(n = 0; n <= nBuffAcf; n ++)
00044 viscAcf[n] = (double *)malloc((nValAcf+1)*sizeof(double ));</pre>
00045
00048
00049
        histRdf = (double *)malloc((sizeHistRdf+1)*sizeof(double));
00050
00051 }
```

References cfOrg, cfVal, histRdf, indexAcf, indexCorr, nBuffAcf, nBuffCorr, nFunCorr, nValAcf, nValCorr, sizeHistRdf, spacetimeCorr, spacetimeCorrAv, viscAcf, viscAcfAv, and viscAcfOrg. Referenced by SetupJob().

Here is the caller graph for this function:



3.7 AllocArrays.c

Go to the documentation of this file.

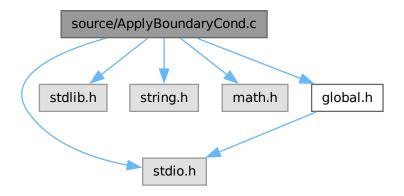
```
00002
       * This file is part of Lamina.
00003
        \star Lamina is free software: you can redistribute it and/or modify \star it under the terms of the GNU General Public License as published by
00004
00005
        * the Free Software Foundation, either version 3 of the License, or
        * (at your option) any later version.
00008
00009 \star Lamina is distributed in the hope that it will be useful,
00010 * but WITHOUT ANY WARRANTY; without even the implied warranty of
       * MERCHANTABILITY or FITNESS FOR A PARTICULAR PURPOSE. See the
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00012 * GNU General Public License for more details.
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00014 * You should have received a copy of the GNU General Public License 00015 * along with Lamina. If not, see <a href="https://www.gnu.org/licenses/">https://www.gnu.org/licenses/</a>.
00016
00017 Copyright (C) 2025 Harish Charan, University of Durham, UK
00018
00019
00020
00021 #include<stdio.h>
00022 #include<stdlib.h>
00023 #include"global.h"
00024
00025 void AllocArrays(){
00026 int n;
00027
        // SPACETIME CORRELATIONS
00028 cfOrg = (double **) malloc ((nBuffCorr+1)*sizeof(double *)); 00029 for (n = 0; n <= nBuffCorr; n++)
        cfOrg[n] = (double *) malloc ((2*nFunCorr+1)*sizeof(double));
00030
00032 cfVal = (double *) malloc ((2*nFunCorr+1)*sizeof(double));
00033 indexCorr = (int *) malloc ((nBuffCorr+1)*sizeof(int));
00034
00035
        spacetimeCorr = (double **) malloc ((nBuffCorr+1)*sizeof(double));
        for (n = 0; n <= nBuffCorr; n++)
00036
00037 spacetimeCorr[n] = (double *) malloc ((nFunCorr*nValCorr+1)*sizeof(double));
00039
        spacetimeCorrAv = (double *) malloc ((nFunCorr*nValCorr+1)*sizeof(double));
00040 // VISCOSITY
00041 indexAcf = (double *)malloc((nBuffAcf+1)*sizeof(double));
00042 viscAcf = (double **)malloc((nBuffAcf+1)*sizeof(double *));
        for (n = 0 ; n \le nBuffAcf ; n ++)
00043
00044
         viscAcf[n] = (double *)malloc((nValAcf+1)*sizeof(double ));
00045
00046 viscAcfOrg = (double *)malloc((nBuffAcf+1)*sizeof(double));
00047 viscAcfAv = (double *)malloc((nValAcf+1)*sizeof(double));
00048
00049
         // RDF
00050
         histRdf = (double *)malloc((sizeHistRdf+1)*sizeof(double));
00051 }
```

3.8 source/ApplyBoundaryCond.c File Reference

```
#include <stdio.h>
#include <stdlib.h>
#include <string.h>
#include <math.h>
```

```
#include "global.h"
```

Include dependency graph for ApplyBoundaryCond.c:



Functions

void ApplyBoundaryCond ()

3.8.1 Function Documentation

3.8.1.1 ApplyBoundaryCond()

```
void ApplyBoundaryCond ( )
Definition at line 27 of file ApplyBoundaryCond.c.
00028
        for(n = 1 ; n <= nAtom ; n ++) {
  if(strcmp(xBoundary, "p") == 0 && strcmp(yBoundary, "p") == 0) {</pre>
00029
                                                                                       // P.B.C along x and y axis
00030
          rx[n] -= region[1]*rint(rx[n]/region[1]);
00031
          ry[n] -= region[2]*rint(ry[n]/region[2]);
00032
00033
          } else if (strcmp(xBoundary, "r") == 0 && strcmp(yBoundary, "r") == 0){ //R.B.C. along x and y
00034
            if((rx[n] + atomRadius[n]) >= regionH[1]){
            rx[n] = 0.999999*regionH[1] - atomRadius[n]; vx[n] = -vx[n];
f((rx[n]-atomRadius[n]) < -regionH[1]) 
00035
00036
00037
              rx[n] = -0.999999 * regionH[1] + atomRadius[n]; vx[n] = -vx[n];
00039
            if((ry[n] + atomRadius[n])>= regionH[2]){
00040
               ry[n] = 0.999999*regionH[2] - atomRadius[n]; vy[n] = -vy[n];
            }if((ry[n]-atomRadius[n]) < -regionH[2]) {
    ry[n] = -0.999999*regionH[2] + atomRadius[n]; vy[n] = -vy[n];</pre>
00041
00042
          } }
00043
           else if (strcmp(xBoundary, "p") == 0 && strcmp(yBoundary, "r") == 0) { //P.B.C. along x and R.B.C
00044
      along y axis
00045
           rx[n] -= region[1]*rint(rx[n]/region[1]);
           if((ry[n] + atomRadius[n]) >= regionH[2]){
   ry[n] = 0.999999*regionH[2] - atomRadius[n]; vy[n] = -vy[n];
00046
00047
00048
            }if((ry[n] - atomRadius[n]) < -regionH[2]){</pre>
               ry[n] = -0.999999*regionH[2] + atomRadius[n]; vy[n] = -vy[n];
00049
00050
00051
           else if(strcmp(xBoundary, "r") == 0 && strcmp(yBoundary, "p") == 0){      //R.B.C. along x and P.B.C
      along y axis
00052
          if((rx[n] + atomRadius[n]) >= regionH[1]){
              rx[n] = 0.999999 * regionH[1] - atomRadius[n]; vx[n] = -vx[n];
00053
            }if((rx[n] - atomRadius[n]) < -regionH[1]){</pre>
00054
00055
             rx[n] = -0.999999*regionH[1] + atomRadius[n]; vx[n] = -vx[n];
00056
00057
            ry[n] -= region[2]*rint(ry[n]/region[2]);
         } else {
00058
00059
           \ensuremath{//} Print error message and exit the program
           fprintf(fpresult, "Error: Invalid boundary configuration: '%s %s'\n", xBoundary, yBoundary);
00060
00061
           exit(EXIT_FAILURE); // Exit with failure status
00062
00063
```

```
ากกรุง เ
```

References atomRadius, fpresult, nAtom, region, regionH, rx, ry, vx, vy, xBoundary, and yBoundary. Referenced by main().

Here is the caller graph for this function:



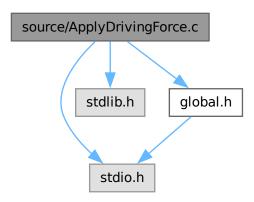
3.9 ApplyBoundaryCond.c

Go to the documentation of this file.

```
00001 /*
00002
      * This file is part of Lamina.
00003
00004
      * Lamina is free software: you can redistribute it and/or modify
00005
       * it under the terms of the GNU General Public License as published by
       * the Free Software Foundation, either version 3 of the License, or
00006
00007
       * (at your option) any later version.
00008
      * Lamina is distributed in the hope that it will be useful,
00009
00010 * but WITHOUT ANY WARRANTY; without even the implied warranty of
00011 * MERCHANTABILITY or FITNESS FOR A PARTICULAR PURPOSE. See the
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00018
00019
00020
00021 #include<stdio.h>
00022 #include<stdlib.h>
00023 #include<string.h>
00024 #include<math.h>
00025 #include"global.h"
00026
00027 void ApplyBoundaryCond() {
00028
       int n;
        if (strcmp(xBoundary, "p") == 0 && strcmp(yBoundary, "p") == 0) {
00029
                                                                                    // P.B.C along x and y axis
          rx[n] -= region[1]*rint(rx[n]/region[1]);
00031
         ry[n] -= region[2]*rint(ry[n]/region[2]);
} else if (strcmp(xBoundary, "r") == 0 && strcmp(yBoundary, "r") == 0){ //R.B.C. along x and y
00032
00033
     axis
           if((rx[n] + atomRadius[n]) >= regionH[1]){
00034
00035
               rx[n] = 0.999999*regionH[1] - atomRadius[n]; vx[n] = -vx[n];
           }if((rx[n]-atomRadius[n]) < -regionH[1]){</pre>
00036
00037
               rx[n] = -0.999999*regionH[1] + atomRadius[n]; vx[n] = -vx[n];
00038
00039
           if((ry[n] + atomRadius[n])>= regionH[2]){
           ry[n] = 0.999999*regionH[2] - atomRadius[n]; vy[n] = -vy[n]; }if((ry[n]-atomRadius[n]) < -regionH[2]){
00040
00041
              ry[n] = -0.999999*regionH[2] + atomRadius[n]; vy[n] = -vy[n];
00042
00043
          else if (strcmp(xBoundary, "p") == 0 && strcmp(yBoundary, "r") == 0){      //P.B.C. along x and R.B.C
00044
      along y axis
00045
          rx[n] -= region[1]*rint(rx[n]/region[1]);
00046
          if((ry[n] + atomRadius[n]) >= regionH[2]){
            ry[n] = 0.999999 * regionH[2] - atomRadius[n]; vy[n] = -vy[n];
00047
00048
           }if((ry[n] - atomRadius[n]) < -regionH[2]){</pre>
00049
              ry[n] = -0.999999*regionH[2] + atomRadius[n]; vy[n] = -vy[n];
00050
          else if (strcmp(xBoundary, "r") == 0 && strcmp(yBoundary, "p") == 0) { //R.B.C. along x and P.B.C
00051
      along y axis
00052
          if((rx[n] + atomRadius[n]) >= regionH[1]){
00053
             rx[n] = 0.999999 * regionH[1] - atomRadius[n]; vx[n] = -vx[n];
00054
           }if((rx[n] - atomRadius[n]) < -regionH[1]){</pre>
            rx[n] = -0.999999*regionH[1] + atomRadius[n]; vx[n] = -vx[n];
00055
00056
```

3.10 source/ApplyDrivingForce.c File Reference

```
#include <stdio.h>
#include <stdlib.h>
#include "global.h"
Include dependency graph for ApplyDrivingForce.c:
```



Functions

• void ApplyDrivingForce ()

3.10.1 Function Documentation

3.10.1.1 ApplyDrivingForce()

```
void ApplyDrivingForce ( )
Definition at line 25 of file ApplyDrivingForce.c.
00025
00026 int n;
00027 double Vxblock, Vyblock;
00028 double Vxsubstrate, Vysubstrate;
00029    Vxblock = 0.0;    Vyblock = 0.0;
00030 Vxsubstrate = 0.0; Vysubstrate = 0.0;
00031 double gammav;
00032 gammav = 0.0;
00033
00034 double count_substrate = 0;
00035 double count_block = 0;
00036
00037 for (n = 1 ; n \le nAtom; n ++) {
        if(atomType[n] == 1 || atomType[n] == 2){
Vxsubstrate += vx[n]; Vysubstrate += vy[n];
00038
00039
00040
        count_substrate++;
00041
00042
         if(atomType[n] == 3 || atomType[n] == 4){
        Vxblock += vx[n]; Vyblock += vy[n];
00043
        count_block++;
00044
00045
        } }
00046
```

```
if(count_substrate > 0) {
               Vxsubstrate /= count_substrate;
Vysubstrate /= count_substrate;
00048
00049
00050
00051
00052
            if (count block > 0) {
              Vxblock /= count_block;
00054
             Vyblock /= count_block;
00055
00056
00057
          for (n = 1 ; n \le nAtom; n ++) {
           if(atomType[n] == 1 || atomType[n] == 2){
ax[n] += -gammav * (vx[n] - Vxsubstrate);
ay[n] += -gammav * (vy[n] - Vysubstrate);
00058
00059
00060
00061
           if(atomType[n] == 3 || atomType[n] == 4){
ax[n] += -gammav * (vx[n] - Vxblock);
ay[n] += -gammav * (vy[n] - Vyblock);
00062
00063
00064
00065 } } }
```

References atomType, ax, ay, nAtom, vx, and vy.

3.11 ApplyDrivingForce.c

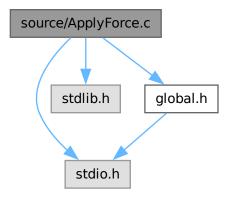
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00018
00019
00020
00021 #include<stdio.h>
00022 #include<stdlib.h>
00023 #include"global.h"
00024
00025 void ApplyDrivingForce(){
00026 int n;
00027 double Vxblock, Vyblock;
        double Vxsubstrate, Vysubstrate;
00029 Vxblock = 0.0; Vyblock = 0.0;
00030 Vxsubstrate = 0.0; Vysubstrate = 0.0;
00031 double gammav;
00032 gammav = 0.0;
00033
00034
        double count_substrate = 0;
00035
        double count_block = 0;
00036
00037
        for (n = 1 ; n \le nAtom; n ++) {
         if(atomType[n] == 1 || atomType[n] == 2){
Vxsubstrate += vx[n]; Vysubstrate += vy[n];
00038
00039
00040
         count substrate++;
00041
00042
          if(atomType[n] == 3 || atomType[n] == 4){
00043
         Vxblock += vx[n]; Vyblock += vy[n];
         count_block++;
00044
00045
00046
00047
         if(count_substrate > 0) {
           Vxsubstrate /= count_substrate;
Vysubstrate /= count_substrate;
00048
00049
00050
00051
00052
         if(count_block > 0) {
          Vxblock /= count_block;
Vyblock /= count_block;
00053
00054
00055
00056
00057
        for(n = 1; n <= nAtom; n ++) {
  if(atomType[n] == 1 || atomType[n] == 2) {
  ax[n] += -gammav * (vx[n] - Vxsubstrate);</pre>
00058
         ay[n] += -gammav * (vy[n] - Vysubstrate);
```

3.12 source/ApplyForce.c File Reference

```
#include <stdio.h>
#include <stdlib.h>
#include "global.h"
```

Include dependency graph for ApplyForce.c:



Functions

• void ApplyForce ()

3.12.1 Function Documentation

3.12.1.1 ApplyForce()

References ax, ay, fx, fxByfy, fy, FyBylx, molID, nAtom, nAtomBlock, and regionH. Referenced by main().

Here is the caller graph for this function:



3.13 ApplyForce.c

Go to the documentation of this file.

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00018
00019 */
00020
00021 #include<stdio.h>
00022 #include<stdlib.h>
00023 #include"global.h"
00024
00025 void ApplyForce(){
00026 int n;
00027 double 1x;
00028 lx = regionH[1];

00029 fy = (FyBylx * lx)/nAtomBlock;

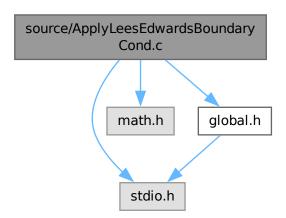
00030 fx = fxByfy * fy;

00031 for(n = 1; n <= nAtom; n ++) {
00032
          if(molID[n] == 2){
00033
           ax[n] += fx;
00034
            ay[n] -= fy;
00035 } }
```

3.14 source/ApplyLeesEdwardsBoundaryCond.c File Reference

```
#include <stdio.h>
#include <math.h>
#include "global.h"
```

Include dependency graph for ApplyLeesEdwardsBoundaryCond.c:



Functions

00040 00041 00042

void ApplyLeesEdwardsBoundaryCond ()

3.14.1 Function Documentation

3.14.1.1 ApplyLeesEdwardsBoundaryCond()

void ApplyLeesEdwardsBoundaryCond ()

//vx[n] += shearVelocity;
ry[n] += region[2];

```
Definition at line 25 of file ApplyLeesEdwardsBoundaryCond.c.
00026 int n;
00027 for (n = 1; n <= nAtom; n++) { 00028 //PBC along x-direction
00029 if(rx[n] >= regionH[1])
00030     rx[n] -= region[1];
00031 else if(rx[n] < -regionH[1])
00032
         rx[n] += region[1];
00033
00034 //LEBC along y-direction
}else if(ry[n] < -regionH[2]) {
  rx[n] += shearDisplacement;
  if(rx[n] >= regionH[1]) rx[n] -= region[1];
```

References nAtom, region, regionH, rx, ry, and shearDisplacement.

3.15 ApplyLeesEdwardsBoundaryCond.c

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00007 \star (at your option) any later version.
```

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00017 Copyright (C) 2025 Harish Charan, University of Durham, UK
00018
00019 */
00020
00021 #include <stdio.h>
00022 #include <math.h>
00023 #include "global.h"
00024
00025 void ApplyLeesEdwardsBoundaryCond() {
00026 int n;
00027 for (n = 1; n <= nAtom; n++) {
00028 //PBC along x-direction
00029 //FBC along x-direction

00029 if(rx[n] >= regionH[1])

00030 rx[n] -= region[1];

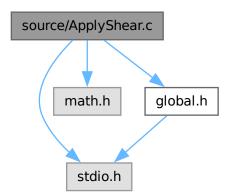
00031 else if(rx[n] < -regionH[1])

00032 rx[n] += region[1];
00034 //LEBC along y-direction
00035 if(ry[n] >= regionH[2]){
00036    rx[n] -= shearDisplacement;
             if(rx[n] < -regionH[1]) rx[n] += region[1];</pre>
00037
            //vx[n] -= shearVelocity;
ry[n] -= region[2];
00038
00039
00040
            }else if(ry[n] < -regionH[2]){</pre>
00041
             rx[n] += shearDisplacement;
             if(rx[n] >= regionH[1]) rx[n] -= region[1];
//vx[n] += shearVelocity;
ry[n] += region[2];
00042
00043
00044
00046
00047 }
00048
```

3.16 source/ApplyShear.c File Reference

```
#include <stdio.h>
#include <math.h>
#include "global.h"
```

Include dependency graph for ApplyShear.c:



3.17 ApplyShear.c 23

Functions

• void ApplyShear ()

3.16.1 Function Documentation

3.16.1.1 ApplyShear()

```
void ApplyShear ( ) Definition at line 25 of file ApplyShear.c. 00025 { 00026 int n; 00027 for (n = 1 ; n <= nAtom ; n ++) { 00028 rx[n] += strain \star ry[n]; 00029 //vx[n] += stranRate \star ry[n]; 00030 } }
```

References nAtom, rx, ry, and strain.

3.17 ApplyShear.c

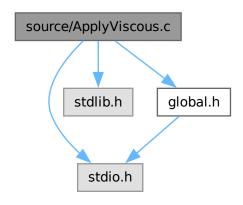
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00018
00019 */
00020
00021 #include<stdio.h>
00022 #include<math.h>
00023 #include"global.h"
00024
00025 void ApplyShear() {
00026 int n;
00027 for(n = 1 ; n <= nAtom ; n ++) {
00028     rx[n] += strain * ry[n];
00029     //vx[n] += stranRate * ry[n];
00030 } }
```

3.18 source/ApplyViscous.c File Reference

```
#include <stdio.h>
#include <stdlib.h>
#include "global.h"
```

Include dependency graph for ApplyViscous.c:



Functions

void ApplyViscous ()

3.18.1 Function Documentation

3.18.1.1 ApplyViscous()

```
void ApplyViscous ( )
Definition at line 25 of file ApplyViscous.c.
00025
00026   int n;
00027   double gammav;
00028   gammav = 1.0;
00029   for (n = 1 ; n <= nAtom; n ++) {
00030    ax[n] += -gammav * vx[n];
00031   ay[n] += -gammav * vy[n];
00032   }
</pre>
```

References ax, ay, nAtom, vx, and vy.

3.19 ApplyViscous.c

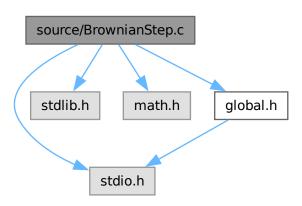
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00018
00019 */
00020
00021 #include<stdio.h>
00022 #include<stdlib.h>
00023 #include"global.h"
```

```
00025 void ApplyViscous() {
00026   int n;
00027   double gammav;
00028   gammav = 1.0;
00029   for(n = 1 ; n <= nAtom; n ++) {
00030    ax[n] += -gammav * vx[n];
00031   ay[n] += -gammav * vy[n];
00032   } }
00033
00034</pre>
```

3.20 source/BrownianStep.c File Reference

```
#include <stdio.h>
#include <stdlib.h>
#include <math.h>
#include "global.h"
```

Include dependency graph for BrownianStep.c:



Functions

void BrownianStep ()

3.20.1 Function Documentation

3.20.1.1 BrownianStep()

```
void BrownianStep ( )
Definition at line 26 of file BrownianStep.c.
00026
          if(stepCount <= stepEquil){</pre>
00027
           double A, S1, S2, T;
00028
00029
           int n;
S1 = 0.; S2 = 0;
00030
           double halfdt = 0.5*deltaT;
00031
00032
           for (n = 1; n <= nAtom; n++) {</pre>
            T = vx[n] + halfdt * ax[n];
S1 += T * ax[n];
00033
00034
             S2 += Sqr(T);
00035
00036
             T = vy[n] + halfdt * ay[n];
S1 += T * ay[n];
00037
00038
              S2 += Sqr(T);
00039
00040
           A = -S1 / S2;
double C = 1 + A*deltaT;
double D = deltaT * (1 + 0.5 * A * deltaT);
00041
00042
00043
           for (n = 1; n <= nAtom; n++) {</pre>
```

```
vx[n] = C * vx[n] + D * ax[n];
             rx[n] += deltaT * vx[n];
vy[n] = C * vy[n] + D * ay[n];
00046
00047
00048
             ry[n] += deltaT * vy[n];
00049
00050
        }else{
             int n;
00052
             //SETTING TEMP = 0.0
00053
            if (stepCount == stepEquil+1){
            for (n = 1; n <= nAtom; n ++) {
vx[n] = 0.0;</pre>
00054
00055
            vy[n] = 0.0;
00056
00057
            } }
00058
            double zeta = 1.0;
00059
            double dx, dy;
00060
            for(n = 1; n <= nAtom; n ++) {
             dx = rx[n];
rx[n] += zeta * ax[n] * deltaT;
00061
00062
             dx = rx[n] - dx;
00063
00064
             vx[n] = dx/deltaT;
00065
             dy = ry[n];
00066
             ry[n] += zeta * ay[n] * deltaT;
             dy = ry[n] - dy;
00067
00068
             vy[n] = dy/deltaT;
00069
00070
00071 }
```

References ax, ay, deltaT, nAtom, rx, ry, Sqr, stepCount, stepEquil, vx, and vy.

3.21 BrownianStep.c

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00018
00019
00020
00021 #include<stdio.h>
00022 #include<stdlib.h>
00023 #include<math.h>
00024 #include"global.h"
00025
00026 void BrownianStep(){
00027
        if(stepCount <= stepEquil){</pre>
00028
           double A, S1, S2, T;
            int n;
S1 = 0.; S2 = 0;
00029
00030
            double halfdt = 0.5*deltaT;
00031
            for (n = 1; n <= nAtom; n++) {</pre>
00032
              T = vx[n] + halfdt * ax[n];
00034
              S1 += T * ax[n];
              S2 += Sqr(T);
00035
00036
               T = vy[n] + halfdt * ay[n];
00037
               S1 += T * ay[n];
00038
               S2 += Sqr(T);
00040
00041
            A = -S1 / S2;
            double C = 1 + A*deltaT;
double D = deltaT * (1 + 0.5 * A * deltaT);
00042
00043
            for (n = 1; n <= nAtom; n++) {
  vx[n] = C * vx[n] + D * ax[n];
00044
00046
               rx[n] += deltaT * vx[n];
00047
               vy[n] = C * vy[n] + D * ay[n];
               ry[n] += deltaT * vy[n];
00048
00049
00050
         }else{
00051
              int n;
               //SETTING TEMP = 0.0
```

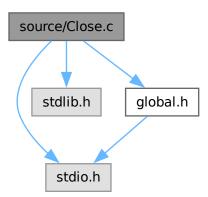
```
if (stepCount == stepEquil+1) {
  for(n = 1 ; n <= nAtom ; n ++) {
    vx[n] = 0.0;
    vy[n] = 0.0;</pre>
00054
00055
00056
00057
00058
                 double zeta = 1.0;
                double dx, dy;

for(n = 1; n <= nAtom; n ++) {
00060
00061
                  dx = rx[n];
                  rx[n] += zeta * ax[n] * deltaT;
dx = rx[n] - dx;
vx[n] = dx/deltaT;
00062
00063
00064
                  dy = ry[n];
ry[n] += zeta * ay[n] * deltaT;
00065
00066
                  dy = ry[n] - dy;
vy[n] = dy/deltaT;
00067
00068
00069
00070 }
00072
```

3.22 source/Close.c File Reference

```
#include <stdio.h>
#include <stdlib.h>
#include "global.h"
```

Include dependency graph for Close.c:



Functions

• void Close ()

3.22.1 Function Documentation

3.22.1.1 Close()

```
void Close ( )
Definition at line 24 of file Close.c.
00024
00025
        int n:
00026
        free(rx);
00027
        free(ry);
00028
       free(vx);
00029
        free(vy);
00030
        free(ax);
00031
        free(ay);
00032
        free (fax);
00033
        free(fay);
```

```
free(cellList);
00035
00036
         free(atomID); free(atomType); free(atomRadius); free(atomMass);
00037
         free (speed);
         free(atom1); free(atom2); free(BondID);
free(BondType); free(kb); free(ro);
free(ImageX); free(ImageY); free(rxUnwrap); free(ryUnwrap);
00038
00039
00041
         free(atomIDInterface);
00042
         free(PairID); free(Pairatom1); free(Pairatom2);
00043
         free(PairXij); free(PairYij);
00044
00045
         free (DeltaXiiOld);
00046
         free (DeltaYijOld);
00047
00048
         free (molID);
00049
         for (n = 0; n <= nAtom; n++) {</pre>
00050
00051
          free(isBonded[n]);
00052
00053
          free(isBonded);
00054
00055
00056
         for(n = 0; n <= nAtom; n++) {
  free(DeltaXijOldPair[n]);</pre>
00057
00058
          free (DeltaYijOldPair[n]);
00059
00060
00061
            free(DeltaXijOldPair);
           free (DeltaYijOldPair);
00062
00063
00064
         for (n = 0; n \le nBuffCorr; n++) {
00065
           free(cfOrg[n]);
00066
            free(spacetimeCorr[n]);
00067
00068
         free(cfOrg);
         free(spacetimeCorr);
00069
00070
         free(cfVal);
00071
         free(indexCorr);
00072
         free(spacetimeCorrAv);
00073
00074
         free(indexAcf);
00075
         free (viscAcfOrg);
00076
         free(viscAcfAv);
for(n = 0 ; n <= nBuffAcf ; n ++)</pre>
00077
00078
           free(viscAcf[n]);
00079
         free(viscAcf);
08000
00081 }
```

References atom1, atom2, atomID, atomIDInterface, atomMass, atomRadius, atomType, ax, ay, BondID, BondType, cellList, cfOrg, cfVal, DeltaXijOld, DeltaXijOldPair, DeltaYijOld, DeltaYijOldPair, fax, fay, ImageX, ImageY, indexAcf, indexCorr, isBonded, kb, molID, nAtom, nBuffAcf, nBuffCorr, Pairatom1, Pairatom2, PairID, PairXij, PairYij, ro, rx, rxUnwrap, ry, ryUnwrap, spacetimeCorr, spacetimeCorrAv, speed, viscAcf, viscAcfAv, viscAcfOrg, vx, and vy.

Referenced by main().

Here is the caller graph for this function:



3.23 Close.c

```
00001 /*
00002 * This file is part of Lamina.
00003 *
00004 * Lamina is free software: you can redistribute it and/or modify
00005 * it under the terms of the GNU General Public License as published by
00006 * the Free Software Foundation, either version 3 of the License, or
```

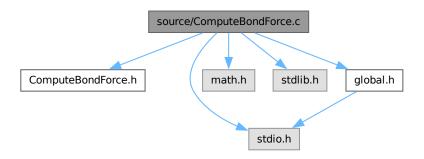
```
00007 * (at your option) any later version.
80000
00009 * Lamina is distributed in the hope that it will be useful,
00010 \, but WITHOUT ANY WARRANTY; without even the implied warranty of 00011 \, \, MERCHANTABILITY or FITNESS FOR A PARTICULAR PURPOSE. See the
00012 * GNU General Public License for more details.
00014 \,\,\star\,\, You should have received a copy of the GNU General Public License
00015 \star along with Lamina. If not, see <a href="https://www.gnu.org/licenses/">https://www.gnu.org/licenses/</a>.
00016
00017 Copyright (C) 2025 Harish Charan, University of Durham, UK
00018
00019
00020
00021 #include<stdio.h>
00022 #include<stdlib.h>
00023 #include"global.h"
00024 void Close() {
00025 int n;
        free(rx);
00027
00028
        free(vx);
00029
        free(vv);
00030
        free (ax):
00031
        free(ay);
00032
        free(fax);
00033
00034
        free(cellList);
00035
00036
        free(atomID); free(atomType); free(atomRadius); free(atomMass);
00037
        free (speed);
00038
        free(atom1); free(atom2); free(BondID);
00039
        free(BondType); free(kb); free(ro);
00040
         free(ImageX); free(ImageY); free(rxUnwrap); free(ryUnwrap);
00041
        free(atomIDInterface);
00042
        free(PairID); free(Pairatom1); free(Pairatom2);
00043
        free(PairXij); free(PairYij);
00044
00045
        free (DeltaXijOld);
00046
        free (DeltaYijOld);
00047
00048
        free (molID):
00049
00050
        for (n = 0; n <= nAtom; n++) {</pre>
00051
         free(isBonded[n]);
00052
00053
         free(isBonded);
00054
00055
00056
        for(n = 0; n <= nAtom; n++) {</pre>
00058
         free (DeltaXijOldPair[n]);
00059
          free (DeltaYijOldPair[n]);
00060
00061
          free(DeltaXijOldPair);
00062
          free (DeltaYijOldPair);
00063
00064
        for (n = 0; n <= nBuffCorr; n++) {</pre>
         free(cfOrg[n]);
00065
00066
          free(spacetimeCorr[n]);
00067
        free(cfOrg);
00068
00069
        free(spacetimeCorr);
00070
00071
        free(indexCorr);
00072
        free(spacetimeCorrAv);
00073
00074
        free(indexAcf);
00075
        free(viscAcfOrg);
        free(viscAcfAv);
for(n = 0 ; n <= nBuffAcf ; n ++)</pre>
00077
00078
          free(viscAcf[n]);
00079
        free(viscAcf);
00080
00081 }
```

3.24 source/ComputeBondForce.c File Reference

```
#include "ComputeBondForce.h"
#include <stdio.h>
#include <math.h>
#include <stdlib.h>
```

#include "global.h"

Include dependency graph for ComputeBondForce.c:



Functions

• void ComputeBondForce ()

3.24.1 Function Documentation

3.24.1.1 ComputeBondForce()

```
void ComputeBondForce ( )
Definition at line 28 of file ComputeBondForce.c.
00028
00029
         int n;
         double dr[NDIM+1], r, rr, ri, roi;
00030
00031
         double uVal, fcVal;
00032
00033
         uVal = 0.0; TotalBondEnergy = 0.0;
00034
         virSumBond = 0.0; virSumBondxx = 0.0; virSumBondyy = 0.0; virSumBondxy = 0.0;
00035
00036
         double vr[NDIM+1], fdVal, rri;
00037
00038
         for (n = 1 ; n \le nAtom ; n ++) {
         nodeDragy[n] = 0.0;
nodeDragy[n] = 0.0;
00039
00040
00041
         } //Important change made on 03Apr2025. Mention it in GitHub
00042
00043
         int atom1TD, atom2TD:
00044
00045
         for (n=1; n<=nBond; n++) {</pre>
00046
         rr = 0.0; rri = 0.0; fcVal = 0.0; fdVal = 0.0; strech = 0.0;
         atom1ID = atom1[n];
atom2ID = atom2[n];
00047
00048
00049
00050
          dr[1] = rx[atom1ID] - rx[atom2ID];
          if(dr[1] >= regionH[1])
00051
00052
           dr[1] -= region[1];
00053
          else if(dr[1] < -regionH[1])</pre>
00054
           dr[1] += region[1];
00055
00056
          dr[2] = ry[atom1ID] - ry[atom2ID];
          if(dr[2] >= regionH[2]) {
  dr[1] -= shearDisplacement;
00057
00058
           if(dr[1] < -regionH[1]) dr[1] += region[1];
dr[2] -= region[2];</pre>
00059
00060
         }else if(dr[2] < -regionH[2]) {
    dr[1] += shearDisplacement;
    if(dr[1] >= regionH[1]) dr[1] -= region[1];
00061
00062
00063
00064
          dr[2] += region[2];
00065
00066
00067
          rr = Sqr(dr[1]) + Sqr(dr[2]);
          r = sqrt(rr);
rri = 1.0/rr;
00068
00069
00070
          ri = 1.0/r;
00071
          roi = 1.0/ro[n];
00072
          strech = (r * roi - 1.0);
```

```
uVal = 0.5 * kb[n] * ro[n] * Sqr(strech);
          fcVal = -kb[n] * strech * ri; //F = -Grad U
00074
00075
          vr[1] = vx[atom1ID] - vx[atom2ID];
vr[2] = vy[atom1ID] - vy[atom2ID];
00076
00077
00078
          fdVal = -gamman * (vr[1]*dr[1] + vr[2]*dr[2]) * rri; //node-node drag
08000
          //DampFlag = 1. LAMMPS version
00081
          if(DampFlag == 1){
00082
          nodeDragx[atom1ID] = fdVal * dr[1]; //node-node drag //Important change made on 03Apr2025.
      Mention it in GitHub
          nodeDragy[atomlID] = fdVal * dr[2]; //node-node drag //Adding the drag forces is wrong. Only add
00083
      the
         nodeDragx[atom2ID] = -fdVal * dr[1]; //node-node drag //total force
nodeDragy[atom2ID] = -fdVal * dr[2]; //node-node drag
00084
00085
00086
          ax[atom1ID] += (fcVal + fdVal) * dr[1];
ay[atom1ID] += (fcVal + fdVal) * dr[2];
ax[atom2ID] += -(fcVal + fdVal) * dr[1];
00087
00088
00089
00090
          ay[atom2ID] += -(fcVal + fdVal) * dr[2];
00091
00092
00093
          //DampFlag = 2. Suzanne notes version
          else if(DampFlag == 2){
00094
          00095
00096
00097
00098
00099
00100
          ax[atom1ID] += (fcVal * dr[1] - gamman * vr[1]);
          ax[atomlID] += (fcVal * dr[1] - gamman * vr[2]);
ax[atom2ID] += -(fcVal * dr[1] - gamman * vr[1]);
00101
00102
00103
          ay[atom2ID] += -(fcVal * dr[2] - gamman * vr[2]);
00104
00105
          //DampFlag = 3. Suzanne PRL, 130, 178203 (2023) version
00106
          else if(DampFlag == 3){
00107
           DeltaXijNew = dr[1];
00108
00109
           DeltaYijNew = dr[2];
00110
00111
            if(stepCount == 0) { // First timestep
            DeltaXijOld[n] = DeltaXijNew;
DeltaYijOld[n] = DeltaYijNew;
00112
00113
00114
00115
00116
            DeltaXij = DeltaXijNew - DeltaXijOld[n];
           DeltaYij = DeltaYijNew - DeltaYijOld[n];
DeltaVXij = DeltaXij / deltaT;
DeltaVYij = DeltaYij / deltaT;
00117
00118
00119
00120
00121
            // Now update for the next timestep
00122
            DeltaXijOld[n] = DeltaXijNew;
00123
            DeltaYijOld[n] = DeltaYijNew;
00124
            nodeDragx[atom1ID] = -gamman * DeltaVXij; //node-node drag
00125
           nodeDragy[atomID] = -gamman * DeltaVYij; //node-node drag
nodeDragy[atom2ID] = -(-gamman * DeltaVXij); //node-node drag
nodeDragy[atom2ID] = -(-gamman * DeltaVYij); //node-node drag
00126
00128
00129
           ax[atom1ID] += (fcVal * dr[1] - gamman * DeltaVXij);
ay[atom1ID] += (fcVal * dr[2] - gamman * DeltaVYij);
ax[atom2ID] += -(fcVal * dr[1] - gamman * DeltaVXij);
00130
00131
00132
            ay[atom2ID] += -(fcVal * dr[2] - gamman * DeltaVYij);
00133
00134
00135
00136
00137
          BondLength[n] = r;
          BondEnergy[n] = uVal; //No 0.5 factor since it is the energy of the bond
00138
00139
          TotalBondEnergy
                                += BondEnergy[n];
00141
                             0.5 * (fcVal + fdVal) * rr;
          virSumBond +=
          virSumBondxx += 0.5 * (fcVal + fdVal) * dr[1] * dr[1];
00142
          virSumBondyy += 0.5 * (fcVal + fdVal) * dr[2] * dr[2];
00143
          virSumBondxy += 0.5 * (fcVal + fdVal) * dr[1] * dr[2];
00144
00145 } }
```

References atom1, atom2, ax, ay, BondEnergy, BondLength, DampFlag, deltaT, DeltaVXij, DeltaVYij, DeltaXij, DeltaXij, DeltaXijNew, DeltaXijOld, DeltaYijNew, DeltaYijOld, gamman, kb, nAtom, nBond, NDIM, nodeDragx, nodeDragy, region, regionH, ro, rx, ry, shearDisplacement, Sqr, stepCount, strech, TotalBondEnergy, virSumBond, virSumBondxx, virSumBondxy, virSumBondyy, vx, and vy. Referenced by main().

Here is the caller graph for this function:



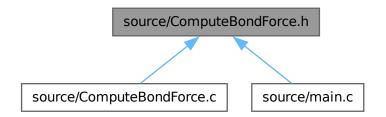
3.25 ComputeBondForce.c

```
00002
       * This file is part of Lamina.
00003
       * Lamina is free software: you can redistribute it and/or modify * it under the terms of the GNU General Public License as published by
00004
00005
        * the Free Software Foundation, either version 3 of the License, or
        * (at your option) any later version.
80000
00009 \star Lamina is distributed in the hope that it will be useful,
00010 * but WITHOUT ANY WARRANTY; without even the implied warranty of
        * MERCHANTABILITY or FITNESS FOR A PARTICULAR PURPOSE. See the
00011
00012
      * GNU General Public License for more details.
00013
* You should have received a copy of the GNU General Public License 00015 * along with Lamina. If not, see <a href="https://www.gnu.org/licenses/">https://www.gnu.org/licenses/</a>.
00016
00017 Copyright (C) 2025 Harish Charan, University of Durham, UK
00018
00019
00020
00021 #include "ComputeBondForce.h"
00022
00023 #include<stdio.h>
00024 #include<math.h>
00025 #include<stdlib.h>
00026 #include"global.h"
00027
00028 void ComputeBondForce(){
00029
        int n;
         double dr[NDIM+1], r, rr, ri, roi;
00030
00031
         double uVal, fcVal;
00032
00033
         uVal = 0.0; TotalBondEnergy = 0.0;
00034
         virSumBond = 0.0; virSumBondxx = 0.0; virSumBondyy = 0.0; virSumBondxy = 0.0;
00035
00036
         double vr[NDIM+1], fdVal, rri;
00037
00038
         for (n = 1 ; n <= nAtom ; n ++) {</pre>
00039
          nodeDragx[n] = 0.0;
00040
         nodeDragy[n] = 0.0;
00041
         } //Important change made on 03Apr2025. Mention it in GitHub
00042
00043
         int atom1ID, atom2ID;
00044
00045
         for (n=1; n<=nBond; n++) {</pre>
00046
          rr = 0.0; rri = 0.0; fcVal = 0.0; fdVal = 0.0; strech = 0.0;
          atom1ID = atom1[n];
atom2ID = atom2[n];
00047
00048
00049
00050
          dr[1] = rx[atom1ID] - rx[atom2ID];
          if(dr[1] >= regionH[1])
dr[1] -= region[1];
00051
00052
00053
          else if(dr[1] < -regionH[1])</pre>
00054
           dr[1] += region[1];
00055
00056
          dr[2] = ry[atom1ID] - ry[atom2ID];
          if(dr[2] >= regionH[2]){
00058
           dr[1] -= shearDisplacement;
00059
           if(dr[1] < -regionH[1]) dr[1] += region[1];</pre>
00060
           dr[2] -= region[2];
         }else if(dr[2] < -regionH[2]){
dr[1] += shearDisplacement;</pre>
00061
00062
00063
           if(dr[1] >= regionH[1]) dr[1] -= region[1];
```

```
dr[2] += region[2];
00065
00066
00067
           rr = Sqr(dr[1]) + Sqr(dr[2]);
           r = sqrt(rr);
00068
00069
           rri = 1.0/rr;
           ri = 1.0/r;
00070
00071
            roi = 1.0/ro[n];
           strech = (r * roi - 1.0);
uVal = 0.5 * kb[n] * ro[n] * Sqr(strech);
00072
00073
00074
           fcVal = -kb[n] * strech * ri; //F = -Grad U
00075
           vr[1] = vx[atom1ID] - vx[atom2ID];
vr[2] = vy[atom1ID] - vy[atom2ID];
00076
00077
00078
            fdVal = -gamman * (vr[1]*dr[1] + vr[2]*dr[2]) * rri; //node-node drag
00079
08000
            //DampFlag = 1. LAMMPS version
            if(DampFlag == 1){
00081
           nodeDragx[atom1ID] = fdVal * dr[1]; //node-node drag //Important change made on 03Apr2025.
00082
       Mention it in GitHub
           nodeDragy[atom1ID] = fdVal * dr[2]; //node-node drag //Adding the drag forces is wrong. Only add
00083
      the
           nodeDragx[atom2ID] = -fdVal * dr[1]; //node-node drag //total force
nodeDragy[atom2ID] = -fdVal * dr[2]; //node-node drag
00084
00085
00086
00087
            ax[atom1ID] += (fcVal + fdVal) * dr[1];
            ax[atom1ID] += (fcVal + fdVal) * dr[2];
ax[atom2ID] += -(fcVal + fdVal) * dr[1];
ay[atom2ID] += -(fcVal + fdVal) * dr[2];
00088
00089
00090
00091
00092
00093
            //DampFlag = 2. Suzanne notes version
00094
            else if(DampFlag == 2){
           nodeDragy[atom1ID] = -gamman * vr[1]; //node-node drag
nodeDragy[atom2ID] = -gamman * vr[2]; //node-node drag
nodeDragy[atom2ID] = -(-gamman * vr[1]); //node-node drag
nodeDragy[atom2ID] = -(-gamman * vr[2]); //node-node drag
00095
00096
00097
00098
           ax[atomlID] += (fcVal * dr[1] - gamman * vr[1]);
ay[atomlID] += (fcVal * dr[2] - gamman * vr[2]);
ax[atom2ID] += -(fcVal * dr[1] - gamman * vr[1]);
00100
00101
00102
            ay[atom2ID] += -(fcVal * dr[2] - gamman * vr[2]);
00103
00104
00105
00106
            //DampFlag = 3. Suzanne PRL, 130, 178203 (2023) version
00107
            else if(DampFlag == 3){
            DeltaXijNew = dr[1];
DeltaYijNew = dr[2];
00108
00109
00110
00111
             if(stepCount == 0) { // First timestep
             DeltaXijOld[n] = DeltaXijNew;
DeltaYijOld[n] = DeltaYijNew;
00112
00113
00114
00115
            DeltaXij = DeltaXijNew - DeltaXijOld[n];
DeltaYij = DeltaYijNew - DeltaYijOld[n];
DeltaVXij = DeltaXij / deltaT;
00116
00117
00118
00119
             DeltaVYij = DeltaYij / deltaT;
00120
00121
             // Now update for the next timestep
            DeltaXijOld[n] = DeltaXijNew;
DeltaYijOld[n] = DeltaYijNew;
00122
00123
00124
            00125
00126
00127
00128
00129
             ax[atom1ID] += (fcVal * dr[1] - gamman * DeltaVXij);
00130
            ax[atomlID] += (fcVal * dr[2] - gamman * DeltaVXij);
ax[atom2ID] += -(fcVal * dr[1] - gamman * DeltaVXij);
ay[atom2ID] += -(fcVal * dr[2] - gamman * DeltaVXij);
00131
00132
00133
00134
00135
00136
00137
            BondLength[n] = r;
00138
            BondEnergy[n] = uVal; //No 0.5 factor since it is the energy of the bond
00139
            TotalBondEnergy += BondEnergy[n];
00140
            virSumBond += 0.5 * (fcVal + fdVal) * rr:
00141
           virSumBondxx += 0.5 * (fcVal + fdVal) * dr[1] * dr[1];
virSumBondyy += 0.5 * (fcVal + fdVal) * dr[2] * dr[2];
00142
           virSumBondxy += 0.5 * (fcVal + fdVal) * dr[1] * dr[2];
00144
00145 } }
```

3.26 source/ComputeBondForce.h File Reference

This graph shows which files directly or indirectly include this file:



Functions

• void ComputeBondForce ()

3.26.1 Function Documentation

3.26.1.1 ComputeBondForce()

```
void ComputeBondForce ( )
Definition at line 28 of file ComputeBondForce.c.
00028
00029
        double dr[NDIM+1], r, rr, ri, roi;
00030
00031
        double uVal, fcVal;
00032
00033
        uVal = 0.0; TotalBondEnergy = 0.0;
        virSumBond = 0.0; virSumBondxx = 0.0; virSumBondyy = 0.0; virSumBondxy = 0.0;
00034
00035
00036
        double vr[NDIM+1], fdVal, rri;
00037
00038
        for (n = 1 ; n <= nAtom ; n ++) {</pre>
00039
         nodeDragx[n] = 0.0;
00040
         nodeDragy[n] = 0.0;
00041
        } //Important change made on 03Apr2025. Mention it in GitHub
00042
00043
        int atom1ID, atom2ID;
00044
00045
        for (n=1; n<=nBond; n++) {</pre>
         rr = 0.0; rri = 0.0; fcVal = 0.0; fdVal = 0.0; strech = 0.0;
atomlID = atoml[n];
atom2ID = atom2[n];
00046
00047
00048
00049
00050
         dr[1] = rx[atom1ID] - rx[atom2ID];
00051
         if(dr[1] >= regionH[1])
         dr[1] -= region[1];
else if(dr[1] < -regionH[1])</pre>
00052
00053
00054
          dr[1] += region[1];
00055
00056
          dr[2] = ry[atom1ID] - ry[atom2ID];
00057
          if(dr[2] >= regionH[2]){
00058
          dr[1] -= shearDisplacement;
           if(dr[1] < -regionH[1]) dr[1] += region[1];</pre>
00059
00060
           dr[2] -= region[2];
        }else if(dr[2] < -regionH[2]) {
  dr[1] += shearDisplacement;</pre>
00061
00062
00063
           if(dr[1] >= regionH[1]) dr[1] -= region[1];
00064
         dr[2] += region[2];
00065
00066
         rr = Sqr(dr[1]) + Sqr(dr[2]);
00067
         r = sqrt(rr);
rri = 1.0/rr;
00068
00070
         ri = 1.0/r;
00071
         roi = 1.0/ro[n];
00072
          strech = (r * roi - 1.0);
```

```
uVal = 0.5 * kb[n] * ro[n] * Sqr(strech);
          fcVal = -kb[n] * strech * ri; //F = -Grad U
00074
00075
          vr[1] = vx[atom1ID] - vx[atom2ID];
vr[2] = vy[atom1ID] - vy[atom2ID];
00076
00077
00078
          fdVal = -gamman * (vr[1]*dr[1] + vr[2]*dr[2]) * rri; //node-node drag
08000
          //DampFlag = 1. LAMMPS version
00081
          if(DampFlag == 1){
00082
          nodeDragx[atom1ID] = fdVal * dr[1]; //node-node drag //Important change made on 03Apr2025.
      Mention it in GitHub
          nodeDragy[atomlID] = fdVal * dr[2]; //node-node drag //Adding the drag forces is wrong. Only add
00083
      the
          nodeDragx[atom2ID] = -fdVal * dr[1]; //node-node drag //total force
nodeDragy[atom2ID] = -fdVal * dr[2]; //node-node drag
00084
00085
00086
          ax[atom1ID] += (fcVal + fdVal) * dr[1];
ay[atom1ID] += (fcVal + fdVal) * dr[2];
ax[atom2ID] += -(fcVal + fdVal) * dr[1];
00087
00088
00089
00090
          ay[atom2ID] += -(fcVal + fdVal) * dr[2];
00091
00092
00093
          //DampFlag = 2. Suzanne notes version
          else if(DampFlag == 2){
00094
          00095
00096
00097
00098
00099
          ax[atomlID] += (fcVal * dr[1] - gamman * vr[1]);
ay[atomlID] += (fcVal * dr[2] - gamman * vr[2]);
ax[atom2ID] += -(fcVal * dr[1] - gamman * vr[1]);
00100
00101
00102
00103
          ay[atom2ID] += -(fcVal * dr[2] - gamman * vr[2]);
00104
00105
          //DampFlag = 3. Suzanne PRL, 130, 178203 (2023) version
00106
          else if(DampFlag == 3){
00107
           DeltaXijNew = dr[1];
00108
00109
           DeltaYijNew = dr[2];
00110
00111
            if(stepCount == 0) { // First timestep
            DeltaXijOld[n] = DeltaXijNew;
DeltaYijOld[n] = DeltaYijNew;
00112
00113
00114
00115
00116
            DeltaXij = DeltaXijNew - DeltaXijOld[n];
           DeltaYij = DeltaYijNew - DeltaYijOld[n];
DeltaVXij = DeltaXij / deltaT;
DeltaVYij = DeltaYij / deltaT;
00117
00118
00119
00120
00121
            // Now update for the next timestep
00122
            DeltaXijOld[n] = DeltaXijNew;
00123
            DeltaYijOld[n] = DeltaYijNew;
00124
            nodeDragx[atom1ID] = -gamman * DeltaVXij; //node-node drag
00125
           nodeDragy[atomID] = -gamman * DeltaVYij; //node-node drag
nodeDragy[atom2ID] = -(-gamman * DeltaVXij); //node-node drag
nodeDragy[atom2ID] = -(-gamman * DeltaVYij); //node-node drag
00126
00128
00129
00130
            ax[atom1ID] += (fcVal * dr[1] - gamman * DeltaVXij);
           ax[atom1ID] += (fcVal * dr[2] - gamman * DeltaVYij);
ax[atom2ID] += -(fcVal * dr[1] - gamman * DeltaVXij);
00131
00132
            ay[atom2ID] += -(fcVal * dr[2] - gamman * DeltaVYij);
00133
00134
00135
00136
00137
          BondLength[n] = r;
          BondEnergy [n] = uVal; //No 0.5 factor since it is the energy of the bond
00138
00139
          TotalBondEnergy
                               += BondEnergy[n];
00141
                             0.5 * (fcVal + fdVal) * rr;
          virSumBond +=
          virSumBondxx += 0.5 * (fcVal + fdVal) * dr[1] * dr[1];
00142
          virSumBondyy += 0.5 * (fcVal + fdVal) * dr[2] * dr[2];
00143
          virSumBondxy += 0.5 * (fcVal + fdVal) * dr[1] * dr[2];
00144
00145 } }
```

References atom1, atom2, ax, ay, BondEnergy, BondLength, DampFlag, deltaT, DeltaVXij, DeltaVYij, DeltaXij, DeltaXij, DeltaXijNew, DeltaXijOld, DeltaYijNew, DeltaYijOld, gamman, kb, nAtom, nBond, NDIM, nodeDragx, nodeDragy, region, regionH, ro, rx, ry, shearDisplacement, Sqr, stepCount, strech, TotalBondEnergy, virSumBond, virSumBondxx, virSumBondxy, virSumBondyy, vx, and vy. Referenced by main().

Here is the caller graph for this function:



3.27 ComputeBondForce.h

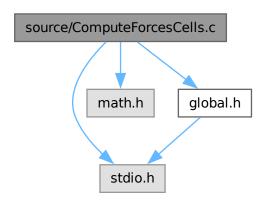
Go to the documentation of this file.

```
00001 #ifndef COMPUTE_BOND_FORCE_H
00002 #define COMPUTE_BOND_FORCE_H
00003
00004 void ComputeBondForce();
00005
00006 #endif
00007
```

3.28 source/ComputeForcesCells.c File Reference

```
#include <stdio.h>
#include <math.h>
#include "global.h"
```

Include dependency graph for ComputeForcesCells.c:



Functions

• void ComputeForcesCells ()

3.28.1 Function Documentation

3.28.1.1 ComputeForcesCells()

void ComputeForcesCells ()

```
Definition at line 25 of file ComputeForcesCells.c.
```

```
double dr[NDIM+1], invWid[NDIM+1], shift[NDIM+1], f, fcVal, rr, ri, r, uVal;
00026
               int c, I, J, m1, m1X, m1Y, m2, m2X, m2Y, n, offset;
int iofX[] = {0, 0, 1, 1, 0, -1, -1, -1, 0, 1},
    iofY[] = {0, 0, 0, 1, 1, 1, 0, -1, -1, -1};
00027
00028
00029
00031
                invWid[1] = cells[1]/region[1];
00032
               invWid[2] = cells[2]/region[2];
00033
00034
               for(n = nAtom+1; n <= nAtom+cells[1]*cells[2]; n++)</pre>
00035
                   cellList[n] = 0;
00036
00037
               for(n = 1 ; n <= nAtom ; n ++) {</pre>
00038
                  c = ((int)((ry[n] + regionH[2])*invWid[2]))*cells[1] + (int)((rx[n] + regionH[1])*invWid[1]) + (int)((ry[n] + regionW[1]) + (int)((r
          nAtom+ 1;
cellList[n] = cellList[c];
00039
00040
                   cellList[c] = n;
00041
00042
00043
                for(n = 1 ; n <= nAtom ; n ++) {</pre>
00044
                  ax[n] = 0.;
00045
                  ay[n] = 0.;
00046
00047
00048
              uSum = 0.0;
00049
               virSum = 0.0;
              rfAtom = 0.0;
00050
00051
              RadiusIJ = 0.0;
00052
00053
               gamman = 1.0:
00054
               double vr[NDIM+1], fd, fdVal, rrinv;
00055
               rrinv = 0.0;
00056
               fd = 0.0;
00057
               fdVal = 0.0;
00058
00059
               int start = 1 + rank*(cells[2]/size);
              int end = (rank+1) * (cells[2]/size);
00061
00062
               for(m1Y = start ; m1Y <= end ; m1Y ++) {</pre>
00063
                   for(m1X = 1 ; m1X <= cells[1] ; m1X ++) {</pre>
                      m1 = (m1Y-1) * cells[1] + m1X + nAtom;

for(offset = 1; offset <= 9; offset ++) {
00064
00065
00066
                   m2X = m1X + iofX[offset]; shift[1] = 0.;
                   if (m2X > cells[1]) {
00067
00068
                       m2X = 1; shift[1] = region[1];
00069
                   else if(m2X == 0){
                      m2X = cells[1]; shift[1] = -region[1];
00070
00071
00072
                   m2Y = m1Y + iofY[offset]; shift[2] = 0.;
                   if (m2Y > cells[2]) {
00074
                      m2Y = 1; shift[2] = region[2];
00075
                   else if(m2Y == 0){
00076
                      m2Y = cells[2]; shift[2] = -region[2];
00077
00078
                   m2 = (m2Y-1)*cells[1] + m2X + nAtom;
                   I = cellList[m1];
08000
                   while (I > 0) {
00081
                      J = cellList[m2];
00082
                       while (J > 0) {
                           if(m1 == m2 && J != I && (atomRadius[I] > 0. && atomRadius[J] > 0.)){
00083
                              dr[1] = rx[I] - rx[J] - shift[1];
dr[2] = ry[I] - ry[J] - shift[2];
00084
00085
00086
                               rr = Sqr(dr[1]) + Sqr(dr[2]);
00087
                               RadiusIJ = atomRadius[I] + atomRadius[J];
                              SqrRadiusIJ = Sqr(RadiusIJ);
00088
00089
                              if(rr < SqrRadiusIJ){</pre>
00090
                           r = sart(rr);
00091
                           ri = 1.0/r;
00092
                                           rrinv = 1.0/rr;
                                           vr[1] = vx[I] - vx[J];
vr[2] = vy[I] - vy[J];
00093
00094
                           RadiusIJInv = 1.0/RadiusIJ;
00095
                           uVal = Sqr(1.0 - r * RadiusIJInv);
fcVal = 2.0 * RadiusIJInv * (1.0 - r * RadiusIJInv) *ri;
00096
00097
00098
                                           fdVal = -gamman * (vr[1]*dr[1] + vr[2]*dr[2]) * rrinv; //disc-disc drag
00099
00100
                           f = fcVal * dr[1];
                                          fd = fdVal * dr[1];
00101
                           ax[I] += (f + fd);
00102
                                          discDragx[I] += fd; //disc-disc drag
00103
00104
00105
                           f = fcVal * dr[2];
00106
                                          fd = fdVal * dr[2];
00107
                           ay[I] += (f + fd);
                                           discDragy[I] += fd; //disc-disc drag
00108
00109
```

```
uSum += 0.5 * uVal;
               virSum += 0.5 * fcVal * rr;
rfAtom += 0.5 * dr[1] * fcVal * dr[2];
00111
00112
00113
00114
               else if (m1 != m2 && (atomRadius[I] > 0. && atomRadius[J] > 0.)){}
                 dr[1] = rx[I] - rx[J] - shift[1];
dr[2] = ry[I] - ry[J] - shift[2];
00115
00116
00117
                 rr = Sqr(dr[1]) + Sqr(dr[2]);
00118
                 RadiusIJ = atomRadius[I] + atomRadius[J];
00119
                 SqrRadiusIJ = Sqr(RadiusIJ);
                 if(rr < SqrRadiusIJ) {</pre>
00120
               r = sqrt(rr);
00121
00122
               ri = 1.0/r;
00123
                       rrinv = 1.0/r;
00124
                        vr[1] = vx[I] - vx[J];
               vr[2] = vy[I] - vy[J];
RadiusIJInv = 1.0/RadiusIJ;
00125
00126
               uVal = Sgr(1.0 - r * RadiusIJInv);
00127
               fcVal = 2.0 * RadiusIJInv * (1.0 - r * RadiusIJInv) *ri;
                        fdVal = -gamman * (vr[1]*dr[1] + vr[2]*dr[2]) * rrinv; //disc-disc drag
00129
00130
00131
               f = fcVal * dr[1];
               fd = fdVal * dr[1];
ax[I] += (f + fd);
00132
00133
00134
                       discDragx[I] += fd; //disc-disc drag
00135
00136
               f = fcVal * dr[2];
00137
                       fd = fdVal * dr[2];
00138
               ay[I] += (f + fd);
                       discDragy[I] += fd; //disc-disc drag
00139
00140
00141
               uSum += 0.5 * uVal;
00142
               virSum += 0.5 * fcVal * rr;
               rfAtom += 0.5 * dr[1] * fcVal * dr[2];
00143
00144
00145
00146
                   J = cellList[J];
             I = cellList[I];
00148
00149
          }
00150
00151
00152
00153 }
```

References atomRadius, ax, ay, cellList, cells, discDragx, discDragy, gamman, nAtom, NDIM, RadiusIJ, RadiusIJInv, rank, region, regionH, rfAtom, rx, ry, size, Sqr, SqrRadiusIJ, uSum, virSum, vx, and vy.

3.29 ComputeForcesCells.c

```
00001 /*
00002 \,\star\, This file is part of Lamina.
00003 *
00004 * Lamina is free software: you can redistribute it and/or modify
       * it under the terms of the GNU General Public License as published by
       \star the Free Software Foundation, either version 3 of the License, or
00007 * (at your option) any later version.
80000
00009 \star Lamina is distributed in the hope that it will be useful,
00010 * but WITHOUT ANY WARRANTY; without even the implied warranty of
00011 * MERCHANTABILITY or FITNESS FOR A PARTICULAR PURPOSE. See the
00012 * GNU General Public License for more details.
00013
00014 \,\, You should have received a copy of the GNU General Public License 00015 \,\, * along with Lamina. If not, see <a href="https://www.gnu.org/licenses/">https://www.gnu.org/licenses/</a>.
00016
00017 Copyright (C) 2025 Harish Charan, University of Durham, UK
00018
00019
00020
00021 #include<stdio.h>
00022 #include<math.h>
00023 #include"global.h"
00025 void ComputeForcesCells(){
00026
      double dr[NDIM+1], invWid[NDIM+1], shift[NDIM+1], f, fcVal, rr, ri, r, uVal;
00027
        int c, I, J, m1, m1X, m1Y, m2, m2X, m2Y, n, offset;
        int iofX[] = \{0, 0, 1, 1, 0, -1, -1, -1, 0, 1\},
00028
             iofY[] = {0, 0, 0, 1 ,1, 1, 0, -1, -1, -1};
00029
00030
        invWid[1] = cells[1]/region[1];
        invWid[2] = cells[2]/region[2];
00032
00033
```

```
for(n = nAtom+1; n <= nAtom+cells[1]*cells[2]; n++)</pre>
00035
                  cellList[n] = 0;
00036
00037
               for(n = 1 ; n <= nAtom ; n ++) {</pre>
                  c = ((int)((ry[n] + regionH[2])*invWid[2]))*cells[1] + (int)((rx[n] + regionH[1])*invWid[1]) + (int)((ry[n] + regionW[1]) + (int)((r
00038
          nAtom+ 1;
00039
               cellList[n] = cellList[c];
00040
                   cellList[c] = n;
00041
00042
00043
               for (n = 1 ; n <= nAtom ; n ++) {</pre>
                ax[n] = 0.;
00044
00045
                  ay[n] = 0.;
00046
00047
00048
               uSum = 0.0;
               virSum = 0.0;
rfAtom = 0.0;
00049
00050
00051
               RadiusIJ = 0.0;
00052
00053
               gamman = 1.0;
00054
               double vr[NDIM+1], fd, fdVal, rrinv;
00055
               rrinv = 0.0:
00056
              fd = 0.0;
00057
               fdVal = 0.0;
00058
00059
               int start = 1 + rank*(cells[2]/size);
00060
              int end = (rank+1) * (cells[2]/size);
00061
00062
               for (m1Y = start ; m1Y <= end ; m1Y ++) {</pre>
00063
                 for (m1X = 1 ; m1X <= cells[1] ; m1X ++) {</pre>
                     m1 = (m1Y-1) * cells[1] + m1X + nAtom;
for(offset = 1; offset <= 9; offset ++){
00064
00065
00066
                   m2X = m1X + iofX[offset]; shift[1] = 0.;
00067
                   if(m2X > cells[1]){
                      m2X = 1; shift[1] = region[1];
00068
00069
                   }else if(m2X == 0){
00070
                     m2X = cells[1]; shift[1] = -region[1];
00071
00072
                   m2Y = m1Y + iofY[offset]; shift[2] = 0.;
00073
                   if(m2Y > cells[2]){
                     m2Y = 1; shift[2] = region[2];
00074
00075
                    else if(m2Y == 0){
00076
                      m2Y = cells[2]; shift[2] = -region[2];
00077
00078
                   m2 = (m2Y-1) * cells[1] + m2X + nAtom;
00079
                   I = cellList[m1];
00080
                   while (I > 0) {
                      J = cellList[m2];
00081
00082
                       while (J > 0) {
                           if (m1 == m2 && J != I && (atomRadius[I] > 0. && atomRadius[J] > 0.)) {
00083
                               dr[1] = rx[I] - rx[J] - shift[1];
dr[2] = ry[I] - ry[J] - shift[2];
00084
00085
                               rr = Sqr(dr[1]) + Sqr(dr[2]);
RadiusIJ = atomRadius[I] + atomRadius[J];
00086
00087
00088
                               SqrRadiusIJ = Sqr(RadiusIJ);
                               if(rr < SqrRadiusIJ) {</pre>
00089
00090
                           r = sqrt(rr);
00091
                           ri = 1.0/r;
                                          rrinv = 1.0/rr;
00092
                                           vr[1] = vx[I] - vx[J];
vr[2] = vy[I] - vy[J];
00093
00094
00095
                           RadiusIJInv = 1.0/RadiusIJ;
                           uVal = Sqr(1.0 - r * RadiusIJInv);
fcVal = 2.0 * RadiusIJInv * (1.0 - r * RadiusIJInv) *ri;
00096
00097
00098
                                           fdVal = -gamman * (vr[1]*dr[1] + vr[2]*dr[2]) * rrinv; //disc-disc drag
00099
00100
                           f = fcVal * dr[1];
                                          fd = fdVal * dr[1];
00101
                           ax[I] += (f + fd);
00102
00103
                                          discDragx[I] += fd; //disc-disc drag
00104
00105
                           f = fcVal * dr[2];
                                          fd = fdVal * dr[2];
00106
                           ay[I] += (f + fd);
00107
                                          discDragy[I] += fd; //disc-disc drag
00108
00109
00110
                           uSum += 0.5 * uVal;
                           virSum += 0.5 * fcVal * rr;
00111
                           rfAtom += 0.5 * dr[1] * fcVal * dr[2];
00112
00113
00114
                           }else if(m1 != m2 && (atomRadius[I] > 0. && atomRadius[J] > 0.)){
                               dr[1] = rx[I] - rx[J] - shift[1];

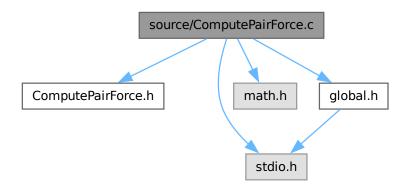
dr[2] = ry[I] - ry[J] - shift[2];
00115
00116
00117
                               rr = Sqr(dr[1]) + Sqr(dr[2]);
                               RadiusIJ = atomRadius[I] + atomRadius[J];
SqrRadiusIJ = Sqr(RadiusIJ);
00118
00119
```

```
if(rr < SqrRadiusIJ){</pre>
00121
               r = sqrt(rr);
00122
               ri = 1.0/r;
                        rrinv = 1.0/r;
00123
                        vr[1] = vx[1] - vx[J];
vr[2] = vy[1] - vy[J];
00124
00125
                RadiusIJInv = 1.0/RadiusIJ;
00126
00127
                uVal = Sqr(1.0 - r * RadiusIJInv);
00128
                fcVal = 2.0 * RadiusIJInv * (1.0 - r * RadiusIJInv) *ri;
                         fdVal = -gamman * (vr[1]*dr[1] + vr[2]*dr[2]) * rrinv; //disc-disc drag
00129
00130
00131
                f = fcVal * dr[1];
                fd = fdVal * dr[1];
ax[I] += (f + fd);
00132
00133
00134
                        discDragx[I] += fd; //disc-disc drag
00135
               f = fcVal * dr[2];
    fd = fdVal * dr[2];
ay[I] += (f + fd);
00136
00137
00138
00139
                        discDragy[I] += fd; //disc-disc drag
00140
               uSum += 0.5 * uVal;
00141
               virSum += 0.5 * fcVal * rr;
rfAtom += 0.5 * dr[1] * fcVal * dr[2];
00142
00143
00144
                  }
00145
00146
                    J = cellList[J];
00147
00148
             I = cellList[I];
00149
          }
00150
00151
00152 }
00153 }
```

3.30 source/ComputePairForce.c File Reference

```
#include "ComputePairForce.h"
#include <stdio.h>
#include <math.h>
#include "global.h"
```

Include dependency graph for ComputePairForce.c:



Functions

• void ComputePairForce (int normFlag)

3.30.1 Function Documentation

3.30.1.1 ComputePairForce()

```
void ComputePairForce (
                 int normFlag )
Definition at line 27 of file ComputePairForce.c.
00028 double dr[NDIM+1], fcVal, rr, ri, r, uVal;
00029 int n, i, j;
00030 uVal = 0.0; uSumPair = 0.0;
00031 virSumPair = 0.0; virSumPairxx = 0.0; virSumPairxy = 0.0; virSumPairxy = 0.0;
00033 for(n = 1 ; n \le nAtom ; n ++) {
00034 ax[n] = 0.0;
00035 ay[n] = 0.0;
00036 discDragx[n] = 0.0;
00037 discDragy[n] = 0.0;
00038 }
00039 for(n = 1; n <= nPairTotal; n ++) {
00040 PairID[n] = 0;
00041 Pairatom1[n] = 0;

00041 Pairatom2[n] = 0;

00042 Pairatom2[n] = 0;

00043 PairXij[n] = 0.0;

00044 PairYij[n] = 0.0;
00045 }
00046
00047
00048 \text{ Kn} = 1.0;
00049 double vr[NDIM+1], fdVal, rri;
00050 nPairActive = 0;
00051 double meff;
00052 \text{ meff} = 0.0;
00053 int atomIDi, atomIDj;
00054 //int processThisPair = 1;
00055
00056 for(i=1;i<=nAtomInterface;i++){</pre>
00057 for(j=i+1;j<=nAtomInterface;j++){
00058 atomIDi = atomIDInterface[i];
        atomIDi = atomIDInterface[i];
00059
         atomIDj = atomIDInterface[j];
00060
         if (isBonded[atomIDi][atomIDj] == 0) { //To exclude pair interaction between bonded atoms
         rr = 0.0; rri = 0.0; fcVal = 0.0; fdVal = 0.0; strech = 0.0; RadiusIJ = 0.0;
00061
00062
00063
00064
         dr[1] = rx[atomIDi] - rx[atomIDj];
         if(dr[1] >= regionH[1])
dr[1] -= region[1];
else if(dr[1] < -regionH[1])</pre>
00065
00066
00067
00068
           dr[1] += region[1];
00069
00070
         dr[2] = ry[atomIDi] - ry[atomIDj];
00071
          if(dr[2] >= regionH[2]){
00072
           dr[1] -= shearDisplacement;
            if(dr[1] < -regionH[1]) dr[1] += region[1];</pre>
00073
           dr[2] -= region[2];
00074
         }else if(dr[2] < -regionH[2]){</pre>
00075
         dr[1] += shearDisplacement;
00076
00077
            if(dr[1] >= regionH[1]) dr[1] -= region[1];
00078
          dr[2] += region[2];
00079
         }
08000
00081
         rr = Sar(dr[1]) + Sar(dr[2]);
         RadiusIJ = atomRadius[atomIDi] + atomRadius[atomIDj];
00083
         SqrRadiusIJ = Sqr(RadiusIJ);
00084
         if(rr < SqrRadiusIJ){</pre>
00085
          r = sqrt(rr);
          ri = 1.0/r;
rri = 1.0/rr;
00086
00087
          RadiusIJInv = 1.0/RadiusIJ;
00088
00089
          strech = (RadiusIJ - r);
00090
           uVal = 0.5 * Kn * Sqr(strech);
00091
           //NormFlag
00092
00093
           if(normFlag == 1){
00094
           strech = strech * RadiusIJInv;
uVal = 0.5 * Kn * RadiusIJ * Sqr(strech);
00095
00096
00097
          fcVal = Kn * strech * ri;
vr[1] = vx[atomIDi] - vx[atomIDj];
vr[2] = vy[atomIDi] - vy[atomIDj];
00098
00099
00100
00101
00102
00103
           PairID[nPairActive] = nPairActive;
00104
           Pairatom1[nPairActive] = atomIDi;
```

```
Pairatom2[nPairActive] = atomIDj;
             PairXij[nPairActive] = dr[1];
00106
00107
             PairYij[nPairActive] = dr[2];
00108
00109
00110
             if (DampFlag == 1) {
             meff = (atomMass[atomIDi]*atomMass[atomIDj])/(atomMass[atomIDi] + atomMass[atomIDj]);
00111
00112
             fdVal = -gamman * meff * (vr[1]*dr[1] + vr[2]*dr[2]) * rri; //disc-disc drag
00113
             discDragx[atomIDi] = fdVal * dr[1]; //disc-disc drag
discDragy[atomIDi] = fdVal * dr[2]; //disc-disc drag
discDragx[atomIDj] = -fdVal * dr[1]; //disc-disc drag
00114
00115
00116
             discDragy[atomIDj] = -fdVal * dr[2]; //disc-disc drag
00117
00118
            discDragx[nPairActive] = discDragx[atomIDi];
discDragy[nPairActive] = discDragy[atomIDi];
00119
00120
00121
00122
             ax[atomIDi] += (fcVal + fdVal) * dr[1];
            ay[atomIDi] += (fcVal + fdVal) * dr[2];
ax[atomIDj] += -(fcVal + fdVal) * dr[1];
ay[atomIDj] += -(fcVal + fdVal) * dr[2];
00124
00125
00126
00127
00128
00129
           //DampFlag = 2
           else if(DampFlag == 2){
00130
            discDragx[atomIDi] = -gamman * vr[1]; //disc-disc drag
discDragy[atomIDi] = -gamman * vr[2]; //disc-disc drag
discDragx[atomIDj] = -(-gamman * vr[1]); //disc-disc drag
discDragy[atomIDj] = -(-gamman * vr[2]); //disc-disc drag
00131
00132
00133
00134
00135
            discDragx[nPairActive] = discDragx[atomIDi];
discDragy[nPairActive] = discDragy[atomIDi];
00136
00137
00138
00139
            ax[atomIDi] += (fcVal * dr[1] - gamman * vr[1]);
ay[atomIDi] += (fcVal * dr[2] - gamman * vr[2]);
ax[atomIDj] += -(fcVal * dr[1] - gamman * vr[1]);
00140
00141
00143
             ay[atomIDj] += -(fcVal * dr[2] - gamman * vr[2]);
00144
00145
00146
           //DampFlag = 3. Suzanne PRL, 130, 178203 (2023) version
           else if(DampFlag == 3){
00147
00148
            //Track compression velocity
           DeltaXijNew = dr[1];
00149
00150
           DeltaYijNew = dr[2];
00151
           if(stepCount == 0) { // Initialization step
            DeltaXijOldPair[atomIDi][atomIDj] = DeltaXijNew;
DeltaYijOldPair[atomIDi][atomIDj] = DeltaYijNew;
00152
00153
00154
00155
00156
             DeltaXij = DeltaXijNew - DeltaXijOldPair[atomIDi][atomIDj];
            DeltaYij = DeltaYijNew - DeltaYijOldPair[atomIDi];
DeltaVXij = DeltaXij / deltaT;
00157
00158
             DeltaVYij = DeltaYij / deltaT;
00159
00160
00161
             // Update history for next step
00162
             DeltaXijOldPair[atomIDi][atomIDj] = DeltaXijNew;
00163
             DeltaYijOldPair[atomIDi][atomIDj] = DeltaYijNew;
00164
            discDragx[atomIDi] = -gamman * DeltaVXij; //disc-disc drag
discDragy[atomIDi] = -gamman * DeltaVYij; //disc-disc drag
discDragx[atomIDj] = -(-gamman * DeltaVXij); //disc-disc drag
00165
00166
00167
             discDragy[atomIDj] = -(-gamman * DeltaVYij); //disc-disc drag
00168
00169
            discDragx[nPairActive] = discDragx[atomIDi];
discDragy[nPairActive] = discDragy[atomIDi];
00170
00171
00172
            ax[atomIDi] += (fcVal * dr[1] - gamman * DeltaVXij);
ay[atomIDi] += (fcVal * dr[2] - gamman * DeltaVYij);
ax[atomIDj] += -(fcVal * dr[1] - gamman * DeltaVXij);
ay[atomIDj] += -(fcVal * dr[2] - gamman * DeltaVYij);
00173
00174
00175
00176
00177
00178
           //In the following, for stress/virial term (fcVal + fdVal) is used since the total pair force =
00179
        Hookean Interaction + relative velocity drag
00180
           uSumPair += 0.5 * uVal;
           virSumPair += 0.5 * (fcVal + fdVal) * rr;
virSumPairxx += 0.5 * (fcVal + fdVal) * dr[1] * dr[1];
virSumPairyy += 0.5 * (fcVal + fdVal) * dr[2] * dr[2];
virSumPairxy += 0.5 * (fcVal + fdVal) * dr[1] * dr[2];
00181
00182
00183
00184
00185
00186
           else { //Resetting the distance between two discs when they are not in contact
00187
            DeltaXijOldPair[atomIDi][atomIDj] = 0.0;
00188
             DeltaYijOldPair[atomIDi][atomIDj] = 0.0;
00189
             DeltaXijOldPair[atomIDj][atomIDi] = 0.0;
00190
             DeltaYijOldPair[atomIDj][atomIDi] = 0.0;
```

```
00191
00192 }
00193 }
00194 }
```

References atomIDInterface, atomMass, atomRadius, ax, ay, DampFlag, deltaT, DeltaVXij, DeltaVYij, DeltaXij, DeltaXij, DeltaXij, DeltaXij, DeltaYijOldPair, discDragx, discDragy, gamman, isBonded, Kn, nAtom, nAtomInterface, NDIM, nPairActive, nPairTotal, Pairatom1, Pairatom2, PairID, PairXij, PairYij, RadiusIJ, RadiusIJInv, region, regionH, rx, ry, shearDisplacement, Sqr, SqrRadiusIJ, stepCount, strech, uSumPair, virSumPairxx, virSumPairxy, virSumPairyy, vx, and vy.

Referenced by main().

Here is the caller graph for this function:



3.31 ComputePairForce.c

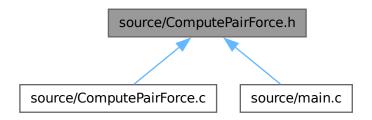
```
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00003
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        \star it under the terms of the GNU General Public License as published by
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00016
00017
        Copyright (C) 2025 Harish Charan, University of Durham, UK
00018
00019
00021 #include "ComputePairForce.h"
00022
00023 #include<stdio.h>
00024 #include<math.h>
00025 #include"global.h"
00026
00027 void ComputePairForce(int normFlag){
00028 double dr[NDIM+1], fcVal, rr, ri, r, uVal;
00029 int n, i, j;
00030 uVal = 0.0;
                     uSumPair = 0.0;
00031 virSumPair = 0.0; virSumPairxx = 0.0; virSumPairxy = 0.0; virSumPairxy = 0.0;
00032
00033 for (n = 1 ; n \le nAtom ; n ++) {
00034 ax[n] = 0.0;
00035 ay[n] = 0.0;
00036 discDragx[n] = 0.0;
       discDragy[n] = 0.0;
00037
00038 }
00039 for(n = 1; n <= nPairTotal; n ++) {
00040 PairID[n] = 0;
00041 Pairatom1[n] = 0;
00042 \quad Pairatom2[n] = 0;
00043 PairXij[n] = 0.0;
00044 PairYij[n] = 0.0;
00045 }
00046
00047
00048 \text{ Kn} = 1.0;
00049 double vr[NDIM+1], fdVal, rri;
```

```
00050 nPairActive = 0;
00051 double meff;
00052 \text{ meff} = 0.0;
00053 int atomIDi, atomIDj;
00054 //int processThisPair = 1;
00055
00056 for(i=1;i<=nAtomInterface;i++) {</pre>
00057 for(j=i+1; j<=nAtomInterface; j++) {
         atomIDi = atomIDInterface[i];
atomIDj = atomIDInterface[j];
00058
00059
         00060
00061
         rr = 0.0; rri = 0.0; fcVal = 0.0; fdVal = 0.0; strech = 0.0;
         RadiusIJ = 0.0;
00062
00063
         dr[1] = rx[atomIDi] - rx[atomIDj];
00064
         if(dr[1] >= regionH[1])
dr[1] -= region[1];
else if(dr[1] < -regionH[1])</pre>
00065
00066
00067
           dr[1] += region[1];
00068
00069
00070
         dr[2] = ry[atomIDi] - ry[atomIDj];
00071
          if(dr[2] >= regionH[2]){
00072
            dr[1] -= shearDisplacement;
           if(dr[1] < -regionH[1]) dr[1] += region[1];
dr[2] -= region[2];</pre>
00073
00074
00075
         }else if(dr[2] < -regionH[2]){</pre>
00076
          dr[1] += shearDisplacement;
00077
           if(dr[1] >= regionH[1]) dr[1] -= region[1];
00078
          dr[2] += region[2];
00079
08000
00081
         rr = Sqr(dr[1]) + Sqr(dr[2]);
00082
         RadiusIJ = atomRadius[atomIDi] + atomRadius[atomIDj];
00083
         SqrRadiusIJ = Sqr(RadiusIJ);
00084
         if(rr < SqrRadiusIJ){</pre>
00085
          r = sqrt(rr);
          ri = 1.0/r;
00086
          rri = 1.0/rr;
00088
          RadiusIJInv = 1.0/RadiusIJ;
00089
           strech = (RadiusIJ - r);
00090
          uVal = 0.5 * Kn * Sqr(strech);
00091
00092
          //NormFlag
00093
          if(normFlag == 1){
00094
            strech = strech * RadiusIJInv;
00095
            uVal = 0.5 * Kn * RadiusIJ * Sqr(strech);
00096
00097
00098
          fcVal = Kn * strech * ri;
          vr[1] = vx[atomIDi] - vx[atomIDj];
vr[2] = vy[atomIDi] - vy[atomIDj];
00099
00100
00101
          nPairActive++;
00102
00103
          PairID[nPairActive] = nPairActive;
          Pairatom1[nPairActive] = atomIDi;
Pairatom2[nPairActive] = atomIDj;
00104
00105
00106
           PairXij[nPairActive] = dr[1];
00107
          PairYij[nPairActive] = dr[2];
00108
00109
          //DampFlag = 1
          if(DampFlag == 1){
00110
          meff = (atomMass[atomIDi]*atomMass[atomIDj])/(atomMass[atomIDi] + atomMass[atomIDj]);
00111
00112
           fdVal = -gamman * meff * (vr[1]*dr[1] + vr[2]*dr[2]) * rri; //disc-disc drag
00113
          discDragx[atomIDi] = fdVal * dr[1]; //disc-disc drag
discDragy[atomIDi] = fdVal * dr[2]; //disc-disc drag
discDragx[atomIDj] = -fdVal * dr[1]; //disc-disc drag
discDragy[atomIDj] = -fdVal * dr[2]; //disc-disc drag
00114
00115
00116
00117
00118
          discDragx[nPairActive] = discDragx[atomIDi];
discDragy[nPairActive] = discDragy[atomIDi];
00119
00120
00121
00122
          ax[atomIDi] += (fcVal + fdVal) * dr[1];
00123
          ay[atomIDi] += (fcVal + fdVal) * dr[1];
ay[atomIDj] += -(fcVal + fdVal) * dr[1];
00124
00125
00126
          ay[atomIDj] += -(fcVal + fdVal) * dr[2];
00127
00128
00129
         //DampFlag = 2
         else if(DampFlag == 2){
00130
          discDragx[atomIDi] = -gamman * vr[1]; //disc-disc drag
discDragy[atomIDi] = -gamman * vr[2]; //disc-disc drag
discDragx[atomIDj] = -(-gamman * vr[1]); //disc-disc drag
00131
00132
00133
          discDragy[atomIDj] = -(-gamman * vr[2]); //disc-disc drag
00134
00135
00136
          discDragx[nPairActive] = discDragx[atomIDi];
```

```
discDragy[nPairActive] = discDragy[atomIDi];
00138
00139
            ax[atomIDi] += (fcVal * dr[1] - gamman * vr[1]);
ay[atomIDi] += (fcVal * dr[2] - gamman * vr[2]);
ax[atomIDj] += -(fcVal * dr[1] - gamman * vr[1]);
00140
00141
00142
            ay[atomIDj] += -(fcVal * dr[2] - gamman * vr[2]);
00143
00144
00145
00146
           //DampFlag = 3. Suzanne PRL, 130, 178203 (2023) version
           else if(DampFlag == 3){
00147
00148
           //{\tt Track\ compression\ velocity}
           DeltaXijNew = dr[1];
DeltaYijNew = dr[2];
00149
00150
00151
           if(stepCount == 0) { // Initialization step
            DeltaXijOldPair[atomIDi][atomIDj] = DeltaXijNew;
DeltaYijOldPair[atomIDi][atomIDj] = DeltaYijNew;
00152
00153
00154
00155
            DeltaXij = DeltaXijNew - DeltaXijOldPair[atomIDi][atomIDj];
DeltaYij = DeltaYijNew - DeltaYijOldPair[atomIDi][atomIDj];
DeltaVXij = DeltaXij / deltaT;
DeltaVYij = DeltaYij / deltaT;
00156
00157
00158
00159
00160
00161
             // Update history for next step
             DeltaXijOldPair[atomIDi][atomIDj] = DeltaXijNew;
00162
00163
             DeltaYijOldPair[atomIDi][atomIDj] = DeltaYijNew;
00164
            discDragx[atomIDi] = -gamman * DeltaVXij; //disc-disc drag
discDragy[atomIDi] = -gamman * DeltaVYij; //disc-disc drag
discDragx[atomIDj] = -(-gamman * DeltaVXij); //disc-disc drag
discDragy[atomIDj] = -(-gamman * DeltaVYij); //disc-disc drag
00165
00166
00167
00168
00169
            discDragx[nPairActive] = discDragx[atomIDi];
discDragy[nPairActive] = discDragy[atomIDi];
00170
00171
00172
            ax[atomIDi] += (fcVal * dr[1] - gamman * DeltaVXij);
ay[atomIDi] += (fcVal * dr[2] - gamman * DeltaVYij);
ax[atomIDj] += -(fcVal * dr[1] - gamman * DeltaVXij);
00173
00174
00175
00176
             ay[atomIDj] += -(fcVal * dr[2] - gamman * DeltaVYij);
00177
00178
           //In the following, for stress/virial term (fcVal + fdVal) is used since the total pair force =
00179
       Hookean Interaction + relative velocity drag
00180
           uSumPair += 0.5 * uVal;
           virSumPair += 0.5 * (fcVal + fdVal) * rr;
virSumPairxx += 0.5 * (fcVal + fdVal) * dr[1] * dr[1];
virSumPairyy += 0.5 * (fcVal + fdVal) * dr[2] * dr[2];
00181
00182
00183
           virSumPairxy += 0.5 * (fcVal + fdVal) * dr[1] * dr[2];
00184
00185
00186
           else { //Resetting the distance between two discs when they are not in contact
00187
           DeltaXijOldPair[atomIDi][atomIDj] = 0.0;
00188
            DeltaYijOldPair[atomIDi][atomIDj] = 0.0;
00189
            DeltaXijOldPair[atomIDj][atomIDi] = 0.0;
00190
            DeltaYijOldPair[atomIDj][atomIDi] = 0.0;
00191
              }
00192
00193
00194 }
00195 }
00196
00197
00198
```

3.32 source/ComputePairForce.h File Reference

This graph shows which files directly or indirectly include this file:



Functions

void ComputePairForce (int normFlag)

3.32.1 Function Documentation

3.32.1.1 ComputePairForce()

```
void ComputePairForce (
                    int normFlag )
Definition at line 27 of file ComputePairForce.c.
00028 double dr[NDIM+1], fcVal, rr, ri, r, uVal;
00029 int n, i,j;
00030 uVal = 0.0;
                          uSumPair = 0.0;
00031 virSumPair = 0.0; virSumPairxx = 0.0; virSumPairxy = 0.0; virSumPairxy = 0.0;
00033 for (n = 1 ; n \le nAtom ; n ++) {
00034 ax[n] = 0.0;
00035 ay[n] = 0.0;
00036 discDragx[n] = 0.0;
00037 discDragy[n] = 0.0;
00038 }
00039 for(n = 1; n <= nPairTotal; n ++) {
00040 PairID[n] = 0;

00041 Pairatom1[n] = 0;

00042 Pairatom2[n] = 0;

00043 PairXij[n] = 0.0;
00044 PairYij[n] = 0.0;
00045 }
00046
00047
00048 \text{ Kn} = 1.0;
00049 double vr[NDIM+1], fdVal, rri;
00050 nPairActive = 0;
00051 double meff;
00052 \text{ meff} = 0.0;
00053 int atomIDi, atomIDj;
00054 //int processThisPair = 1;
00055
00056 for(i=1;i<=nAtomInterface;i++) {</pre>
00057 for(j=i+1; j<=nAtomInterface; j++) {
00058    atomIDi = atomIDInterface[i];
00059    atomIDj = atomIDInterface[j];</pre>
          if (isBonded[atomIDi][atomIDj] == 0) { //To exclude pair interaction between bonded atoms
rr = 0.0; rri = 0.0; fcVal = 0.0; fdVal = 0.0; strech = 0.0;
RadiusIJ = 0.0;
00060
00061
00062
00063
00064
           dr[1] = rx[atomIDi] - rx[atomIDj];
          if(dr[1] >= regionH[1])
dr[1] -= region[1];
else if(dr[1] < -regionH[1])</pre>
00065
00066
00067
00068
             dr[1] += region[1];
00069
```

```
dr[2] = ry[atomIDi] - ry[atomIDj];
00071
           if(dr[2] >= regionH[2]){
             dr[1] -= shearDisplacement;
00072
00073
             if(dr[1] < -regionH[1]) dr[1] += region[1];</pre>
00074
             dr[2] -= region[2];
00075
          }else if(dr[2] < -regionH[2]){</pre>
           dr[1] += shearDisplacement;
00076
00077
             if(dr[1] >= regionH[1]) dr[1] -= region[1];
00078
           dr[2] += region[2];
00079
00080
00081
          rr = Sqr(dr[1]) + Sqr(dr[2]);
          RadiusIJ = atomRadius[atomIDi] + atomRadius[atomIDj];
00082
          SqrRadiusIJ = Sqr(RadiusIJ);
00083
00084
          if(rr < SqrRadiusIJ) {</pre>
           r = sqrt(rr);
ri = 1.0/r;
00085
00086
00087
           rri = 1.0/rr;
           RadiusIJInv = 1.0/RadiusIJ;
00089
            strech = (RadiusIJ - r);
00090
            uVal = 0.5 * Kn * Sqr(strech);
00091
00092
            //NormFlag
00093
            if(normFlag == 1){
            strech = strech * RadiusIJInv;
uVal = 0.5 * Kn * RadiusIJ * Sqr(strech);
00094
00095
00096
00097
           fcVal = Kn * strech * ri;
vr[1] = vx[atomIDi] - vx[atomIDj];
vr[2] = vy[atomIDi] - vy[atomIDj];
00098
00099
00100
00101
00102
            nPairActive++;
00103
            PairID[nPairActive] = nPairActive;
            Pairatom1[nPairActive] = atomIDi;
Pairatom2[nPairActive] = atomIDj;
00104
00105
           PairXij[nPairActive] = dr[1];
PairYij[nPairActive] = dr[2];
00106
00108
00109
            //DampFlag = 1
00110
            if(DampFlag == 1) {
            meff = (atomMass[atomIDi]*atomMass[atomIDj])/(atomMass[atomIDi] + atomMass[atomIDj]);
00111
00112
            fdVal = -qamman * meff * (vr[1]*dr[1] + vr[2]*dr[2]) * rri; //disc-disc drag
00113
            discDragx[atomIDi] = fdVal * dr[1]; //disc-disc drag
00114
            discDragy[atomIDi] = fdVal * dr[2]; //disc-disc drag
00115
           discDragx[atomIDj] = -fdVal * dr[1]; //disc-disc drag
discDragy[atomIDj] = -fdVal * dr[2]; //disc-disc drag
00116
00117
00118
00119
            discDragx[nPairActive] = discDragx[atomIDi];
            discDragy[nPairActive] = discDragy[atomIDi];
00120
00121
00122
           ax[atomIDi] += (fcVal + fdVal) * dr[1];
ay[atomIDi] += (fcVal + fdVal) * dr[2];
ax[atomIDj] += -(fcVal + fdVal) * dr[1];
ay[atomIDj] += -(fcVal + fdVal) * dr[2];
00123
00124
00125
00126
00127
00128
00129
          //DampFlag = 2
          else if(DampFlag == 2){
00130
           discDragx[atomIDi] = -gamman * vr[1]; //disc-disc drag
discDragy[atomIDi] = -gamman * vr[2]; //disc-disc drag
discDragx[atomIDj] = -(-gamman * vr[1]); //disc-disc drag
00131
00132
00133
            discDragy[atomIDj] = -(-gamman * vr[2]); //disc-disc drag
00134
00135
00136
           discDragx[nPairActive] = discDragx[atomIDi];
discDragy[nPairActive] = discDragy[atomIDi];
00137
00138
00139
           ax[atomIDi] += (fcVal * dr[1] - gamman * vr[1]);
ay[atomIDi] += (fcVal * dr[2] - gamman * vr[2]);
ax[atomIDj] += -(fcVal * dr[1] - gamman * vr[1]);
ay[atomIDj] += -(fcVal * dr[2] - gamman * vr[2]);
00140
00141
00142
00143
00144
00145
00146
          //DampFlag = 3. Suzanne PRL, 130, 178203 (2023) version
00147
           else if(DampFlag == 3){
00148
           //Track compression velocity
00149
          DeltaXijNew = dr[1];
          DeltaYijNew = dr[2];
00150
           if(stepCount == 0) { // Initialization step
00151
           DeltaXijOldPair[atomIDi][atomIDj] = DeltaXijNew;
DeltaYijOldPair[atomIDi][atomIDj] = DeltaYijNew;
00152
00153
00154
00155
00156
            DeltaXij = DeltaXijNew - DeltaXijOldPair[atomIDi][atomIDj];
```

```
DeltaYij = DeltaYijNew - DeltaYijOldPair[atomIDi][atomIDj];
00158
           DeltaVXij = DeltaXij / deltaT;
           DeltaVYij = DeltaYij / deltaT;
00159
00160
00161
           // Update history for next step
           DeltaXijOldPair[atomIDi] [atomIDj] = DeltaXijNew;
00162
           DeltaYijOldPair[atomIDi][atomIDj] = DeltaYijNew;
00163
00164
00165
           discDragx[atomIDi] = -gamman * DeltaVXij; //disc-disc drag
          discDragy[atomIDi] = -gamman * DeltaVYij; //disc-disc drag
discDragx[atomIDj] = -(-gamman * DeltaVXij); //disc-disc drag
discDragy[atomIDj] = -(-gamman * DeltaVYij); //disc-disc drag
00166
00167
00168
00169
00170
           discDragx[nPairActive] = discDragx[atomIDi];
00171
           discDragy[nPairActive] = discDragy[atomIDi];
00172
          ax[atomIDi] += (fcVal * dr[1] - gamman * DeltaVXij);
ay[atomIDi] += (fcVal * dr[2] - gamman * DeltaVYij);
ax[atomIDj] += -(fcVal * dr[1] - gamman * DeltaVXij);
00173
00174
00176
          ay[atomIDj] += -(fcVal * dr[2] - gamman * DeltaVYij);
00177
00178
         //{\rm In~the~following,~for~stress/virial~term~(fcVal~+~fdVal)}~is~used~since~the~total~pair~force=
00179
       Hookean Interaction + relative velocity drag
00180
         uSumPair += 0.5 * uVal;
         virSumPair += 0.5 * (fcVal + fdVal) * rr;
00181
00182
         virSumPairxx += 0.5 * (fcVal + fdVal) * dr[1] * dr[1];
         virSumPairyy += 0.5 * (fcVal + fdVal) * dr[2] * dr[2];
virSumPairxy += 0.5 * (fcVal + fdVal) * dr[1] * dr[2];
00183
00184
00185
00186
         else { //Resetting the distance between two discs when they are not in contact
00187
          DeltaXijOldPair[atomIDi][atomIDj] = 0.0;
00188
           DeltaYijOldPair[atomIDi][atomIDj] = 0.0;
00189
           DeltaXijOldPair[atomIDj][atomIDi] = 0.0;
00190
          DeltaYijOldPair[atomIDj][atomIDi] = 0.0;
00191
00192
00194
00195 }
```

References atomIDInterface, atomMass, atomRadius, ax, ay, DampFlag, deltaT, DeltaVXij, DeltaVYij, DeltaXij, DeltaXij, DeltaXij, DeltaXij, DeltaXij, DeltaXij, DeltaYij, DeltaYij, DeltaYijOldPair, discDragx, discDragy, gamman, isBonded, Kn, nAtom, nAtomInterface, NDIM, nPairActive, nPairTotal, Pairatom1, Pairatom2, PairID, PairXij, PairYij, RadiusIJ, RadiusIJInv, region, regionH, rx, ry, shearDisplacement, Sqr, SqrRadiusIJ, stepCount, strech, uSumPair, virSumPairxy, virSumPairxy, virSumPairyy, vx, and vy.

Referenced by main().

Here is the caller graph for this function:



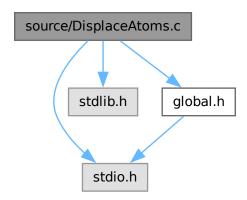
3.33 ComputePairForce.h

```
00001 #ifndef COMPUTE_PAIR_FORCE_H
00002 #define COMPUTE_PAIR_FORCE_H
00003
00004 void ComputePairForce(int normFlag);
00005
00006 #endif
```

3.34 source/DisplaceAtoms.c File Reference

```
#include <stdio.h>
#include <stdlib.h>
#include "global.h"
```

Include dependency graph for DisplaceAtoms.c:



Functions

void DisplaceAtoms ()

3.34.1 Function Documentation

3.34.1.1 DisplaceAtoms()

References DeltaX, DeltaY, molID, nAtom, rx, and ry.

Referenced by main().

Here is the caller graph for this function:



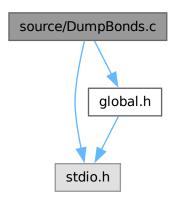
3.35 DisplaceAtoms.c

```
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00016
00017 Copyright (C) 2025 Harish Charan, University of Durham, UK
00018
00019 */
00020
00021 #include<stdio.h>
00022 #include<stdlib.h>
00023 #include"global.h"
00024
00025 void DisplaceAtoms(){
00026 int n;
          for(n = 1; n <= nAtom; n ++) {
00027
           if(molID[n] == 2) {
00028
            rx[n] += DeltaX;
ry[n] += DeltaY;
00029
00030
00031 } }
```

3.36 source/DumpBonds.c File Reference

```
#include <stdio.h>
#include "global.h"
```

Include dependency graph for DumpBonds.c:



Functions

• void DumpBonds ()

3.36.1 Function Documentation

3.36.1.1 DumpBonds()

```
void DumpBonds ( )

Definition at line 24 of file DumpBonds.c.
```

3.37 DumpBonds.c 51

```
int n;
          //Trajectory file in LAMMPS dump format for OVITO visualization fprintf(fpbond, "ITEM: TIMESTEP\n"); fprintf(fpbond, "%lf\n",timeNow);
00026
00027
00028
          fprintf(fpbond, "ITEM: NUMBER OF ENTRIES\n");
fprintf(fpbond, "%d\n", nBond);
00029
00030
           fprintf(fpbond, "ITEM: BOX BOUNDS pp ff pp\n");
          fprintf(fpbond, "%lf %lf xlo xhi\n", -regionH[1], regionH[1]);
fprintf(fpbond, "%lf %lf ylo yhi\n", -regionH[2], regionH[2]);
fprintf(fpbond, "%lf %lf zlo zhi\n", -0.1, 0.1);
00032
00033
00034
          fprintf(fpbond, "ITEM: ENTRIES BondID, BondType, atom1 atom2 BondLength BondLengthEqul nodeDragx1
00035
        \verb|nodeDragy1\n"|;
00036
00037
           for (n=1; n<=nBond; n++)</pre>
00038
            fprintf(fpbond, "%d %d %d %d %0.161f %0.161f %0.161f %0.161f\n", BondID[n], BondType[n], atom1[n],
        atom2[n],
00039
            BondLength[n], ro[n], nodeDragx[atom1[n]], nodeDragy[atom1[n]]);
00040
```

References atom1, atom2, BondID, BondLength, BondType, fpbond, nBond, nodeDragx, nodeDragy, regionH, ro, and timeNow.

Referenced by main().

Here is the caller graph for this function:

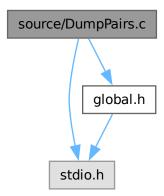


3.37 DumpBonds.c

```
00001 /*
00002
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00003
00004
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00005
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00006
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00007
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00016
00017
        Copyright (C) 2025 Harish Charan, University of Durham, UK
00018
00019
00020
00021 #include<stdio.h>
00022 #include"global.h"
00023
00024 void DumpBonds() {
00025
          //Trajectory file in LAMMPS dump format for OVITO visualization fprintf(fpbond, "ITEM: TIMESTEP\n");
00026
00027
          fprintf(fpbond, "%lf\n",timeNow);
00028
          fprintf(fpbond, "ITEM: NUMBER OF ENTRIES\n");
00029
          fprintf(fpbond, "%d\n", nBond);
00030
         iprintf(fpbond, "aa\n",nsond);
fprintf(fpbond, "ITEM: BOX BOUNDS pp ff pp\n");
fprintf(fpbond, "%lf %lf xlo xhi\n", -regionH[1], regionH[1]);
fprintf(fpbond, "%lf %lf ylo yhi\n", -regionH[2], regionH[2]);
fprintf(fpbond, "%lf %lf zlo zhi\n", -0.1, 0.1);
00031
00032
00033
00035
          fprintf(fpbond, "ITEM: ENTRIES BondID, BondType, atom1 atom2 BondLength BondLengthEqul nodeDragx1
       nodeDragy1n");
00036
00037
          for(n=1; n<=nBond; n++)
fprintf(fpbond, "%d %d %d %d %0.161f %0.161f %0.161f %0.161f \n", BondID[n], BondType[n], atoml[n],</pre>
00038
```

3.38 source/DumpPairs.c File Reference

```
#include <stdio.h>
#include "global.h"
Include dependency graph for DumpPairs.c:
```



Functions

• void DumpPairs ()

3.38.1 Function Documentation

3.38.1.1 DumpPairs()

```
void DumpPairs ( )
Definition at line 25 of file DumpPairs.c.
00025
                int n;
//Trajectory file in LAMMPS dump format for OVITO visualization
fprintf(fppair, "ITEM: TIMESTEP\n");
fprintf(fppair, "%lf\n",timeNow);
fprintf(fppair, "ITEM: NUMBER OF ENTRIES\n");
fprintf(fppair, "%d\n",nPairActive);
fprintf(fppair, "ITEM: BOX BOUNDS pp ff pp\n");
fprintf(fppair, "%lf %lf xlo xhi\n", -regionH[1], regionH[1]);
fprintf(fppair, "%lf %lf ylo yhi\n", -regionH[2], regionH[2]);
fprintf(fppair, "%lf %lf zlo zhi\n", -0.1, 0.1);
fprintf(fppair, "ITEM: ENTRIES index, atom1 atom2 xij yij discDragx1 discDragy1\n");
00026
00027
00030
00031
00032
00033
00034
00035
00036
00037
                for(n=1; n<=nPairActive; n++)
fprintf(fppair, "%d %d %0 .161f %0 .161f %0 .161f %0 .161f \n", PairID[n], Pairatom1[n],</pre>
00038
00039
00040
                   PairXij[n], PairYij[n], discDragx[n], discDragy[n]);
00041
00042
```

References discDragx, discDragy, fppair, nPairActive, Pairatom1, Pairatom2, PairID, PairXij, PairYij, regionH, and timeNow.

Referenced by main().

3.39 DumpPairs.c 53

Here is the caller graph for this function:



3.39 DumpPairs.c

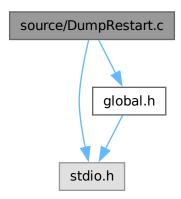
Go to the documentation of this file.

```
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00016
00017 Copyright (C) 2025 Harish Charan, University of Durham, UK
00018
00019
00020
00021
00022 #include<stdio.h>
00023 #include"global.h"
00024
 00025 void DumpPairs(){
00026
                  int n;
                       //Trajectory file in LAMMPS dump format for OVITO visualization
fprintf(fppair, "ITEM: TIMESTEP\n");
fprintf(fppair, "%lf\n",timeNow);
fprintf(fppair, "ITEM: NUMBER OF ENTRIES\n");
fprintf(fppair, "%d\n",nPairActive);
00027
00028
00029
00030
00031
                        fprintf(fppair, %u(n', hraffxctive);
fprintf(fppair, "ITEM: BOX BOUNDS pp ff pp\n");
fprintf(fppair, "%lf %lf xlo xhi\n", -regionH[1], regionH[1]);
fprintf(fppair, "%lf %lf ylo yhi\n", -regionH[2], regionH[2]);
fprintf(fppair, "%lf %lf zlo zhi\n", -0.1, 0.1);
fprintf(fppair, "ITEM: ENTRIES index, atoml atom2 xij yij discDragxl discDragyl\n");
00032
00033
00034
00035
00036
00037
                        for(n=1; n<=nPairActive; n++)</pre>
00039
                            fprintf(fppair, "%d %d %0.161f %0.161f %0.161f %0.161f \n", PairID[n], Pairatom1[n], Pairatom1[n],
                 Pairatom2[n],
00040
                          PairXij[n], PairYij[n], discDragx[n], discDragy[n]);
00041
00042
00043
00044
00045
```

3.40 source/DumpRestart.c File Reference

```
#include <stdio.h>
#include "global.h"
```

Include dependency graph for DumpRestart.c:



Functions

void DumpRestart ()

3.40.1 Function Documentation

3.40.1.1 DumpRestart()

```
void DumpRestart ( )
Definition at line 25 of file DumpRestart.c.
00025
00026
         char DUMP[256];
        FILE *fpDUMP;
00028 sprintf(DUMP, "%s.Restart", prefix);
00029 fpDUMP = fopen(DUMP, "w");
00030 if(fpDUMP == NULL) {
00031 fprintf(stderr, "Error opening file %s for writing\n", DUMP);
00032
          return:
00033 }
00034
          fprintf(fpDUMP,
fprintf(fpDUMP,
    "natom %d\n", natom);
fprintf(fpDUMP, "nBond %d\n", nBond);
fprintf(fpDUMP, "nAtomType %d\n", nAtomType);
fprintf(fpDUMP, "nBondType %d\n", nBondType);
fprintf(fpDUMP, "region[1] %0.141f\n", region[1]);
fprintf(fpDUMP, "region[2] %0.141f\n", region[2]);
00035
00036
00037
00038
00039
00040
          fprintf(fpDUMP, "region[2] %0.14lf\n", region[2]);
00041
00042
00043
00044
          fprintf(fpDUMP, "Atoms\n");
          for (n = 1; n <= nAtom; n ++)</pre>
00045
            fprintf(fpDUMP, "%d %d %d %0.21f %0.161f %0.161f %0.161f %0.161f\n", atomID[n], molID[n],
00046
       atomType[n], atomRadius[n], rx[n], ry[n], vx[n], vy[n]);
00047
00048
          fprintf(fpDUMP, "Bonds\n");
00049
          for(n=1; n<=nBond; n++)

fprintf(fpDUMP, "%d %d %d %d %0.21f %0.161f\n", BondID[n], BondType[n], atom1[n], atom2[n], kb[n],
00050
00051
        <u>ro[n]);</u>
00052
00053
          fclose(fpDUMP);
00054 }
```

References atom1, atom2, atomID, atomRadius, atomType, BondID, BondType, kb, molID, nAtom, nAtomType, nBond, nBondType, prefix, region, ro, rx, ry, timeNow, vx, and vy. Referenced by main().

3.41 DumpRestart.c 55

Here is the caller graph for this function:

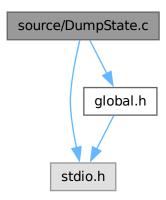


3.41 DumpRestart.c

```
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00017 Copyright (C) 2025 Harish Charan, University of Durham, UK
00018
00019
00020
00021
00022 #include <stdio.h>
00023 #include "global.h"
00024
00025 void DumpRestart() {
00026 char DUMP[256];
00027 FILE *fpDUMP;
00028 sprintf(DUMP, "%s.Restart", prefix);
00029 fpDUMP = fopen(DUMP, "w");
                 if (fpDUMP == NULL) {
00030
                   fprintf(stderr, "Error opening file %s for writing\n", DUMP);
00031
00032
                    return;
00033 }
00034
                   fprintf(fpDUMP, "timeNow %lf\n", timeNow);
fprintf(fpDUMP, "nAtom %d\n", nAtom);
fprintf(fpDUMP, "nBond %d\n", nBond);
00035
00036
00037
                   fprintf(fpDUMP, "nBond %d\n", nBond);
fprintf(fpDUMP, "nAtomType %d\n", nAtomType);
fprintf(fpDUMP, "nBondType %d\n", nBondType);
fprintf(fpDUMP, "region[1] %0.141f\n", region[1]);
fprintf(fpDUMP, "region[2] %0.141f\n", region[2]);
00039
00040
00041
00042
00043
00044
                    fprintf(fpDUMP, "Atoms\n");
00045
                    for(n = 1; n <= nAtom; n ++)</pre>
00046
                       \label{eq:fpdump} \texttt{fprintf(fpDUMP, "%d %d %d %0.21f %0.161f %0.161f %0.161f %0.161f %0.161f n", atomID[n], molID[n], molID[
              atomType[n], atomRadius[n], rx[n], ry[n], vx[n], vy[n]);
00047
00048
00049
                    fprintf(fpDUMP, "Bonds\n");
                  for(n=1; n<=nBond; n++) fprintf(fpDUMP, "%d %d %d %d %0.21f %0.161f\n", BondID[n], BondType[n], atom1[n], atom2[n], kb[n],
00050
00051
               ro[n]);
00052
00053
                    fclose(fpDUMP);
00054 }
```

3.42 source/DumpState.c File Reference

```
#include <stdio.h>
#include "global.h"
Include dependency graph for DumpState.c:
```



Functions

· void DumpState ()

3.42.1 Function Documentation

3.42.1.1 DumpState()

```
void DumpState ( )
Definition at line 25 of file DumpState.c.
00026
          char DUMP[256];
00027 FILE *fpDUMP;
00028 sprintf(DUMP, "%s.STATE", prefix);
00020 sprincr(Down, %s.STATE, 00029 fpDUMP = fopen(DUMP, "w"); 00030 if(fpDUMP == NULL) {
           fprintf(stderr, "Error opening file %s for writing\n", DUMP);
00032
00033 }
00034
           fprintf(fpDUMP, "ITEM: TIMESTEP\n");
00035
           fprintf(fpDUMP, "%lf\n",timeNow);
fprintf(fpDUMP, "ITEM: NUMBER OF ATOMS\n");
00036
           fprintf(fpDUMP, "liem: NUMBER OF ATOMS(n");
fprintf(fpDUMP, "%d\n", nAtom);
fprintf(fpDUMP, "ITEM: BOX BOUNDS pp pp pp\n");
fprintf(fpDUMP, "%lf %lf xlo xhi\n", -regionH[1], regionH[1]);
fprintf(fpDUMP, "%lf %lf ylo yhi\n", -regionH[2], regionH[2]);
fprintf(fpDUMP, "%lf %lf zlo zhi\n", -0.1, 0.1);
00038
00039
00040
00041
00042
            fprintf(fpDUMP, "ITEM: ATOMS id mol type radius x y vx vy fx fy\n");
00043
            int n;
            for (n = 1; n <= nAtom; n++) {
    fprintf(fpDUMP, "%d\t %d\t %d\t %0.21f\t %0.161f\t %0.161f\t %0.161f\t %0.161f\t %0.161f\t %0.161f\t
00045
00046
        %0.16lf\n'
00047
             atomID[n], molID[n], atomType[n], atomRadius[n], rx[n], ry[n], vx[n], vy[n], ax[n], ay[n]);
00048
00049
            fclose(fpDUMP);
00050 }
```

References atomID, atomRadius, atomType, ax, ay, moIID, nAtom, prefix, regionH, rx, ry, timeNow, vx, and vy. Referenced by main().

3.43 DumpState.c 57

Here is the caller graph for this function:



3.43 DumpState.c

Go to the documentation of this file.

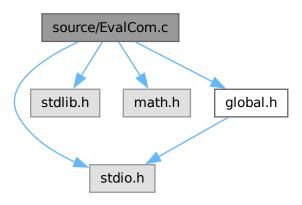
```
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00016
00017 Copyright (C) 2025 Harish Charan, University of Durham, UK
00018
00019 */
00020
00021
00022 #include <stdio.h>
00023 #include "global.h"
00024
00025 void DumpState() {
00026 char DUMP[256];
00027 FILE *fpDUMP;
00028 sprintf(DUMP, "%s.STATE", prefix);
00029 fpDUMP = fopen(DUMP, "w");
00030 if(fpDUMP == NULL) {
           fprintf(stderr, "Error opening file %s for writing\n", DUMP);
00031
00032
            return;
00033 }
00034
           fprintf(fpDUMP, "ITEM: TIMESTEP\n");
fprintf(fpDUMP, "%lf\n",timeNow);
00035
00036
           fprintf(fpDUMP, "%lf\n",timeNow);
fprintf(fpDUMP, "ITEM: NUMBER OF ATOMS\n");
fprintf(fpDUMP, "%d\n",nAtom);
fprintf(fpDUMP, "TEM: BOX BOUNDS pp pp pp\n");
fprintf(fpDUMP, "%lf %lf xlo xhi\n", -regionH[1], regionH[1]);
fprintf(fpDUMP, "%lf %lf ylo yhi\n", -regionH[2], regionH[2]);
fprintf(fpDUMP, "%lf %lf zlo zhi\n", -0.1, 0.1);
fprintf(fpDUMP, "ITEM: ATOMS id mol type radius x y vx vy fx fy\n");
int n.
00037
00039
00040
00041
00042
00043
00044
            int n;
           for (n = 1; n <= nAtom; n++) {
    fprintf(fpDUMP, "%d\t %d\t %d\t %0.21f\t %0.161f\t %0.161f\t %0.161f\t %0.161f\t %0.161f\t</pre>
00045
00046
        %0.16lf\n",
00047
             atomID[n], molID[n], atomType[n], atomRadius[n], rx[n], ry[n], vx[n], vy[n], ax[n], ay[n]);
00048
            fclose(fpDUMP);
00050 }
00051
```

3.44 source/EvalCom.c File Reference

```
#include <stdio.h>
#include <stdlib.h>
#include <math.h>
```

```
#include "global.h"
```

Include dependency graph for EvalCom.c:



Functions

• void EvalCom ()

3.44.1 Function Documentation

3.44.1.1 EvalCom()

```
void EvalCom ( )
Definition at line 27 of file EvalCom.c.
00027
00028 int n;

00029 ComX = 0.0; ComY = 0.0; ComXRatio = 0.0; ComYRatio = 0.0;

00030 TotalMass = 0.0;
00031
00032  for(n=1; n<=nAtom; n++){
00032   if(molID[n] == 2){
00034        ComX += atomMass[n] * rxUnwrap[n];
00035        ComY += atomMass[n] * ryUnwrap[n];</pre>
00036
          TotalMass += atomMass[n];
00037
           } }
00038
          ComX = ComX/TotalMass;
ComY = ComY/TotalMass;
00039
00040
00041
00042
           if(timeNow == 0.0){
          ComX0 = ComX; ComY0 = ComY;
00043
00044
00045
           ComXRatio = ComX/ComX0; ComYRatio = ComY/ComY0;
00046 }
```

References atomMass, ComX, ComX0, ComXRatio, ComY, ComY0, ComYRatio, molID, nAtom, rxUnwrap, ryUnwrap, timeNow, and TotalMass.

Referenced by main().

3.45 EvalCom.c 59

Here is the caller graph for this function:



3.45 EvalCom.c

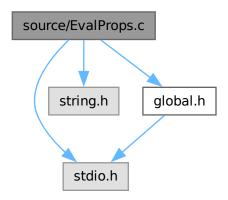
Go to the documentation of this file.

```
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00016
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00018
00019 */
00020
00021
00022 #include<stdio.h>
00023 #include<stdlib.h>
00024 #include<math.h>
00025 #include"global.h"
00027 void EvalCom() {
00028 int n;
00029 ComX = 0.0; ComY = 0.0; ComXRatio = 0.0; ComYRatio = 0.0;
00030 TotalMass = 0.0;
00031
00032
         for (n=1; n<=nAtom; n++) {</pre>
00033
         if(molID[n] == 2){
          ComX += atomMass[n] * rxUnwrap[n];
ComY += atomMass[n] * ryUnwrap[n];
00034
00035
00036
          TotalMass += atomMass[n];
00037
          } }
00038
00039
          ComX = ComX/TotalMass;
00040
          ComY = ComY/TotalMass;
00041
00042
          if(timeNow == 0.0){
00043
          ComX0 = ComX; ComY0 = ComY;
00044
00045
          ComXRatio = ComX/ComX0; ComYRatio = ComY/ComY0;
00046
00047
00048
00049
```

3.46 source/EvalProps.c File Reference

```
#include <stdio.h>
#include <string.h>
#include "global.h"
```

Include dependency graph for EvalProps.c:



Functions

· void EvalProps ()

3.46.1 Function Documentation

3.46.1.1 EvalProps()

```
void EvalProps ( )
Definition at line 26 of file EvalProps.c.
00026
00027
       real v, vv;
00028
       virSum = 0.0;
00029 vSumX = 0.0; vSumY = 0.0; vSum = 0.0; 00030 vvSum = 0.;
00031 int n;
00032
00033 for (n = 1; n \le nAtom; n++) {
00034
        vv = 0.;
00035
        // Initialize v with a default value to avoid "uninitialized" warning.
        v = 0.0;
// X direction velocity
00036
00037
        if (strcmp(solver, "Verlet") == 0) {
00038
        v = vx[n];
} else if (strcmp(solver, "LeapFrog") == 0) {
v = vx[n] - 0.5 * deltaT * ax[n];
00039
00040
00041
00042
00043
         vSum += v;
         vv += Sqr(v);
00044
         vSumX += v;
00045
         // Y direction velocity
00046
00047
          if (strcmp(solver, "Verlet") == 0) {
         v = vy[n];
} else if (strcmp(solver, "LeapFrog") == 0) {
00048
00049
         v = vy[n] - 0.5 * deltaT * ay[n];
00050
00051
00052
          vSum += v;
          vSumY += v;
00053
00054
          vv += Sqr(v);
00055
         vvSum += vv;
00056
00057
00058
        kinEnergy = 0.5 * vvSum / nAtom ;
        uSumPairPerAtom = uSumPair / nAtom ;
00059
00060 BondEnergyPerAtom = TotalBondEnergy / (0.5*nAtom); //Factor of 0.5 since each atom has one half the
     bond energy
00061 potEnergy = uSumPairPerAtom + BondEnergyPerAtom;
00062
        totEnergy = kinEnergy + potEnergy;
virSumxx = virSumPairxx + virSumBondxx;
virSumyy = virSumPairyy + virSumBondyy;
00063
00064
00065
        virSumxy = virSumPairxy + virSumBondxy;
```

3.47 EvalProps.c 61

```
00066  virSum = virSumPair + virSumBond;
00067  pressure = density * (vvSum + virSum) / (nAtom * NDIM);
00068
00069 }
```

References ax, ay, BondEnergyPerAtom, deltaT, density, kinEnergy, nAtom, NDIM, potEnergy, pressure, solver, Sqr, TotalBondEnergy, totEnergy, uSumPair, uSumPairPerAtom, virSum, virSumBond, virSumBondxx, virSumBondxy, virSumBondyy, virSumPair, virSumPairx, virSumPairxy, virSumPairyy, virSumxx, virSumxy, virSumyy, vSum, vSumX, vSumY, vvSum, vx, and vy.

Referenced by main().

Here is the caller graph for this function:



3.47 EvalProps.c

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00016
00017 Copyright (C) 2025 Harish Charan, University of Durham, UK
00018
00019
00020
00021
00022 #include <stdio.h>
00023 #include <string.h>
00024 #include "global.h"
00025
00026 void EvalProps() {
00027 real v, vv;

00028 virSum = 0.0;

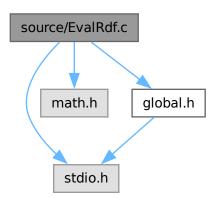
00029 vSumX = 0.0; vSumY = 0.0; vSum = 0.0;
00030 vvSum = 0.;
00031
        int n;
00033
        for (n = 1; n <= nAtom; n++) {</pre>
00034
        vv = 0.;
00035
         // Initialize v with a default value to avoid "uninitialized" warning.
00036
        v = 0.0:
00037
         // X direction velocity
         if (strcmp(solver, "Verlet") == 0) {
00038
00039
           v = vx[n];
00040
         } else if (strcmp(solver, "LeapFrog") == 0) {
00041
           v = vx[n] - 0.5 * deltaT * ax[n];
00042
00043
          vSum += v;
00044
          vv += Sqr(v);
00046
          // Y direction velocity
00047
          if (strcmp(solver, "Verlet") == 0) {
00048
          v = vy[n];
          } else if (strcmp(solver, "LeapFrog") == 0) {
v = vy[n] - 0.5 * deltaT * ay[n];
00049
00050
00051
```

```
vSum += v;
           vSumY += v;
vv += Sqr(v);
00053
00054
           vvSum += vv;
00055
00056
00057
00058
          kinEnergy = 0.5 * vvSum / nAtom ;
00059
          uSumPairPerAtom = uSumPair / nAtom ;
00060 BondEnergyPerAtom = TotalBondEnergy / (0.5*nAtom); //Factor of 0.5 since each atom has one half the
       bond energy
00061
          potEnergy = uSumPairPerAtom + BondEnergyPerAtom ;
          totEnergy = kinEnergy + potEnergy;
virSumxx = virSumPairxx + virSumBondxx;
virSumyy = virSumPairxy + virSumBondxy;
virSumxy = virSumPairxy + virSumBondxy;
00062
00063
00064
00065
          virSum = virSumPair + virSumBond;
pressure = density * (vvSum + virSum) / (nAtom * NDIM);
00066
00067
00068
00069 }
00070
```

3.48 source/EvalRdf.c File Reference

```
#include <stdio.h>
#include <math.h>
#include "global.h"
```

Include dependency graph for EvalRdf.c:



Functions

• void EvalRdf ()

3.48.1 Function Documentation

3.48.1.1 EvalRdf()

```
void EvalRdf ( )
Definition at line 26 of file EvalRdf.c.
00026
00027
         real dr[NDIM+1], deltaR, normFac, rr, rrRange;
         int j1, j2, n;
countRdf ++;
00028
00029
00030
         if(countRdf == 1) {
          for (n = 1 ; n <= sizeHistRdf ; n ++)
  histRdf[n] = 0.;</pre>
00031
00032
00033
00034
         rrRange = Sqr(rangeRdf);
         deltaR = rangeRdf / sizeHistRdf;
```

3.49 EvalRdf.c 63

```
for(j1 = 1; j1 <= nAtom - 1; j1 ++) {
  for(j2 = j1 + 1; j2 <= nAtom; j2 ++) {</pre>
00037
00038
               dr[1] = rx[j1] - rx[j2];
if(fabs(dr[1]) > regionH[1])
00039
00040
            dr[1] -= SignR(region[1], dr[1]);
00041
00042
00043
               dr[2] = ry[j1] - ry[j2];
if(fabs(dr[2]) > regionH[2])
00044
00045
            dr[2] -= SignR(region[2], dr[2]);
00046
00047
               rr = Sqr(dr[1]) + Sqr(dr[2]);
00048
00049
               if(rr < rrRange){</pre>
00050
            n = (int)(sqrt(rr)/deltaR) + 1;
00051
            histRdf[n] ++;
00052
               }
00053
00054
00055
00056
          if(countRdf == limitRdf){
00057
            normFac = region[1]*region[2] / (M_PI*Sqr(deltaR)*nAtom*nAtom*countRdf );
            for(n = 1; n <= sizeHistRdf; n ++)
histRdf[n] *= normFac/(n-0.5);
// PRINT THE RADIAL DISTRIBUTION DATA ON TO DISK FILE</pre>
00058
00059
00060
00061
            real rBin;
00062
             int n;
00063
             fprintf(fprdf,"rdf @ timeNow %lf\n", timeNow);
            for (n = 1; n <= sizeHistRdf; n ++) {
    rBin = (n - 0.5)*rangeRdf/sizeHistRdf;</pre>
00064
00065
               fprintf(fprdf, "%lf %lf\n", rBin, histRdf[n]);
00066
00067
00068
00069
00070 }
```

References countRdf, fprdf, histRdf, limitRdf, nAtom, NDIM, rangeRdf, region, regionH, rx, ry, SignR, sizeHistRdf, Sqr, and timeNow.

3.49 EvalRdf.c

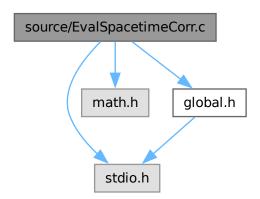
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00016
00017 Copyright (C) 2025 Harish Charan, University of Durham, UK
00018
00019 */
00020
00021
00022 #include<stdio.h>
00023 #include<math.h>
00024 #include"global.h"
00025
00026 void EvalRdf(){
00027
        real dr[NDIM+1], deltaR, normFac, rr, rrRange;
        int j1, j2, n;
countRdf ++;
00028
00029
        if(countRdf == 1) {
  for(n = 1; n <= sizeHistRdf; n ++)
   histRdf[n] = 0.;</pre>
00030
00031
00032
00033
00034
        rrRange = Sqr(rangeRdf);
        deltaR = rangeRdf / sizeHistRdf;
for(j1 = 1; j1 <= nAtom - 1; j1 ++) {</pre>
00035
00036
           for(j2 = j1 + 1; j2 <= nAtom; j2 ++) {
00037
00038
00039
             dr[1] = rx[j1] - rx[j2];
              if(fabs(dr[1]) > regionH[1])
00041
           dr[1] -= SignR(region[1], dr[1]);
00042
```

```
dr[2] = ry[j1] - ry[j2];
if(fabs(dr[2]) > regionH[2])
00044
00045
               dr[2] -= SignR(region[2], dr[2]);
00046
                   rr = Sqr(dr[1]) + Sqr(dr[2]);
00047
00048
                  if(rr < rrRange){</pre>
00050
               n = (int) (sqrt(rr)/deltaR) + 1;
00051
               histRdf[n] ++;
00052
00053
00054
00055
00056
            if(countRdf == limitRdf){
               normFac = region[1]*region[2] / (M_PI*Sqr(deltaR)*nAtom*nAtom*countRdf);
for(n = 1; n <= sizeHistRdf; n ++)
histRdf[n] *= normFac/(n-0.5);
// PRINT THE RADIAL DISTRIBUTION DATA ON TO DISK FILE</pre>
00057
00058
00059
00060
00061
               real rBin;
00062
               fprintf(fprdf, "rdf @ timeNow %lf\n", timeNow);
for(n = 1; n <= sizeHistRdf; n ++) {
  rBin = (n - 0.5) *rangeRdf/sizeHistRdf;
  fprintf(fprdf, "%lf %lf\n", rBin, histRdf[n]);</pre>
00063
00064
00065
00066
00067
00068 }
00069
00070 }
00071
```

3.50 source/EvalSpacetimeCorr.c File Reference

```
#include <stdio.h>
#include <math.h>
#include "global.h"
```

Include dependency graph for EvalSpacetimeCorr.c:



Functions

void EvalSpacetimeCorr ()

3.50.1 Function Documentation

3.50.1.1 EvalSpacetimeCorr()

```
void EvalSpacetimeCorr ( )
Definition at line 26 of file EvalSpacetimeCorr.c.
00026
```

```
real cosV=0., cosV0=0., cosV1=0., cosV2=0., sinV=0., sinV1=0., sinV2=0.;
00028
          real COSA, SINA, COSV, SINV;
          int j, m, n, nb, ni, nv;
real kMin = 2. * M_PI / region[1];
real kMax = M_PI;
00029
00030
00031
00032
          real deltaK = (kMax - kMin) / nFunCorr;
00034
          for (j = 1; j <= 2*nFunCorr; j++)</pre>
00035
            cfVal[j] = 0.;
00036
00037
          for (n = 1; n <= nAtom; n++) {</pre>
00038
            j = 1;
            COSA = cos(kMin*rx[n]);
SINA = sin(kMin*rx[n]);
00039
00040
            for (m = 1; m <= nFunCorr; m++) {
  if (m == 1) {</pre>
00041
00042
            cosV = cos(deltaK*rx[n]);
00043
00044
            sinV = sin(deltaK*rx[n]);
00045
            cosV0 = cosV;
               }else if(m == 2){
00046
00047
             cosV1 = cosV;
             sinV1 = sinV;
00048
            cosV = 2.*cosV0*cosV1-1;
sinV = 2.*cosV0*sinV1;
00049
00050
00051
               }else{
             cosV2 = cosV1;
00052
00053
            sinV2 = sinV1;
00054
             cosV1 = cosV;
00055
             sinV1 = sinV;
            cosV = 2.*cosV0*cosV1-cosV2;
00056
00057
            sinV = 2.*cosV0*sinV1-sinV2;
00058
00059
               COSV = COSA*cosV - SINA*sinV;
00060
               SINV = SINA*cosV + COSA*sinV;
               cfVal[j] += COSV;
cfVal[j+1] += SINV;
00061
00062
00063
               j += 2;
00064
00065
00066
00067
          for (nb = 1; nb <= nBuffCorr; nb++) {</pre>
00068
           indexCorr[nb] += 1;
             if (indexCorr[nb] <= 0) continue;</pre>
00069
00070
            ni = nFunCorr * (indexCorr[nb] - 1);
            if (indexCorr[nb] == 1) {
    for (j = 1; j <= 2*nFunCorr; j++)
    cfOrg[nb][j] = cfVal[j];
00071
00072
00073
00074
00075
             for (j = 1; j <= nFunCorr; j++)</pre>
00076
              spacetimeCorr[nb][ni + j] = 0.;
00078
00079
00080
             for (m = 1; m <= nFunCorr; m++) {</pre>
00081
              nv = m + ni;
00082
                spacetimeCorr[nb][nv] \ += \ cfVal[j] \ \star \ cfOrg[nb][j] \ + \ cfVal[j+1] \ \star \ cfOrg[nb][j+1]; 
               j += 2;
00084
00085
00086
00087
00088
          // ACCUMULATE SPACETIME CORRELATIONS
00089
          for (nb = 1; nb <= nBuffCorr; nb++) {</pre>
           if (indexCorr[nb] == nValCorr) {
  for (j = 1; j <= nFunCorr*nValCorr; j++)
    spacetimeCorrAv[j] += spacetimeCorr[nb][j];</pre>
00090
00091
00092
00093
              indexCorr[nb] = 0.;
00094
              countCorrAv ++;
00095
              if (countCorrAv == limitCorrAv) {
                 for (j = 1; j <= nFunCorr*nValCorr; j++)</pre>
00097
              spacetimeCorrAv[j] /= (nAtom*limitCorrAv);
                fprintf(fpdnsty, "NDIM %d\n", NDIM);
fprintf(fpdnsty, "nAtom %d\n", nAtom);
fprintf(fpdnsty, "region %lf\n", region[1]);
fprintf(fpdnsty, "region %d\n", nFunCorr);
fprintf(fpdnsty, "limitCorrAv %d\n", limitCorrAv);
00098
00099
00100
00101
                 fprintf(fpdnsty, "stepCorr %d\n", stepCorr);
fprintf(fpdnsty, "nValCorr %d\n", nValCorr);
fprintf(fpdnsty, "deltaT %lf\n", deltaT);
00103
00104
00105
00106
                 real tVal;
                 for (n = 1; n <= nValCorr; n++) {</pre>
00107
              tVal = (n-1) *stepCorr*deltaT;
00108
              fprintf (fpdnsty, "%e\t", tVal);
00109
              00110
00111
00112
00113
```

References cfOrg, cfVal, countCorrAv, deltaT, fpdnsty, indexCorr, limitCorrAv, nAtom, nBuffCorr, NDIM, nFunCorr, nValCorr, region, rx, spacetimeCorr, spacetimeCorrAv, and stepCorr.

3.51 EvalSpacetimeCorr.c

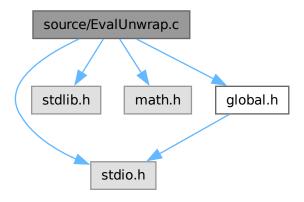
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00017
        Copyright (C) 2025 Harish Charan, University of Durham, UK
00018
00019
00020
00021
00022 #include<stdio.h>
00023 #include<math.h>
00024 #include"global.h"
00025
00026 void EvalSpacetimeCorr (){
00027    real cosV=0., cosV0=0., cosV1=0., cosV2=0., sinV=0., sinV1=0., sinV2=0.;
00028    real COSA, SINA, COSV, SINV;
         int j, m, n, nb, ni, nv;
        real kMin = 2. * M_PI / region[1];
real kMax = M_PI;
00030
00031
        real deltaK = (kMax - kMin) / nFunCorr;
00032
00033
00034
        for (j = 1; j <= 2*nFunCorr; j++)
  cfVal[j] = 0.;</pre>
00036
00037
         for (n = 1; n <= nAtom; n++) {</pre>
00038
           j = 1;
           COSA = cos(kMin*rx[n]);
SINA = sin(kMin*rx[n]);
00039
00040
00041
           for (m = 1; m <= nFunCorr; m++) {</pre>
00042
              if (m == 1) {
00043
           cosV = cos(deltaK*rx[n]);
           sinV = sin(deltaK*rx[n]);
00044
           cosV0 = cosV;
00045
00046
             }else if(m == 2){
           cosV1 = cosV;
00048
           sinV1 = sinV;
           cosV = 2.*cosV0*cosV1-1;
sinV = 2.*cosV0*sinV1;
00049
00050
             }else{
00051
00052
           cosV2 = cosV1;
           sinV2 = sinV1;
00053
           cosV1 = cosV;
00054
00055
           sinV1 = sinV;
00056
           cosV = 2.*cosV0*cosV1-cosV2;
           sinV = 2.*cosV0*sinV1-sinV2;
00057
00058
00059
              COSV = COSA*cosV - SINA*sinV;
00060
             SINV = SINA*cosV + COSA*sinV;
00061
              cfVal[j] += COSV;
00062
             cfVal[j+1] += SINV;
00063
              j += 2;
00064
00065
00066
00067
         for (nb = 1; nb <= nBuffCorr; nb++) {</pre>
00068
          indexCorr[nb] += 1;
```

```
if (indexCorr[nb] <= 0) continue;</pre>
00070
               ni = nFunCorr * (indexCorr[nb] - 1);
               if (indexCorr[nb] == 1) {
   for (j = 1; j <= 2*nFunCorr; j++)
   cfOrg[nb][j] = cfVal[j];</pre>
00071
00072
00073
00074
00075
00076
                for (j = 1; j <= nFunCorr; j++)</pre>
00077
                 spacetimeCorr[nb][ni + j] = 0.;
00078
00079
                j = 1;
               for (m = 1; m <= nFunCorr; m++) {</pre>
00080
                 nv = m + ni;
00081
00082
                  spacetimeCorr[nb][nv] += cfVal[j] * cfOrg[nb][j] + cfVal[j + 1] * cfOrg[nb][j + 1];
00083
00084
00085
00086
            }
00087
00088
            // ACCUMULATE SPACETIME CORRELATIONS
00089
            for (nb = 1; nb <= nBuffCorr; nb++) {</pre>
00090
              if (indexCorr[nb] == nValCorr) {
                 for (j = 1; j <= nFunCorr*nValCorr; j++)
    spacetimeCorrAv[j] += spacetimeCorr[nb][j];
indexCorr[nb] = 0.;</pre>
00091
00092
00093
00094
                 countCorrAv ++;
00095
                     (countCorrAv == limitCorrAv) {
00096
                    for (j = 1; j <= nFunCorr*nValCorr; j++)</pre>
                for (j = 1; j <= nFunCorr*nValCorr; j++)
spacetimeCorrAv[j] /= (nAtom*limitCorrAv);
fprintf(fpdnsty, "NDIM %d\n", NDIM);
fprintf(fpdnsty, "nAtom %d\n", nAtom);
fprintf(fpdnsty, "region %lf\n", region[1]);
fprintf(fpdnsty, "nFunCorr %d\n", nFunCorr);
fprintf(fpdnsty, "limitCorrAv %d\n", limitCorrAv);
fprintf(fpdnsty, "stepCorr %d\n", stepCorr);
fprintf(fpdnsty, "nValCorr %d\n", nValCorr);
fprintf(fpdnsty, "deltaT %lf\n", deltaT);
real tVal;</pre>
00097
00098
00099
00100
00101
00102
00103
00104
00105
                    real tVal;
00107
                    for (n = 1; n <= nValCorr; n++) {</pre>
                00108
00109
00110
00111
00112
00113
00114
00115
                 countCorrAv = 0.;
  for (j = 1; j <= nFunCorr*nValCorr; j++)
spacetimeCorrAv[j] = 0.;</pre>
00116
00117
00118
00119
00120
00121
            }
00122 }
```

3.52 source/EvalUnwrap.c File Reference

```
#include <stdio.h>
#include <stdlib.h>
#include <math.h>
#include "global.h"
```

Include dependency graph for EvalUnwrap.c:



Functions

· void EvalUnwrap ()

3.52.1 Function Documentation

3.52.1.1 EvalUnwrap()

References ImageX, ImageY, nAtom, region, rx, rxUnwrap, ry, and ryUnwrap. Referenced by main().

Here is the caller graph for this function:



3.53 EvalUnwrap.c

```
00001 /*
00002 * This file is part of Lamina.
00003 *
00004 * Lamina is free software: you can redistribute it and/or modify
00005 * it under the terms of the GNU General Public License as published by
00006 * the Free Software Foundation, either version 3 of the License, or
```

```
00007 * (at your option) any later version.
 00008 *
 00009 \,\star\, Lamina is distributed in the hope that it will be useful,
00010 * but WITHOUT ANY WARRANTY; without even the implied warranty of the war
 00014 \,\,\star\,\, You should have received a copy of the GNU General Public License
00015 \star along with Lamina. If not, see <a href="https://www.gnu.org/licenses/">https://www.gnu.org/licenses/</a>.
 00016
00017 Copyright (C) 2025 Harish Charan, University of Durham, UK
 00018
 00019 */
 00020
 00021
00022 #include <stdio.h>
00023 #include <stdlib.h>
00024 #include <math.h>
00025 #include "global.h"
 00026
 00027 void EvalUnwrap() {
00028 int n;

00029 for (n = 1; n <= nAtom; n++) {

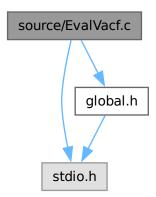
00030 rxUnwrap[n] = rx[n] + ImageX[n] * region[1];

00031 ryUnwrap[n] = ry[n] + ImageY[n] * region[2];

00032 }
 00033 }
00034
```

3.54 source/EvalVacf.c File Reference

```
#include <stdio.h>
#include "global.h"
Include dependency graph for EvalVacf.c:
```



Functions

- void AccumVacf ()
- void EvalVacf ()

3.54.1 Function Documentation

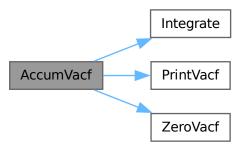
3.54.1.1 AccumVacf()

```
00029 int j, nb;
00030 for(nb = 1 ; nb <= nBuffAcf ; nb ++) {
         if(indexAcf[nb] == nValAcf;
for(j = 1; j <= nValAcf; j ++){
  viscAcfAv[j] += viscAcf[nb][j];
00031
00032
00033
00034
         indexAcf[nb] = 0;
00036
         countAcfAv ++;
         if(countAcfAv == limitAcfAv){
00037
          fac = 1./(kinEnergy*region[1]*region[2]*limitAcfAv);
00038
          viscAcfInt = fac*stepAcf*deltaT*Integrate(viscAcfAv, nValAcf);
00039
          PrintVacf();
00040
00041
          ZeroVacf();
00042 } } }
```

References countAcfAv, deltaT, indexAcf, Integrate(), kinEnergy, limitAcfAv, nBuffAcf, nValAcf, PrintVacf(), region, stepAcf, viscAcf, viscAcfAv, viscAcfInt, and ZeroVacf().

Referenced by EvalVacf().

Here is the call graph for this function:



Here is the caller graph for this function:

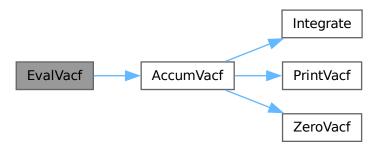


3.54.1.2 EvalVacf()

```
void EvalVacf ( )
Definition at line 26 of file EvalVacf.c.
00026
00027
           int n, nb, ni;
           double viscVec = 0.;
00028
00029
           double v[3];
00030
           for (n = 1 ; n <= nAtom ; n ++) {</pre>
             v[1] = vx[n] - 0.5*ax[n]*deltaT;
v[2] = vy[n] - 0.5*ay[n]*deltaT;
viscVec += v[1]*v[2];
00031
00032
00033
00034
00035
           viscVec += rfAtom;
           for(nb = 1 ; nb <= nBuffAcf ; nb ++) {</pre>
00036
             indexAcf[nb] ++;
if(indexAcf[nb] <= 0)continue;
if(indexAcf[nb] == 1){</pre>
00037
00038
00039
```

3.55 EvalVacf.c 71

References AccumVacf(), ax, ay, deltaT, indexAcf, nAtom, nBuffAcf, rfAtom, viscAcf, viscAcfOrg, vx, and vy. Here is the call graph for this function:



3.55 EvalVacf.c

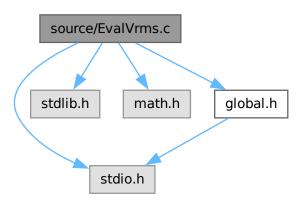
```
00002
        * This file is part of Lamina.
00003
00004 \,\, \star Lamina is free software: you can redistribute it and/or modify 00005 \,\, \star it under the terms of the GNU General Public License as published by
         * the Free Software Foundation, either version 3 of the License, or
00006
         * (at your option) any later version.
80000
00009
        \star Lamina is distributed in the hope that it will be useful,
00010 \,\, but WITHOUT ANY WARRANTY; without even the implied warranty of 00011 \,\, MERCHANTABILITY or FITNESS FOR A PARTICULAR PURPOSE. See the
00012 * GNU General Public License for more details.
00013 *
00014 * You should have received a copy of the GNU General Public License 00015 * along with Lamina. If not, see <a href="https://www.gnu.org/licenses/">https://www.gnu.org/licenses/</a>.
00016
00017 Copyright (C) 2025 Harish Charan, University of Durham, UK
00018
00019
00020
00021
00022 #include<stdio.h>
00023 #include"global.h"
00024
00025 void AccumVacf();
00026 void EvalVacf() {
00027
         int n, nb, ni;
00028
          double viscVec = 0.;
          double v[3];
for(n = 1; n <= nAtom; n ++) {</pre>
00029
00030
            v[1] = vx[n] - 0.5*ax[n]*deltaT;
v[2] = vy[n] - 0.5*ay[n]*deltaT;
00031
00032
00033
             viscVec += v[1]*v[2];
00034
00035
          viscVec += rfAtom;
00036
          for(nb = 1 ; nb <= nBuffAcf ; nb ++) {</pre>
            indexAcf[nb] ++;
if(indexAcf[nb] <= 0)continue;
if(indexAcf[nb] == 1){</pre>
00037
00038
00039
00040
                viscAcfOrg[nb] = viscVec;
00041
             ni = indexAcf[nb];
viscAcf[nb][ni] = viscAcfOrg[nb]*viscVec;
00042
00043
00044
00045
          AccumVacf();
```

00046 }

3.56 source/EvalVrms.c File Reference

```
#include <stdio.h>
#include <stdlib.h>
#include <math.h>
#include "global.h"
```

Include dependency graph for EvalVrms.c:



Functions

• void EvalVrms ()

3.56.1 Function Documentation

3.56.1.1 EvalVrms()

References nAtom, Sqr, VMeanSqr, VRootMeanSqr, VSqr, vx, and vy. Referenced by main().

3.57 EvalVrms.c 73

Here is the caller graph for this function:



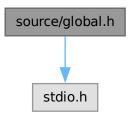
3.57 EvalVrms.c

```
00002
        * This file is part of Lamina.
00003
          \star Lamina is free software: you can redistribute it and/or modify \star it under the terms of the GNU General Public License as published by
00004
00005
          * the Free Software Foundation, either version 3 of the License, or
00007
          * (at your option) any later version.
80000
00000 * Lamina is distributed in the hope that it will be useful,
00010 * but WITHOUT ANY WARRANTY; without even the implied warranty of
00011 * MERCHANTABILITY or FITNESS FOR A PARTICULAR PURPOSE. See the
00012 * GNU General Public License for more details.
00013 *
00014 * You should have received a copy of the GNU General Public License 00015 * along with Lamina. If not, see <a href="https://www.gnu.org/licenses/">https://www.gnu.org/licenses/</a>.
00016
00017 Copyright (C) 2025 Harish Charan, University of Durham, UK
00018
00019
00020
00021
00022 #include<stdio.h>
00023 #include<stdlib.h>
00024 #include<math.h>
00025 #include"global.h"
00026
00027 void EvalVrms(){
00028
           int n;
VSqr = 0.0;
00029
           VMeanSqr = 0.0;
00030
           VRootMeanSqr = 0.0;
00031
00032
00033
            for(n = 1 ; n <= nAtom ; n ++) {</pre>
00034
           VSqr += Sqr(vx[n]) + Sqr(vy[n]);
00035
           VMeanSqr = VSqr/nAtom;
00036
00037
           VRootMeanSqr = sqrt (VMeanSqr);
00039
00040
00041
```

3.58 source/global.h File Reference

#include <stdio.h>

Include dependency graph for global.h:



This graph shows which files directly or indirectly include this file:

Macros

- #define EXTERN extern
- #define NDIM 2
- #define Sqr(x) ((x) * (x))
- #define SignR(x, y) (((y) >= 0) ? (x) : (- (x)))

Typedefs

• typedef double real

Variables

- double * rx
- double * ry
- double * vx
- double * vy
- double * ax
- double * ay
- double * speed
- double region [2+1]
- double regionH [2+1]
- double deltaT
- double timeNow
- double potEnergy
- double kinEnergy
- double totEnergy
- double density
- double pressure
- double rCut
- double kappa
- double uSum

- · double virSum
- · double svirSum
- double vSum
- double vSumX
- double vSumY
- double vvSum
- double sPotEnergy
- double sKinEnergy
- double sTotEnergy
- double sPressure
- double ssPotEnergy
- double ssKinEnergy
- double ssTotEnergy
- double ssPressure
- int initUcell [2+1]
- · int moreCycles
- int nAtom
- · int stepAvg
- int stepCount
- · int stepEquil
- · int stepLimit
- int stepTraj
- · int stepDump
- double RadiusIJ
- double SqrRadiusIJ
- · double RadiusIJInv
- int nAtomType
- int * atomType
- int * atomID
- double * atomRadius
- double * atomMass
- double TotalMass
- int nBond
- int nBondType
- int * atom1
- int * atom2
- int * BondID
- int * BondType
- double * kb
- double * ro
- double * BondEnergy
- double * BondLength
- double TotalBondEnergy
- double BondEnergyPerAtom
- double gamman
- double * discDragx
- double * discDragy
- double * nodeDragx
- double * nodeDragy
- · double strain
- double strainRate
- · double shearDisplacement
- · double shearVelocity
- double VSqr
- double VMeanSqr

- double VRootMeanSqr
- double ComX
- double ComY
- double ComX0
- · double ComY0
- · double ComXRatio
- · double ComYRatio
- double HaltCondition
- double DeltaY
- · double DeltaX
- int * ImageX
- int * ImageY
- double * rxUnwrap
- double * ryUnwrap
- int nAtomInterface
- · int nDiscInterface
- int nAtomBlock
- int * atomIDInterface
- double Kn
- double fx
- · double fy
- · double FyBylx
- double fxByfy
- · int DampFlag
- · double strech
- int dumpPairFlag
- int nPairTotal
- int nPairActive
- int * PairID
- int * Pairatom1
- int * Pairatom2
- double * PairXii
- double * PairYij
- · char solver [128]
- char xBoundary [10]
- char yBoundary [10]
- char thermo
- double * DeltaXijOld
- double * DeltaYijOld
- · double DeltaXijNew
- double DeltaYijNew
- double DeltaXij
- · double DeltaYij
- double DeltaVXij
- double DeltaVYij
- double ** DeltaXijOldPair
- double ** DeltaYijOldPair
- int * molID
- int ** isBonded
- int * cellList
- int cells [2+1]
- int rank
- int size
- int master
- double * fax

- double * fay
- · double fuSum
- · double fvirSum
- double frfAtom
- · double uSumPair
- · double uSumPairPerAtom
- · double virSumPair
- double virSumPairxx
- double virSumPairyy
- double virSumPairxy
- double virSumBond
- double virSumBondxx
- double virSumBondyy
- double virSumBondxy
- double virSumxx
- double virSumyy
- · double virSumxy
- int freezeAtomType
- double ** cfOrg
- double ** spacetimeCorr
- double * cfVal
- double * spacetimeCorrAv
- int * indexCorr
- · int countCorrAv
- int limitCorrAv
- int nBuffCorr
- int nFunCorr
- int nValCorr
- int stepCorr
- double rfAtom
- double * indexAcf
- double ** viscAcf
- double * viscAcfOrg
- double * viscAcfAv
- double viscAcfInt
- int nValAcf
- int nBuffAcf
- int stepAcf
- · int countAcfAv
- · int limitAcfAv
- double * histRdf
- · double rangeRdf
- · int countRdf
- int limitRdf
- int sizeHistRdf
- int stepRdf
- char * prefix
- char result [250]
- FILE * fpresult
- char xyz [256]
- FILE * fpxyz
- char bond [256]
- FILE * fpbond
- char dump [256]
- FILE * fpdump

- char dnsty [256]
- FILE * fpdnsty
- char visc [256]
- FILE * fpvisc
- char rdf [256]
- FILE * fprdf
- char vrms [256]FILE * fpvrms
- · char stress [256]
- FILE * fpstress
- char momentum [256]
- FILE * fpmomentum
- char com [256]
- FILE * fpcom
- char pair [256]
- FILE * fppair

3.58.1 Macro Definition Documentation

3.58.1.1 EXTERN

```
#define EXTERN extern

Definition at line 8 of file global.h.
```

3.58.1.2 NDIM

#define NDIM 2

Definition at line 13 of file global.h.

Referenced by ComputeBondForce(), ComputeForcesCells(), ComputePairForce(), EvalProps(), EvalRdf(), EvalSpacetimeCorr(), and LeapfrogStep().

3.58.1.3 SignR

Definition at line 15 of file global.h.

Referenced by EvalRdf(), and LeapfrogStep().

3.58.1.4 Sqr

```
#define Sqr( x ) ((x) * (x))
```

Definition at line 14 of file global.h.

Referenced by AccumProps(), BrownianStep(), ComputeBondForce(), ComputeForcesCells(), ComputePairForce(), EvalProps(), EvalRdf(), EvalVrms(), and LeapfrogStep().

3.58.2 Typedef Documentation

3.58.2.1 real

typedef double real

Definition at line 11 of file global.h.

3.58.3 Variable Documentation

3.58.3.1 atom1

```
int* atom1 [extern]
```

Referenced by Close(), ComputeBondForce(), DumpBonds(), DumpRestart(), and Init().

3.58.3.2 atom2

int * atom2

Definition at line 34 of file global.h.

Referenced by Close(), ComputeBondForce(), DumpBonds(), DumpRestart(), and Init().

3.58.3.3 atomID

int* atomID [extern]

Referenced by Close(), DumpRestart(), DumpState(), Init(), and Trajectory().

3.58.3.4 atomIDInterface

int* atomIDInterface [extern]

Referenced by Close(), ComputePairForce(), and Init().

3.58.3.5 atomMass

double* atomMass [extern]

Referenced by Close(), ComputePairForce(), EvalCom(), and Init().

3.58.3.6 atomRadius

double* atomRadius [extern]

Referenced by ApplyBoundaryCond(), Close(), ComputeForcesCells(), ComputePairForce(), DumpRestart(), DumpState(), Init(), and Trajectory().

3.58.3.7 atomType

int* atomType [extern]

Referenced by ApplyDrivingForce(), Close(), DumpRestart(), DumpState(), Init(), Trajectory(), and VelocityVerletStep().

3.58.3.8 ax

double * ax

Definition at line 17 of file global.h.

Referenced by ApplyDrivingForce(), ApplyForce(), ApplyViscous(), BrownianStep(), Close(), ComputeBondForce(), ComputeForcesCells(), ComputePairForce(), DumpState(), EvalProps(), EvalVacf(), Init(), LeapfrogStep(), Trajectory(), and VelocityVerletStep().

3.58.3.9 av

double * ay

Definition at line 17 of file global.h.

Referenced by ApplyDrivingForce(), ApplyForce(), ApplyViscous(), BrownianStep(), Close(), ComputeBondForce(), ComputeForcesCells(), ComputePairForce(), DumpState(), EvalProps(), EvalVacf(), Init(), LeapfrogStep(), Trajectory(), and VelocityVerletStep().

3.58.3.10 bond

char bond[256] [extern]

Referenced by main().

3.58.3.11 BondEnergy

double* BondEnergy [extern]

Referenced by ComputeBondForce(), and Init().

3.58.3.12 BondEnergyPerAtom

double BondEnergyPerAtom

Definition at line 38 of file global.h.

Referenced by EvalProps(), and PrintSummary().

3.58.3.13 BondID

int* BondID [extern]
Referenced by Close(), DumpBonds(), DumpRestart(), and Init().

3.58.3.14 BondLength

double * BondLength

Definition at line 37 of file global.h.

Referenced by ComputeBondForce(), DumpBonds(), and Init().

3.58.3.15 BondType

int * BondType
Definition at line 35 of file global.h.
Referenced by Close(), DumpBonds(), DumpRestart(), and Init().

3.58.3.16 cellList

int* cellList [extern]
Referenced by Close(), ComputeForcesCells(), and Init().

3.58.3.17 cells

int cells[2+1]

Definition at line 78 of file global.h.

Referenced by ComputeForcesCells(), and Init().

3.58.3.18 cfOrg

double** cfOrg [extern]
Referenced by AllocArrays(), Close(), and EvalSpacetimeCorr().

3.58.3.19 cfVal

double * cfVal

Definition at line 89 of file global.h.

Referenced by AllocArrays(), Close(), and EvalSpacetimeCorr().

3.58.3.20 com

char com[256] [extern] Referenced by main().

3.58.3.21 ComX

double ComX [extern]
Referenced by EvalCom(), and PrintCom().

3.58.3.22 ComX0

double ComX0

Definition at line 44 of file global.h.

Referenced by EvalCom().

3.58.3.23 ComXRatio

double ComXRatio

Definition at line 44 of file global.h.

Referenced by EvalCom().

3.58.3.24 ComY

double ComY

Definition at line 44 of file global.h.

Referenced by EvalCom(), and PrintCom().

3.58.3.25 ComY0

double ComY0

Definition at line 44 of file global.h.

Referenced by EvalCom().

3.58.3.26 ComYRatio

double ComYRatio

Definition at line 44 of file global.h.

Referenced by EvalCom().

3.58.3.27 countAcfAv

int countAcfAv

Definition at line 95 of file global.h.

Referenced by AccumVacf(), and ZeroVacf().

3.58.3.28 countCorrAv

int countCorrAv

Definition at line 90 of file global.h.

Referenced by EvalSpacetimeCorr(), and SetupJob().

3.58.3.29 countRdf

int countRdf [extern]

Referenced by EvalRdf(), and SetupJob().

3.58.3.30 DampFlag

int DampFlag [extern]

Referenced by ComputeBondForce(), ComputePairForce(), and Init().

3.58.3.31 deltaT

double deltaT

Definition at line 20 of file global.h.

Referenced by AccumVacf(), BrownianStep(), ComputeBondForce(), ComputePairForce(), EvalProps(), EvalSpacetimeCorr(), EvalVacf(), Init(), LeapfrogStep(), main(), PrintVacf(), and VelocityVerletStep().

3.58.3.32 DeltaVXij

double DeltaVXij

Definition at line 70 of file global.h.

Referenced by ComputeBondForce(), and ComputePairForce().

3.58.3.33 DeltaVYij

double DeltaVYij

Definition at line 70 of file global.h.

Referenced by ComputeBondForce(), and ComputePairForce().

3.58.3.34 DeltaX

double DeltaX

Definition at line 46 of file global.h.

Referenced by DisplaceAtoms(), and Init().

3.58.3.35 DeltaXij

double DeltaXij [extern]

Referenced by ComputeBondForce(), and ComputePairForce().

3.58.3.36 DeltaXijNew

double DeltaXijNew [extern]

Referenced by ComputeBondForce(), and ComputePairForce().

3.58.3.37 DeltaXijOld

double* DeltaXijOld [extern]

Referenced by Close(), ComputeBondForce(), and Init().

3.58.3.38 DeltaXijOldPair

double** DeltaXijOldPair [extern]

Referenced by Close(), ComputePairForce(), and Init().

3.58.3.39 DeltaY

double DeltaY [extern]

Referenced by DisplaceAtoms(), and Init().

3.58.3.40 DeltaYij

double DeltaYij

Definition at line 70 of file global.h.

Referenced by ComputeBondForce(), and ComputePairForce().

3.58.3.41 DeltaYijNew

double DeltaYijNew

Definition at line 69 of file global.h.

Referenced by ComputeBondForce(), and ComputePairForce().

3.58.3.42 DeltaYijOld

double * DeltaYijOld

Definition at line 68 of file global.h.

Referenced by Close(), ComputeBondForce(), and Init().

3.58.3.43 DeltaYijOldPair

double ** DeltaYijOldPair

Definition at line 71 of file global.h.

Referenced by Close(), ComputePairForce(), and Init().

3.58.3.44 density

```
double density
```

Definition at line 21 of file global.h.

Referenced by EvalProps(), and Init().

3.58.3.45 discDragx

```
double* discDragx [extern]
```

Referenced by ComputeForcesCells(), ComputePairForce(), DumpPairs(), and Init().

3.58.3.46 discDragy

```
double * discDragy
```

Definition at line 40 of file global.h.

Referenced by ComputeForcesCells(), ComputePairForce(), DumpPairs(), and Init().

3.58.3.47 dnsty

```
char dnsty[256] [extern]
```

3.58.3.48 dump

```
char dump[256] [extern]
```

3.58.3.49 dumpPairFlag

```
int dumpPairFlag [extern]
```

3.58.3.50 fax

```
double* fax [extern]
```

Referenced by Close(), and Init().

3.58.3.51 fay

```
double * fay
```

Definition at line 80 of file global.h.

Referenced by Close(), and Init().

3.58.3.52 fpbond

```
FILE* fpbond [extern]
```

Referenced by DumpBonds(), and main().

3.58.3.53 fpcom

```
FILE* fpcom [extern]
```

Referenced by Init(), main(), and PrintCom().

3.58.3.54 fpdnsty

```
FILE* fpdnsty [extern]
```

Referenced by EvalSpacetimeCorr().

3.58.3.55 fpdump

```
FILE* fpdump [extern]
```

3.58.3.56 fpmomentum

```
FILE* fpmomentum [extern]
Referenced by PrintMomentum().
```

3.58.3.57 fppair

```
FILE* fppair [extern]
Referenced by DumpPairs(), and main().
```

3.58.3.58 fprdf

```
FILE* fprdf [extern]
Referenced by EvalRdf().
```

3.58.3.59 fpresult

```
FILE* fpresult [extern]

Referenced by ApplyBoundaryCond(), HaltConditionCheck(), Init(), main(), and PrintSummary().
```

3.58.3.60 fpstress

```
FILE* fpstress [extern]
Referenced by PrintStress().
```

3.58.3.61 fpvisc

```
FILE* fpvisc [extern]
Referenced by PrintVacf().
```

3.58.3.62 fpvrms

```
FILE* fpvrms [extern]
Referenced by Init(), main(), and PrintVrms().
```

3.58.3.63 fpxyz

```
FILE* fpxyz [extern]
Referenced by main(), and Trajectory().
```

3.58.3.64 freezeAtomType

```
int freezeAtomType [extern]
Referenced by Init(), and VelocityVerletStep().
```

3.58.3.65 frfAtom

```
double frfAtom
```

Definition at line 80 of file global.h.

3.58.3.66 fuSum

double fuSum

Definition at line 80 of file global.h.

3.58.3.67 fvirSum

```
double fvirSum
```

Definition at line 80 of file global.h.

3.58.3.68 fx

```
double fx [extern]
Referenced by ApplyForce().
```

3.58.3.69 fxByfy

double fxByfy

Definition at line 52 of file global.h.

Referenced by ApplyForce(), and Init().

3.58.3.70 fy

double fy

Definition at line 52 of file global.h.

Referenced by ApplyForce().

3.58.3.71 FyBylx

double FyBylx

Definition at line 52 of file global.h.

Referenced by ApplyForce(), and Init().

3.58.3.72 gamman

double gamman [extern]

Referenced by ComputeBondForce(), ComputeForcesCells(), ComputePairForce(), and Init().

3.58.3.73 HaltCondition

double HaltCondition [extern]

Referenced by HaltConditionCheck(), and Init().

3.58.3.74 histRdf

double* histRdf [extern]

Referenced by AllocArrays(), and EvalRdf().

3.58.3.75 ImageX

int* ImageX [extern]

Referenced by Close(), EvalUnwrap(), Init(), and VelocityVerletStep().

3.58.3.76 ImageY

int * ImageY

Definition at line 47 of file global.h.

Referenced by Close(), EvalUnwrap(), Init(), and VelocityVerletStep().

3.58.3.77 indexAcf

double* indexAcf [extern]

Referenced by AccumVacf(), AllocArrays(), Close(), EvalVacf(), and InitVacf().

3.58.3.78 indexCorr

int* indexCorr [extern]

Referenced by AllocArrays(), Close(), EvalSpacetimeCorr(), and SetupJob().

3.58.3.79 initUcell

```
int initUcell[2+1] [extern]
```

3.58.3.80 isBonded

```
int** isBonded [extern]
```

Referenced by Close(), ComputePairForce(), and Init().

3.58.3.81 kappa

double kappa

Definition at line 21 of file global.h.

Referenced by Init(), and LeapfrogStep().

3.58.3.82 kb

double* kb [extern]

Referenced by Close(), ComputeBondForce(), DumpRestart(), and Init().

3.58.3.83 kinEnergy

double kinEnergy

Definition at line 20 of file global.h.

Referenced by AccumProps(), AccumVacf(), EvalProps(), and PrintSummary().

3.58.3.84 Kn

double Kn [extern]

Referenced by ComputePairForce().

3.58.3.85 limitAcfAv

int limitAcfAv

Definition at line 95 of file global.h.

Referenced by AccumVacf(), and Init().

3.58.3.86 limitCorrAv

int limitCorrAv

Definition at line 90 of file global.h.

Referenced by EvalSpacetimeCorr(), and Init().

3.58.3.87 limitRdf

int limitRdf

Definition at line 99 of file global.h.

Referenced by EvalRdf(), and Init().

3.58.3.88 master

int master

Definition at line 79 of file global.h.

3.58.3.89 molID

int* molID [extern]

Referenced by ApplyForce(), Close(), DisplaceAtoms(), DumpRestart(), DumpState(), EvalCom(), Init(), and Trajectory().

3.58.3.90 momentum

char momentum[256] [extern]

3.58.3.91 moreCycles

int moreCycles

Definition at line 24 of file global.h.

Referenced by main().

3.58.3.92 nAtom

int nAtom

Definition at line 24 of file global.h.

Referenced by ApplyBoundaryCond(), ApplyDrivingForce(), ApplyForce(), ApplyLeesEdwardsBoundaryCond(), ApplyShear(), ApplyViscous(), BrownianStep(), Close(), ComputeBondForce(), ComputeForcesCells(), ComputePairForce(), DisplaceAtoms(), DumpRestart(), DumpState(), EvalCom(), EvalProps(), EvalRdf(), EvalSpacetimeCorr(), EvalUnwrap(), EvalVacf(), EvalVrms(), Init(), LeapfrogStep(), Trajectory(), and VelocityVerletStep().

3.58.3.93 nAtomBlock

int nAtomBlock

Definition at line 49 of file global.h.

Referenced by ApplyForce(), and Init().

3.58.3.94 nAtomInterface

int nAtomInterface [extern]
Referenced by ComputePairForce(), and Init().

3.58.3.95 nAtomType

int nAtomType [extern]

Referenced by DumpRestart(), and Init().

3.58.3.96 nBond

int nBond [extern]

Referenced by ComputeBondForce(), DumpBonds(), DumpRestart(), and Init().

3.58.3.97 nBondType

int nBondType

Definition at line 33 of file global.h.

Referenced by DumpRestart(), and Init().

3.58.3.98 nBuffAcf

int nBuffAcf

Definition at line 95 of file global.h.

Referenced by AccumVacf(), AllocArrays(), Close(), EvalVacf(), Init(), and InitVacf().

3.58.3.99 nBuffCorr

int nBuffCorr

Definition at line 90 of file global.h.

Referenced by AllocArrays(), Close(), EvalSpacetimeCorr(), Init(), and SetupJob().

3.58.3.100 nDiscInterface

int nDiscInterface

Definition at line 49 of file global.h.

Referenced by Init().

3.58.3.101 nFunCorr

int nFunCorr

Definition at line 90 of file global.h.

Referenced by AllocArrays(), EvalSpacetimeCorr(), Init(), and SetupJob().

3.58.3.102 nodeDragx

double * nodeDragx

Definition at line 40 of file global.h.

Referenced by ComputeBondForce(), DumpBonds(), and Init().

3.58.3.103 nodeDragy

double * nodeDragy

Definition at line 40 of file global.h.

Referenced by ComputeBondForce(), DumpBonds(), and Init().

3.58.3.104 nPairActive

int nPairActive

Definition at line 58 of file global.h.

Referenced by ComputePairForce(), and DumpPairs().

3.58.3.105 nPairTotal

int nPairTotal [extern]

Referenced by ComputePairForce(), and Init().

3.58.3.106 nValAcf

int nValAcf [extern]

Referenced by AccumVacf(), AllocArrays(), Init(), InitVacf(), PrintVacf(), and ZeroVacf().

3.58.3.107 nValCorr

int nValCorr

Definition at line 90 of file global.h.

Referenced by AllocArrays(), EvalSpacetimeCorr(), Init(), and SetupJob().

3.58.3.108 pair

char pair[256] [extern]

Referenced by main().

3.58.3.109 Pairatom1

int * Pairatom1

Definition at line 59 of file global.h.

Referenced by Close(), ComputePairForce(), DumpPairs(), and Init().

3.58.3.110 Pairatom2

int * Pairatom2

Definition at line 59 of file global.h.

Referenced by Close(), ComputePairForce(), DumpPairs(), and Init().

3.58.3.111 PairID

int* PairID [extern]

Referenced by Close(), ComputePairForce(), DumpPairs(), and Init().

3.58.3.112 PairXij

double* PairXij [extern]

Referenced by Close(), ComputePairForce(), DumpPairs(), and Init().

3.58.3.113 PairYij

double * PairYij

Definition at line 60 of file global.h.

Referenced by Close(), ComputePairForce(), DumpPairs(), and Init().

3.58.3.114 potEnergy

double potEnergy

Definition at line 20 of file global.h.

Referenced by AccumProps(), EvalProps(), and PrintSummary().

3.58.3.115 prefix

char* prefix [extern]

Definition at line 13 of file main.c.

Referenced by DumpRestart(), and DumpState().

3.58.3.116 pressure

double pressure

Definition at line 21 of file global.h.

Referenced by AccumProps(), EvalProps(), PrintStress(), and PrintSummary().

3.58.3.117 RadiusIJ

double RadiusIJ [extern]

Referenced by ComputeForcesCells(), and ComputePairForce().

3.58.3.118 RadiusIJInv

double RadiusIJInv

Definition at line 26 of file global.h.

Referenced by ComputeForcesCells(), and ComputePairForce().

3.58.3.119 rangeRdf

double rangeRdf

Definition at line 98 of file global.h.

Referenced by EvalRdf(), and Init().

3.58.3.120 rank

int rank [extern]

Referenced by ComputeForcesCells().

3.58.3.121 rCut

double rCut

Definition at line 21 of file global.h.

Referenced by Init(), and LeapfrogStep().

3.58.3.122 rdf

char rdf[256] [extern]

3.58.3.123 region

```
double region[2+1] [extern]
```

Referenced by AccumVacf(), ApplyBoundaryCond(), ApplyLeesEdwardsBoundaryCond(), ComputeBondForce(), ComputeForcesCells(), ComputePairForce(), DumpRestart(), EvalRdf(), EvalSpacetimeCorr(), EvalUnwrap(), Init(), LeapfrogStep(), and VelocityVerletStep().

3.58.3.124 regionH

double regionH[2+1]

Definition at line 20 of file global.h.

Referenced by ApplyBoundaryCond(), ApplyForce(), ApplyLeesEdwardsBoundaryCond(), ComputeBondForce(), ComputeForcesCells(), ComputePairForce(), DumpBonds(), DumpPairs(), DumpState(), EvalRdf(), Init(), LeapfrogStep(), Trajectory(), and VelocityVerletStep().

3.58.3.125 result

```
char result[250] [extern] Referenced by main().
```

3.58.3.126 rfAtom

double rfAtom [extern]

Referenced by ComputeForcesCells(), and EvalVacf().

3.58.3.127 ro

double * ro

Definition at line 36 of file global.h.

Referenced by Close(), ComputeBondForce(), DumpBonds(), DumpRestart(), and Init().

3.58.3.128 rx

```
double* rx [extern]
```

Referenced by ApplyBoundaryCond(), ApplyLeesEdwardsBoundaryCond(), ApplyShear(), BrownianStep(), Close(), ComputeBondForce(), ComputeForcesCells(), ComputePairForce(), DisplaceAtoms(), DumpRestart(), DumpState(), EvalRdf(), EvalSpacetimeCorr(), EvalUnwrap(), Init(), LeapfrogStep(), Trajectory(), and VelocityVerletStep().

3.58.3.129 rxUnwrap

```
double* rxUnwrap [extern]
```

Referenced by Close(), EvalCom(), EvalUnwrap(), and Init().

3.58.3.130 ry

double * ry

Definition at line 17 of file global.h.

Referenced by ApplyBoundaryCond(), ApplyLeesEdwardsBoundaryCond(), ApplyShear(), BrownianStep(), Close(), ComputeBondForce(), ComputeForcesCells(), ComputePairForce(), DisplaceAtoms(), DumpRestart(), DumpState(), EvalRdf(), EvalUnwrap(), Init(), LeapfrogStep(), Trajectory(), and VelocityVerletStep().

3.58.3.131 ryUnwrap

```
double * ryUnwrap
Definition at line 48 of file global.h.
Referenced by Close(), EvalCom(), EvalUnwrap(), and Init().
```

3.58.3.132 shearDisplacement

```
double shearDisplacement [extern] Referenced by ApplyLeesEdwardsBoundaryCond(), ComputeBondForce(), ComputePairForce(), and Init().
```

3.58.3.133 shearVelocity

```
double shearVelocity
Definition at line 42 of file global.h.
Referenced by Init().
```

3.58.3.134 size

int size

Definition at line 79 of file global.h. Referenced by ComputeForcesCells().

3.58.3.135 sizeHistRdf

int sizeHistRdf
Definition at line 99 of file global.h.
Referenced by AllocArrays(), EvalRdf(), and Init().

3.58.3.136 sKinEnergy

double skinEnergy
Definition at line 21 of file global.h.
Referenced by AccumProps().

3.58.3.137 solver

```
char solver[128] [extern]
Referenced by EvalProps(), and Init().
```

3.58.3.138 spacetimeCorr

```
double ** spacetimeCorr

Definition at line 89 of file global.h.

Referenced by AllocArrays(), Close(), and EvalSpacetimeCorr().
```

3.58.3.139 spacetimeCorrAv

```
double * spacetimeCorrAv

Definition at line 89 of file global.h.

Referenced by AllocArrays(), Close(), EvalSpacetimeCorr(), and SetupJob().
```

3.58.3.140 speed

```
double* speed [extern]
Referenced by Close(), and Init().
```

3.58.3.141 sPotEnergy

double sPotEnergy

Definition at line 21 of file global.h.

Referenced by AccumProps().

3.58.3.142 sPressure

double sPressure

Definition at line 22 of file global.h.

Referenced by AccumProps().

3.58.3.143 SqrRadiusIJ

double SqrRadiusIJ

Definition at line 26 of file global.h.

Referenced by ComputeForcesCells(), and ComputePairForce().

3.58.3.144 ssKinEnergy

double ssKinEnergy

Definition at line 22 of file global.h.

Referenced by AccumProps().

3.58.3.145 ssPotEnergy

double ssPotEnergy

Definition at line 22 of file global.h.

Referenced by AccumProps().

3.58.3.146 ssPressure

double ssPressure

Definition at line 22 of file global.h.

Referenced by AccumProps().

3.58.3.147 ssTotEnergy

double ssTotEnergy

Definition at line 22 of file global.h.

Referenced by AccumProps().

3.58.3.148 stepAcf

int stepAcf

Definition at line 95 of file global.h.

Referenced by AccumVacf(), Init(), and PrintVacf().

3.58.3.149 stepAvg

int stepAvg

Definition at line 24 of file global.h.

Referenced by AccumProps(), Init(), and main().

3.58.3.150 stepCorr

int stepCorr

Definition at line 90 of file global.h.

Referenced by EvalSpacetimeCorr(), and Init().

3.58.3.151 stepCount

int stepCount

Definition at line 24 of file global.h.

Referenced by BrownianStep(), ComputeBondForce(), ComputePairForce(), HaltConditionCheck(), LeapfrogStep(), main(), and SetupJob().

3.58.3.152 stepDump

int stepDump

Definition at line 24 of file global.h.

Referenced by Init(), and main().

3.58.3.153 stepEquil

int stepEquil

Definition at line 24 of file global.h.

Referenced by BrownianStep(), Init(), and LeapfrogStep().

3.58.3.154 stepLimit

int stepLimit

Definition at line 24 of file global.h.

Referenced by Init(), and main().

3.58.3.155 stepRdf

int stepRdf

Definition at line 99 of file global.h.

Referenced by Init().

3.58.3.156 stepTraj

int stepTraj

Definition at line 24 of file global.h.

Referenced by Init(), and main().

3.58.3.157 sTotEnergy

double sTotEnergy

Definition at line 22 of file global.h.

Referenced by AccumProps().

3.58.3.158 strain

double strain [extern]

Referenced by ApplyShear(), and Init().

3.58.3.159 strainRate

double strainRate

Definition at line 41 of file global.h.

Referenced by Init().

3.58.3.160 strech

double strech [extern]

Referenced by ComputeBondForce(), and ComputePairForce().

3.58.3.161 stress

char stress[256] [extern]

3.58.3.162 svirSum

double svirSum

Definition at line 21 of file global.h.

Referenced by AccumProps().

3.58.3.163 thermo

char thermo [extern]

Referenced by Init(), and LeapfrogStep().

3.58.3.164 timeNow

double timeNow

Definition at line 20 of file global.h.

Referenced by DumpBonds(), DumpPairs(), DumpRestart(), DumpState(), EvalCom(), EvalRdf(), Init(), main(), PrintCom(), PrintMomentum(), PrintStress(), PrintSummary(), PrintVrms(), and Trajectory().

3.58.3.165 TotalBondEnergy

double TotalBondEnergy [extern]

Referenced by ComputeBondForce(), and EvalProps().

3.58.3.166 TotalMass

double TotalMass

Definition at line 31 of file global.h.

Referenced by EvalCom().

3.58.3.167 totEnergy

 $\verb"double totEnergy"$

Definition at line 20 of file global.h.

Referenced by AccumProps(), EvalProps(), and PrintSummary().

3.58.3.168 uSum

double uSum

Definition at line 21 of file global.h.

Referenced by ComputeForcesCells().

3.58.3.169 uSumPair

double uSumPair [extern]

Referenced by ComputePairForce(), and EvalProps().

3.58.3.170 uSumPairPerAtom

double uSumPairPerAtom

Definition at line 83 of file global.h.

Referenced by EvalProps(), and PrintSummary().

3.58.3.171 virSum

double virSum

Definition at line 21 of file global.h.

Referenced by AccumProps(), ComputeForcesCells(), EvalProps(), and PrintSummary().

3.58.3.172 virSumBond

double virSumBond [extern]

Referenced by ComputeBondForce(), and EvalProps().

3.58.3.173 virSumBondxx

double virSumBondxx

Definition at line 84 of file global.h.

Referenced by ComputeBondForce(), and EvalProps().

3.58.3.174 virSumBondxy

double virSumBondxy

Definition at line 84 of file global.h.

Referenced by ComputeBondForce(), and EvalProps().

3.58.3.175 virSumBondyy

double virSumBondyy

Definition at line 84 of file global.h.

Referenced by ComputeBondForce(), and EvalProps().

3.58.3.176 virSumPair

double virSumPair

Definition at line 83 of file global.h.

Referenced by ComputePairForce(), and EvalProps().

3.58.3.177 virSumPairxx

double virSumPairxx

Definition at line 83 of file global.h.

Referenced by ComputePairForce(), and EvalProps().

3.58.3.178 virSumPairxy

double virSumPairxy

Definition at line 83 of file global.h.

Referenced by ComputePairForce(), and EvalProps().

3.58.3.179 virSumPairyy

double virSumPairyy

Definition at line 83 of file global.h.

Referenced by ComputePairForce(), and EvalProps().

3.58.3.180 virSumxx

double virSumxx [extern]

Referenced by EvalProps(), and PrintStress().

3.58.3.181 virSumxy

double virSumxy

Definition at line 85 of file global.h.

Referenced by EvalProps(), and PrintStress().

3.58.3.182 virSumyy

double virSumyy

Definition at line 85 of file global.h.

Referenced by EvalProps(), and PrintStress().

3.58.3.183 visc

char visc[256] [extern]

3.58.3.184 viscAcf

double ** viscAcf

Definition at line 94 of file global.h.

Referenced by AccumVacf(), AllocArrays(), Close(), and EvalVacf().

3.58.3.185 viscAcfAv

double * viscAcfAv

Definition at line 94 of file global.h.

Referenced by AccumVacf(), AllocArrays(), Close(), PrintVacf(), and ZeroVacf().

3.58.3.186 viscAcfInt

double viscAcfInt

Definition at line 94 of file global.h.

Referenced by AccumVacf(), and PrintVacf().

3.58.3.187 viscAcfOrg

double * viscAcfOrg

Definition at line 94 of file global.h.

Referenced by AllocArrays(), Close(), and EvalVacf().

3.58.3.188 VMeanSqr

double VMeanSqr

Definition at line 43 of file global.h.

Referenced by EvalVrms().

3.58.3.189 vrms

char vrms[256] [extern]

Referenced by main().

3.58.3.190 VRootMeanSqr

double VRootMeanSqr

Definition at line 43 of file global.h.

Referenced by EvalVrms(), main(), and PrintVrms().

3.58.3.191 VSqr

double VSqr [extern]

Referenced by EvalVrms().

3.58.3.192 vSum

double vSum

Definition at line 21 of file global.h.

Referenced by EvalProps(), and PrintSummary().

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3.58.3.193 vSumX

```
double vSumX
```

Definition at line 21 of file global.h.

Referenced by EvalProps(), and PrintMomentum().

3.58.3.194 vSumY

```
double vSumY
```

Definition at line 21 of file global.h.

Referenced by EvalProps(), and PrintMomentum().

3.58.3.195 vvSum

```
double vvSum
```

Definition at line 21 of file global.h.

Referenced by EvalProps(), and LeapfrogStep().

3.58.3.196 vx

```
double * vx
```

Definition at line 17 of file global.h.

Referenced by ApplyBoundaryCond(), ApplyDrivingForce(), ApplyViscous(), BrownianStep(), Close(), ComputeBondForce(), ComputeForcesCells(), ComputePairForce(), DumpRestart(), DumpState(), EvalProps(), EvalVacf(), EvalVrms(), Init(), LeapfrogStep(), Trajectory(), and VelocityVerletStep().

3.58.3.197 vy

```
double * vy
```

Definition at line 17 of file global.h.

Referenced by ApplyBoundaryCond(), ApplyDrivingForce(), ApplyViscous(), BrownianStep(), Close(), ComputeBondForce(), ComputeForcesCells(), ComputePairForce(), DumpRestart(), DumpState(), EvalProps(), EvalVacf(), EvalVrms(), Init(), LeapfrogStep(), Trajectory(), and VelocityVerletStep().

3.58.3.198 xBoundary

```
char xBoundary[10] [extern]
```

Referenced by ApplyBoundaryCond(), and Init().

3.58.3.199 xyz

```
char xyz[256] [extern] Referenced by main().
```

3.58.3.200 yBoundary

```
char yBoundary[10]
```

Definition at line 64 of file global.h.

Referenced by ApplyBoundaryCond(), and Init().

3.59 global.h

Go to the documentation of this file.

```
00001 #ifndef GLOBAL_H
00002 #define GLOBAL_H
00003 #include <stdio.h> // Required for FILE*
00005 #ifdef DEFINE_GLOBALS
00006 #define EXTERN
00007 #else
00008 #define EXTERN extern
00009 #endif
```

```
00011 typedef double real;
00012
00013 #define NDTM 2
00014 #define Sqr(x) ((x) * (x))
00015 #define SignR(x, y) (((y) >= 0) ? (x) : (- (x)))
00017 EXTERN double *rx, *ry, *vx, *vy, *ax, *ay;
00018 EXTERN double *speed;
00019
00020 EXTERN double region[NDIM+1], regionH[NDIM+1], deltaT, timeNow, potEnergy, kinEnergy, totEnergy, 00021 density, pressure, rCut, kappa, uSum, virSum, svirSum, vSum, vSumX, vSumY, vvSum, sPotEnergy,
      sKinEnergy,
00022 sTotEnergy, sPressure, ssPotEnergy, ssKinEnergy, ssTotEnergy, ssPressure;
00023
00024 EXTERN int
                      initUcell[NDIM+1], moreCycles, nAtom, stepAvg, stepCount, stepEquil, stepLimit,
      stepTraj, stepDump;
00025
00026 EXTERN double RadiusIJ, SqrRadiusIJ, RadiusIJInv;
00027 EXTERN int
                      nAtomType;
00028 EXTERN int
                       *atomType;
                       *atomID;
00029 EXTERN int
00030 EXTERN double *atomRadius;
00031 EXTERN double *atomMass, TotalMass;
00032
00033 EXTERN int
                      nBond, nBondType;
                      *atom1, *atom2;
00034 EXTERN int
00035 EXTERN int
                      *BondID, *BondType;
00036 EXTERN double
                      *kb, *ro;
00037 EXTERN double *BondEnergy, *BondLength;
00038 EXTERN double
                      TotalBondEnergy, BondEnergyPerAtom;
00039 EXTERN double
                      gamman;
00040 EXTERN double
                      *discDragx, *discDragy, *nodeDragx, *nodeDragy;
00041 EXTERN double
                      strain, strainRate;
00042 EXTERN double
                      shearDisplacement, shearVelocity;
                      VSqr, VMeanSqr, VRootMeanSqr;
ComX, ComY, ComX0, ComY0, ComXRatio, ComYRatio;
00043 EXTERN double
00044 EXTERN double
00045 EXTERN double
                      HaltCondition;
00046 EXTERN double DeltaY, DeltaX;
00047 EXTERN int
                      *ImageX, *ImageY;
00048 EXTERN double
                      *rxUnwrap, *ryUnwrap;
00049 EXTERN int
                      nAtomInterface, nDiscInterface, nAtomBlock;
00050 EXTERN int.
                       *atomIDInterface:
00051 EXTERN double Kn;
00052 EXTERN double fx, fy, FyBylx, fxByfy;
00053 EXTERN int
                      DampFlag;
00054 EXTERN double strech:
00055
00056 //For dumping the pair interaction data
00057 EXTERN int dumpPairFlag;
00058 EXTERN int
                      nPairTotal, nPairActive;
00059 EXTERN int
                      *PairID, *Pairatom1, *Pairatom2;
00060 EXTERN double *PairXij, *PairYij;
00061
00062
                   solver[128];
xBoundary[10], yBoundary[10];
00063 EXTERN char
00064 EXTERN char
00065 EXTERN char
                       thermo:
00066
00067 //For damping as in PRL, 130, 178203 (2023)
00068 EXTERN double *DeltaXijOld, *DeltaYijOld; 00069 EXTERN double DeltaXijNew, DeltaYijNew;
00070 EXTERN double DeltaXij, DeltaYij, DeltaVXij, DeltaVYij;
00071 EXTERN double **DeltaXijOldPair, **DeltaYijOldPair;
00072
00073 //For molecule-ID as per LAMMPS, helpful!
00074 EXTERN int *molID;
00075 EXTERN int
                     **isBonded:
00077 //Following three for MPI only
00078 EXTERN int *cellList, cells[NDIM+1];
00079 EXTERN int
                      rank, size, master;
00080 EXTERN double *fax, *fay, fuSum, fvirSum, frfAtom;
00081
00082 //For thermodynamic properties
00083 EXTERN double uSumPair, uSumPairPerAtom, virSumPair, virSumPairxx, virSumPairyy, virSumPairxy;
00084 EXTERN double virSumBond, virSumBondxx, virSumBondyy, virSumBondxy;
00085 EXTERN double virSumxx, virSumyy, virSumxy;
00086 EXTERN int
                      freezeAtomType;
00087
00088 // Spacetime Correlations
00089 EXTERN double **cfOrg, **spacetimeCorr, *cfVal, *spacetimeCorrAv;
                      *indexCorr, countCorrAv, limitCorrAv, nBuffCorr, nFunCorr, nValCorr, stepCorr;
00090 EXTERN int
00091
00092 // Viscosity
00093 EXTERN double rfAtom, frfAtom;
00094 EXTERN double *indexAcf, **viscAcfOrg, *viscAcfAv, viscAcfInt;
```

```
00095 EXTERN int
                       nValAcf, nBuffAcf, stepAcf, countAcfAv, limitAcfAv;
00096
00097 // Radial distribution function
00098 EXTERN double *histRdf, rangeRdf;
00099 EXTERN int countRdf, limitRdf, sizeHistRdf, stepRdf;
00100
00102 // Output files prefixes
00103 EXTERN char
00104
00105 EXTERN char
                      result[250];
                    *fpresult;
00106 EXTERN FILE
00107
00108 EXTERN char
                      xyz[256];
00109 EXTERN FILE
                      *fpxyz;
00110
00111 EXTERN char
                      bond[256];
00112 EXTERN FILE
                      *fpbond;
00115 EXTERN char
                      dump[256];
00116 EXTERN FILE
                      *fpdump;
00117
00118 EXTERN char
                      dnstv[256];
00119 EXTERN FILE
                      *fpdnsty;
00120
00121 EXTERN char
                      visc[256];
00122 EXTERN FILE
                      *fpvisc;
00123
00124 EXTERN char
                      rdf[2561;
00125 EXTERN FILE
                      *fprdf;
00126
00127 EXTERN char
                      vrms[256];
00128 EXTERN FILE
                      *fpvrms;
00129
00130 EXTERN char
                      stress[256];
00131 EXTERN FILE
                      *fpstress;
00133 EXTERN char
                      momentum[256];
00134 EXTERN FILE
                      *fpmomentum;
00135
00136 EXTERN char
                      com[256];
00137 EXTERN FILE
                      *fpcom;
00138
00139 EXTERN char
                      pair[256];
00140 EXTERN FILE
00141
00142 #endif // GLOBALEXTERN_H
```

3.60 source/globalExtern.h File Reference

Macros

- #define NDIM 2
- #define Sqr(x) ((x) * (x))
- #define SignR(x, y) (((y) \geq = 0) ? (x) : (- (x)))

Typedefs

· typedef double real

Variables

- double * rx
- double * ry
- double * vx
- double * vv
- double * ax
- double * ay
- double * speed
- double region [2+1]
- double regionH [2+1]
- · double deltaT

- double timeNow
- double potEnergy
- · double kinEnergy
- double totEnergy
- · double density
- · double pressure
- double rCut
- · double kappa
- double uSum
- · double virSum
- double svirSum
- double vSum
- double vSumX
- double vSumY
- double vvSum
- double sPotEnergy
- · double sKinEnergy
- double sTotEnergy
- double sPressure
- double ssPotEnergy
- double ssKinEnergy
- double ssTotEnergy
- double ssPressure
- int initUcell [2+1]
- · int moreCycles
- int nAtom
- int stepAvg
- · int stepCount
- · int stepEquil
- · int stepLimit
- int stepTraj
- int stepDump
- double RadiusIJ
- · double SqrRadiusIJ
- double RadiusIJInv
- int nAtomType
- int * atomType
- int * atomID
- double * atomRadius
- double * atomMass
- double TotalMass
- int nBond
- int nBondType
- int * atom1
- int * atom2
- int * BondID
- int * BondType
- double * kb
- double * ro
- double * BondEnergy
- double * BondLength
- double TotalBondEnergy
- double BondEnergyPerAtom
- double gamman
- double * discDragx

- double * discDragy
- double * nodeDragx
- double * nodeDragy
- · double strain
- · double strainRate
- · double shearDisplacement
- · double shearVelocity
- · double VSqr
- double VMeanSqr
- double VRootMeanSqr
- double ComX
- double ComY
- double ComX0
- double ComY0
- double ComXRatio
- double ComYRatio
- · double HaltCondition
- · double DeltaY
- double DeltaX
- int * ImageX
- int * ImageY
- double * rxUnwrap
- double * ryUnwrap
- int nAtomInterface
- · int nDiscInterface
- int nAtomBlock
- int * atomIDInterface
- double Kn
- double fx
- · double fy
- · double FyBylx
- · double fxByfy
- · int DampFlag
- · double strech
- · int dumpPairFlag
- int nPairTotal
- int nPairActive
- int * PairID
- int * Pairatom1
- int * Pairatom2
- double * PairXii
- double * PairYij
- · char solver [128]
- char xBoundary [10]
- char yBoundary [10]
- double * DeltaXijOld
- double * DeltaYijOld
- double DeltaXijNew
- · double DeltaYijNew
- double DeltaXij
- double DeltaYij
- · double DeltaVXij
- double DeltaVYij
- double ** DeltaXijOldPairdouble ** DeltaYijOldPair

- int * molID
- int ** isBonded
- int * cellList
- int cells [2+1]
- · int rank
- int size
- · int master
- double * fax
- double * fay
- · double fuSum
- double fvirSum
- double frfAtom
- double uSumPair
- double uSumPairPerAtom
- double virSumPair
- double virSumPairxx
- double virSumPairyy
- double virSumPairxy
- double virSumBond
- double virSumBondxx
- double virSumBondyy
- · double virSumBondxy
- double virSumxx
- · double virSumyy
- · double virSumxy
- int freezeAtomType
- double ** cfOrg
- double ** spacetimeCorr
- double * cfVal
- double * spacetimeCorrAv
- int * indexCorr
- · int countCorrAv
- int limitCorrAv
- · int nBuffCorr
- int nFunCorr
- int nValCorr
- int stepCorr
- double rfAtom
- double * indexAcf
- double ** viscAcf
- double * viscAcfOrg
- double * viscAcfAv
- double viscAcfInt
- int nValAcf
- int nBuffAcf
- · int stepAcf
- int countAcfAv
- · int limitAcfAv
- double * histRdf
- · double rangeRdf
- · int countRdf
- int limitRdf
- int sizeHistRdf
- int stepRdf
- char * prefix

- · char result [250]
- FILE * fpresult
- char xyz [256]
- FILE * fpxyz
- char bond [256]
- FILE * fpbond
- char dump [256]
- FILE * fpdump
- char dnsty [256]
- FILE * fpdnsty
- char visc [256]
- FILE * fpvisc
- char rdf [256]
- FILE * fprdf
- char vrms [256]
- FILE * fpvrms
- char stress [256]
- FILE * fpstress
- char momentum [256]
- FILE * fpmomentum
- char com [256]
- FILE * fpcom
- char pair [256]
- FILE * fppair

3.60.1 Macro Definition Documentation

3.60.1.1 NDIM

```
#define NDIM 2
```

Definition at line 6 of file globalExtern.h.

3.60.1.2 SignR

Definition at line 8 of file globalExtern.h.

3.60.1.3 Sqr

```
 \begin{tabular}{ll} \#define & Sqr ( & x & ) & ((x) & * & (x)) \\ \hline Definition & at line & 7 & of file & global Extern.h. \\ \end{tabular}
```

3.60.2 Typedef Documentation

3.60.2.1 real

```
typedef double real
```

Definition at line 4 of file globalExtern.h.

3.60.3 Variable Documentation

3.60.3.1 atom1

```
int* atom1 [extern]
```

3.60.3.2 atom2

int * atom2

Definition at line 27 of file globalExtern.h.

3.60.3.3 atomID

int* atomID [extern]

3.60.3.4 atomIDInterface

int* atomIDInterface [extern]

3.60.3.5 atomMass

double* atomMass [extern]

3.60.3.6 atomRadius

double* atomRadius [extern]

3.60.3.7 atomType

int* atomType [extern]

3.60.3.8 ax

double * ax

Definition at line 10 of file globalExtern.h.

3.60.3.9 ay

double * ay

Definition at line 10 of file globalExtern.h.

3.60.3.10 bond

char bond[256] [extern]

3.60.3.11 BondEnergy

double* BondEnergy [extern]

3.60.3.12 BondEnergyPerAtom

double BondEnergyPerAtom

Definition at line 31 of file globalExtern.h.

3.60.3.13 BondID

int* BondID [extern]

3.60.3.14 BondLength

double * BondLength

Definition at line 30 of file globalExtern.h.

3.60.3.15 BondType

int * BondType

Definition at line 28 of file globalExtern.h.

3.60.3.16 cellList

int* cellList [extern]

3.60.3.17 cells

int cells[2+1]

Definition at line 70 of file globalExtern.h.

3.60.3.18 cfOrg

double** cfOrg [extern]

3.60.3.19 cfVal

double * cfVal

Definition at line 81 of file globalExtern.h.

3.60.3.20 com

char com[256] [extern]

3.60.3.21 ComX

double ComX [extern]

3.60.3.22 ComX0

double ComX0

Definition at line 37 of file globalExtern.h.

3.60.3.23 ComXRatio

double ComXRatio

Definition at line 37 of file globalExtern.h.

3.60.3.24 ComY

double ComY

Definition at line 37 of file globalExtern.h.

3.60.3.25 ComY0

double ComY0

Definition at line 37 of file globalExtern.h.

3.60.3.26 ComYRatio

double ComYRatio

Definition at line 37 of file globalExtern.h.

3.60.3.27 countAcfAv

int countAcfAv

Definition at line 87 of file globalExtern.h.

3.60.3.28 countCorrAv

int countCorrAv

Definition at line 82 of file globalExtern.h.

3.60.3.29 countRdf

int countRdf [extern]

3.60.3.30 DampFlag

int DampFlag [extern]

3.60.3.31 deltaT

double deltaT

Definition at line 13 of file globalExtern.h.

3.60.3.32 DeltaVXij

double DeltaVXij

Definition at line 62 of file globalExtern.h.

3.60.3.33 DeltaVYij

double DeltaVYij

Definition at line 62 of file globalExtern.h.

3.60.3.34 DeltaX

double DeltaX

Definition at line 39 of file globalExtern.h.

3.60.3.35 DeltaXij

double DeltaXij [extern]

3.60.3.36 DeltaXijNew

double DeltaXijNew [extern]

3.60.3.37 DeltaXijOld

double* DeltaXijOld [extern]

3.60.3.38 DeltaXijOldPair

double** DeltaXijOldPair [extern]

3.60.3.39 DeltaY

double DeltaY [extern]

3.60.3.40 DeltaYij

double DeltaYij

Definition at line 62 of file globalExtern.h.

3.60.3.41 DeltaYijNew

double DeltaYijNew

Definition at line 61 of file globalExtern.h.

3.60.3.42 DeltaYijOld

double * DeltaYijOld

Definition at line 60 of file globalExtern.h.

3.60.3.43 DeltaYijOldPair

double ** DeltaYijOldPair

Definition at line 63 of file globalExtern.h.

3.60.3.44 density

double density

Definition at line 14 of file globalExtern.h.

3.60.3.45 discDragx

double* discDragx [extern]

3.60.3.46 discDragy

double * discDragy

Definition at line 33 of file globalExtern.h.

3.60.3.47 dnsty

char dnsty[256] [extern]

3.60.3.48 dump

char dump[256] [extern]

3.60.3.49 dumpPairFlag

int dumpPairFlag [extern]

3.60.3.50 fax

double* fax [extern]

3.60.3.51 fay

double * fav

Definition at line 72 of file globalExtern.h.

3.60.3.52 fpbond

FILE* fpbond [extern]

3.60.3.53 fpcom

FILE* fpcom [extern]

3.60.3.54 fpdnsty

FILE* fpdnsty [extern]

3.60.3.55 fpdump

FILE* fpdump [extern]

3.60.3.56 fpmomentum

FILE* fpmomentum [extern]

3.60.3.57 fppair

FILE* fppair [extern]

3.60.3.58 fprdf

FILE* fprdf [extern]

3.60.3.59 fpresult

FILE* fpresult [extern]

3.60.3.60 fpstress

FILE* fpstress [extern]

3.60.3.61 fpvisc

FILE* fpvisc [extern]

3.60.3.62 fpvrms

FILE* fpvrms [extern]

3.60.3.63 fpxyz

FILE* fpxyz [extern]

3.60.3.64 freezeAtomType

int freezeAtomType [extern]

3.60.3.65 frfAtom

double frfAtom

Definition at line 72 of file globalExtern.h.

3.60.3.66 fuSum

double fuSum

Definition at line 72 of file globalExtern.h.

3.60.3.67 fvirSum

double fvirSum

Definition at line 72 of file globalExtern.h.

3.60.3.68 fx

double fx [extern]

3.60.3.69 fxByfy

double fxByfy

Definition at line 45 of file globalExtern.h.

3.60.3.70 fy

double fy

Definition at line 45 of file globalExtern.h.

3.60.3.71 FyBylx

double FyBylx

Definition at line 45 of file globalExtern.h.

3.60.3.72 gamman

double gamman [extern]

3.60.3.73 HaltCondition

double HaltCondition [extern]

3.60.3.74 histRdf

double* histRdf [extern]

3.60.3.75 ImageX

int* ImageX [extern]

3.60.3.76 ImageY

int * ImageY

Definition at line 40 of file globalExtern.h.

3.60.3.77 indexAcf

double* indexAcf [extern]

3.60.3.78 indexCorr

int* indexCorr [extern]

3.60.3.79 initUcell

int initUcell[2+1] [extern]

3.60.3.80 isBonded

int** isBonded [extern]

3.60.3.81 kappa

double kappa

Definition at line 14 of file globalExtern.h.

3.60.3.82 kb

double* kb [extern]

3.60.3.83 kinEnergy

double kinEnergy

Definition at line 13 of file globalExtern.h.

3.60.3.84 Kn

double Kn [extern]

3.60.3.85 limitAcfAv

int limitAcfAv

Definition at line 87 of file globalExtern.h.

3.60.3.86 limitCorrAv

int limitCorrAv

Definition at line 82 of file globalExtern.h.

3.60.3.87 limitRdf

int limitRdf

Definition at line 91 of file globalExtern.h.

3.60.3.88 master

int master

Definition at line 71 of file globalExtern.h.

3.60.3.89 molID

int* molID [extern]

3.60.3.90 momentum

char momentum[256] [extern]

3.60.3.91 moreCycles

int moreCycles

Definition at line 17 of file globalExtern.h.

3.60.3.92 nAtom

int nAtom

Definition at line 17 of file globalExtern.h.

3.60.3.93 nAtomBlock

int nAtomBlock

Definition at line 42 of file globalExtern.h.

3.60.3.94 nAtomInterface

int nAtomInterface [extern]

3.60.3.95 nAtomType

int nAtomType [extern]

3.60.3.96 nBond

int nBond [extern]

3.60.3.97 nBondType

int nBondType

Definition at line 26 of file globalExtern.h.

3.60.3.98 nBuffAcf

int nBuffAcf

Definition at line 87 of file globalExtern.h.

3.60.3.99 nBuffCorr

int nBuffCorr

Definition at line 82 of file globalExtern.h.

3.60.3.100 nDiscInterface

int nDiscInterface

Definition at line 42 of file globalExtern.h.

3.60.3.101 nFunCorr

int nFunCorr

Definition at line 82 of file globalExtern.h.

3.60.3.102 nodeDragx

double * nodeDragx

Definition at line 33 of file globalExtern.h.

3.60.3.103 nodeDragy

double * nodeDragy

Definition at line 33 of file globalExtern.h.

3.60.3.104 nPairActive

int nPairActive

Definition at line 51 of file globalExtern.h.

3.60.3.105 nPairTotal

int nPairTotal [extern]

3.60.3.106 nValAcf

int nValAcf [extern]

3.60.3.107 nValCorr

int nValCorr

Definition at line 82 of file globalExtern.h.

3.60.3.108 pair

char pair[256] [extern]

3.60.3.109 Pairatom1

int * Pairatom1

Definition at line 52 of file globalExtern.h.

3.60.3.110 Pairatom2

int * Pairatom2

Definition at line 52 of file globalExtern.h.

3.60.3.111 PairID

int* PairID [extern]

3.60.3.112 PairXij

double* PairXij [extern]

3.60.3.113 PairYij

double * PairYij

Definition at line 53 of file globalExtern.h.

3.60.3.114 potEnergy

double potEnergy

Definition at line 13 of file globalExtern.h.

3.60.3.115 prefix

char* prefix [extern]

Definition at line 13 of file main.c.

Referenced by main().

3.60.3.116 pressure

double pressure

Definition at line 14 of file globalExtern.h.

3.60.3.117 RadiusIJ

double RadiusIJ [extern]

3.60.3.118 RadiuslJInv

double RadiusIJInv

Definition at line 19 of file globalExtern.h.

3.60.3.119 rangeRdf

double rangeRdf

Definition at line 90 of file globalExtern.h.

3.60.3.120 rank

int rank [extern]

3.60.3.121 rCut

double rCut

Definition at line 14 of file globalExtern.h.

3.60.3.122 rdf

char rdf[256] [extern]

3.60.3.123 region

double region[2+1] [extern]

3.60.3.124 regionH

double regionH[2+1]

Definition at line 13 of file globalExtern.h.

3.60.3.125 result

char result[250] [extern]

3.60.3.126 rfAtom

double rfAtom [extern]

3.60.3.127 ro

double * ro

Definition at line 29 of file globalExtern.h.

3.60.3.128 rx

double* rx [extern]

3.60.3.129 rxUnwrap

double* rxUnwrap [extern]

3.60.3.130 ry

double * ry

Definition at line 10 of file globalExtern.h.

3.60.3.131 ryUnwrap

double * ryUnwrap

Definition at line 41 of file globalExtern.h.

3.60.3.132 shearDisplacement

double shearDisplacement [extern]

3.60.3.133 shearVelocity

double shearVelocity

Definition at line 35 of file globalExtern.h.

3.60.3.134 size

int size

Definition at line 71 of file globalExtern.h.

3.60.3.135 sizeHistRdf

int sizeHistRdf

Definition at line 91 of file globalExtern.h.

3.60.3.136 sKinEnergy

double sKinEnergy

Definition at line 14 of file globalExtern.h.

3.60.3.137 solver

char solver[128] [extern]

3.60.3.138 spacetimeCorr

double ** spacetimeCorr

Definition at line 81 of file globalExtern.h.

3.60.3.139 spacetimeCorrAv

double * spacetimeCorrAv

Definition at line 81 of file globalExtern.h.

3.60.3.140 speed

double* speed [extern]

3.60.3.141 sPotEnergy

double sPotEnergy

Definition at line 14 of file globalExtern.h.

3.60.3.142 sPressure

double sPressure

Definition at line 15 of file globalExtern.h.

3.60.3.143 SqrRadiusIJ

double SqrRadiusIJ

Definition at line 19 of file globalExtern.h.

3.60.3.144 ssKinEnergy

double ssKinEnergy

Definition at line 15 of file globalExtern.h.

3.60.3.145 ssPotEnergy

double ssPotEnergy

Definition at line 15 of file globalExtern.h.

3.60.3.146 ssPressure

double ssPressure

Definition at line 15 of file globalExtern.h.

3.60.3.147 ssTotEnergy

double ssTotEnergy

Definition at line 15 of file globalExtern.h.

3.60.3.148 stepAcf

int stepAcf

Definition at line 87 of file globalExtern.h.

3.60.3.149 stepAvg

int stepAvg

Definition at line 17 of file globalExtern.h.

3.60.3.150 stepCorr

int stepCorr

Definition at line 82 of file globalExtern.h.

3.60.3.151 stepCount

int stepCount

Definition at line 17 of file globalExtern.h.

3.60.3.152 stepDump

int stepDump

Definition at line 17 of file globalExtern.h.

3.60.3.153 stepEquil

int stepEquil

Definition at line 17 of file globalExtern.h.

3.60.3.154 stepLimit

int stepLimit

Definition at line 17 of file globalExtern.h.

3.60.3.155 stepRdf

int stepRdf

Definition at line 91 of file globalExtern.h.

3.60.3.156 stepTraj

 $\verb"int stepTraj"$

Definition at line 17 of file globalExtern.h.

3.60.3.157 sTotEnergy

 $\verb"double sTotEnergy"$

Definition at line 15 of file globalExtern.h.

3.60.3.158 strain

double strain [extern]

3.60.3.159 strainRate

double strainRate

Definition at line 34 of file globalExtern.h.

3.60.3.160 strech

double strech [extern]

3.60.3.161 stress

char stress[256] [extern]

3.60.3.162 svirSum

double svirSum

Definition at line 14 of file globalExtern.h.

3.60.3.163 timeNow

double timeNow

Definition at line 13 of file globalExtern.h.

3.60.3.164 TotalBondEnergy

double TotalBondEnergy [extern]

3.60.3.165 TotalMass

double TotalMass

Definition at line 24 of file globalExtern.h.

3.60.3.166 totEnergy

double totEnergy

Definition at line 13 of file globalExtern.h.

3.60.3.167 uSum

double uSum

Definition at line 14 of file globalExtern.h.

3.60.3.168 uSumPair

double uSumPair [extern]

3.60.3.169 uSumPairPerAtom

double uSumPairPerAtom

Definition at line 75 of file globalExtern.h.

3.60.3.170 virSum

double virSum

Definition at line 14 of file globalExtern.h.

3.60.3.171 virSumBond

double virSumBond [extern]

3.60.3.172 virSumBondxx

double virSumBondxx

Definition at line 76 of file globalExtern.h.

3.60.3.173 virSumBondxy

double virSumBondxy

Definition at line 76 of file globalExtern.h.

3.60.3.174 virSumBondyy

double virSumBondyy

Definition at line 76 of file globalExtern.h.

3.60.3.175 virSumPair

double virSumPair

Definition at line 75 of file globalExtern.h.

3.60.3.176 virSumPairxx

double virSumPairxx

Definition at line 75 of file globalExtern.h.

3.60.3.177 virSumPairxy

double virSumPairxy

Definition at line 75 of file globalExtern.h.

3.60.3.178 virSumPairyy

double virSumPairyy

Definition at line 75 of file globalExtern.h.

3.60.3.179 virSumxx

double virSumxx [extern]

3.60.3.180 virSumxy

double virSumxy

Definition at line 77 of file globalExtern.h.

3.60.3.181 virSumyy

double virSumyy

Definition at line 77 of file globalExtern.h.

3.60.3.182 visc

char visc[256] [extern]

3.60.3.183 viscAcf

double ** viscAcf

Definition at line 86 of file globalExtern.h.

3.60.3.184 viscAcfAv

 $\verb"double" * \verb"viscAcfAv"$

Definition at line 86 of file globalExtern.h.

3.60.3.185 viscAcfInt

double viscAcfInt

Definition at line 86 of file globalExtern.h.

3.60.3.186 viscAcfOrg

double * viscAcfOrg

Definition at line 86 of file globalExtern.h.

3.60.3.187 VMeanSqr

double VMeanSqr

Definition at line 36 of file globalExtern.h.

3.60.3.188 vrms

char vrms[256] [extern]

3.60.3.189 VRootMeanSqr

double VRootMeanSqr

Definition at line 36 of file globalExtern.h.

3.60.3.190 VSqr

double VSqr [extern]

3.60.3.191 vSum

double vSum

Definition at line 14 of file globalExtern.h.

3.60.3.192 vSumX

double vSumX

Definition at line 14 of file globalExtern.h.

3.60.3.193 vSumY

double vSumY

Definition at line 14 of file globalExtern.h.

3.60.3.194 vvSum

double vvSum

Definition at line 14 of file globalExtern.h.

3.60.3.195 vx

double * vx

Definition at line 10 of file globalExtern.h.

3.60.3.196 vy

double * vy

Definition at line 10 of file globalExtern.h.

3.60.3.197 xBoundary

char xBoundary[10] [extern]

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3.60.3.198 xyz

```
char xyz[256] [extern]
```

3.60.3.199 yBoundary

char yBoundary[10]

Definition at line 57 of file globalExtern.h.

3.61 globalExtern.h

Go to the documentation of this file.

```
00001 #ifndef GLOBALEXTERN_H
00002 #define GLOBALEXTERN H
00003
00004 typedef double real;
00005
00006 #define NDIM 2
00007 #define Sqr(x) ((x) * (x))
00008 #define SignR(x, y) (((y) >= 0) ? (x) : (- (x)))
00009
00010 extern double
                     *rx, *ry, *vx, *vy, *ax, *ay;
                     *speed;
00011 extern double
00013 extern double region[NDIM+1], regionH[NDIM+1], deltaT, timeNow, potEnergy, kinEnergy, totEnergy,
00014 density, pressure, rCut, kappa, uSum, virSum, svirSum, vSum, vSumX, vSumY, vvSum, sPotEnergy,
     sKinEnergy,
00015 sTotEnergy, sPressure, ssPotEnergy, ssKinEnergy, ssTotEnergy, ssPressure;
00016
                     initUcell[NDIM+1], moreCycles, nAtom, stepAvg, stepCount, stepEquil, stepLimit,
00017 extern int
      stepTraj, stepDump;
00018
00019 extern double RadiusIJ, SqrRadiusIJ, RadiusIJInv;
00020 extern int
                     nAtomType;
00021 extern int
                     *atomType;
00022 extern int
                     *atomID;
00023 extern double *atomRadius;
00024 extern double *atomMass, TotalMass;
00025
00026 extern int
00027 extern int
                     nBond, nBondType;
                     *atom1, *atom2;
*BondID, *BondType;
00028 extern int
00029 extern double *kb, *ro;
00030 extern double *BondEnergy, *BondLength;
00031 extern double
                     TotalBondEnergy, BondEnergyPerAtom;
00032 extern double
                     gamman;
                    *discDragx, *discDragy, *nodeDragx, *nodeDragy;
00033 extern double
00034 extern double
                     strain, strainRate;
00035 extern double
                     shearDisplacement, shearVelocity;
00036 extern double
                     VSqr, VMeanSqr, VRootMeanSqr;
00037 extern double
                    ComX, ComY, ComX0, ComY0, ComXRatio, ComYRatio;
HaltCondition;
00038 extern double
00039 extern double DeltaY, DeltaX;
00040 extern int
                     *ImageX, *ImageY;
00041 extern double
                     *rxUnwrap, *ryUnwrap;
00042 extern int
                     nAtomInterface, nDiscInterface, nAtomBlock;
00043 extern int
                     *atomIDInterface;
00044 extern double Kn;
00045 extern double fx, fy, FyBylx, fxByfy;
00046 extern int
                     DampFlag;
00047 extern double strech;
00048
00049 //For dumping the pair interaction data
00050 extern int dumpPairFlag;
00054
00055
00056 extern char solver[128];
00057 extern char xBoundary[10], yBoundary[10];
00058
00059 //For damping as in PRL, 130, 178203 (2023)
00060 extern double *DeltaXijOld, *DeltaYijOld;
00061 extern double DeltaXijNew, DeltaYijNew;
00062 extern double DeltaXij, DeltaYij, DeltaVXij, DeltaVYij;
00063 extern double **DeltaXijOldPair, **DeltaYijOldPair;
00064
00065 //For molecule-ID as per LAMMPS, helpful!
00066 extern int *molID;
00067 extern int
                    **isBonded;
```

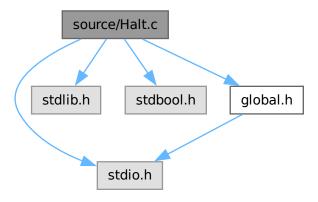
```
00069 //Following three for MPI only
00070 extern int *cellList, cells[NDIM+1];
00071 extern int
                    rank, size, master;
00072 extern double *fax, *fay, fuSum, fvirSum, frfAtom;
00073
00074 //For thermodynamic properties
00075 extern double uSumPair, uSumPairPerAtom, virSumPair, virSumPairxx, virSumPairxy, virSumPairxy;
00076 extern double virSumBond, virSumBondxx, virSumBondyy, virSumBondxy;
00077 extern double virSumxx, virSumyy, virSumxy;
00078 extern int
                    freezeAtomType;
00079
00080 // Spacetime Correlations
00081 extern double ***cfOrg, **spacetimeCorr, *cfVal, *spacetimeCorrAv;
00082 extern int *indexCorr, countCorrAv, limitCorrAv, nBuffCorr, nFunCorr, nValCorr, stepCorr;
00083
00084 // Viscosity
00085 extern double rfAtom, frfAtom;
00086 extern double *indexAcf, **viscAcf, *viscAcfOrg, *viscAcfAv, viscAcfInt;
00087 extern int
                     nValAcf, nBuffAcf, stepAcf, countAcfAv, limitAcfAv;
00088
00089 // Radial distribution function
00091 extern int
00092
00093
00094 // Output files prefixes
00095 extern char
00096
00097 extern char
                     result[250];
00098 extern FILE
                     *fpresult;
00099
00100 extern char
                     xyz[256];
00101 extern FILE
                     *fpxyz;
00102
                    bond[256];
00103 extern char
00104 extern FILE
                     *fpbond;
00106
00107 extern char
                     dump[256];
00108 extern FILE
                     *fpdump;
00109
00110 extern char
                     dnstv[256]:
00111 extern FILE
                     *fpdnsty;
00112
00113 extern char
                     visc[256];
00114 extern FILE
                     *fpvisc;
00115
00116 extern char
                     rdf[2561;
00117 extern FILE
                     *fordf:
00118
00119 extern char
                     vrms[256];
00120 extern FILE
                     *fpvrms;
00121
00122 extern char
                     stress[256];
00123 extern FILE
                     *fpstress;
00125 extern char
                     momentum[256];
00126 extern FILE
                     *fpmomentum;
00127
00128 extern char
                     com[256]:
00129 extern FILE
                     *fpcom;
00130
00131 extern char
                     pair[256];
00132 extern FILE
                     *fppair;
00133
00134 #endif // GLOBALEXTERN_H
```

3.62 source/Halt.c File Reference

```
#include <stdio.h>
#include <stdlib.h>
#include <stdbool.h>
#include "global.h"
```

3.63 Halt.c 121

Include dependency graph for Halt.c:



Functions

bool HaltConditionCheck (double value, int stepCount)

3.62.1 Function Documentation

3.62.1.1 HaltConditionCheck()

References fpresult, HaltCondition, and stepCount.

Referenced by main().

Here is the caller graph for this function:



3.63 Halt.c

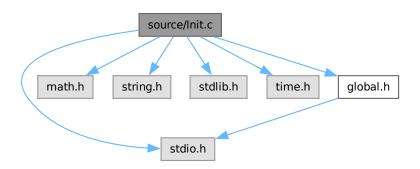
```
Go to the documentation of this file.
```

```
00001 /* 00002 * This file is part of Lamina.
```

```
00003
       * Lamina is free software: you can redistribute it and/or modify * it under the terms of the GNU General Public License as published by
00004
00005
       * the Free Software Foundation, either version 3 of the License, or
00006
00007
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80000
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00010
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00016
00017 Copyright (C) 2025 Harish Charan, University of Durham, UK
00018
00019 */
00020
00021
00022 #include <stdio.h>
00023 #include <stdlib.h>
00024 #include <stdbool.h>
00025 #include "global.h"
00026
00027 bool HaltConditionCheck(double value, int stepCount) {
00028
00029
         if(value <= HaltCondition && value != 0) {</pre>
00030 fprintf(fpresult, "Halt condition met at step = %d with Vrms = %.10f\n", stepCount, value);
00031
         return true;
                               // Signal that the halt condition is met
00032
00033 return false; // Halt condition not met
00034 }
00035
```

3.64 source/Init.c File Reference

```
#include <stdio.h>
#include <math.h>
#include <string.h>
#include <stdlib.h>
#include <time.h>
#include "global.h"
Include dependency graph for Init.c:
```



Functions

• void Init ()

3.64.1 Function Documentation

3.64.1.1 Init()

```
void Init ( )
Definition at line 29 of file Init.c.
00029
00030
                      char dummy[128];
00031
                     char inputConfig[128];
FILE *fp;
00032
                   FILE *fp;
fp = fopen("input-data","r");
fscanf(fp, "%s %s", dummy, inputConfig);
fscanf(fp, "%s %s", dummy, solver);
fscanf(fp, "%s %s %s", dummy, xBoundary, yBoundary);
fscanf(fp, "%s %d", dummy, &DampFlag);
fscanf(fp, "%s %d", dummy, &Tcut);
fscanf(fp, "%s %lf", dummy, &gamman);
fscanf(fp, "%s %lf", dummy, &kappa);
fscanf(fp, "%s %lf", dummy, &deltaT);
fscanf(fp, "%s %lf", dummy, &fxBylx);
fscanf(fp, "%s %lf", dummy, &FxBylx);
fscanf(fp, "%s %lf", dummy, &FxBylx);
fscanf(fp, "%s %lf", dummy, &DeltaY);
fscanf(fp, "%s %lf", dummy, &DeltaX);
fscanf(fp, "%s %lf", dummy, &HaltCondition);
fscanf(fp, "%s %d", dummy, &stepAvg);
fscanf(fp, "%s %d", dummy, &stepEquil);
fscanf(fp, "%s %d", dummy, &stepEquil);
fscanf(fp, "%s %d", dummy, &stepLimit);
fscanf(fp, "%s %d", dummy, &stepDump);
                      fp = fopen("input-data", "r");
00034
00035
00036
00037
00038
00040
00041
00042
00043
00044
00045
00046
00047
00048
00049
00050
00051
                    fscanf(fp, "%s %d",
f
00052
00053
00054
00055
00056
00057
                    fscanf(fp, "%s %d", dummy, &limitAcfAv fscanf(fp, "%s %d", dummy, &nBuffAcf); fscanf(fp, "%s %d", dummy, &nValAcf); fscanf(fp, "%s %d", dummy, &stepAcf);
00059
                                                                               dummy, &limitAcfAv);
00060
00061
00062
                     fscanf(fp, "%s %lf", dummy, &rangeRdf);
00063
                    fscanf(fp, "%s %d", dummy, &limitRdf);
fscanf(fp, "%s %d", dummy, &sizeHistRdf);
fscanf(fp, "%s %d", dummy, &stepRdf);
 00064
00065
00066
00067
00068
                     fclose(fp):
00069
                     FILE *fpSTATE;
                      if((fpSTATE = fopen(inputConfig, "r")) ==NULL) {
00071
                      printf("Error occurred: Could not open STATE file\n Exiting now..\n");
00072
00073
00074
                     fscanf(fpSTATE, "%s %lf", dummy, &timeNow);
fscanf(fpSTATE, "%s %d", dummy, &nAtom);
00075
00076
                      fscanf(fpSTATE, "%s %d", dummy, &nBond);
00077
                     fscanf(fpSTATE, "%s %d", dummy, &nAtomType); fscanf(fpSTATE, "%s %d", dummy, &nBondType); fscanf(fpSTATE, "%s %d", dummy, &nBondType); fscanf(fpSTATE, "%s %lf", dummy, &region[1]); fscanf(fpSTATE, "%s %lf", dummy, &region[2]);
00078
00079
08000
00081
00082
 00083
                      region[2] *= 1.5; //Remove this when put on GitHub
00084
00085
                     density = nAtom/(region[1]*region[2]);
                     cells[1] = region[1] / rCut;
cells[2] = region[2] / rCut;
00086
00087
00088
                      cellList = (int*)malloc((nAtom + cells[1] * cells[2] + 1) * sizeof(int));
                     regionH[1] = 0.5*region[1];
00090
                     regionH[2] = 0.5*region[2];
00091
00092
                     //strain information
00093
                     strainRate = strain/deltaT;
shearDisplacement = strain * region[2];
00094
00095
                      shearVelocity = strainRate * region[2];
00096
00097
00098
                     rx = (double*)malloc((nAtom + 1) * sizeof(double));
00099
                      ry = (double*)malloc((nAtom + 1) * sizeof(double));
                      vx = (double*)malloc((nAtom + 1) * sizeof(double));
00100
                     vy = (double*)malloc((nAtom + 1) * sizeof(double));
ax = (double*)malloc((nAtom + 1) * sizeof(double));
00101
 00102
00103
                      ay = (double*)malloc((nAtom + 1) * sizeof(double));
                      fax = (double*)malloc((nAtom + 1) * sizeof(double));
fay = (double*)malloc((nAtom + 1) * sizeof(double));
00104
00105
                      atomID = (int*)malloc((nAtom+1) * sizeof(int));
00106
00107
                      atomType = (int*)malloc((nAtom+1) * sizeof(int));
                     atomRadius = (double*)malloc((nAtom + 1) * sizeof(double));
```

```
atomMass = (double*)malloc((nAtom + 1) * sizeof(double));
         speed = (double*)malloc((nAtom + 1) * sizeof(double));
00110
00111
         atom1 = (int*)malloc((nBond+1)*sizeof(int));
         atom2 = (int*)malloc((nBond+1)*sizeof(int));
00112
         BondID = (int*)malloc((nBond+1)*sizeof(int));
00113
00114
         BondType = (int*)malloc((nBond+1)*sizeof(int));
         kb = (double*)malloc((nBond+1)*sizeof(double));
00115
00116
         ro = (double*)malloc((nBond+1)*sizeof(double));
         BondEnergy = (double*)malloc((nBond+1)*sizeof(double));
BondLength = (double*)malloc((nBond+1)*sizeof(double));
00117
00118
         discDragx = (double*)malloc((nAtom + 1) * sizeof(double));
discDragy = (double*)malloc((nAtom + 1) * sizeof(double));
00119
00120
         nodeDragx = (double*)malloc((nAtom + 1) * sizeof(double));
00121
         nodeDragy = (double*)malloc((nAtom + 1) * sizeof(double));
00122
         ImageX = (int*)malloc((nAtom+1) * sizeof(int));
ImageY = (int*)malloc((nAtom+1) * sizeof(int));
00123
00124
         rxUnwrap = (double*)malloc((nAtom + 1) * sizeof(double));
ryUnwrap = (double*)malloc((nAtom + 1) * sizeof(double));
00125
00126
         DeltaXijOld = (double*)malloc((nBond+1)*sizeof(double));
         DeltaYijOld = (double*)malloc((nBond+1)*sizeof(double));
00128
         DeltaXijOldPair = (double**)malloc((nAtom+1) * sizeof(double*));
DeltaYijOldPair = (double**)malloc((nAtom+1) * sizeof(double*));
00129
00130
         for(int n = 0; n <= nAtom; n++) {
  DeltaXijOldPair[n] = (double*)malloc((nAtom+1) * sizeof(double));</pre>
00131
00132
00133
          DeltaYijOldPair[n] = (double*)malloc((nAtom+1) * sizeof(double));
00134
00135
         molID = (int*)malloc((nAtom+1) * sizeof(int));
00136
         for(n = 1; n <= nAtom; n ++) {
  atomMass[n] = 1.0;</pre>
00137
00138
00139
00140
00141
         fscanf(fpSTATE, "%s\n", dummy);
00142
         for (n = 1; n \le nAtom; n ++)
          fscanf (fpSTATE, "%d %d %d %lf %lf %lf %lf %lf %lf n", &atomID[n], &molID[n], &atomType[n], \\
00143
      &atomRadius[n], &rx[n], &ry[n], &vx[n], &vy[n]);
00144
00145
00146
         fscanf(fpSTATE, "%s\n", dummy);
         for(n=1; n<=nBond; n++) fscanf(fpSTATE, "%d %d %d %d %lf %lf\n", &BondID[n], &BondType[n], &atom1[n], &atom2[n], &kb[n],
00147
00148
      &ro[n]);
00149
00150
         fclose(fpSTATE);
00151
00152
        //2D-List of bonded atoms. This is used to remove pair interaction
00153
        //calculation for the bonded atoms
00154
          isBonded = (int**)malloc((nAtom + 1) * sizeof(int*));
00155
         for (int i = 0; i <= nAtom; i++) {</pre>
          isBonded[i] = (int*)malloc((nAtom + 1) * sizeof(int));
00156
           for (int j = 0; j <= nAtom; j++) {
   isBonded[i][j] = 0;</pre>
00157
00158
00159
00160
        }
00161
00162
         for (n = 1; n \le nBond; n++) {
          int i = atom1[n];
            int j = atom2[n];
00164
           isBonded[i][j] = 1;
isBonded[j][i] = 1; // symmetric
00165
00166
00167 }
00168
00169
        //For thermostate, update in final version
00170 thermo = 'C';
00171
00172
00173 // List the interface atoms
00174 nAtomInterface = 0:
00175 nAtomBlock = 0;
00176 nDiscInterface = 0;
00177
        double InterfaceWidth, bigDiameter;
        bigDiameter = 2.8;
00178
00179
        InterfaceWidth = 5.0 * bigDiameter;
00180
00181
        for (n = 1; n <= nAtom; n++) {</pre>
        if(fabs(ry[n]) < InterfaceWidth){</pre>
00182
00183
         nAtomInterface++;
00184
         if(molID[n] == 2){
00185
00186
         nAtomBlock++:
00187
00188
         if(atomRadius[n] != 0.0){
00189
         nDiscInterface++;
00190
00191
          atomIDInterface = (int*)malloc((nAtomInterface+1)*sizeof(int));
00192
00193
```

```
00194
           int m;
00195
00196
           for (n=1; n<=nAtom; n++) {</pre>
00197
            if(fabs(ry[n]) < InterfaceWidth){</pre>
00198
            atomIDInterface[m] = atomID[n];
00199
            m++;
00201
00202
           nPairTotal = 0.5 * nAtomInterface * (nAtomInterface-1);
00203
           PairID = (int*)malloc((nPairTotal+1) * sizeof(int));
           Pairatom1 = (int*)malloc((nPairTotal+1) * sizeof(int));
00204
           Pairatom2 = (int*)malloc((nPairTotal+1) * sizeof(int));
00205
00206
           PairXij = (double*)malloc((nPairTotal+1) * sizeof(double));
           PairYij = (double*)malloc((nPairTotal+1) * sizeof(double));
00207
00208
00209
              fprintf(fpresult, "-----
              fprintf(fpresult, "-----PARAMETERS----\n");
00210
              00211
00212
              fprintf(fpresult, "nBond\t\t\t\d\n",
00213
                                                                  nBond);
              fprintf(fpresult, "nAtomBlock\t\t\t\d\n", nAtomBlock);
fprintf(fpresult, "nAtomInterface\t\d\n", nAtomInter
00214
                                                                         nAtomInterface);
00215
              fprintf(fpresult, "nDiscInterface\t\t\%d\n", nDiscInterface);
00216
              fprintf(fpresult, "gamman\t\t\t\0.6g\n", gamman);
fprintf(fpresult, "strain\t\t\t\0.6g\n", strain);
fprintf(fpresult, "strainRate\t\t\0.6g\n", strainRate);
00217
00218
              fprintf(fpresult, "FyBylx\t\t\t\80.6g\n", FyBylx);
fprintf(fpresult, "fxByfy\t\t\80.6g\n", fxByfy);
fprintf(fpresult, "DeltaY\t\t\80.6g\n", DeltaY);
00220
00221
00222
              fprintf(fpresult, "DeltaX\t\t\t\80.6g\n", DeltaX\;
fprintf(fpresult, "BeltaX\t\t\80.6g\n", DeltaX\;
fprintf(fpresult, "HaltCondition\t\t\80.6g\n", HaltCondition);
fprintf(fpresult, "kappa\t\t\t\$g\n", kappa);
00223
00224
00225
              fprintf(fpresult, "density\t\t\t\g\n", density);
00226
00227
              fprintf(fpresult, "rCut\t\t\t\t\g\n", rCut);
              fprintf(fpresult, "deltaT\t\t\t\g\n", deltaT);
fprintf(fpresult, "stepEquil\t\t\t\g\n", stepEquil);
00228
00229
              fprintf(fpresult, "stepLimit);
fprintf(fpresult, "stepLimit);
fprintf(fpresult, "region[1]\t\t\t\0.161f\n", region[1]);
00230
              fprintf(fpresult, "region[2])t\t\t\s\0.161f\n", region[2]);
fprintf(fpresult, "cells[1]\t\t\s\n", cells[1]);
fprintf(fpresult, "cells[2]\t\t\s\d\n", cells[2]);
00232
00233
00234
              fprintf(fpresult, "solver\t\t\t\s\n", solver);
fprintf(fpresult, "boundary\t\t\t\s\n", xBoundary, yBoundary);
00235
00236
              fprintf(fpresult, "DampFlag\t\t\d\n", DampFlag);
00237
00238
00239
              00240
00241
       PairEnergyPerAtom BondEnergyPerAtom Press VirialSum\n");
fprintf(fpvrms, "#timeNow\tVrms \n");
fprintf(fpcom, "#timeNow\tComX\tComY\n");
00242
00243
00244
00245 /\star //Uncomment the following as per your acquirement
                                                     %lf\n", strain);
%lf\n", region[1]);
              fprintf(fpstress, "strain
fprintf(fpstress, "region[1]
00246
00247
              fprintf(fpstress, "region[2] % lf\n", region[2]);
fprintf(fpstress, "#timeNow virSumxx virSumyy virSumxy pressure\n");
00248
              fprintf(fpmomentum, "#timeNow Px Py\n");
00250
00251 */
00252
            if((strcmp(xBoundary, "p") != 0 && strcmp(xBoundary, "r") != 0) ||
(strcmp(yBoundary, "p") != 0 && strcmp(yBoundary, "r") != 0)) {
00253
00254
             (strcmp(yBoundary, "p") != 0 && strcmp(yBoundary, "r") != 0)) {
fprintf(fpresult, "Error: Invalid boundary value detected: '%s %s'. Only 'p' or 'r' are
00255
             owed.\n", xBoundary, yBoundary);
exit(EXIT_FAILURE); // Exit with failure status
00256
00257
00258
00259
```

References atom1, atom2, atomID, atomIDInterface, atomMass, atomRadius, atomType, ax, ay, BondEnergy, BondID, BondLength, BondType, cellList, cells, DampFlag, deltaT, DeltaX, DeltaXijOld, DeltaXijOldPair, DeltaY, DeltaYijOld, DeltaYijOldPair, density, discDragx, discDragy, fax, fay, fpcom, fpresult, fpvrms, freezeAtomType, fxByfy, FyBylx, gamman, HaltCondition, ImageX, ImageY, isBonded, kappa, kb, limitAcfAv, limitCorrAv, limitRdf, molID, nAtom, nAtomBlock, nAtomInterface, nAtomType, nBond, nBondType, nBuffAcf, nBuffCorr, nDiscInterface, nFunCorr, nodeDragx, nodeDragy, nPairTotal, nValAcf, nValCorr, Pairatom1, Pairatom2, PairID, PairXij, PairYij, rangeRdf, rCut, region, regionH, ro, rx, rxUnwrap, ry, ryUnwrap, shearDisplacement, shearVelocity, sizeHistRdf, solver, speed, stepAcf, stepAvg, stepCorr, stepDump, stepEquil, stepLimit, stepRdf, stepTraj, strain, strainRate, thermo, timeNow, vx, vy, xBoundary, and yBoundary.

Referenced by main().

Here is the caller graph for this function:



3.65 Init.c

Go to the documentation of this file.

```
00002
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00003
00004
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00014
00015
00016
00017
               Copyright (C) 2025 Harish Charan, University of Durham, UK
00018
00019
00020
00021
00022 #include<stdio.h>
00023 #include<math.h>
00024 #include<string.h>
00025 #include<stdlib.h>
00026 #include <time.h>
00027 #include"global.h"
00028
00029 void Init() {
                 char dummy[128];
char inputConfig[128];
00030
00031
00032
                  FILE *fp;
                 fribe *tp;
fp = fopen("input-data","r");
fscanf(fp, "%s %s", dummy, inputConfig);
fscanf(fp, "%s %s", dummy, solver);
fscanf(fp, "%s %s", dummy, xBoundary, yBoundary);
00033
00034
00035
00036
                 fscanf(fp, "%s %s %s", dummy, xBoundary, yBoundary, grant f(fp, "%s %d", dummy, &DampFlag);
fscanf(fp, "%s %d", dummy, &freezeAtomType);
fscanf(fp, "%s %lf", dummy, &rCut);
fscanf(fp, "%s %lf", dummy, &gamman);
fscanf(fp, "%s %lf", dummy, &kappa);
fscanf(fp, "%s %lf", dummy, &ctall;
fscanf(fp, "%s %lf", dummy, &freeleft, "%s %lf", dummy, &freeleft, "%s %lf", dummy, &freeleft, grant f(fp, "%s %lf", dummy, &freeleft, grant f(fp, "%s %lf", dummy, &freeleft, grant f(fp, grant f(fp
00037
00039
00040
00041
00042
00043
00044
                  fscanf(fp, "%s %lf", dummy, &FyBylx);
                  fscanf(fp, "%s %lf", dummy, &fxByfy);
00045
                  fscanf(fp, "%s %lf", dummy, &DeltaY);
fscanf(fp, "%s %lf", dummy, &DeltaX);
00046
00047
                  fscanf(fp, "%s %lf", dummy, &HaltCondition);
00048
                  fscanf(fp, "%s %d", dummy, &stepAvg);
00049
                  fscanf(fp, "%s %d", dummy, &stepEquil);
00050
                  fscanf(fp, "%s %d", fscanf(fp, "%s %d",
00051
                                                                    dummy, &stepLimit);
00052
                                                                   dummy, &stepDump);
                  fscanf(fp, "%s %d", fscanf(fp, "%s %d", fscanf(fp, "%s %d",
00053
                                                                    dummy, &stepTraj);
00054
                                                                    dummy, &limitCorrAv);
00055
                                                                    dummy, &nBuffCorr);
                  fscanf(fp, "%s %d", fscanf(fp, "%s %d",
00056
                                                                    dummy, &nFunCorr);
                                                                    dummy, &nValCorr);
                  fscanf(fp, "%s %d", fscanf(fp, "%s %d",
00058
                                                                    dummy, &stepCorr);
00059
                                                                    dummy, &limitAcfAv);
                  fscanf(fp, "%s %d",
00060
                                                                    dummy, &nBuffAcf);
                  fscanf(fp, "%s %d", dummy, &nWalAcf);
fscanf(fp, "%s %d", dummy, &stepAcf);
fscanf(fp, "%s %d", dummy, &rangeRdf);
00061
00062
00063
```

3.65 Init.c 127

```
fscanf(fp, "%s %d", dummy, &limitRdf);
fscanf(fp, "%s %d", dummy, &sizeHistRdf);
fscanf(fp, "%s %d", dummy, &stepRdf);
00065
00066
00067
00068
         fclose(fp);
00069
         FILE *fpSTATE:
         if((fpSTATE = fopen(inputConfig, "r")) == NULL) {
00070
00071
         printf("Error occurred: Could not open STATE file\n Exiting now..\n");
00072
00073
00074
         fscanf(fpSTATE, "%s %lf", dummy, &timeNow);
fscanf(fpSTATE, "%s %d", dummy, &nAtom);
fscanf(fpSTATE, "%s %d", dummy, &nBond);
00075
00076
00077
          fscanf(fpSTATE, "%s %d",
00078
                                         dummy, &nAtomType);
         fscanf(fpSTATE, "%s %d", dummy, &nBondType);
fscanf(fpSTATE, "%s %lf", dummy, &region[1]);
fscanf(fpSTATE, "%s %lf", dummy, &region[2]);
00079
08000
00081
00082
00083
         region[2] *= 1.5; //Remove this when put on GitHub
00084
00085
         density = nAtom/(region[1]*region[2]);
         cells[1] = region[1] / rCut;
cells[2] = region[2] / rCut;
00086
00087
00088
         cellList = (int*)malloc((nAtom + cells[1] * cells[2] + 1) * sizeof(int));
         regionH[1] = 0.5*region[1];
00089
00090
         regionH[2] = 0.5*region[2];
00091
00092
         //strain information
00093
         strainRate = strain/deltaT;
00094
         shearDisplacement = strain * region[2];
00095
         shearVelocity = strainRate * region[2];
00096
00097
         rx = (double*)malloc((nAtom + 1) * sizeof(double));
ry = (double*)malloc((nAtom + 1) * sizeof(double));
00098
00099
         vx = (double*)malloc((nAtom + 1) * sizeof(double));
00100
         vy = (double*)malloc((nAtom + 1) * sizeof(double));
00102
         ax = (double*)malloc((nAtom + 1) * sizeof(double));
         ay = (double*)malloc((nAtom + 1) * sizeof(double));
00103
         fax = (double*)malloc((nAtom + 1) * sizeof(double));
fay = (double*)malloc((nAtom + 1) * sizeof(double));
00104
00105
         atomTD = (int*)malloc((nAtom+1) * sizeof(int)):
00106
00107
         atomType = (int*)malloc((nAtom+1) * sizeof(int));
         atomRadius = (double*)malloc((nAtom + 1) * sizeof(double));
00108
00109
         atomMass = (double*)malloc((nAtom + 1) * sizeof(double));
00110
         speed = (double*)malloc((nAtom + 1) * sizeof(double));
00111
         atom1 = (int*)malloc((nBond+1)*sizeof(int));
         atom2 = (int*)malloc((nBond+1)*sizeof(int));
00112
00113
         BondID = (int*)malloc((nBond+1)*sizeof(int));
00114
         BondType = (int*)malloc((nBond+1)*sizeof(int));
00115
         kb = (double*)malloc((nBond+1)*sizeof(double));
00116
          ro = (double*)malloc((nBond+1)*sizeof(double));
         BondEnergy = (double*)malloc((nBond+1)*sizeof(double));
BondLength = (double*)malloc((nBond+1)*sizeof(double));
00117
00118
         discDragx = (double*)malloc((nAtom + 1) * sizeof(double));
discDragy = (double*)malloc((nAtom + 1) * sizeof(double));
00119
         nodeDragx = (double*)malloc((nAtom + 1) * sizeof(double));
00121
         nodeDragy = (double*)malloc((nAtom + 1) * sizeof(double));
00122
         ImageX = (int*)malloc((nAtom+1) * sizeof(int));
ImageY = (int*)malloc((nAtom+1) * sizeof(int));
00123
00124
         rxUnwrap = (double*)malloc((nAtom + 1) * sizeof(double));
ryUnwrap = (double*)malloc((nAtom + 1) * sizeof(double));
00125
00126
         DeltaXijOld = (double*)malloc((nBond+1)*sizeof(double));
00127
         DeltaYijOld = (double*)malloc((nBond+1)*sizeof(double));
00128
         DeltaXijOldPair = (double**)malloc((nAtom+1) * sizeof(double*));
DeltaYijOldPair = (double**)malloc((nAtom+1) * sizeof(double*));
00129
00130
         for (int n = 0; n <= nAtom; n++) {</pre>
00131
          DeltaXijOldPair[n] = (double*)malloc((nAtom+1) * sizeof(double));
00132
          DeltaYijOldPair[n] = (double*)malloc((nAtom+1) * sizeof(double));
00133
00134
00135
         molID = (int*)malloc((nAtom+1) * sizeof(int));
00136
00137
         for (n = 1; n \le nAtom; n ++) {
00138
          atomMass[n] = 1.0;
00139
00140
00141
          fscanf(fpSTATE, "%sn", dummy);
         00142
00143
       &atomRadius[n], &rx[n], &ry[n], &vx[n], &vy[n]);
00144
00145
00146
          fscanf(fpSTATE, "%s\n", dummy);
         for(n=1; n<=nBond; n++)
fscanf(fpSTATE, "%d %d %d %d %lf %lf\n", &BondID[n], &BondType[n], &atom1[n], &atom2[n], &kb[n],</pre>
00147
00148
       &ro[n]);
```

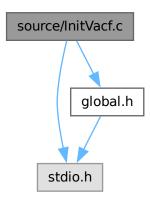
```
00150
            fclose(fpSTATE);
00151
00152
          //2D-List of bonded atoms. This is used to remove pair interaction
00153
           //calculation for the bonded atoms
             isBonded = (int**)malloc((nAtom + 1) * sizeof(int*));
00154
            for (int i = 0; i <= nAtom; i++) {</pre>
00156
               isBonded[i] = (int*)malloc((nAtom + 1) * sizeof(int));
00157
               for (int j = 0; j <= nAtom; j++) {</pre>
                      isBonded[i][j] = 0;
00158
00159
00160
00161
            for (n = 1; n <= nBond; n++) {</pre>
00162
              int i = atom1[n];
int j = atom2[n];
00163
00164
               isBonded[i][j] = 1;
isBonded[j][i] = 1; // symmetric
00165
00166
00167 }
00168
00169 //For thermostate, update in final version
00170 thermo = 'C';
00171
00172
00173 // List the interface atoms
00174 nAtomInterface = 0;
00175 nAtomBlock = 0;
00176 nDiscInterface = 0;
00177 double InterfaceWidth, bigDiameter;
00178 bigDiameter = 2.8;
00179 InterfaceWidth = 5.0 * bigDiameter;
00180
00181 for(n = 1; n <= nAtom; n++){
00182
           if(fabs(ry[n]) < InterfaceWidth){</pre>
00183
           nAtomInterface++;
00184
            if (molID[n] == 2) {
00185
00186
           nAtomBlock++;
00187
00188
             if (atomRadius[n] != 0.0) {
00189
            nDiscInterface++;
00190
            } }
00191
00192
             atomIDInterface = (int*)malloc((nAtomInterface+1)*sizeof(int));
00193
00194
00195
            m = 1;
00196
            for (n=1; n<=nAtom; n++) {</pre>
00197
             if(fabs(ry[n]) < InterfaceWidth){</pre>
00198
             atomIDInterface[m] = atomID[n];
00199
             m++;
00200
00201
           nPairTotal = 0.5 * nAtomInterface * (nAtomInterface-1);
PairID = (int*)malloc((nPairTotal+1) * sizeof(int));
00202
00203
00204
            Pairatom1 = (int*)malloc((nPairTotal+1) * sizeof(int));
            Pairatom2 = (int*)malloc((nPairTotal+1) * sizeof(int));
00206
            PairXij = (double*)malloc((nPairTotal+1) * sizeof(double));
00207
            PairYij = (double*)malloc((nPairTotal+1) * sizeof(double));
00208
                fprintf(fpresult, "----\n");
00209
               00210
00211
               fprintf(fpresult, "nAtom\t\t\t\t\d\n", nAtom);
fprintf(fpresult, "nBond\t\t\t\d\n", nBond);
00212
00213
               fprintf(fpresult, "nBond\t\t\t\d\n", nBond);
fprintf(fpresult, "nAtomBlock\t\t\d\n", nAtomBlock);
fprintf(fpresult, "nAtomInterface\t\d\n", nAtomInterface);
fprintf(fpresult, "nDiscInterface\t\t\d\n", nDiscInterface);
fprintf(fpresult, "gamman\t\t\t\d\0.6g\n", gamman);
fprintf(fpresult, "strain\t\t\t\0.6g\n", strain);
fprintf(fpresult, "strainRate\t\t\0.6g\n", strainRate);
fprintf(fpresult, "FyBylx\t\t\0.6g\n", FyBylx);
fprintf(fpresult, "FyBylx\t\t\0.6g\n", fxByfy);
00214
00215
00216
00217
00218
00219
00220
               fprintf(fpresult, "FyBylx\t\t\t\0.6g\n", FyBylx);
fprintf(fpresult, "fxByfy\t\t\t\0.6g\n", fxByfy);
fprintf(fpresult, "DeltaY\t\t\0.6g\n", DeltaY);
fprintf(fpresult, "DeltaX\t\t\0.6g\n", DeltaX);
fprintf(fpresult, "HaltCondition\t\t\0.6g\n", HaltCondition);
00221
00222
00223
00224
               fprintf(fpresult, "HaltCondition\t\t\\0,0\n", HaltCon
fprintf(fpresult, "kappa\t\t\t\\0,0\n", kappa);
fprintf(fpresult, "density\t\t\\0,0\n", density);
fprintf(fpresult, "rCut\t\t\t\\0,0\n", rCut);
fprintf(fpresult, "deltaT\t\t\\0,0\n", deltaT);
fprintf(fpresult, "stepEquil\t\t\\0,0\n", stepEquil);
fprintf(fpresult, "stepLimit\t\t\\0,0\n", stepLimit);
00225
00226
00227
00228
00229
00230
               fprintf(fpresult, "stepLimit\t\t\t\0.161f\n", region[1]);
fprintf(fpresult, "region[2]\t\t\t\0.161f\n", region[2]);
fprintf(fpresult, "region[2]\t\t\t\0.161f\n", region[2]);
fprintf(fpresult, "cells[1]\t\t\t\0.161f\n", cells[1]);
fprintf(fpresult, "cells[2]\t\t\t\0.161f\n", cells[2]);
fprintf(fpresult, "solver\t\t\t\0.181f\n", solver);
00231
00232
00233
00234
00235
```

```
 fprintf(fpresult, "boundary\t\t\t\s \s.\n", xBoundary, yBoundary); fprintf(fpresult, "DampFlag\t\t\t\n", DampFlag); 
00237
00238
00239
                fprintf(fpresult, "-----\n");
fprintf(fpresult, "#TimeNow TotalMomentum PotEngyPerAtom KinEngyPerAtom TotEngyPerAtom
00240
00241
         PairEnergyPerAtom BondEnergyPerAtom Press VirialSum\n");
fprintf(fpvrms, "#timeNow\tVrms \n");
fprintf(fpcom, "#timeNow\tComX\tComY\n");
00242
00243
00244
00245 /\star // \text{Uncomment} the following as per your acquirement
               //uncomment the following as per your acquirement
fprintf(fpstress, "strain %lf\n", strain);
fprintf(fpstress, "region[1] %lf\n", region[1]);
fprintf(fpstress, "region[2] %lf\n", region[2]);
fprintf(fpstress, "#timeNow virSumxx virSumxy virSumxy pressure\n");
00246
00247
00248
00249
00250
                fprintf(fpmomentum, "#timeNow Px Pyn");
00251 */
00252
               if((strcmp(xBoundary, "p") != 0 && strcmp(xBoundary, "r") != 0) ||
  (strcmp(yBoundary, "p") != 0 && strcmp(yBoundary, "r") != 0)) {
               (strcmp(yBoundary, "p") != 0 && strcmp(yBoundary, "r") != 0)) {
fprintf(fpresult, "Error: Invalid boundary value detected: '%s %s'. Only 'p' or 'r' are
00254
00255
allowed.\n", xBoundary, yBoundary);
00256 exit(EXIT_FAILURE); // Exit wit
               exit(EXIT_FAILURE); // Exit with failure status
00257
00258
00259 }
```

3.66 source/InitVacf.c File Reference

```
#include <stdio.h>
#include "global.h"
```

Include dependency graph for InitVacf.c:



Functions

- void ZeroVacf ()
- void InitVacf ()

3.66.1 Function Documentation

3.66.1.1 InitVacf()

```
00030 ZeroVacf();
```

References indexAcf, nBuffAcf, nValAcf, and ZeroVacf().

Referenced by SetupJob().

Here is the call graph for this function:



Here is the caller graph for this function:



3.66.1.2 ZeroVacf()

Referenced by InitVacf().

Here is the caller graph for this function:



3.67 InitVacf.c

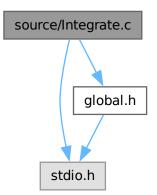
Go to the documentation of this file.

```
00001 /*
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00006 * the Free Software Foundation, either version 3 of the License, or
00007 * (at your option) any later version.
```

```
* 80000
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00013 *
00014 * You should have received a copy of the GNU General Public License 00015 * along with Lamina. If not, see <a href="https://www.gnu.org/licenses/">https://www.gnu.org/licenses/</a>.
00016
00017 Copyright (C) 2025 Harish Charan, University of Durham, UK
00018
00019 */
00020
00021
00022 #include<stdio.h>
00023 #include"global.h"
00024
00025 void ZeroVacf();
00026 void InitVacf() {
00027 int nb;
for(nb = 1; nb <= nBuffAcf; nb ++)
00029    indexAcf[nb] = -(nb-1)*nValAcf/nBuffAcf;</pre>
00030 ZeroVacf();
00031 }
```

3.68 source/Integrate.c File Reference

```
#include <stdio.h>
#include "global.h"
Include dependency graph for Integrate.c:
```



Functions

• double Integrate (double *f, int nf)

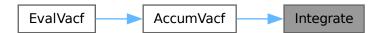
3.68.1 Function Documentation

3.68.1.1 Integrate()

```
00030 s += f[i];
00031 return(s);
00032 }
```

Referenced by AccumVacf().

Here is the caller graph for this function:



3.69 Integrate.c

Go to the documentation of this file.

```
00002 * This file is part of Lamina.
00003
00004 \,\star\, Lamina is free software: you can redistribute it and/or modify
00005 \, \, \, it under the terms of the GNU General Public License as published by
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00006
00007
         * (at your option) any later version.
80000
00009
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00016
00017 Copyright (C) 2025 Harish Charan, University of Durham, UK
00018
00019
00020
00021
00022 #include<stdio.h>
00023 #include"global.h"
00024
00025 double Integrate(double *f, int nf){
00025 double into
00026 double s;
00027 int i;
00022 int i;

00028 s = 0.5*(f[1] + f[nf]);

00029 for(i = 2; i <= nf - 1; i ++)

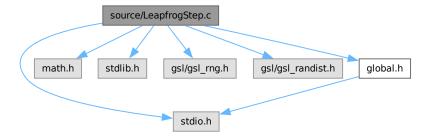
00030 s += f[i];

00031 return(s);
00032 }
00033
```

3.70 source/LeapfrogStep.c File Reference

```
#include <stdio.h>
#include <math.h>
#include <stdlib.h>
#include <gsl/gsl_rng.h>
#include <gsl/gsl_randist.h>
#include "global.h"
```

Include dependency graph for LeapfrogStep.c:



Functions

void LeapfrogStep (char thermo, gsl_rng *rnd)

3.70.1 Function Documentation

3.70.1.1 LeapfrogStep()

```
void LeapfrogStep (
                char thermo,
                gsl_rng * rnd )
Definition at line 28 of file LeapfrogStep.c.
00029 double temperature, GAMMA;
00030 \text{ GAMMA} = 100;
00031
00032 double *TValSum;
00033 TValSum = (double*)malloc((nAtom + 1) * sizeof(double));
00034
00036
        if(stepCount <= stepEquil){</pre>
00037
          double gSum, varS, massS;
00038
           temperature = 1./GAMMA;
00039
00040
          if(stepCount == 1) varS = 0.;
          double A, S1, S2, T;
00041
00042
           int n;
00043
           S1 = 0.; S2 = 0.; gSum = 0.; massS = 0.1;
00044
           vvSum = 0.;
double halfdt = 0.5*deltaT;
for (n = 1; n <= nAtom; n++) {
    T = vx[n] + halfdt * ax[n];</pre>
00045
00046
00047
00048
00049
            S1 += T * ax[n];
            S2 += Sqr(T);
00050
00051
00052
             T = vy[n] + halfdt * ay[n];
            S1 += T * ay[n];
00053
00054
             S2 += Sqr(T);
00055
            vvSum += (Sqr(vx[n]) + Sqr(vy[n]));
00056
00057
00058
           A = -S1 / S2;
           S2 = vvSum;
00059
00060
00061
           double C = 1 + A*deltaT;
00062
           double D = deltaT * (1 + 0.5 * A * deltaT);
00063
00064
           int i,j;
           real dr[NDIM+1], r, rr, ri, rrCut;
00065
00066
           double vv;
00067
00068
           double uVal, AA, AASum;
00069
00070
           double TVal;
00071
           double deno, VVSum;
00072
           deno = 0.;
00073
           VVSum = 0.;
```

```
00074
            AASum = 0.;
00075
00076
         for (n=1; n<=nAtom; n++)</pre>
00077
             TValSum[n] = 0.;
00078
00079
         rrCut = Sgr(rCut);
00081 /*****Calculating Configarational temperature*****/
00082 //Solving the equation of motion here 00083 if(thermo == 'C') {
00084 for(i = 1; i <= nAtom; i ++) {
          for(j = i+1; j <= nAtom; j ++) {
  dr[1] = rx[i] - rx[j];</pre>
00085
00086
00087
               if(fabs(dr[1]) > regionH[1])
00088
            dr[1] -= SignR(region[1], dr[1]);
00089
            dr[2] = ry[i] - ry[j];
  if(fabs(dr[2]) > regionH[2])
dr[2] -= SignR(region[2], dr[2]);
00090
00091
00092
00093
00094
              rr = Sqr(dr[1]) + Sqr(dr[2]);
00095
             if(rr < rrCut ) {</pre>
00096
             r = sqrt(rr);
ri = 1/r;
00097
00098
             uVal = ri*exp(-kappa*r);
00099
00100
              TVal = (1./rr + Sqr(kappa) + kappa/r) *uVal;
00101
              TValSum[i] += TVal;
              TValSum[j] += TVal;
00102
00103
00104
             AA = Sqr(ax[i]) + Sqr(ay[i]);
00105
             AASum += AA;
00106
              vv = Sqr(vx[i]) + Sqr(vy[i]);
00107
              VVSum += vv;
00108
              deno += TValSum[i];
00109 }
00110
00111
             double gSumconfig, varSconfig, massSconfig;
00112
              if(stepCount == 1) varSconfig = 0.;
00113
              gSumconfig = 0.; massSconfig = 2.0;
00114
              gSumconfig = (AASum/temperature - deno)/massSconfig;
varSconfig += deltaT*gSumconfig;
00115
00116
00117
00118
               /*****Configarational Nose-Hoover thermostat****/
00119
           for (n = 1; n <= nAtom; n++) {
              r (n - 1; n <- natom; n++)(
vx[n] += deltaT * ax[n];
rx[n] += deltaT * (vx[n] + varSconfig * ax[n]);
vy[n] += deltaT * ay[n];
ry[n] += deltaT * (vy[n] + varSconfig * ay[n]);</pre>
00120
00121
00122
00123
00124
            }
00125
               /****Kinetic Nose-Hoover thermostat****/
         }else if(thermo == 'N') {
  gSum = (0.5*S2 - (nAtom + 1)*temperature)/massS;
00126
00127
            varS += deltaT*gSum;
00128
00129
           for (n = 1; n <= nAtom; n++) {</pre>
              vx[n] += deltaT * (ax[n] - varS *vx[n]);
rx[n] += deltaT * vx[n];
00130
00131
               vy[n] += deltaT * (ay[n] - varS *vy[n]);
ry[n] += deltaT * vy[n];
00132
00133
00134
           }
00135
               /****for Gaussian thermostat****/
00136
        }else if(thermo == 'G'){
              for (n = 1; n <= nAtom; n++) {</pre>
00137
00138
               vx[n] = C * vx[n] + D * ax[n];
              rx[n] += deltaT * vx[n];
vy[n] = C * vy[n] + D * ay[n];
00139
00140
              ry[n] += deltaT * vy[n];
00141
00142
00143
           }else if (thermo == 'L') {
         double nu = 0.03066;
double var = sqrt(2*nu/(GAMMA*deltaT));
00144
00145
         double scale = 1. + nu*deltaT/2.;
double scale_v = 2./scale - 1.;
00146
00147
00148
         double scale_f = deltaT/scale;
00149
         int n;
00150
          for (n = 1 ; n <= nAtom ; n ++) {</pre>
00151
               vx[n] = scale_v*vx[n] + scale_f*(ax[n] + var*gsl_ran_gaussian(rnd,1));
               rx[n] += deltaT * vx[n];
00152
               vy[n] = scale_v*vy[n] + scale_f*(ay[n] + var*gsl_ran_gaussian(rnd,1));
00153
               ry[n] += deltaT * vy[n];
00154
00155
            }
00156
00157
         }else{
           int n;
for(n = 1 ; n <= nAtom ; n ++) {
  vx[n] += deltaT * ax[n];</pre>
00158
00159
00160
```

3.71 LeapfrogStep.c 135

References ax, ay, deltaT, kappa, nAtom, NDIM, rCut, region, regionH, rx, ry, SignR, Sqr, stepCount, stepEquil, thermo, vvSum, vx, and vy.

3.71 LeapfrogStep.c

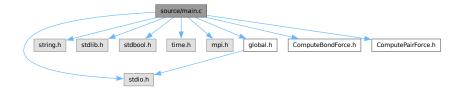
Go to the documentation of this file.

```
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00016
00017 Copyright (C) 2025 Harish Charan, University of Durham, UK
00018
00020 #include<stdio.h>
00021 #include<math.h>
00022 #include<stdlib.h>
00023 #include <gsl/gsl_rng.h>
00024 #include <gsl/gsl_randist.h>
00025 #include"global.h"
00026
00027
00028 void LeapfrogStep(char thermo, gsl_rng * rnd){
00029 double temperature, GAMMA;
00030 \text{ GAMMA} = 100;
00031
00032 double *TValSum;
00033 TValSum = (double*)malloc((nAtom + 1) * sizeof(double));
00034
00035
00036
         if(stepCount <= stepEquil){</pre>
           double gSum, varS, massS;
temperature = 1./GAMMA;
00037
00039
00040
           if(stepCount == 1) varS = 0.;
            double A, S1, S2, T;
00041
00042
            int n;
            S1 = 0.; S2 = 0.; gSum = 0.; massS = 0.1;
00043
00044
00045
            vvSum = 0.;
00046
            double halfdt = 0.5*deltaT;
00047
            for (n = 1; n <= nAtom; n++) {</pre>
00048
              T = vx[n] + halfdt * ax[n];
00049
              S1 += T * ax[n];
              S2 += Sqr(T);
00050
00051
00052
              T = vy[n] + halfdt * ay[n];
00053
              S1 += T * ay[n];
              S2 += Sqr(T);
00054
             vvSum += (Sqr(vx[n]) + Sqr(vy[n]));
00055
00056
00057
00058
            A = -S1 / S2;
00059
            S2 = vvSum;
00060
00061
            double C = 1 + A*deltaT ;
double D = deltaT * (1 + 0.5 * A * deltaT);
00062
00063
00064
00065
            real dr[NDIM+1], r, rr, ri, rrCut;
00066
            double vv;
00067
00068
            double uVal, AA, AASum;
00069
            double TVal;
00070
00071
            double deno, VVSum;
```

```
00072
            deno = 0.;
00073
            VVSum = 0.;
           AASum = 0.;
00074
00075
00076
         for (n=1; n<=nAtom; n++)</pre>
00077
             TValSum[n] = 0.;
00078
00079
         rrCut = Sqr(rCut);
08000
00081 /*****Calculating Configarational temperature*****/
00082 //Solving the equation of motion here
00083 if(thermo == 'C') {
00084 for(i = 1; i <= nAtom; i ++) {
           for(j = i+1; j <= nAtom; j ++) {
  dr[1] = rx[i] - rx[j];</pre>
00085
00086
00087
               if(fabs(dr[1]) > regionH[1])
00088
            dr[1] -= SignR(region[1], dr[1]);
00089
             dr[2] = ry[i] - ry[j];
00091
               if(fabs(dr[2]) > regionH[2])
00092
            dr[2] -= SignR(region[2], dr[2]);
00093
00094
             rr = Sqr(dr[1]) + Sqr(dr[2]);
00095
             if(rr < rrCut ) {</pre>
00096
             r = sqrt(rr);
             ri = 1/r;
00097
00098
             uVal = ri*exp(-kappa*r);
00099
             TVal = (1./rr + Sqr(kappa) + kappa/r)*uVal;
TValSum[i] += TVal;
TValSum[j] += TVal;
00100
00101
00102
00103
           } }
00104
             AA = Sqr(ax[i]) + Sqr(ay[i]);
00105
             AASum += AA;
00106
             vv = Sqr(vx[i]) + Sqr(vy[i]);
             VVSum += vv;
00107
             deno += TValSum[i];
00108
00109 }
00110
             double gSumconfig, varSconfig, massSconfig;
if(stepCount == 1) varSconfig = 0.;
gSumconfig = 0.; massSconfig = 2.0;
00111
00112
00113
00114
00115
             gSumconfig = (AASum/temperature - deno)/massSconfig;
             varSconfig += deltaT*gSumconfig;
00116
00117
00118
              /*****Configarational Nose-Hoover thermostat****/
00119
           for (n = 1; n \le nAtom; n++) {
              rx[n] += deltaT * ax[n];
rx[n] += deltaT * (vx[n] + varSconfig * ax[n]);
00120
00121
               vy[n] += deltaT * ay[n];
00122
00123
              ry[n] += deltaT * (vy[n] + varSconfig * ay[n]);
00124
00125
              /*****Kinetic Nose-Hoover thermostat****/\\
         }else if(thermo == 'N') {
   gSum = (0.5*S2 - (nAtom + 1)*temperature)/massS;
00126
00127
            varS += deltaT*gSum;
00128
00129
           for (n = 1; n <= nAtom; n++) {</pre>
              vx[n] += deltaT * (ax[n] - varS *vx[n]);
rx[n] += deltaT * vx[n];
00130
00131
              vy[n] += deltaT * (ay[n] - varS *vy[n]);
00132
              ry[n] += deltaT * vy[n];
00133
00134
          }
00135
              /*****for \ {\tt Gaussian} \ {\tt thermostat} *****/
00136
        }else if(thermo == 'G'){
              for (n = 1; n <= nAtom; n++) {</pre>
00137
              vx[n] = C * vx[n] + D * ax[n];
rx[n] += deltaT * vx[n];
00138
00139
              vy[n] = C * vy[n] + D * ay[n];
00140
              ry[n] += deltaT * vy[n];
00141
00142
          }else if (thermo == 'L') {
00143
         double nu = 0.03066;
double var = sqrt(2*nu/(GAMMA*deltaT));
double scale = 1. + nu*deltaT/2.;
double scale_v = 2./scale - 1.;
00144
00145
00146
00147
00148
         double scale_f = deltaT/scale;
         int n;
for(n = 1 ; n <= nAtom ; n ++) {</pre>
00149
00150
00151
              vx[n] = scale_v*vx[n] + scale_f*(ax[n] + var*gsl_ran_gaussian(rnd,1));
              rx[n] += deltaT * vx[n];
vy[n] = scale_v*vy[n] + scale_f*(ay[n] + var*gsl_ran_gaussian(rnd,1));
00152
00153
00154
               ry[n] += deltaT * vy[n];
00155
00156
00157
        }else{
00158
           int n:
```

3.72 source/main.c File Reference

```
#include <stdio.h>
#include <string.h>
#include <stdlib.h>
#include <stdbool.h>
#include <time.h>
#include <mpi.h>
#include "global.h"
#include "ComputeBondForce.h"
#include "ComputePairForce.h"
Include dependency graph for main.c:
```



Macros

• #define DEFINE_GLOBALS

Functions

- void Init ()
- void SetupJob ()
- void EvalSpacetimeCorr ()
- void Trajectory ()
- void DumpState ()
- void ComputeForcesCells ()
- void LeapfrogStep ()
- void BrownianStep ()
- void ApplyBoundaryCond ()
- void EvalProps ()
- void EvalVacf ()
- void EvalRdf ()
- void AccumProps (int icode)
- void PrintSummary ()
- void PrintVrms ()
- void DumpBonds ()
- void VelocityVerletStep (int icode)
- void ApplyForce ()
- void ApplyDrivingForce ()
- · void ApplyShear ()

```
· void ApplyLeesEdwardsBoundaryCond ()
```

- · void PrintStress ()
- void Close ()
- void PrintMomentum ()
- void DisplaceAtoms ()
- void DumpRestart ()
- bool HaltConditionCheck (double value, int stepCount)
- void EvalCom ()
- void PrintCom ()
- void EvalVrms ()
- void EvalUnwrap ()
- void DumpPairs ()
- void ApplyViscous ()
- int main (int argc, char **argv)

Variables

char * prefix = NULL

3.72.1 Macro Definition Documentation

3.72.1.1 DEFINE_GLOBALS

#define DEFINE_GLOBALS
Definition at line 7 of file main.c.

3.72.2 Function Documentation

3.72.2.1 AccumProps()

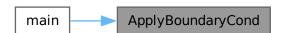
3.72.2.2 ApplyBoundaryCond()

```
void ApplyBoundaryCond ( )
Definition at line 27 of file ApplyBoundaryCond.c.
00027
00028    int n;
00029    for(n = 1 ; n <= nAtom ; n ++) {</pre>
```

```
if(strcmp(xBoundary, "p") == 0 && strcmp(yBoundary, "p") == 0){
                                                                            // P.B.C along x and y axis
         rx[n] -= region[1]*rint(rx[n]/region[1]);
ry[n] -= region[2]*rint(ry[n]/region[2]);
00031
00032
        } else if (strcmp(xBoundary, "r") == 0 && strcmp(yBoundary, "r") == 0){    //R.B.C. along x and y
00033
     axis
00034
          if((rx[n] + atomRadius[n]) >= regionH[1]){
             rx[n] = 0.999999 * regionH[1] - atomRadius[n]; vx[n] = -vx[n];
00035
00036
          }if((rx[n]-atomRadius[n]) < -regionH[1]){</pre>
00037
            rx[n] = -0.999999*regionH[1] + atomRadius[n]; vx[n] = -vx[n];
00038
00039
          if ((ry[n] + atomRadius[n]) >= regionH[2]) {
          ry[n] = 0.999999*regionH[2] - atomRadius[n]; vy[n] = -vy[n];
}if((ry[n]-atomRadius[n]) < -regionH[2]){</pre>
00040
00041
00042
            ry[n] = -0.999999*regionH[2] + atomRadius[n]; vy[n] = -vy[n];
00043
00044
         along y axis
00045
         rx[n] -= region[1]*rint(rx[n]/region[1]);
         if((ry[n] + atomRadius[n]) >= regionH[2]){
00046
           ry[n] = 0.999999*regionH[2] - atomRadius[n]; vy[n] = -vy[n];
00047
00048
          }if((ry[n] - atomRadius[n]) < -regionH[2]){</pre>
00049
             ry[n] = -0.999999*regionH[2] + atomRadius[n]; vy[n] = -vy[n];
00050
         else if(strcmp(xBoundary, "r") == 0 && strcmp(yBoundary, "p") == 0){      //R.B.C. along x and P.B.C
00051
     along y axis
00052
        if((rx[n] + atomRadius[n]) >= regionH[1]){
          00053
00054
00055
           rx[n] = -0.999999*regionH[1] + atomRadius[n]; vx[n] = -vx[n];
00056
00057
          ry[n] -= region[2] *rint(ry[n]/region[2]);
00058
       } else
00059
        // Print error message and exit the program
00060
          fprintf (fpresult, "Error: Invalid boundary configuration: '%s %s' \n", xBoundary, yBoundary); \\
00061
         exit(EXIT_FAILURE); // Exit with failure status
00062
00063
      }
00064 }
```

References atomRadius, fpresult, nAtom, region, regionH, rx, ry, vx, vy, xBoundary, and yBoundary. Referenced by main().

Here is the caller graph for this function:



3.72.2.3 ApplyDrivingForce()

void ApplyDrivingForce ()

Definition at line 25 of file ApplyDrivingForce.c.

```
00026
       int n;
00027 double Vxblock, Vyblock;
      double Vxsubstrate, Vysubstrate;
00029
       Vxblock = 0.0; Vyblock = 0.0;
00030 Vxsubstrate = 0.0; Vysubstrate = 0.0;
00031 double gammav;
00032 gammav = 0.0;
00033
00034
       double count_substrate = 0;
00035
       double count_block = 0;
00036
00037
       for (n = 1 ; n \le nAtom; n ++) {
        if(atomType[n] == 1 || atomType[n] == 2){
Vxsubstrate += vx[n]; Vysubstrate += vy[n];
00038
00039
        count_substrate++;
00040
00041
00042
         if(atomType[n] == 3 || atomType[n] == 4){
        Vxblock += vx[n]; Vyblock += vy[n];
00043
        count_block++;
00044
```

```
00045
          } }
00046
00047
          if(count_substrate > 0) {
             Vxsubstrate /= count_substrate;
Vysubstrate /= count_substrate;
00048
00049
00050
00051
00052
          if(count_block > 0) {
          Vxblock /= count_block;
Vyblock /= count_block;
00053
00054
00055
00056
00057
         for (n = 1 ; n \le nAtom; n ++) {
00058
          if(atomType[n] == 1 || atomType[n] == 2){
          ax[n] += -gammav * (vx[n] - Vxsubstrate);
ay[n] += -gammav * (vy[n] - Vysubstrate);
00059
00060
00061
00062
          if(atomType[n] == 3 || atomType[n] == 4){
ax[n] += -gammav * (vx[n] - Vxblock);
00063
          ay[n] += -gammav * (vy[n] - Vyblock);
00064
00065 } } }
```

References atomType, ax, ay, nAtom, vx, and vy.

3.72.2.4 ApplyForce()

```
void ApplyForce ( )
```

Definition at line 25 of file ApplyForce.c.

```
00025 {
00026 int n;
00027 double lx;
00028 lx = regionH[1];
00029 fy = (FyBylx * lx)/nAtomBlock;
00030 fx = fxByfy * fy;
00031 for(n = 1; n <= nAtom; n ++) {
00032 if(molID[n] == 2) {
00033 ax[n] += fx;
00034 ay[n] -= fy;
```

 $References\ ax,\ ay,\ fx,\ fxByfy,\ fy,\ FyBylx,\ molID,\ nAtom,\ nAtomBlock,\ and\ regionH.$

Referenced by main().

Here is the caller graph for this function:



3.72.2.5 ApplyLeesEdwardsBoundaryCond()

void ApplyLeesEdwardsBoundaryCond ()

Definition at line 25 of file ApplyLeesEdwardsBoundaryCond.c.

```
00025
00026 int n;
          for (n = 1; n <= nAtom; n++) {</pre>
00027
00028 //PBC along x-direction
00029 if(rx[n] >= regionH[1])
00030 rx[n] -= region[1];
00031 else if(rx[n] < -regionH[1])
00032
            rx[n] += region[1];
00033
00034 //LEBC along y-direction
           if(ry[n] >= regionH[2]) {
  rx[n] -= shearDisplacement;
00035
00036
00037
             if(rx[n] < -regionH[1]) rx[n] += region[1];</pre>
          //vx[n] -= shearVelocity;
ry[n] -= region[2];
}else if(ry[n] < -regionH[2]) {
rx[n] += shearDisplacement;</pre>
00038
00039
00040
00041
             if(rx[n] >= regionH[1]) rx[n] -= region[1];
```

References nAtom, region, regionH, rx, ry, and shearDisplacement.

3.72.2.6 ApplyShear()

References nAtom, rx, ry, and strain.

3.72.2.7 ApplyViscous()

3.72.2.8 BrownianStep()
void BrownianStep ()

Definition at line 26 of file BrownianStep.c.

```
if(stepCount <= stepEquil){</pre>
00027
00028
           double A, S1, S2, T;
00029
           int n;
00030
           S1 = 0.; S2 = 0;
00031
           double halfdt = 0.5*deltaT;
00032
           for (n = 1; n <= nAtom; n++) {</pre>
            T = vx[n] + halfdt * ax[n];
00033
00034
             S1 += T * ax[n];
             S2 += Sqr(T);
00035
00036
00037
             T = vy[n] + halfdt * ay[n];
S1 += T * ay[n];
00039
              S2 += Sqr(T);
00040
           A = -S1 / S2;
00041
           double C = 1 + A*deltaT;
double D = deltaT * (1 + 0.5 * A * deltaT);
00042
00043
00044
           for (n = 1; n <= nAtom; n++) {</pre>
00045
             vx[n] = C * vx[n] + D * ax[n];
              rx[n] += deltaT * vx[n];
vy[n] = C * vy[n] + D * ay[n];
ry[n] += deltaT * vy[n];
00046
00047
00048
00049
00050
         }else{
00051
             //SETTING TEMP = 0.0
if (stepCount == stepEquil+1) {
00052
00053
             for(n = 1; n <= nAtom; n ++) {
vx[n] = 0.0;</pre>
00054
00055
00056
             vy[n] = 0.0;
00057
00058
             double zeta = 1.0;
00059
             double dx, dy;
00060
             for(n = 1; n <= nAtom; n ++) {
00061
              dx = rx[n];
00062
              rx[n] += zeta * ax[n] * deltaT;
00063
              dx = rx[n] - dx;
00064
              vx[n] = dx/deltaT;
00065
              dy = ry[n];
              ry[n] += zeta * ay[n] * deltaT;
00066
00067
              dy = ry[n] - dy;
```

References ax, ay, deltaT, nAtom, rx, ry, Sqr, stepCount, stepEquil, vx, and vy.

3.72.2.9 Close()

```
void Close ( )
Definition at line 24 of file Close.c.
00024
00025
        int n;
00026
        free(rx);
00027
        free(ry);
00028
        free(vx);
00029
        free(vy);
00030
        free (ax):
00031
        free(av);
00032
        free (fax);
00033
        free(fay);
00034
        free(cellList);
00035
00036
        free(atomID); free(atomType); free(atomRadius); free(atomMass);
00037
        free (speed):
00038
        free(atom1); free(atom2); free(BondID);
00039
        free(BondType); free(kb); free(ro);
00040
        free(ImageX); free(ImageY); free(rxUnwrap); free(ryUnwrap);
00041
        free(atomIDInterface);
        free(PairID); free(Pairatom1); free(Pairatom2);
00042
00043
        free(PairXij); free(PairYij);
00044
00045
        free(DeltaXijOld);
00046
        free (DeltaYijOld);
00047
00048
        free (molID);
00049
00050
        for (n = 0; n <= nAtom; n++) {</pre>
         free(isBonded[n]);
00052
00053
         free (isBonded);
00054
00055
00056
00057
        for(n = 0; n <= nAtom; n++) {</pre>
00058
         free (DeltaXijOldPair[n]);
00059
         free (DeltaYijOldPair[n]);
00060
00061
          free(DeltaXijOldPair);
00062
          free(DeltaYijOldPair);
00063
00064
        for (n = 0; n <= nBuffCorr; n++) {</pre>
00065
          free(cfOrg[n]);
00066
          free(spacetimeCorr[n]);
00067
       free(cfOrg);
00068
00069
        free(spacetimeCorr);
00070
        free(cfVal);
00071
        free(indexCorr);
00072
        free(spacetimeCorrAv);
00073
00074
        free (indexAcf):
00075
        free(viscAcfOrg);
00076
        free(viscAcfAv);
00077
        for(n = 0 ; n \le nBuffAcf ; n ++)
00078
          free(viscAcf[n]);
00079
        free(viscAcf);
00080
00081 }
```

References atom1, atom2, atomID, atomIDInterface, atomMass, atomRadius, atomType, ax, ay, BondID, BondType, cellList, cfOrg, cfVal, DeltaXijOld, DeltaXijOldPair, DeltaYijOld, DeltaYijOldPair, fax, fay, ImageX, ImageY, indexAcf, indexCorr, isBonded, kb, molID, nAtom, nBuffAcf, nBuffCorr, Pairatom1, Pairatom2, PairID, PairXij, PairYij, ro, rx, rxUnwrap, ry, ryUnwrap, spacetimeCorr, spacetimeCorrAv, speed, viscAcf, viscAcfAv, viscAcfOrg, vx, and vy.

Referenced by main().

Here is the caller graph for this function:



3.72.2.10 ComputeForcesCells()

```
void ComputeForcesCells ( )
Definition at line 25 of file ComputeForcesCells.c.
 00025
                         double dr[NDIM+1], invWid[NDIM+1], shift[NDIM+1], f, fcVal, rr, ri, r, uVal;
 00026
                         int c, I, J, m1, m1x, m1y, m2, m2x, m2y, n, offset; int iofX[] = {0, 0, 1, 1, 0, -1, -1, -1, 0, 1}, iofY[] = {0, 0, 0, 1, 1, 1, 0, -1, -1, -1, -1};
 00027
 00028
 00029
 00030
 00031
                         invWid[1] = cells[1]/region[1];
                        invWid[2] = cells[2]/region[2];
00032
00033
 00034
                        for(n = nAtom+1; n <= nAtom+cells[1]*cells[2]; n++)</pre>
                              cellList[n] = 0;
 00035
 00036
00037
                         for(n = 1 ; n <= nAtom ; n ++) {</pre>
00038
                            \texttt{c} = ((\texttt{int})((\texttt{ry}[\texttt{n}] + \texttt{regionH}[\texttt{2}]) * \texttt{invWid}[\texttt{2}])) * \texttt{cells}[\texttt{1}] + (\texttt{int})((\texttt{rx}[\texttt{n}] + \texttt{regionH}[\texttt{1}]) * \texttt{invWid}[\texttt{1}]) + (\texttt{int})((\texttt{ry}[\texttt{n}] + \texttt{regionH}[\texttt{1}]) * \texttt{invWid}[\texttt{1}]) + (\texttt{int})((\texttt{ry}[\texttt{n}] + \texttt{regionH}[\texttt{1}]) * \texttt{invWid}[\texttt{1}]) + (\texttt{int})((\texttt{ry}[\texttt{n}] + \texttt{regionH}[\texttt{1}]) * \texttt{invWid}[\texttt{1}])) + (\texttt{int})((\texttt{ry}[\texttt{n}] + \texttt{regionH}[\texttt{1}])) + (\texttt{int})((\texttt{n})(\texttt{n})) + (\texttt{int})((\texttt{n})(\texttt{n})) + (\texttt{int})((\texttt{n})(\texttt{n}))) + (\texttt{int})((\texttt{n})(\texttt{n})(\texttt{n})(\texttt{n})) + (\texttt{int})((\texttt{n})(\texttt{n})(\texttt{n}))) + (\texttt{int})((\texttt{n})(\texttt{n})(\texttt{n})(\texttt{n})) + (\texttt{int})((\texttt{n})(\texttt{n})(\texttt{n})(\texttt{n})) + (\texttt{int})((\texttt{n})(\texttt{n})(\texttt{n})(\texttt{n})) + (\texttt{int})((\texttt{n})(\texttt{n})(\texttt{n})(\texttt{n})(\texttt{n})) + (\texttt{int})((\texttt{n})(\texttt{n})(\texttt{n})(\texttt{n})(\texttt{n})) + (\texttt{int})((\texttt{n})(\texttt{n})(\texttt{n})(\texttt{n})(\texttt{n})) + (\texttt{int})((\texttt{n})(\texttt{n})(\texttt{n})(\texttt{n})(\texttt{n})(\texttt{n})(\texttt{n})) + (\texttt{int})((\texttt{n})(\texttt{n})(\texttt{n})(\texttt{n})(\texttt{n})(\texttt{n})) + (\texttt{int})((\texttt{n})(\texttt{n
                  nAtom+ 1;
 00039
                         cellList[n] = cellList[c];
 00040
                               cellList[c] = n;
 00041
 00042
                         for(n = 1; n <= nAtom; n ++) {
  ax[n] = 0.;</pre>
 00043
 00044
 00045
                              ay[n] = 0.;
 00046
 00047
 00048
                         uSum = 0.0;
                        virSum = 0.0;
rfAtom = 0.0;
 00049
00050
                         RadiusIJ = 0.0;
 00051
 00052
 00053
 00054
                         double vr[NDIM+1], fd, fdVal, rrinv;
00055
                        rrinv = 0.0;
                         fd = 0.0;
 00056
 00057
                         fdVal = 0.0;
 00058
 00059
                         int start = 1 + rank*(cells[2]/size);
 00060
                        int end = (rank+1) * (cells[2]/size);
 00061
 00062
                         for(m1Y = start ; m1Y \le end ; m1Y ++) {
                               for (m1X = 1; m1X <= cells[1]; m1X ++) {
    m1 = (m1Y-1) * cells[1] + m1X + nAtom;
    for (offset = 1; offset <= 9; offset ++) {
 00063
 00064
 00065
 00066
                                m2X = m1X + iofX[offset]; shift[1] = 0.;
                               if(m2X > cells[1]) {
  m2X = 1; shift[1] = region[1];
 00067
 00068
                               }else if(m2X == 0){
  m2X = cells[1]; shift[1] = -region[1];
00069
 00070
 00071
 00072
                                m2Y = m1Y + iofY[offset]; shift[2] = 0.;
 00073
                                if(m2Y > cells[2]){
00074
                                    m2Y = 1; shift[2] = region[2];
 00075
                                else if (m2Y == 0) {
 00076
                                     m2Y = cells[2]; shift[2] = -region[2];
 00077
 00078
                               m2 = (m2Y-1)*cells[1] + m2X + nAtom;
 00079
                                I = cellList[m1];
 08000
                                while (I > 0) {
                                     J = cellList[m2];
 00081
 00082
                                      while (J > 0) {
 00083
                                             if(m1 == m2 && J != I && (atomRadius[I] > 0. && atomRadius[J] > 0.)){
                                                   dr[1] = rx[I] - rx[J] - shift[1];
                                                   dr[2] = ry[I] - ry[J] - shift[2];
00085
```

```
rr = Sqr(dr[1]) + Sqr(dr[2]);
00087
                  RadiusIJ = atomRadius[I] + atomRadius[J];
                  SqrRadiusIJ = Sqr(RadiusIJ);
00088
                  if(rr < SqrRadiusIJ) {</pre>
00089
00090
                r = sqrt(rr);
00091
                ri = 1.0/r;
00092
                        rrinv = 1.0/rr;
                        vr[1] = vx[I] - vx[J];
vr[2] = vy[I] - vy[J];
00093
00094
00095
                RadiusIJInv = 1.0/RadiusIJ;
                uVal = Sqr(1.0 - r * RadiusIJInv);
00096
                fcVal = 2.0 * RadiusIJInv * (1.0 - r * RadiusIJInv) *ri;
00097
00098
                         fdVal = -gamman * (vr[1]*dr[1] + vr[2]*dr[2]) * rrinv; //disc-disc drag
00099
00100
                f = fcVal * dr[1];
                fd = fdVal * dr[1];
ax[I] += (f + fd);
00101
00102
                        discDragx[I] += fd; //disc-disc drag
00103
00104
00105
                f = fcVal * dr[2];
00106
                        fd = fdVal * dr[2];
                ay[I] += (f + fd);
00107
                        discDragy[I] += fd; //disc-disc drag
00108
00109
                uSum += 0.5 * uVal;
00110
                virSum += 0.5 * fcVal * rr;
rfAtom += 0.5 * dr[1] * fcVal * dr[2];
00111
00112
00113
00114
                }else if(m1 != m2 && (atomRadius[I] > 0. && atomRadius[J] > 0.)){
                  dr[1] = rx[1] - rx[3] - shift[1];
dr[2] = ry[1] - ry[3] - shift[2];
rr = Sqr(dr[1]) + Sqr(dr[2]);
00115
00116
00117
00118
                  RadiusIJ = atomRadius[I] + atomRadius[J];
00119
                  SqrRadiusIJ = Sqr(RadiusIJ);
00120
                  if(rr < SqrRadiusIJ){</pre>
00121
                r = sqrt(rr);
               ri = 1.0/r;
00122
                        rrinv = 1.0/r;
                         vr[1] = vx[I] - vx[J];
vr[2] = vy[I] - vy[J];
00124
00125
00126
                RadiusIJInv = 1.0/RadiusIJ;
                uVal = Sqr(1.0 - r * RadiusIJInv);
fcVal = 2.0 * RadiusIJInv * (1.0 - r * RadiusIJInv) *ri;
00127
00128
                         fdVal = -gamman * (vr[1]*dr[1] + vr[2]*dr[2]) * rrinv; //disc-disc drag
00129
00130
00131
                f = fcVal * dr[1];
                fd = fdVal * dr[1];

ax[I] += (f + fd);
00132
00133
                        discDragx[I] += fd; //disc-disc drag
00134
00135
00136
                f = fcVal * dr[2];
00137
                        fd = fdVal * dr[2];
00138
                ay[I] += (f + fd);
00139
                        discDragy[I] += fd; //disc-disc drag
00140
00141
                uSum += 0.5 * uVal;
                virSum += 0.5 * fcVal * rr;
00143
                rfAtom += 0.5 * dr[1] * fcVal * dr[2];
00144
                 }
               }
00145
                    J = cellList[J];
00146
00147
00148
             I = cellList[I];
00149
           }
00150
00151
00152
00153 }
```

References atomRadius, ax, ay, cellList, cells, discDragx, discDragy, gamman, nAtom, NDIM, RadiusIJ, RadiusIJInv, rank, region, regionH, rfAtom, rx, ry, size, Sqr, SqrRadiusIJ, uSum, virSum, vx, and vy.

3.72.2.11 DisplaceAtoms()

Referenced by main().

Here is the caller graph for this function:



3.72.2.12 DumpBonds()

```
void DumpBonds ( )
```

Definition at line 24 of file DumpBonds.c.

```
00025
            //Trajectory file in LAMMPS dump format for OVITO visualization fprintf(fpbond, "ITEM: TIMESTEP\n"); fprintf(fpbond, "%lf\n",timeNow);
00026
00027
00028
            fprintf(fpbond, "ITEM: NUMBER OF ENTRIES\n");
            fprintf(fpbond, "%d\n", nBond);
00030
           fprintf(fpbond, "%d\n",nBond);
fprintf(fpbond, "ITEM: BOX BOUNDS pp ff pp\n");
fprintf(fpbond, "%lf %lf xlo xhi\n", -regionH[1], regionH[1]);
fprintf(fpbond, "%lf %lf ylo yhi\n", -regionH[2], regionH[2]);
fprintf(fpbond, "%lf %lf zlo zhi\n", -0.1, 0.1);
00031
00032
00033
00034
00035
            fprintf(fpbond, "ITEM: ENTRIES BondID, BondType, atom1 atom2 BondLength BondLengthEqul nodeDragx1
         nodeDragy1\n");
00036
            for(n=1; n<=nBond; n++)
fprintf(fpbond, "%d %d %d %d %0.16lf %0.16lf %0.16lf %0.16lf \n", BondID[n], BondType[n], atoml[n],</pre>
00037
00038
         atom2[n],
00039
             BondLength[n], ro[n], nodeDragx[atom1[n]], nodeDragy[atom1[n]]);
```

References atom1, atom2, BondID, BondLength, BondType, fpbond, nBond, nodeDragx, nodeDragy, regionH, ro, and timeNow.

Referenced by main().

Here is the caller graph for this function:



3.72.2.13 DumpPairs()

```
void DumpPairs ( )
```

```
Definition at line 25 of file DumpPairs.c.
00025
00026
            int n;
            //Trajectory file in LAMMPS dump format for OVITO visualization
00027
            fprintf(fppair, "ITEM: NUMBER OF ENTRIES\n");
fprintf(fppair, "$1f\n",timeNow);
fprintf(fppair, "ITEM: NUMBER OF ENTRIES\n");
00028
00029
00030
            fprintf(fppair, "%d\n", nPairActive);
00031
            fprintf(fppair, "ITEM: BOX BOUNDS pp ff pp\n");
fprintf(fppair, "%lf %lf xlo xhi\n", -regionH[1], regionH[1]);
fprintf(fppair, "%lf %lf ylo yhi\n", -regionH[2], regionH[2]);
00032
00033
00034
00035
            fprintf(fppair, "%lf %lf zlo zhi\n", -0.1, 0.1);
```

References discDragx, discDragy, fppair, nPairActive, Pairatom1, Pairatom2, PairID, PairXij, PairYij, regionH, and timeNow.

Referenced by main().

Here is the caller graph for this function:



3.72.2.14 DumpRestart()

```
void DumpRestart ( )
Definition at line 25 of file DumpRestart.c.
00025
00026
         char DUMP[256];
00027 FILE *fpDUMP;
        sprintf(DUMP, "%s.Restart", prefix);
00029
         fpDUMP = fopen(DUMP, "w");
00030
         if(fpDUMP == NULL) {
00031
          fprintf(stderr, "Error opening file %s for writing\n", DUMP);
00032
          return;
00033
00034
00035
          fprintf(fpDUMP, "timeNow %lf\n", timeNow);
          fprintf(fpDUMP, "nAtom %d\n", nAtom);
fprintf(fpDUMP, "nBond %d\n", nBond);
00036
00037
          fprintf(fpDUMP, "nBond %d\n", nBond);
fprintf(fpDUMP, "nAtomType %d\n", nAtomType);
fprintf(fpDUMP, "nBondType %d\n", nBondType);
fprintf(fpDUMP, "region[1] %0.141f\n", region[1]);
fprintf(fpDUMP, "region[2] %0.141f\n", region[2]);
00038
00039
00040
00041
00042
00043
          fprintf(fpDUMP, "Atoms\n");
for(n = 1; n <= nAtom; n ++)</pre>
00044
00045
           fprintf(fpDUMP, "%d %d %d %0.21f %0.161f %0.161f %0.161f %0.161f\n", atomID[n], molID[n],
00046
       atomType[n], atomRadius[n], rx[n], ry[n], vx[n], vy[n]);
00047
00048
00049
         fprintf(fpDUMP, "Bonds\n");
          for (n=1; n<=nBond; n++)</pre>
00050
                                "%d %d %d %0.21f %0.161f\n", BondID[n], BondType[n], atom1[n], atom2[n], kb[n],
00051
         fprintf(fpDUMP,
       ro[n]);
00052
00053
          fclose(fpDUMP);
00054 }
```

References atom1, atom2, atomID, atomRadius, atomType, BondID, BondType, kb, molID, nAtom, nAtomType, nBond, nBondType, prefix, region, ro, rx, ry, timeNow, vx, and vy. Referenced by main().

Here is the caller graph for this function:



3.72.2.15 DumpState()

```
void DumpState ( )
Definition at line 25 of file DumpState.c.
00025
           char DUMP[256];
00026
00027
           FILE *fpDUMP;
           sprintf(DUMP, "%s.STATE", prefix);
00028
00029
           fpDUMP = fopen(DUMP, "w");
00030 if(fpDUMP == NULL) {
00031
            fprintf(stderr, "Error opening file %s for writing\n", DUMP);
00032
00033 }
00034
00035
            fprintf(fpDUMP, "ITEM: TIMESTEP\n");
           fprintf(fpDUMP, "TTEM: TIMESTEP\n");
fprintf(fpDUMP, "%lf\n",timeNow);
fprintf(fpDUMP, "ITEM: NUMBER OF ATOMS\n");
fprintf(fpDUMP, "%d\n", nAtom);
fprintf(fpDUMP, "ITEM: BOX BOUNDS pp pp pp\n");
fprintf(fpDUMP, "%lf %lf xlo xhi\n", -regionH[1], regionH[1]);
fprintf(fpDUMP, "%lf %lf ylo yhi\n", -regionH[2], regionH[2]);
fprintf(fpDUMP, "%lf %lf zlo zhi\n", -0.1, 0.1);
fprintf(fpDUMP, "ITEM: ATOMS id mol type radius x y vx vy fx fy\n");
int n:
00036
00037
00038
00039
00040
00041
00042
00043
00044
            int n;
            for (n = 1; n <= nAtom; n++) {
    fprintf(fpDUMP, "%d\t %d\t %d\t %0.21f\t %0.161f\t %0.161f\t %0.161f\t %0.161f\t %0.161f\t</pre>
00045
00046
         %0.161f\n"
00047
             atomID[n], molID[n], atomType[n], atomRadius[n], rx[n], ry[n], vx[n], vy[n], ax[n], ay[n]);
00048
00049
            fclose(fpDUMP);
00050 }
```

References atomID, atomRadius, atomType, ax, ay, molID, nAtom, prefix, regionH, rx, ry, timeNow, vx, and vy. Referenced by main().

Here is the caller graph for this function:



3.72.2.16 EvalCom()

```
00033 if(molID[n] == 2){
        ComX += atomMass[n] * rxUnwrap[n];
ComY += atomMass[n] * ryUnwrap[n];
00034
00035
00036
        TotalMass += atomMass[n];
00037
00038
        ComX = ComX/TotalMass;
00039
00040
        ComY = ComY/TotalMass;
00041
        if(timeNow == 0.0){
00042
00043
        ComX0 = ComX; ComY0 = ComY;
00044
00045
        ComXRatio = ComX/ComX0;
                                    ComYRatio = ComY/ComY0;
00046 }
```

References atomMass, ComX, ComX0, ComXRatio, ComY, ComY0, ComYRatio, molID, nAtom, rxUnwrap, ryUnwrap, timeNow, and TotalMass.

Referenced by main().

Here is the caller graph for this function:



3.72.2.17 EvalProps()

```
void EvalProps ( )
Definition at line 26 of file EvalProps.c.
00026
00027
00027 real v, vv;
00028 virSum = 0.0;
00029 vSumX = 0.0; vSumY = 0.0; vSum = 0.0;
00030 vvSum = 0.;
00031
        int n;
00032
00033 for (n = 1; n \le nAtom; n++) {
00034
        vv = 0.;
00035
        // Initialize v with a default value to avoid "uninitialized" warning.
00036
        v = 0.0;
00037
        // X direction velocity
00038
        if (strcmp(solver, "Verlet") == 0) {
        v = vx[n];
} else if (strcmp(solver, "LeapFrog") == 0) {
v = vx[n] - 0.5 * deltaT * ax[n];
00039
00040
00041
00042
         }
00043
          vSum += v;
00044
          vv += Sqr(v);
          vSumX += v;
00045
00046
          // Y direction velocity
          if (strcmp(solver, "Verlet") == 0) {
00047
          v = vy[n];
00048
00049
          } else if (strcmp(solver, "LeapFrog") == 0) {
00050
          v = vy[n] - 0.5 * deltaT * ay[n];
00051
00052
          vSum += v:
00053
          vSumY += v;
00054
          vv += Sqr(v);
00055
          vvSum += vv;
00056
00057
        kinEnergy = 0.5 * vvSum / nAtom ;
uSumPairPerAtom = uSumPair / nAtom ;
00058
00059
00060
         BondEnergyPerAtom = TotalBondEnergy / (0.5*nAtom); //Factor of 0.5 since each atom has one half the
      bond energy
00061
         potEnergy = uSumPairPerAtom + BondEnergyPerAtom ;
00062
         totEnergy = kinEnergy + potEnergy;
         virSumxy = virSumPairxx + virSumBondxx;
virSumyy = virSumPairxy + virSumBondxy;
virSumxy = virSumPairxy + virSumBondxy;
00063
00064
00065
        virSum = virSumPair + virSumBond;
00067
        pressure = density * (vvSum + virSum) / (nAtom * NDIM);
```

```
00068
```

References ax, ay, BondEnergyPerAtom, deltaT, density, kinEnergy, nAtom, NDIM, potEnergy, pressure, solver, Sqr, TotalBondEnergy, totEnergy, uSumPair, uSumPairPerAtom, virSum, virSumBond, virSumBondxx, virSumBondxy, virSumBondyy, virSumPair, virSumPairx, virSumPairxy, virSumPairy, virSumPairy, virSumxx, virSumxy, virSumy, vSum, vSumX, vSumY, vvSum, vx, and vy.

Referenced by main().

Here is the caller graph for this function:



3.72.2.18 EvalRdf()

```
void EvalRdf ( )
Definition at line 26 of file EvalRdf.c.
          real dr[NDIM+1], deltaR, normFac, rr, rrRange;
         int j1, j2, n;
countRdf ++;
00028
00029
         if (countRdf == 1) {
00030
00031
           for (n = 1 ; n \le sizeHistRdf ; n ++)
              histRdf[n] = 0.;
00033
00034
          rrRange = Sqr(rangeRdf);
         deltaR = rangeRdf / sizeHistRdf;
for(j1 = 1; j1 <= nAtom - 1; j1 ++) {
  for(j2 = j1 + 1; j2 <= nAtom; j2 ++) {</pre>
00035
00036
00037
00038
              dr[1] = rx[j1] - rx[j2];
if(fabs(dr[1]) > regionH[1])
00039
00040
00041
            dr[1] -= SignR(region[1], dr[1]);
00042
              dr[2] = ry[j1] - ry[j2];
if(fabs(dr[2]) > regionH[2])
00043
00044
00045
            dr[2] -= SignR(region[2], dr[2]);
00046
00047
              rr = Sqr(dr[1]) + Sqr(dr[2]);
00048
00049
              if(rr < rrRange){
00050
            n = (int) (sqrt(rr)/deltaR) + 1;
            histRdf[n] ++;
00052
00053
00054
         }
00055
00056
         if(countRdf == limitRdf){
           normFac = region[1]*region[2] / (M_PI*Sqr(deltaR)*nAtom*nAtom*countRdf );
00058
            for (n = 1 ; n \le sizeHistRdf ; n ++)
00059
            histRdf[n] *= normFac/(n-0.5);
// PRINT THE RADIAL DISTRIBUTION DATA ON TO DISK FILE
00060
00061
            real rBin;
00062
            int n;
00063
            fprintf(fprdf,"rdf @ timeNow %lf\n", timeNow);
00064
            for (n = 1 ; n <= sizeHistRdf ; n ++) {</pre>
              rBin = (n - 0.5)*rangeRdf/sizeHistRdf;
fprintf(fprdf, "%lf %lf\n", rBin, histRdf[n]);
00065
00066
00067
00068
         }
00069
```

References countRdf, fprdf, histRdf, limitRdf, nAtom, NDIM, rangeRdf, region, regionH, rx, ry, SignR, sizeHistRdf, Sqr, and timeNow.

3.72.2.19 EvalSpacetimeCorr()

```
void EvalSpacetimeCorr ( )
Definition at line 26 of file EvalSpacetimeCorr.c.
          real cosV=0., cosV0=0., cosV1=0., cosV2=0., sinV=0., sinV1=0., sinV2=0.;
real COSA, SINA, COSV, SINV;
00027
00028
         int j, m, n, nb, ni, nv;
real kMin = 2. * M_PI / region[1];
00029
00030
00031
          real kMax = M_PI;
00032
          real deltaK = (kMax - kMin) / nFunCorr;
00033
         for (j = 1; j <= 2*nFunCorr; j++)
  cfVal[j] = 0.;</pre>
00034
00035
00036
00037
          for (n = 1; n <= nAtom; n++) {</pre>
00038
            j = 1;
00039
             COSA = cos(kMin*rx[n]);
             SINA = sin(kMin*rx[n]);
00040
            for (m = 1; m <= nFunCorr; m++) {
  if (m == 1) {
00041
00042
00043
            cosV = cos(deltaK*rx[n]);
00044
            sinV = sin(deltaK*rx[n]);
00045
             cosV0 = cosV;
              }else if(m == 2){
00046
00047
             cosV1 = cosV;
            sinV1 = sinV;
00048
            cosV = 2.*cosV0*cosV1-1;
00049
            sinV = 2.*cosV0*sinV1;
00050
00051
              }else{
00052
             cosV2 = cosV1;
00053
            sinV2 = sinV1;
             cosV1 = cosV;
00054
00055
             sinV1 = sinV;
             cosV = 2.*cosV0*cosV1-cosV2;
00056
            sinV = 2.*cosV0*sinV1-sinV2;
00057
00058
               COSV = COSA*cosV - SINA*sinV;
00059
               SINV = SINA*cosV + COSA*sinV;
00060
               cfVal[j] += COSV;
cfVal[j+1] += SINV;
00061
00062
00063
               j += 2;
00064
00065
00066
00067
          for (nb = 1; nb <= nBuffCorr; nb++) {</pre>
           indexCorr[nb] += 1;
             if (indexCorr[nb] <= 0) continue;</pre>
00070
             ni = nFunCorr * (indexCorr[nb] - 1);
00071
             if (indexCorr[nb] == 1) {
             for (j = 1; j <= 2*nFunCorr; j++)
cfOrg[nb][j] = cfVal[j];</pre>
00072
00073
00074
00075
00076
             for (j = 1; j <= nFunCorr; j++)</pre>
00077
              spacetimeCorr[nb][ni + j] = 0.;
00078
00079
             \dot{1} = 1:
             for (m = 1; m <= nFunCorr; m++) {</pre>
08000
               nv = m + ni;
00081
00082
               spacetimeCorr[nb][nv] += cfVal[j] * cfOrg[nb][j] + cfVal[j + 1] * cfOrg[nb][j + 1];
00083
               j += 2;
00084
             }
00085
00086
00088
          // ACCUMULATE SPACETIME CORRELATIONS
00089
          for (nb = 1; nb <= nBuffCorr; nb++) {</pre>
00090
           if (indexCorr[nb] == nValCorr){
              for (j = 1; j <= nFunCorr*nValCorr; j++)
   spacetimeCorrAv[j] += spacetimeCorr[nb][j];</pre>
00091
00092
              indexCorr[nb] = 0.;
00093
00094
              countCorrAv ++;
00095
              if (countCorrAv == limitCorrAv) {
00096
                for (j = 1; j <= nFunCorr*nValCorr; j++)</pre>
              spacetimeCorrAv[j] /= (nAtom*limitCorrAv);
fprintf(fpdnsty, "NDIM %d\n", NDIM);
fprintf(fpdnsty, "nAtom %d\n", nAtom);
00097
00098
00099
                iprintf(fpdnsty, "nAtom %q\n", nAtom);
fprintf(fpdnsty, "region %lf\n", region[1]);
fprintf(fpdnsty, "nFunCorr %d\n", nFunCorr);
fprintf(fpdnsty, "limitCorrAv %d\n", limitCorrAv);
fprintf(fpdnsty, "stepCorr %d\n", stepCorr);
fprintf(fpdnsty, "nValCorr %d\n", nValCorr);
00100
00101
00102
00103
00104
                 fprintf(fpdnsty, "deltaT %lf\n", deltaT);
00105
                 real tVal;
                 for (n = 1; n <= nValCorr; n++) {</pre>
00107
```

```
tVal = (n-1) *stepCorr*deltaT;
          00109
00110
00111
00112
00113
00114
00115
          countCorrAv = 0.;
for (j = 1; j <= nFunCorr*nValCorr; j++)
spacetimeCorrAv[j] = 0.;</pre>
00116
00117
00118
00119
00120
        }
00121
00122 }
```

References cfOrg, cfVal, countCorrAv, deltaT, fpdnsty, indexCorr, limitCorrAv, nAtom, nBuffCorr, NDIM, nFunCorr, nValCorr, region, rx, spacetimeCorr, spacetimeCorrAv, and stepCorr.

3.72.2.20 EvalUnwrap()

References ImageX, ImageY, nAtom, region, rx, rxUnwrap, ry, and ryUnwrap.

Referenced by main().

Here is the caller graph for this function:

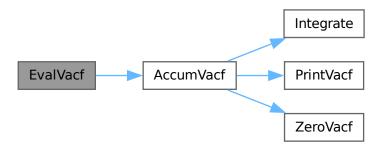


3.72.2.21 EvalVacf()

```
void EvalVacf ( )
Definition at line 26 of file EvalVacf.c.
00026
         int n, nb, ni;
00028
         double viscVec = 0.;
00029
         double v[3];
00030
         for (n = 1 ; n <= nAtom ; n ++) {</pre>
          v[1] = vx[n] - 0.5*ax[n]*deltaT;
v[2] = vy[n] - 0.5*ay[n]*deltaT;
00031
00032
00033
           viscVec += v[1]*v[2];
00034
00035
         viscVec += rfAtom;
00036
        for (nb = 1; nb <= nBuffAcf; nb ++) {
           indexAcf[nb] ++;
00037
           if (indexAcf[nb] <= 0) continue;
if (indexAcf[nb] == 1) {</pre>
00038
00039
00040
             viscAcfOrg[nb] = viscVec;
00041
00042
           ni = indexAcf[nb];
           viscAcf[nb][ni] = viscAcfOrg[nb] *viscVec;
00043
00044
00045
         AccumVacf():
00046 }
```

 $References\ AccumVacf(),\ ax,\ ay,\ deltaT,\ indexAcf,\ nAtom,\ nBuffAcf,\ rfAtom,\ viscAcf,\ viscAcfOrg,\ vx,\ and\ vy.$

Here is the call graph for this function:



3.72.2.22 EvalVrms()

```
void EvalVrms ( )
Definition at line 27 of file EvalVrms.c.
00027
          int n;
         VSqr = 0.0;
VMeanSqr = 0.0;
00029
00030
00031
         VRootMeanSqr = 0.0;
00032
         for(n = 1 ; n <= nAtom ; n ++) {
VSqr += Sqr(vx[n]) + Sqr(vy[n]);</pre>
00033
00034
00035
00036
         VMeanSqr = VSqr/nAtom;
         VRootMeanSqr = sqrt (VMeanSqr);
00037
00038
```

References nAtom, Sqr, VMeanSqr, VRootMeanSqr, VSqr, vx, and vy.

Referenced by main().

Here is the caller graph for this function:



3.72.2.23 HaltConditionCheck()

References fpresult, HaltCondition, and stepCount. Referenced by main().

Here is the caller graph for this function:



3.72.2.24 Init() void Init ()

```
Definition at line 29 of file Init.c.
                   char dummy[128];
00031
                   char inputConfig[128];
00032
                   FILE *fp;
                  fp = fopen("input-data","r");
fscanf(fp, "%s %s", dummy, inputConfig);
fscanf(fp, "%s %s", dummy, solver);
00033
00034
00035
                  fscanf(fp, "%s %s %s", dummy, xBoundary, yBoundary);
00036
                 fscanf(fp, "%s %d", dummy, &DampFlag);
fscanf(fp, "%s %d", dummy, &freezeAtomType);
fscanf(fp, "%s %lf", dummy, &rCut);
fscanf(fp, "%s %lf", dummy, &gamman);
fscanf(fp, "%s %lf", dummy, &kappa);
00037
00038
00039
00040
                fscanf(fp, "%s %lf", dummy, &kappa);
fscanf(fp, "%s %lf", dummy, &deltaT);
fscanf(fp, "%s %lf", dummy, &strain);
fscanf(fp, "%s %lf", dummy, &FyBylx);
fscanf(fp, "%s %lf", dummy, &fxByfy);
fscanf(fp, "%s %lf", dummy, &DeltaY);
fscanf(fp, "%s %lf", dummy, &DeltaX);
fscanf(fp, "%s %lf", dummy, &HaltCondition);
fscanf(fp, "%s %d", dummy, &stepAvg);
fscanf(fp, "%s %d", dummy, &stepEquil);
fscanf(fp, "%s %d", dummy, &stepLimit);
fscanf(fp, "%s %d", dummy, &stepTraj);
fscanf(fp, "%s %d", dummy, &stepTraj);
fscanf(fp, "%s %d", dummy, &nBuffCorr);
00041
00042
00043
00044
00045
00046
00047
00048
00049
00050
00051
00052
00053
00054
00055
                 fscanf(fp, "%s %d", dummy, &nFunCorr);
fscanf(fp, "%s %d", dummy, &nValCorr);
fscanf(fp, "%s %d", dummy, &stepCorr);
fscanf(fp, "%s %d", dummy, &imitAcfAv
fscanf(fp, "%s %d", dummy, &nBuffAcf);
00056
00057
00058
                                                                    dummy, &limitAcfAv);
00059
00060
                 fscanf(fp, "%s %d", dummy, &nBuffAcf);
fscanf(fp, "%s %d", dummy, &nValAcf);
fscanf(fp, "%s %d", dummy, &stepAcf);
fscanf(fp, "%s %d", dummy, &rangeRdf);
fscanf(fp, "%s %d", dummy, &limitRdf);
fscanf(fp, "%s %d", dummy, &sizeHistRdf);
fscanf(fp, "%s %d", dummy, &stepRdf);
00061
00062
00063
00064
00065
00066
00067
00068
                   fclose(fp);
00069
                   FILE *fpSTATE;
                   if((fpSTATE = fopen(inputConfig,"r")) ==NULL) {
00070
00071
                   printf("Error occurred: Could not open STATE file\n Exiting now..\n");
00072
                   exit(0);
00073
00074
                  fscanf(fpSTATE, "%s %lf", dummy, &timeNow);
fscanf(fpSTATE, "%s %d", dummy, &nAtom);
fscanf(fpSTATE, "%s %d", dummy, &nBond);
fscanf(fpSTATE, "%s %d", dummy, &nAtomType);
00075
00076
00077
00078
00079
                  fscanf(fpSTATE, "%s %d", dummy, &nBondType);
                  fscanf(fpSTATE, "%s %lf", dummy, &region[1]);
fscanf(fpSTATE, "%s %lf", dummy, &region[2]);
08000
00081
00082
00083
                   region[2] *= 1.5; //Remove this when put on GitHub
```

density = nAtom/(region[1]*region[2]);

cells[1] = region[1] / rCut;

cells[2] = region[2] / rCut;

```
cellList = (int*)malloc((nAtom + cells[1] * cells[2] + 1) * sizeof(int));
         regionH[1] = 0.5*region[1];
regionH[2] = 0.5*region[2];
00089
00090
00091
00092
         //strain information
00093
         strainRate = strain/deltaT;
         shearDisplacement = strain * region[2];
00094
00095
         shearVelocity = strainRate * region[2];
00096
00097
00098
         rx = (double*)malloc((nAtom + 1) * sizeof(double));
         ry = (double*)malloc((nAtom + 1) * sizeof(double));
00099
         vx = (double*)malloc((nAtom + 1) * sizeof(double));
00100
         vy = (double*)malloc((nAtom + 1) * sizeof(double));
00101
00102
         ax = (double*)malloc((nAtom + 1) * sizeof(double));
         ay = (double*)malloc((nAtom + 1) * sizeof(double));
00103
         fax = (double*)malloc((nAtom + 1) * sizeof(double));
fay = (double*)malloc((nAtom + 1) * sizeof(double));
00104
00105
         atomID = (int*)malloc((nAtom+1) * sizeof(int));
         atomType = (int*)malloc((nAtom+1) * sizeof(int));
00107
00108
         atomRadius = (double*)malloc((nAtom + 1) * sizeof(double));
00109
         atomMass = (double*)malloc((nAtom + 1) * sizeof(double));
         speed = (double*)malloc((nAtom + 1) * sizeof(double));
00110
         atom1 = (int*)malloc((nBond+1)*sizeof(int));
00111
00112
         atom2 = (int*)malloc((nBond+1)*sizeof(int));
         BondID = (int*)malloc((nBond+1)*sizeof(int));
00113
00114
         BondType = (int*)malloc((nBond+1)*sizeof(int));
         kb = (double*)malloc((nBond+1)*sizeof(double));
00115
         ro = (double*)malloc((nBond+1)*sizeof(double));
00116
         BondEnergy = (double*)malloc((nBond+1)*sizeof(double));
00117
         BondLength = (double*) malloc((nBond+1)*sizeof(double));
00118
         discPragx = (double*)malloc((nAtom + 1) * sizeof(double));
discPragy = (double*)malloc((nAtom + 1) * sizeof(double));
00119
00120
00121
         nodeDragx = (double*)malloc((nAtom + 1) * sizeof(double));
         nodeDragy = (double*)malloc((nAtom + 1) * sizeof(double));
00122
         ImageX = (int*)malloc((nAtom+1) * sizeof(int));
ImageY = (int*)malloc((nAtom+1) * sizeof(int));
00123
00124
         rxUnwrap = (double*)malloc((nAtom + 1) * sizeof(double));
ryUnwrap = (double*)malloc((nAtom + 1) * sizeof(double));
00126
         DeltaXijOld = (double*)malloc((nBond+1)*sizeof(double));
DeltaYijOld = (double*)malloc((nBond+1)*sizeof(double));
00127
00128
         DeltaXijOldPair = (double**)malloc((nAtom+1) * sizeof(double*));
DeltaYijOldPair = (double**)malloc((nAtom+1) * sizeof(double*));
00129
00130
00131
         for (int n = 0; n \le nAtom; n++) {
          DeltaXijOldPair[n] = (double*)malloc((nAtom+1) * sizeof(double));
DeltaYijOldPair[n] = (double*)malloc((nAtom+1) * sizeof(double));
00132
00133
00134
00135
         molID = (int*)malloc((nAtom+1) * sizeof(int));
00136
00137
         for(n = 1; n \le nAtom; n ++) {
          atomMass[n] = 1.0;
00138
00139
00140
         fscanf(fpSTATE, "%s\n", dummy);
00141
         for (n = 1; n \le nAtom; n ++)
00142
          fscanf(fpSTATE, "%d %d %d %lf %lf %lf %lf %lf %lf\n", &atomID[n], &molID[n], &atomType[n],
00143
      &atomRadius[n], &rx[n], &ry[n], &vx[n], &vy[n]);
00144
00145
         fscanf(fpSTATE, "%s\n", dummy);
00146
         for(n=1; n<=nBond; n++)
fscanf(fpSTATE, "%d %d %d %d %lf %lf\n", &BondID[n], &BondType[n], &atom1[n], &atom2[n], &kb[n],</pre>
00147
00148
      &ro[n]);
00149
00150
         fclose(fpSTATE);
00151
00152
        //2D-List of bonded atoms. This is used to remove pair interaction
00153
        //calculation for the bonded atoms
00154
          isBonded = (int**)malloc((nAtom + 1) * sizeof(int*));
00155
         for (int i = 0; i <= nAtom; i++) {</pre>
            isBonded[i] = (int*)malloc((nAtom + 1) * sizeof(int));
for (int j = 0; j <= nAtom; j++) {</pre>
00156
00157
                isBonded[i][j] = 0;
00158
00159
00160
         }
00161
00162
         for (n = 1; n <= nBond; n++) {</pre>
          int i = atom1[n];
int j = atom2[n];
00163
00164
            isBonded[i][j] = 1;
isBonded[j][i] = 1; // symmetric
00165
00166
00167 }
00168
00169
        //For thermostate, update in final version
00170 thermo = 'C';
00171
00172
```

```
00173 // List the interface atoms
00174 nAtomInterface = 0;
00175 nAtomBlock = 0;
00176 nDiscInterface = 0;
00177
             double InterfaceWidth, bigDiameter;
00178 bigDiameter = 2.8;
00179
             InterfaceWidth = 5.0 * bigDiameter;
00180
00181 for (n = 1; n \le nAtom; n++) {
00182
              if(fabs(ry[n]) < InterfaceWidth){</pre>
              nAtomInterface++;
00183
00184
00185
               if (molID[n] == 2) {
00186
               nAtomBlock++;
00187
00188
                if(atomRadius[n] != 0.0){
               nDiscInterface++:
00189
00190
               } }
00191
00192
                 atomIDInterface = (int*)malloc((nAtomInterface+1)*sizeof(int));
00193
00194
00195
               m = 1;
               for(n=1; n<=nAtom; n++) {
  if(fabs(ry[n]) < InterfaceWidth) {</pre>
00196
00197
00198
                 atomIDInterface[m] = atomID[n];
00199
00200
                } }
00201
00202
               nPairTotal = 0.5 * nAtomInterface * (nAtomInterface-1);
00203
               PairID = (int*)malloc((nPairTotal+1) * sizeof(int));
               Pairatom1 = (int*)malloc((nPairTotal+1) * sizeof(int));
Pairatom2 = (int*)malloc((nPairTotal+1) * sizeof(int));
00204
00205
00206
               PairXij = (double*)malloc((nPairTotal+1) * sizeof(double));
               PairYij = (double*)malloc((nPairTotal+1) * sizeof(double));
00207
00208
00209
                    fprintf(fpresult, "-----
                   fprintf(fpresult, "-----\n");
fprintf(fpresult, "-----\n");
fprintf(fpresult, "-----\n");
fprintf(fpresult, "nAtom\t\t\t\t\d\n", nAtom);
fprintf(fpresult, "nBond\t\t\t\d\n", nBond);
fprintf(fpresult, "nAtomBlock\t\t\d\n", nAtomBlock);
fprintf(fpresult, "nAtomInterface\t\d\n", nAtomInterface);
fprintf(fpresult, "nDiscInterface\t\t\d\n", nDiscInterface);
00211
00212
00213
00214
00215
00216
                   fprintf(fpresult, "nDiscInterface\t\t%d\n", nDiscInterface);
fprintf(fpresult, "gamman\t\t\t%0.6g\n", gamman);
fprintf(fpresult, "strainRate\t\t%0.6g\n", strain);
fprintf(fpresult, "strainRate\t\t\t%0.6g\n", strainRate);
fprintf(fpresult, "FyBylx\t\t\t%0.6g\n", FyBylx);
fprintf(fpresult, "fxByfy\t\t\t%0.6g\n", fxByfy);
fprintf(fpresult, "DeltaY\t\t\t%0.6g\n", DeltaY);
fprintf(fpresult, "BeltaY\t\t\t%0.6g\n", DeltaY);
fprintf(fpresult, "HaltCondition\t\t%0.6g\n", HaltCondition);
fprintf(fpresult, "kappa\t\t\t\t\t\g\n", kappa);
00217
00218
00219
00220
00221
00222
                  fprintf(fpresult, "HaltCondition)t\t\%0.6g\n", HaltCondition);
fprintf(fpresult, "kappa\t\t\t\$g\n", kappa);
fprintf(fpresult, "density\t\t\$g\n", density);
fprintf(fpresult, "rCut\t\t\t\$g\n", rCut);
fprintf(fpresult, "deltaT\t\t\$g\n", rCut);
fprintf(fpresult, "stepEquil\t\t\t\$d\n", stepEquil);
fprintf(fpresult, "stepLimit\t\t\$d\n", stepLimit);
fprintf(fpresult, "region[1]\t\t\t\$0.161f\n", region[1]);
fprintf(fpresult, "region[2]\t\t\$0.161f\n", region[2]);
fprintf(fpresult, "cells[1]\t\t\$d\n", cells[1]);
fprintf(fpresult, "cells[2]\t\t\$d\n", cells[2]);
fprintf(fpresult, "solver\t\t\$\$\n", solver);
fprintf(fpresult, "boundary\t\t\$\$\s\n", xBoundary, yBoundary);
fprintf(fpresult, "DampFlag\t\t\$d\n", DampFlag);
00224
00225
00226
00227
00228
00230
00231
00232
00233
00234
00235
00236
00237
00238
00239
                   fprintf(fpresult, "-----\n");
fprintf(fpresult, "#TimeNow TotalMomentum PotEngyPerAtom KinEngyPerAtom TotEngyPerAtom
00240
00241
          PairEnergyPerAtom BondEnergyPerAtom Press VirialSum\n");
                   fprintf(fpvrms, "#timeNow\tVrms \n");
fprintf(fpcom, "#timeNow\tComX\tComY\n");
00242
00243
00244
00245 /\star //Uncomment the following as per your acquirement
                  00246
00247
                   fprintf(fpstress, "region[2] %lf\n", region[2]);
fprintf(fpstress, "#timeNow virSumxx virSumyy virSumxy pressure\n");
fprintf(fpmomentum, "#timeNow Px Py\n");
00248
00249
00250
00251 */
00252
          if((strcmp(xBoundary, "p") != 0 && strcmp(xBoundary, "r") != 0) ||
  (strcmp(yBoundary, "p") != 0 && strcmp(yBoundary, "r") != 0)) {
  fprintf(fpresult, "Error: Invalid boundary value detected: '%s %s'. Only 'p' or 'r' are
  allowed.\n", xBoundary, yBoundary);
  exit(EXIT_FAILURE); // Exit with failure status
00254
00255
00256
00257
```

```
00258
```

References atom1, atom2, atomID, atomIDInterface, atomMass, atomRadius, atomType, ax, ay, BondEnergy, BondID, BondLength, BondType, cellList, cells, DampFlag, deltaT, DeltaX, DeltaXijOld, DeltaXijOldPair, DeltaY, DeltaYijOld, DeltaYijOldPair, density, discDragx, discDragy, fax, fay, fpcom, fpresult, fpvrms, freezeAtomType, fxByfy, FyBylx, gamman, HaltCondition, ImageX, ImageY, isBonded, kappa, kb, limitAcfAv, limitCorrAv, limitRdf, molID, nAtom, nAtomBlock, nAtomInterface, nAtomType, nBond, nBondType, nBuffAcf, nBuffCorr, nDiscInterface, nFunCorr, nodeDragx, nodeDragy, nPairTotal, nValAcf, nValCorr, Pairatom1, Pairatom2, PairID, PairXij, PairYij, rangeRdf, rCut, region, regionH, ro, rx, rxUnwrap, ry, ryUnwrap, shearDisplacement, shearVelocity, sizeHistRdf, solver, speed, stepAcf, stepAvg, stepCorr, stepDump, stepEquil, stepLimit, stepRdf, stepTraj, strain, strainRate, thermo, timeNow, vx, vy, xBoundary, and yBoundary.

Referenced by main().

Here is the caller graph for this function:



3.72.2.25 LeapfrogStep()

```
void LeapfrogStep ( )
```

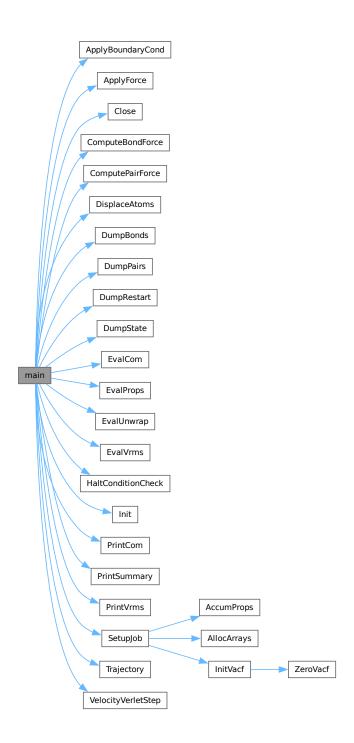
3.72.2.26 main()

```
int main (
                  int arac.
                  char ** argv )
Definition at line 51 of file main.c.
00051
00052
        time_t t1 = 0, t2;
00053
         if (argc < 2) {
        fprintf(stderr, "Usage: %s <output_prefix>\n", argv[0]);
00054
00055
00056
         int prefix_size = snprintf(NULL, 0, "../output/%s", argv[1]) + 1; // +1 for the null terminator
00057
00058
         prefix = malloc(prefix_size);
if(prefix == NULL) {
00059
00060
          fprintf(stderr, "Memory allocation failed\n");
00061
           return 1;
00062
00063
         // Write the formatted string into the allocated space
snprintf(prefix, prefix_size, "../output/%s", argv[1]);
sprintf(result, "%s.result", prefix);
00064
00065
00066
          fpresult = fopen(result, "w");
00067
         sprintf(xyz, "%s.xyz", prefix);
fpxyz = fopen(xyz, "w");
00068
00069
         sprintf(vrms, "%s.vrms", prefix);
fpvrms = fopen(vrms, "w");
sprintf(bond, "%s.bond", prefix);
00070
00071
00072
         fpbond = fopen(bond, "w");
00073
00074
          sprintf(com, "%s.com", prefix);
         fpcom = fopen(com, "w");
sprintf(pair, "%s.pair", prefix);
00075
00076
         fppair = fopen(pair, "w");
00077
00078
00079
          /* //Uncomment the following as per your acquirement
08000
          sprintf(dnsty, "%s.curr-dnsty", prefix);
00081
          fpdnsty = fopen(dnsty, "w");
00082
          sprintf(visc, "%s.viscosity", prefix);
         fpvisc = fopen(visc, "w");
sprintf(rdf, "%s.rdf", prefix);
00083
00084
00085
         fprdf = fopen(rdf, "w");
00086
         sprintf(stress, "%s.stress", prefix);
```

```
fpstress = fopen(stress, "w");
         sprintf(momentum, "%s.momentum", prefix);
fpmomentum = fopen(momentum, "w");
00088
00089
00090
00091
00092
         Init();
         SetupJob();
00093
00094
         t1 = time(NULL);
00095
         moreCycles = 1;
         timeNow = 0.0;
if(timeNow == 0.0) {
00096
00097
         DisplaceAtoms();
00098
00099
          ComputePairForce(1);
00100
          ComputeBondForce();
00101
          ApplyForce();
00102
          DumpBonds();
          DumpPairs();
Trajectory();
00103
00104
          EvalUnwrap();
00105
00106
          ApplyBoundaryCond();
00107
          EvalProps();
00108
          EvalVrms();
          EvalCom();
00109
          PrintVrms();
00110
00111
          PrintCom();
00112
          PrintSummary();
00113
00114
00115 //Here starts the main loop of the program
00116
         while (moreCycles) {
00117
         if(stepLimit == 0){
00118
          exit(0);
00119
00120
00121
          stepCount ++;
          timeNow = stepCount * deltaT; //for adaptive step size: timeNow += deltaT
00122
00123
00124
          VelocityVerletStep(1);
00125
          EvalUnwrap();
00126
          ApplyBoundaryCond();
00127
          ComputePairForce(1);
          ComputeBondForce();
00128
          ApplyForce();
00129
          VelocityVerletStep(2);
00130
00131
          ApplyBoundaryCond();
00132
          EvalProps();
          EvalVrms();
00133
00134
          EvalCom();
          if(stepCount % stepAvg == 0){
00135
00136
           PrintSummary();
           PrintVrms();
00137
00138
           PrintCom();
00139
00140
          if(stepCount % stepTraj == 0){
00141
           Trajectory();
00142
           DumpBonds();
00143
           DumpPairs();
00144
00145
          if(stepCount % stepDump == 0){
                             // Save the current state for input
// Save the current state for config
00146
           DumpRestart();
00147
           DumpState();
00148
00149
          if(HaltConditionCheck(VRootMeanSqr, stepCount)) {
           DumpRestart(); // Save the current state for input
DumpState(); // Save the current state for config
00150
00151
           DumpState();
00152
           break; // Exit the loop when the halt condition is met
00153
00154
00155
           if(stepCount >= stepLimit)
            moreCycles = 0;
00156
00157
00158
00159
         t2 = time(NULL);
00160
        fprintf(fpresult, "#Execution time %lf secs\n", difftime(t2,t1));
fprintf(fpresult, "#Execution speed %lf steps per secs\n", stepLimit/difftime(t2,t1));
00161
00162
00163
00164
         fclose(fpresult);
00165
        fclose(fpxyz);
00166
        fclose(fpvrms);
00167
         fclose(fpbond);
00168
         fclose(fppair);
00169
         fclose(fpcom);
00170
00171 /\star//\mathrm{Uncomment} the following as per your acquirement
00172
        fclose(fpdnstv);
00173
        fclose(fpvisc);
```

References ApplyBoundaryCond(), ApplyForce(), bond, Close(), com, ComputeBondForce(), ComputePairForce(), deltaT, DisplaceAtoms(), DumpBonds(), DumpPairs(), DumpRestart(), DumpState(), EvalCom(), EvalProps(), EvalUnwrap(), EvalVrms(), fpbond, fpcom, fppair, fpresult, fpvrms, fpxyz, HaltConditionCheck(), Init(), moreCycles, pair, prefix, PrintCom(), PrintSummary(), PrintVrms(), result, SetupJob(), stepAvg, stepCount, stepDump, stepLimit, stepTraj, timeNow, Trajectory(), VelocityVerletStep(), vrms, VRootMeanSqr, and xyz.

Here is the call graph for this function:



3.72.2.27 PrintCom()

References ComX, ComY, fpcom, and timeNow.

Referenced by main().

Here is the caller graph for this function:



3.72.2.28 PrintMomentum()

3.72.2.29 PrintStress()

References fpstress, pressure, timeNow, virSumxx, virSumxy, and virSumyy.

3.72.2.30 PrintSummary()

References BondEnergyPerAtom, fpresult, kinEnergy, potEnergy, pressure, timeNow, totEnergy, uSumPairPerAtom, virSum, and vSum.

Referenced by main().

Here is the caller graph for this function:



3.72.2.31 PrintVrms()

```
void PrintVrms ( )
```

Definition at line 27 of file PrintVrms.c.

```
00027 {
00028 fprintf(fpvrms, "%0.41f\t%0.161f\n", timeNow, VRootMeanSqr);
00029 fflush(fpvrms);
00030 }
```

References fpvrms, timeNow, and VRootMeanSqr.

Referenced by main().

Here is the caller graph for this function:



3.72.2.32 SetupJob()

void SetupJob ()

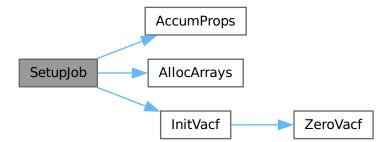
Definition at line 27 of file SetupJob.c.

```
00027
          AllocArrays();
00028
         AccumProps(0);
InitVacf();
stepCount = 0;
00029
00030
00031
00032
          // INITIALISE SPACETIME CORRELATIONS
00033
         for (n = 1; n <= nBuffCorr; n++)
  indexCorr[n] = -(n - 1)*nValCorr/nBuffCorr;</pre>
00034
00035
00036
00037
         countCorrAv = 0.;
00038
00039
         for (n = 1; n <= nFunCorr*nValCorr; n++)</pre>
00040
00041
            spacetimeCorrAv[n] = 0.;
00042
         //RDF
00043
         countRdf = 0;
```

References AccumProps(), AllocArrays(), countCorrAv, countRdf, indexCorr, InitVacf(), nBuffCorr, nFunCorr, nValCorr, spacetimeCorrAv, and stepCount.

Referenced by main().

Here is the call graph for this function:



Here is the caller graph for this function:



3.72.2.33 Trajectory()

References atomID, atomRadius, atomType, ax, ay, fpxyz, moIID, nAtom, regionH, rx, ry, timeNow, vx, and vy. Referenced by main().

Here is the caller graph for this function:



3.72.2.34 VelocityVerletStep()

```
void VelocityVerletStep (
                        int icode )
Definition at line 26 of file VelocityVerletStep.c.
00026
00027 int n;
00028 if(icode == 1){
00029
           for (n= 1; n <= nAtom; n++) {</pre>
            if (n- 1; n <- hatom; n++) {
   if (atomType[n] != freezeAtomType) {
    vx[n] += ax[n] * 0.5 * deltaT;
   vy[n] += ay[n] * 0.5 * deltaT;
   rx[n] += vx[n] * deltaT;
   ry[n] += vy[n] * deltaT;</pre>
00031
00032
00033
00034
00035
00036
             //Calculating the image flags here
            if (rx[n] >= regionH[1]) {
  rx[n] -= region[1];
  ImageX[n]++;
00037
00038
00039
```

3.73 main.c 163

```
} else if (rx[n] < -regionH[1]) {</pre>
00041
          rx[n] += region[1];
00042
         ImageX[n]--;
00043
         if (ry[n] >= regionH[2]) {
00044
         ry[n] -= region[2];
ImageY[n]++;
00045
00047
          } else if (ry[n] < -regionH[2]) {</pre>
00048
          ry[n] += region[2];
00049
          ImageY[n]--;
00050
          } } }
00051
         else if(icode == 2){
00052
         for(n = 1; n <= nAtom; n++) {</pre>
         if (atomType[n] != freezeAtomType) {
  vx[n] += ax[n] * 0.5 * deltaT;
00053
00054
          vy[n] += ay[n] * 0.5 * deltaT;
00056 } } } }
```

References atomType, ax, ay, deltaT, freezeAtomType, ImageX, ImageY, nAtom, region, regionH, rx, ry, vx, and vy.

Referenced by main().

Here is the caller graph for this function:



3.72.3 Variable Documentation

3.72.3.1 prefix

char* prefix = NULL

Definition at line 13 of file main.c.

Referenced by DumpRestart(), DumpState(), and main().

3.73 main.c

Go to the documentation of this file.

```
00001 #include<stdio.h>
00002 #include<string.h>
00003 #include<stdlib.h>
00004 #include <stdbool.h>
00005 #include <time.h>
00006 #include <mpi.h>
00007 #define DEFINE_GLOBALS
00008 #include "global.h"
00009 #include "ComputeBondForce.h"
00010 #include "ComputePairForce.h"
00011
00012
00013 char *prefix = NULL; // Definition of prefix
00014
00015 void Init();
00016 void SetupJob();
00017 void EvalSpacetimeCorr();
00018 void Trajectory();
00019 void DumpState();
00020 void ComputeForcesCells();
00021 void LeapfrogStep();
00022 void BrownianStep();
00023 void ApplyBoundaryCond();
00024 void EvalProps();
00025 void EvalVacf();
00026 void EvalRdf();
00027 void AccumProps(int icode);
00028 void PrintSummary();
00029 void PrintVrms():
00030 //void ComputeBondForce();
```

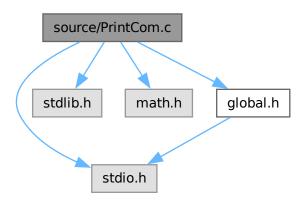
```
00031 void DumpBonds();
00032 void VelocityVerletStep(int icode);
00033 void ApplyForce();
00034 void ApplyDrivingForce();
00035 void ApplyShear();
00036 void ApplyLeesEdwardsBoundaryCond();
00037 void PrintStress();
00038 void Close();
00039 //void ComputePairForce(int normFlag);
00040 void PrintMomentum();
00041 void DisplaceAtoms();
00042 void DumpRestart();
00043 bool HaltConditionCheck(double value, int stepCount);
00044 void EvalCom();
00045 void PrintCom();
00046 void EvalVrms();
00047 void EvalUnwrap();
00048 void DumpPairs();
00049 void ApplyViscous();
00050
00051 int main(int argc, char **argv) {
00052 time_t t1 = 0, t2;
00053 if (argc < 2) {
00054 fprintf(stderr, "Usage: %s <output_prefix>\n", argv[0]);
00055
         return 1;
00056
00057
          int prefix_size = snprintf(NULL, 0, "../output/%s", argv[1]) + 1; // +1 for the null terminator
00058
          prefix = malloc(prefix_size);
           if(prefix == NULL) {
  fprintf(stderr, "Memory allocation failed\n");
00059
00060
00061
           return 1:
00062
00063
          // Write the formatted string into the allocated space
snprintf(prefix, prefix_size, "../output/%s", argv[1]);
sprintf(result, "%s.result", prefix);
fpresult = fopen(result, "w");
sprintf(xyz, "%s.xyz", prefix);
fpxyz = fopen(xyz, "w");
00064
00065
00066
00067
00069
          fpxyz = fopen(xyz, "w");
sprintf(vrms, "%s.vrms", prefix);
fpvrms = fopen(vrms, "w");
sprintf(bond, "%s.bond", prefix);
fpbond = fopen(bond, "w");
sprintf(com, "%s.com", prefix);
fpcom = fopen(com, "w");
sprintf(pair, "%s.pair", prefix);
fppair = fopen(pair, "w");
00070
00071
00072
00073
00074
00075
00076
00077
00078
00079
          /* //Uncomment the following as per your acquirement
          sprintf(dnsty, "%s.curr-dnsty", prefix);
08000
          fpdnsty = fopen(dnsty, "w");
00082
          sprintf(visc, "%s.viscosity", prefix);
          fpvisc = fopen(visc, "w");
sprintf(rdf, "%s.rdf", prefix);
fprdf = fopen(rdf, "w");
sprintf(stress, "%s.stress", prefix);
fpstress = fopen(stress, "w");
00083
00084
00085
00086
00087
00088
          sprintf(momentum, "%s.momentum", prefix);
           fpmomentum = fopen(momentum, "w");
00089
00090
00091
          Init();
SetupJob();
00092
00093
00094
          t1 = time(NULL);
00095
          moreCycles = 1;
          timeNow = 0.0;
00096
          if (timeNow == 0.0) {
  DisplaceAtoms();
00097
00098
00099
           ComputePairForce(1);
00100
            ComputeBondForce();
00101
            ApplyForce();
00102
            DumpBonds();
00103
            DumpPairs();
            Trajectory();
00104
00105
            EvalUnwrap();
00106
            ApplyBoundaryCond();
00107
            EvalProps();
00108
            EvalVrms();
00109
            EvalCom();
00110
            PrintVrms():
00111
            PrintCom();
00112
            PrintSummary();
00113
00114
00115 //Here starts the main loop of the program
          while (moreCycles) {
  if (stepLimit == 0) {
00116
00117
```

```
00118
         exit(0);
00119
00120
00121
         stepCount ++;
00122
         timeNow = stepCount * deltaT; //for adaptive step size: timeNow += deltaT
00123
00124
         VelocityVerletStep(1);
00125
         EvalUnwrap();
00126
         ApplyBoundaryCond();
00127
         ComputePairForce(1);
         ComputeBondForce();
00128
         ApplyForce();
00129
          VelocityVerletStep(2);
00130
00131
         ApplyBoundaryCond();
00132
         EvalProps();
00133
         EvalVrms();
00134
         EvalCom();
00135
         if(stepCount % stepAvg == 0){
          PrintSummary();
00136
00137
          PrintVrms();
00138
          PrintCom();
00139
         if(stepCount % stepTraj == 0){
00140
00141
          Trajectory();
00142
          DumpBonds();
00143
          DumpPairs();
00144
00145
         if(stepCount % stepDump == 0){
          DumpRestart(); // Save the current state for input
DumpState(); // Save the current state for config
00146
00147
00148
00149
         if (HaltConditionCheck(VRootMeanSqr, stepCount)) {
          DumpRestart(); // Save the current state for input
DumpState(); // Save the current state for config
00150
00151
          DumpState();
00152
          break; // Exit the loop when the halt condition is met
00153
00154
00155
          if(stepCount >= stepLimit)
00156
           moreCycles = 0;
00157
00158
00159
        t2 = time(NULL);
00160
        fprintf(fpresult, "#Execution time %lf secs\n", difftime(t2,t1));
00161
00162
        fprintf(fpresult, "#Execution speed %lf steps per secs\n", stepLimit/difftime(t2,t1));
00163
00164
        fclose(fpresult);
00165
        fclose(fpxyz);
00166
        fclose(fpvrms);
00167
        fclose(fpbond);
00168
        fclose(fppair);
00169
00170
00171 /*//Uncomment the following as per your acquirement 00172 \, fclose(fpdnsty);
00173
        fclose(fpvisc);
        fclose(fprdf);
00175
        fclose(fpstress);
00176 fclose(fpmomentum);
00177 */
00178
00179
        free (prefix);
00180
        Close();
00181
        return 0;
00182 }
```

3.74 source/PrintCom.c File Reference

```
#include <stdio.h>
#include <stdlib.h>
#include <math.h>
#include "global.h"
```

Include dependency graph for PrintCom.c:



Functions

void PrintCom ()

3.74.1 Function Documentation

3.74.1.1 PrintCom()

Referenced by main().

Here is the caller graph for this function:



3.75 PrintCom.c

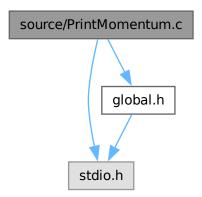
Go to the documentation of this file.

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00017 Copyright (C) 2025 Harish Charan, University of Durham, UK
00018
00019 */
00020
00021
00022
00023 #include<stdio.h>
00024 #include<stdlib.h>
00025 #include<math.h>
00026 #include"global.h"
00027
00028 void PrintCom(){
00029 fprintf(fpcom, "%0.41f\t%0.161f\t%0.161f\n", timeNow, ComX, ComY);
00030 fflush(fpcom);
00031
00032
00033
00034
```

3.76 source/PrintMomentum.c File Reference

```
#include <stdio.h>
#include "global.h"
Include dependency graph for PrintMomentum.c:
```



Functions

void PrintMomentum ()

3.76.1 Function Documentation

3.76.1.1 PrintMomentum()

```
void PrintMomentum ( )
Definition at line 25 of file PrintMomentum.c.
00025
00026 fprintf(fpmomentum, "%0.41f\t%0.161f\t%0.161f\n", timeNow, vSumX, vSumY);
00027 fflush(fpmomentum);
00028 }
```

References fpmomentum, timeNow, vSumX, and vSumY.

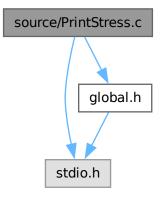
3.77 PrintMomentum.c

Go to the documentation of this file.

```
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00003
00004
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00016
00017 Copyright (C) 2025 Harish Charan, University of Durham, UK
00018
00019 */
00020
00021
00022 #include<stdio.h>
00023 #include"global.h"
00024
00025 void PrintMomentum(){
00026 fprintf(fpmomentum, "%0.41f\t%0.161f\t%0.161f\n", timeNow, vSumX, vSumY); 00027 fflush(fpmomentum);
00028 }
```

3.78 source/PrintStress.c File Reference

```
#include <stdio.h>
#include "global.h"
Include dependency graph for PrintStress.c:
```



Functions

• void PrintStress ()

3.78.1 Function Documentation

3.78.1.1 PrintStress()

void PrintStress ()

Definition at line 25 of file PrintStress.c.

3.79 PrintStress.c 169

References fpstress, pressure, timeNow, virSumxx, virSumxy, and virSumyy.

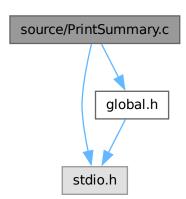
3.79 PrintStress.c

```
Go to the documentation of this file.
```

```
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00007
        \star (at your option) any later version.
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00016
00017 Copyright (C) 2025 Harish Charan, University of Durham, UK
00018
00019
00020
00021
00022 #include<stdio.h>
00023 #include"global.h"
00024
00025 void PrintStress(){
00026
        fprintf(fpstress, "%0.41f\t%0.161f\t%0.161f\t%0.161f\t%0.161f\t%0.161f\n", timeNow, virSumxx, virSumyy,
virSumxy, pressure);
00027 fflush(fpstress);
00028 }
```

3.80 source/PrintSummary.c File Reference

```
#include <stdio.h>
#include "global.h"
Include dependency graph for PrintSummary.c:
```



Functions

• void PrintSummary ()

3.80.1 Function Documentation

3.80.1.1 PrintSummary()

References BondEnergyPerAtom, fpresult, kinEnergy, potEnergy, pressure, timeNow, totEnergy, uSumPairPerAtom, virSum, and vSum.

Referenced by main().

Here is the caller graph for this function:



3.81 PrintSummary.c

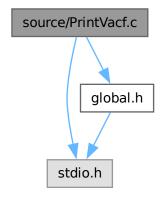
Go to the documentation of this file.

3.82 source/PrintVacf.c File Reference

```
#include <stdio.h>
#include "global.h"
```

3.83 PrintVacf.c 171

Include dependency graph for PrintVacf.c:



Functions

void PrintVacf ()

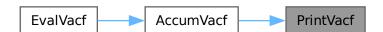
3.82.1 Function Documentation

3.82.1.1 PrintVacf()

References deltaT, fpvisc, nValAcf, stepAcf, viscAcfAv, and viscAcfInt.

Referenced by AccumVacf().

Here is the caller graph for this function:



3.83 PrintVacf.c

Go to the documentation of this file.

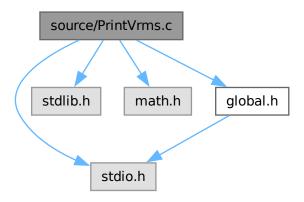
```
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```

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00016
00017 Copyright (C) 2025 Harish Charan, University of Durham, UK
00018
00019 */
00020
00021
00022 #include<stdio.h>
00023 #include"global.h"
00024
00025 void PrintVacf(){
00026 double tVal;
00027 int j;
00028 fprintf(fpvisc, "viscosity acf\n");
00029 for(j = 1; j <= nValAcf; j ++){
00030 tVal = (j-1)*stepAcf*deltaT;
00031 fprintf(fpvisc, "%lf\t %lf\t", tVal, viscAcfAv[j], viscAcfAv[j]/viscAcfAv[1]);
00032
00033 fprintf(fpvisc, "viscosity acf integral: %lf\n", viscAcfInt);
00034 }
00035
00036
```

3.84 source/PrintVrms.c File Reference

```
#include <stdio.h>
#include <stdlib.h>
#include <math.h>
#include "global.h"
```

Include dependency graph for PrintVrms.c:



Functions

• void PrintVrms ()

3.85 PrintVrms.c 173

3.84.1 Function Documentation

3.84.1.1 PrintVrms()

References fpvrms, timeNow, and VRootMeanSqr.

Referenced by main().

Here is the caller graph for this function:



3.85 PrintVrms.c

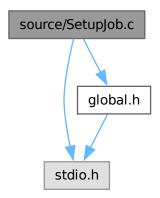
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00018
00020
00021
00022 #include<stdio.h>
00023 #include<stdlib.h>
00024 #include<math.h>
00025 #include"global.h"
00026
00027 void PrintVrms(){
00028 fprintf(fpvrms, "%0.41f\t%0.161f\n", timeNow, VRootMeanSqr);
00029 fflush(fpvrms);
00030 }
00032
00033
```

3.86 source/SetupJob.c File Reference

```
#include <stdio.h>
#include "global.h"
```

Include dependency graph for SetupJob.c:



Functions

- · void AllocArrays ()
- void AccumProps (int icode)
- void InitVacf ()
- void SetupJob ()

3.86.1 Function Documentation

3.86.1.1 AccumProps()

References kinEnergy, potEnergy, pressure, sKinEnergy, sPotEnergy, sPressure, Sqr, ssKinEnergy, ssPotEnergy, ssPressure, ssTotEnergy, stepAvg, sTotEnergy, svirSum, totEnergy, and virSum. Referenced by SetupJob().

Here is the caller graph for this function:



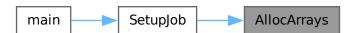
3.86.1.2 AllocArrays()

```
void AllocArrays ( )
Definition at line 25 of file AllocArrays.c.
00025
00026
      // SPACETIME CORRELATIONS
00027
00028
      cfOrg = (double **) malloc ((nBuffCorr+1)*sizeof(double *));
           (n = 0; n \le nBuffCorr; n++)
00030
       cfOrg[n] = (double *) malloc ((2*nFunCorr+1)*sizeof(double));
00031
00032 cfVal = (double *) malloc ((2*nFunCorr+1)*sizeof(double));
00033 indexCorr = (int *) malloc ((nBuffCorr+1)*sizeof(int));
00034
00035 spacetimeCorr = (double **) malloc ((nBuffCorr+1)*sizeof(double));
00036 for (n = 0; n <= nBuffCorr; n++)
00037 spacetimeCorr[n] = (double *) malloc ((nFunCorr*nValCorr+1)*sizeof(double));
00038
00039 spacetimeCorrAy = (double *) malloc ((nFunCorr*nValCorr+1)*sizeof(double));
00040 // VISCOSITY
00041 indexAcf = (double *)malloc((nBuffAcf+1)*sizeof(double));
00042 viscAcf = (double **)malloc((nBuffAcf+1)*sizeof(double *));
00043 for (n = 0 ; n \le nBuffAcf ; n ++)
00044
       viscAcf[n] = (double *)malloc((nValAcf+1)*sizeof(double));
00045
00046 viscAcfOrg = (double *)malloc((nBuffAcf+1)*sizeof(double));
00047 viscAcfAv = (double *) malloc((nValAcf+1) *sizeof(double));
00048
00049
       histRdf = (double *)malloc((sizeHistRdf+1)*sizeof(double));
00050
00051 }
```

References cfOrg, cfVal, histRdf, indexAcf, indexCorr, nBuffAcf, nBuffCorr, nFunCorr, nValAcf, nValCorr, sizeHistRdf, spacetimeCorr, spacetimeCorrAv, viscAcf, viscAcfAv, and viscAcfOrg.

Referenced by SetupJob().

Here is the caller graph for this function:



3.86.1.3 InitVacf()

```
00030 ZeroVacf();
```

References indexAcf, nBuffAcf, nValAcf, and ZeroVacf().

Referenced by SetupJob().

Here is the call graph for this function:



Here is the caller graph for this function:



3.86.1.4 SetupJob()

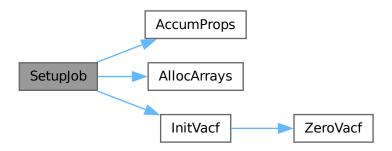
```
void SetupJob ( )
Definition at line 27 of file SetupJob.c.
00027
00028
         AllocArrays();
00029
         AccumProps(0);
00030
         InitVacf();
         stepCount = 0;
00031
00032
         \//\  INITIALISE SPACETIME CORRELATIONS
         int n;
for (n = 1; n <= nBuffCorr; n++)</pre>
00033
00034
00035
           indexCorr[n] = -(n - 1)*nValCorr/nBuffCorr;
00036
00037
         countCorrAv = 0.;
00038
         for (n = 1; n <= nFunCorr*nValCorr; n++)
    spacetimeCorrAv[n] = 0.;</pre>
00039
00041
00042
         countRdf = 0;
00044 }
```

References AccumProps(), AllocArrays(), countCorrAv, countRdf, indexCorr, InitVacf(), nBuffCorr, nFunCorr, nValCorr, spacetimeCorrAv, and stepCount.

Referenced by main().

3.87 SetupJob.c 177

Here is the call graph for this function:



Here is the caller graph for this function:



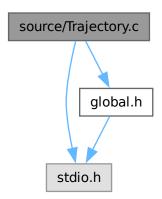
3.87 SetupJob.c

Go to the documentation of this file.

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00018
00020
00021 #include<stdio.h>
00022 #include"global.h"
00023
00024 void AllocArrays();
00025 void AccumProps(int icode);
00026 void InitVacf();
00027 void SetupJob(){
00028
         AllocArrays();
00029
         AccumProps(0);
         InitVacf();
stepCount = 0;
00030
00031
00032
         // INITIALISE SPACETIME CORRELATIONS
00033
00034
         for (n = 1; n <= nBuffCorr; n++)</pre>
           indexCorr[n] = -(n - 1)*nValCorr/nBuffCorr;
00035
```

3.88 source/Trajectory.c File Reference

```
#include <stdio.h>
#include "global.h"
Include dependency graph for Trajectory.c:
```



Functions

· void Trajectory ()

3.88.1 Function Documentation

3.88.1.1 Trajectory()

References atomID, atomRadius, atomType, ax, ay, fpxyz, moIID, nAtom, regionH, rx, ry, timeNow, vx, and vy. Referenced by main().

3.89 Trajectory.c 179

Here is the caller graph for this function:



3.89 Trajectory.c

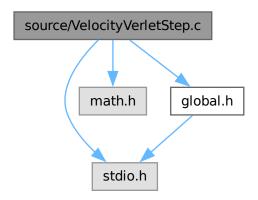
Go to the documentation of this file.

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00018
00019
00020
00021
00022 #include<stdio.h>
00023 #include"global.h"
00024
 00025 void Trajectory(){
00026 int n;
00027
           //Trajectory file in LAMMPS dump format for OVITO visualization
00027 //Trajectory file in LAMMPS dump format for OVITO visualization
00028 fprintf(fpxyz, "ITEM: TIMESTEP\n");
00029 fprintf(fpxyz, "%lf\n",timeNow);
00030 fprintf(fpxyz, "ITEM: NUMBER OF ATOMS\n");
00031 fprintf(fpxyz, "%d\n",nAtom);
00032 fprintf(fpxyz, "%d\n",nAtom);
00033 fprintf(fpxyz, "%lf %lf xlo xhi\n", -regionH[1], regionH[1]);
00034 fprintf(fpxyz, "%lf %lf ylo yhi\n", -regionH[2], regionH[2]);
00035 fprintf(fpxyz, "%lf %lf zlo zhi\n", -0.1, 0.1);
00036 fprintf(fpxyz, "ITEM: ATOMS id mol type radius x y vx vy fx fy\n");
00037 for(n=1: n<=nAtom: n++)</pre>
00037 for (n=1; n<=nAtom; n++)
            fprintf(fpxyz, "%d\t %d\t %0.21f\t %0.161f\t %0.161f\t %0.161f\t %0.161f\t %0.161f\t
00039
           atomID[n], molID[n], atomType[n], atomRadius[n], rx[n], ry[n], vx[n], vy[n], ax[n], ay[n]);
00040 }
00041
00042
00043
```

3.90 source/VelocityVerletStep.c File Reference

```
#include <stdio.h>
#include <math.h>
#include "global.h"
```

Include dependency graph for VelocityVerletStep.c:



Functions

void VelocityVerletStep (int icode)

3.90.1 Function Documentation

3.90.1.1 VelocityVerletStep()

void VelocityVerletStep (

```
int icode )
Definition at line 26 of file VelocityVerletStep.c.
00026
00027 int n;
00028 if (icode == 1) {
00029 for (n= 1; n <= nAtom; n++) {
          if(atomType[n] != freezeAtomType) {
          00031
00032
00033
00034
00035
          //Calculating the image flags here
if (rx[n] >= regionH[1]) {
  rx[n] -= region[1];
00037
00038
00039
            ImageX[n]++;
           } else if (rx[n] < -regionH[1]) {
rx[n] += region[1];</pre>
00040
00041
00042
           ImageX[n]--;
00043
00044
           if (ry[n] >= regionH[2]) {
           ry[n] -= region[2];
ImageY[n]++;
} else if (ry[n] < -regionH[2]) {
ry[n] += region[2];</pre>
00045
00046
00047
00048
00049
            ImageY[n]--;
00050
            } } }
00051
           else if(icode == 2){
          for(n = 1; n <= nAtom; n++) {
   if(atomType[n] != freezeAtomType) {
    vx[n] += ax[n] * 0.5 * deltaT;
   vy[n] += ay[n] * 0.5 * deltaT;</pre>
00052
00053
00054
00056 } } }
```

References atomType, ax, ay, deltaT, freezeAtomType, ImageX, ImageY, nAtom, region, regionH, rx, ry, vx, and vy.

Referenced by main().

Here is the caller graph for this function:



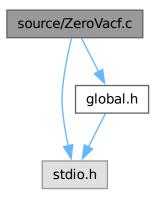
3.91 VelocityVerletStep.c

Go to the documentation of this file.

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00018
00019
00020
00021
00022 #include<stdio.h>
00023 #include<math.h>
00024 #include"global.h"
00026 void VelocityVerletStep(int icode){
00027 int n;
00029 for (n= 1; n <= nAtom; n++) {
         if (atomType[n] != freezeAtomType) {
  vx[n] += ax[n] * 0.5 * deltaT;
00030
          vy[n] += ay[n] * 0.5 * deltaT;
rx[n] += vx[n] * deltaT;
00032
00033
          ry[n] += vy[n] * deltaT;
00034
00035
          ^{\prime} //Calculating the image flags here
00036
00037
          if (rx[n] >= regionH[1]) {
          rx[n] -= region[1];
00039
           ImageX[n]++;
          } else if (rx[n] < -regionH[1]) {
rx[n] += region[1];</pre>
00040
00041
          ImageX[n]--;
00042
00043
00044
          if (ry[n] >= regionH[2]) {
00045
          ry[n] -= region[2];
00046
           ImageY[n]++;
           } else if (ry[n] < -regionH[2]) {
ry[n] += region[2];</pre>
00047
00048
00049
           ImageY[n]--;
00050
           } } }
00051
          else if(icode == 2){
00052
          for(n = 1; n <= nAtom; n++) {</pre>
          if(atomType[n] != freezeAtomType){
  vx[n] += ax[n] * 0.5 * deltaT;
  vy[n] += ay[n] * 0.5 * deltaT;
00053
00054
00055
00056 } } }
```

3.92 source/ZeroVacf.c File Reference

```
#include <stdio.h>
#include "global.h"
Include dependency graph for ZeroVacf.c:
```



Functions

• void ZeroVacf ()

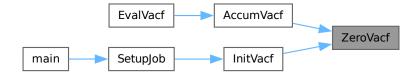
3.92.1 Function Documentation

3.92.1.1 ZeroVacf()

References countAcfAv, nValAcf, and viscAcfAv.

Referenced by AccumVacf(), and InitVacf().

Here is the caller graph for this function:



3.93 ZeroVacf.c

Go to the documentation of this file.

3.93 ZeroVacf.c 183

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00018
00019 */
00020
00021
00022 #include<stdio.h>
00023 #include"global.h"
00024
00025 void ZeroVacf(){
00026 int j;
00027 countAcfAv= 0;
```

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