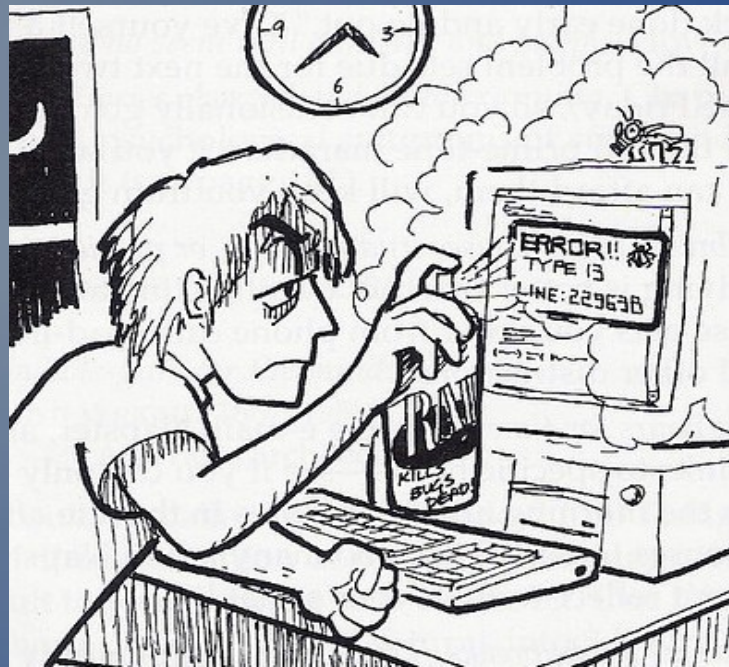


The Survival



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Workshop on Computer Programming and Advanced
Tools for Scientific Research Work & Quantum
ESPRESSO Developer Training

Outline

- Project Management & distribution of task
- Refactor the Code and provide python interface
- Project Documentation
- Changes in Simulation Infrastructure

Project Management & distribution of task

- Development (2)
- Changes in Simulation Infrastructure (1)
- Unit testing (1)
- Documentation (1)

Modules of Fortran program & Python Interface

Modules: cell.f90,force.f90,getekin.f90, f90, io.f90, kinds.f90, ljmd.f90,mdsys.f90, physconst.f90,utils.f90, veverlet.f90

Python Interface

```
import ljmd
import numpy as np
pyfname = np.array('argon_108')

ljmd.interface.callopen(pyfname)

#ljmd.interface.set_natoms(108)          # natoms
#ljmd.interface.set_mass(39.948)        # mass in AMU
#ljmd.interface.set_epsilon(0.2379)    # epsilon in kcal/mol
#ljmd.interface.set_sigma(3.405)       # sigma in angstrom
#ljmd.interface.set_rcut(8.5)          # rcut in angstrom
#ljmd.interface.set_box(17.1580)       # box length (in angstrom)
#ljmd.interface.set_nsteps(10000)      # nr MD steps
#ljmd.interface.set_dt(5.0)            # MD time step (in fs)
#ljmd.interface.set_nprint(100)       # output print frequency

ljmd.interface.run()
```

Modules of Fortran program & Python Interface

Fortran interface

```
MODULE interface
```

```
  USE ljmd
```

```
IMPLICIT NONE
```

```
CONTAINS
```

```
  SUBROUTINE run
```

```
    IMPLICIT NONE
```

```
      CALL ljmd
```

```
  END SUBROUTINE
```

```
  SUBROUTINE callopen(pyfname)
```

```
    IMPLICIT NONE
```

```
    CHARACTER*(*) :: pyfname
```

```
      CALL openfiles(pyfname)
```

```
  END SUBROUTINE
```

```
  SUBROUTINE set_natoms(myatoms)
```

```
    IMPLICIT NONE
```

```
    INTEGER :: myatoms
```

```
    natoms = myatoms
```

```
  END SUBROUTINE set_natoms
```

Modules of Fortran program & Python Interface

ljmd

```
MODULE ljmd
```

```
  USE kinds
```

```
  USE io
```

```
  USE utils
```

```
  USE mdsys
```

```
  USE cell
```

```
  INTEGER :: nprint
```

```
CONTAINS
```

```
SUBROUTINE lljmd
```

```
IMPLICIT NONE
```

```
  INTEGER :: i, j
```

```
  INTEGER, EXTERNAL :: omp_get_num_threads
```

```
  nthreads = 1
```

```
  !$OMP parallel shared(nthreads)
```

```
  !$OMP master
```

```
  !$ nthreads = omp_get_num_threads()
```

```
  !$ WRITE(stdout, '(A,I2,A)') 'Running OpenMP version
```

```
using ', nthreads, ' thread(s).'
```

```
  !$OMP end master
```

```
  !$OMP end parallel
```

```
  ! allocate storage for simulation data.
```

```
  ALLOCATE(pos(natoms,3), vel(natoms,3), frc(natoms,3, nthreads))
```

```
DO i=1, natoms
```

```
  READ(12, *) (pos(i,j), j=1,3)
```

```
END DO
```

```
DO i=1, natoms
```

```
  READ(12, *) (vel(i,j), j=1,3)
```

```
END DO
```

```
CLOSE(12)
```

```
! set up cell list
```

```
CALL mkcell
```

```
CALL updcell
```

```
! initialize forces and energies
```

```
nfi=0
```

```
frc(:, :, :) = 0.0_dbl
```

```
CALL force
```

```
CALL getekin
```

```
WRITE(stdout, *) 'Starting simulation with ', natoms,  
' atoms for', nsteps, ' steps'
```

```
WRITE(stdout, *) '      NFI           TEMP
```

```
EKIN           EPOT&  
      &           ETOT'
```

```
CALL output
```

```
! main MD loop
```

```
DO nfi=1, nsteps
```

```
  ! write output, if requested
```

```
  IF (mod(nfi, nprint) == 0) THEN
```

```
    CALL output
```

```
  END IF
```

Modules of Fortran program & Python Interface

```
! propagate system and recompute energies
CALL updcell
CALL velverlet
CALL getekin
END DO

! clean up: close files, free memory
WRITE(stdout,'(A)') 'Simulation Done.'
CALL rmccl
CALL ioclose
DEALLOCATE(pos,vel,frc)
END SUBROUTINE ljmd

SUBROUTINE openfiles(pyfname)
  IMPLICIT NONE
  CHARACTER(LEN=30) :: ifname,restfname, trajfname,
  ergfname, logfname,fname
  CHARACTER*(*) :: pyfname
  INTEGER :: iosl,iosR, length!, nprint
  fname = TRIM( pyfname )
  length = LEN_TRIM( fname )
  ifname = TRIM(fname)
  ifname( length+1:length+4 ) = ".inp"
  OPEN(UNIT=11,FILE=ifname,IOSTAT=iosl,STATUS='OLD')
  IF( iosl /= 0 ) THEN
    WRITE(6,*) " "
    WRITE(6,*) "No input file ", ifname, " found."
    WRITE(6,*) " "
  ELSE
    READ(11,*) natoms
    READ(11,*) mass
    READ(11,*) epsilon
    READ(11,*) sigma
```

```
READ(11,*) rcut
  READ(11,*) box
  READ(11,*) nsteps
  READ(11,*) dt
  READ(11,*) nprint
  rewind(11)
END IF
restfname = TRIM(fname)
restfname( length+1:length+5 ) = ".rest"
OPEN( UNIT=12, FILE=restfname, IOSTAT=iosR, STATUS="OLD" )
IF( iosR /= 0 ) THEN
  WRITE(6,*) " "
  WRITE(6,*) "No restart file ", restfname, " found."
  WRITE(6,*) "Program STOP "
  WRITE(6,*) " "
  STOP
END IF

trajfname = TRIM(fname)
trajfname( length+1:length+4 ) = ".xyz"
OPEN( UNIT=13, FILE=trajfname, STATUS="UNKNOWN" )

ergfname = TRIM(fname)
ergfname( length+1:length+4 ) = ".dat"
OPEN( UNIT=14, FILE=ergfname, STATUS="UNKNOWN" )

logfname = TRIM(fname)
logfname( length+1:length+4 ) = ".log"
OPEN( UNIT=15, FILE=logfname, STATUS="UNKNOWN" )

END SUBROUTINE openfiles

END MODULE ljmd
```

Modules of Fortran program & Python Interface

```
# -*- Makefile -*-
SHELL=/bin/sh
FC=gfortran
FFLAGS=-Wall -g -std=f95 -O3 -ffast-math -fomit-frame-pointer
AR=ar

# list of source files
SRC=kinds.f90 physconst.f90 mdsys.f90 getekin.f90 utils.f90
io.f90 cell.f90 force.f90 velverlet.f90 ljmd.f90 interface.f90
VPATH=../src
OBJ=$(SRC:%.f90=%.o)

default: ljmd.so

clean:
    rm -f *.mod *.o *.a *.so *.pyf

# linker rule
../ljmd-serial.x: $(OBJ)
    $(FC) -o $@ $(FFLAGS) $^
```

```
# compilation pattern rule for objects
%.o: %.f90
    $(FC) -c -fPIC $(FFLAGS) $<

# dependencies
ljmd.o: ljmd.f90

# signature file creation
ljmd.pyf : ../src/interface.f90
    f2py -m ljmd -h $@ $<

# concatenation of libraries
libljmd.a : $(OBJ)
    ar crvs $@ $^

# compilation of share library
ljmd.so : ljmd.pyf libljmd.a
    f2py -m ljmd --fcompiler=gfortran -c $^
```


Documentation

- 2 versions of documentation was created. Doxygen was used to extract structure from the code
- 1st version is based on original code
- 2nd version is based on our modified code

Kinds.f90

!> Return the kind value of a real data type. Define the precision of double and single precision.

```
MODULE kinds
```

```
  IMPLICIT NONE
```

```
  INTEGER, PARAMETER :: dbl = selected_real_kind(14,200) ! double precision float
```

```
  INTEGER, PARAMETER :: sgl = selected_real_kind(6,30)   ! single precision float
```

```
  INTEGER, PARAMETER :: sln = 200                       ! length of I/O input line
```

```
  PRIVATE
```

```
  PUBLIC :: sgl, dbl, sln
```

```
END MODULE kinds
```

Chapter 1

Data Type Index

1.1 Data Types List

Here are the data types with brief descriptions:

[kinds](#)

Return the kind value of a real data type. Define the precision of double and single precision 3

[mdsys](#)

Define all the variables used in the molecular dynamics 3

[physconst](#)

Define the physics constants: boltzman constant and $m \cdot v^2$. It is dependent of module kinds 4

Doc: Unit Testing

Test-case ID:

Test-case Statement:

Purpose (Prerequisite):

Test (Input) Data:

Steps to be executed:

Expected Result:

Actual Result:

Pass/Fail:

Comments:

Changes in Simulation Infrastructure

Extended the simulation infrastructure so it can handle N type of atoms instead of one atom
Tried to do it with minimum change to original code
And it work!

Demo

Link: <https://github.com/chrismacdermaid/ljmd-f>

