

```
1      subroutine gaussfg_init(natom,nbonds,ib,jb,nquant,labels,nlink)
2
3 !    callするとき:
4 !      gaussfg_init(natom,mbona,ix(i18),ix(i20),nquant,labels,nlink)
5
6 !      natom      : 全原子数（リンク原子含まず）
7 !      nbonds     : 水素を含まないbondの数
8 !      ib(*)      : (bondする原子iのインデックス-1)x3
9 !      jb(*)      : (bondする原子jのインデックス-1)x3
10 !      nquant     : QM原子数（リンク原子含まず）
11 !      labels(*)  : QM原子のインデックスリスト
12 !      nlink      : リンク原子の数
13 !
14 !      nqm_gfg    : - QM原子数
15 !      nln_gfg    : - リンク原子数
16 !      nqmln_gfg  : - QM原子数+リンク原子数
17 !      iqmln_gfg(*) : - リンクしているQM原子インデックス
18 !      immln_gfg(*) : - リンクしているMM原子インデックス
19 !
20 !      gh_route_gfg : - Gaussian route section
21 !      gh_molsp_gfg : - Gaussian molecule specification section
22
23 use iflport ! Intel Fortran 7.0
24 !      use ifport ! Intel Fortran 9.0
25
26 implicit none
27 integer natom,nbonds,ib(*),jb(*),nquant,labels(*),nlink
28
29 integer natom_gfg, nqm_gfg, nln_gfg, nqmln_gfg,
30 &      iqmln_gfg(1000), immln_gfg(1000)
31 character gh_route_gfg*400, gh_molsp_gfg*80
32
33 common /gfg/ natom_gfg, nqm_gfg, nln_gfg, nqmln_gfg,
34 &      iqmln_gfg, immln_gfg,
35 &      gh_route_gfg, gh_molsp_gfg
36
37 integer i, j, ii, jj, mm
38 integer io1, ierr
39 parameter (io1=70)
40
41 ! gaussian.headerがないときはリターン
42 inquire(file='gaussian.header', exist=ierr)
43 if(.not. ierr) return
44
45 ! 定数
46 natom_gfg = natom
47 nqm_gfg = nquant
48 nln_gfg = nlink
49 nqmln_gfg = nqm_gfg+nln_gfg
50
51 ! Gaussianヘッダーの読み込み
52 open(io1,file='gaussian.header',status='old',action='read')
53 read(io1,'(a)') gh_route_gfg ! forceやchargeが入っていてはいけない
54 read(io1,'(a)') gh_molsp_gfg
55 close(io1)
56
57 ierr = runqq('rm', '-f gaussian-debug.log') ! log
58
59 ! リンク原子に結合しているQM原子,MM原子の取得
60 mm = 0
61 do i=1, nbonds
62   ii = 0
63   jj = 0
64   do j=1, nquant
65     if((ib(i)/3+1)==labels(j)) ii=1
```

```
66      if((jb(i)/3+1)==labels(j)) jj=1
67      enddo
68      if((ii+jj)==1) then
69          if(ii==1) then
70              iqmln_gfg(mm+1) = ib(i)/3+1
71              immln_gfg(mm+1) = jb(i)/3+1
72          else
73              iqmln_gfg(mm+1) = jb(i)/3+1
74              immln_gfg(mm+1) = ib(i)/3+1
75          endif
76          mm = mm + 1
77      endif
78  enddo
79
80  end
81
82 ! -----
83
84 SUBROUTINE gaussfg(XPARAM,INT,ESCF,FULSCF,GRAD,LGRAD,
85 .                      coord,nquant,crdsmm,chrags,iread,dxyzqm,
86 .                      dxyzcl,icalcf,iqmres,mark,jqatms)
87 C
88 C Routine has been modified for qm-mm coupling and includes COSMO.
89 C
90 C Variables for qm-mm:
91 C
92 C coord(3,numatm) - Cartesian coordinates of qm atoms. This was
93 C originally a MOPAC array defined locally in
94 C COMPFG and filled by GMETRY. Now Cartesian
95 C coordinates are passed directly.
96 C nquant           - Number of qm atoms.
97 C crdsmm(*)       - Main AMBER array containing coordinates for
98 C all atoms (used here for mm atoms).
99 C chrags(*)       - Atomic charges for mm atoms.
100 C iread            - Flag for reading in density matrix 0=no, 1=yes.
101 C dxyzqm(3,*)     - Quantum mechanical derivatives from qm-mm
102 C interactions.
103 C dxyzcl(3,*)     - Classical derivatives from qm-mm interactions.
104 C icalcf           - Flag for calculating atomic forces 0=yes, 1=no.
105 C iqmres(*)        - Atom-based interaction list.
106 C mark(*)          - Center of geometry atom marker list.
107 C jqatms(*)        - Atom numbers for qm atoms (based on AMBER scheme).
108 C
109 C
110 C          -- S. Dixon (8/11/94)
111 C
112 IMPLICIT DOUBLE PRECISION (A-H,O-Z)
113 *COMDECK SIZES
114 ****
115 * THIS FILE CONTAINS ALL THE ARRAY SIZES FOR USE IN MOPAC.
116 *
117 * THERE ARE ONLY 5 PARAMETERS THAT THE PROGRAMMER NEED SET:
118 * MAXHEV = MAXIMUM NUMBER OF HEAVY ATOMS (HEAVY: NON-HYDROGEN ATOMS)
119 * MAXLIT = MAXIMUM NUMBER OF HYDROGEN ATOMS.
120 * MAXTIM = DEFAULT TIME FOR A JOB. (SECONDS)
121 * MAXDMP = DEFAULT TIME FOR AUTOMATIC RESTART FILE GENERATION (SECS)
122 * ISYBYL = 1 IF MOPAC IS TO BE USED IN THE SYBYL PACKAGE, =0 OTHERWISE
123 * SEE ALSO NMECI, NPULAY AND MESP AT THE END OF THIS FILE
124 *
125 PARAMETER (MAXHEV=40, MAXLIT=40)
126 PARAMETER (MAXTIM=3600, MAXDMP=3600)
127 PARAMETER (ISYBYL=1)
128 *
129 ****
130 *
```

```
131 * THE FOLLOWING CODE DOES NOT NEED TO BE ALTERED BY THE PROGRAMMER
132 *
133 ****
134 *
135 * ALL OTHER PARAMETERS ARE DERIVED FUNCTIONS OF THESE TWO PARAMETERS
136 *
137 *      NAME          DEFINITION
138 *      NUMATM        MAXIMUM NUMBER OF ATOMS ALLOWED.
139 *      MAXORB        MAXIMUM NUMBER OF ORBITALS ALLOWED.
140 *      MAXPAR        MAXIMUM NUMBER OF PARAMETERS FOR OPTIMISATION.
141 *      N2ELEC        MAXIMUM NUMBER OF TWO ELECTRON INTEGRALS ALLOWED.
142 *      MPACK         AREA OF LOWER HALF TRIANGLE OF DENSITY MATRIX.
143 *      MORB2         SQUARE OF THE MAXIMUM NUMBER OF ORBITALS ALLOWED.
144 *      MAXHES        AREA OF HESSIAN MATRIX
145 *      MAXALL        LARGER THAN MAXORB OR MAXPAR.
146 ****
147      PARAMETER (VERSON=7.00D0)
148      PARAMETER (NUMATM=MAXHEV+MAXLIT)
149      PARAMETER (MAXORB=4*MAXHEV+MAXLIT)
150      PARAMETER (MAXPAR=3*NUMATM)
151      PARAMETER (MAXBIG=MAXORB*MAXORB**2)
152      PARAMETER (N2ELEC=(50*MAXHEV*(MAXHEV-1)+10*MAXHEV*MAXLIT
153      + (MAXLIT*(MAXLIT-1))/2))
154      PARAMETER (MAXHES=(MAXPAR*(MAXPAR+1))/2,MORB2=MAXORB**2)
155      PARAMETER (MPACK=(MAXORB*(MAXORB+1))/2)
156      PARAMETER (MAXPR=6*MAXORB)
157      PARAMETER (MAXALL=4*MAXHEV+3*MAXLIT)
158      PARAMETER (NMECI=11, NPULAY=MPACK, MMCI=60)
159      PARAMETER (MESP=10)
160      PARAMETER (LENABC=600)
161      PARAMETER (LENAB2=LENABC*(LENABC+5))
162      PARAMETER (NPPA = 1082, MAXNSS = 500)
163      PARAMETER (MAXDEN=10*MAXHEV+MAXLIT)
164 ****
165 *DECK MOPAC
166      dimension crdsmm(*),chrgs(*),dxyzqm(3,*),dxyzcl(3,*),iqmres(*),
167      . mark(*),jqatms(*)
168      . DIMENSION XPARAM(MAXPAR),GRAD(MAXPAR)
169      LOGICAL LGRAD, FULSCF
170      COMMON/KTLLBTL/KILLIT
171      COMMON /GEOVAR/ NVAR,LOC(2,MAXPAR),IDUMY,DUMY(MAXPAR)
172      COMMON /GEOSYM/ NDEP,LOCPAR(MAXPAR),IDEPFN(MAXPAR),LOCDEP(MAXPAR)
173      COMMON /GEOM / GEO(3,NUMATM)
174      COMMON /ATHLET/ ATHEAT
175      COMMON /WMATRX/ WJ(N2ELEC), WK(N2ELEC)
176      COMMON /ENUCLR/ ENUCLR
177      COMMON /NATYPE/ NZTYPE(107),MTYPE(30),LTYPE
178      COMMON /ELECT / ELECT
179      PARAMETER (MDUMY=MAXPAR**2-MPACK)
180      COMMON /SCRACH/ RXYZ(MPACK), XDUMY(MDUMY)
181      COMMON /HMATRX/ H(MPACK)
182      COMMON /GEOKST/ NATOMS,LABELS(NUMATM),
183      . NA(NUMATM), NB(NUMATM), NC(NUMATM)
184      COMMON /ERRFN / ERRFN(MAXPAR), AICORR(MAXPAR)
185      COMMON /VECTOR/ C(MORB2),EIGS(MAXORB),CBETA(MORB2),EIGB(MAXORB)
186      COMMON /LAST / LAST
187      COMMON /NUMCAL/ NUMCAL
188      COMMON /SCFTYP/ EMIN, LIMSCF
189      COMMON /MOLMEC/ HTYPE(4),NHCO(4,20),NNHCO,ITYPE
190      . /MOLKST/ NUMAT,NAT(NUMATM),NFIRST(NUMATM),NMIDDLE(NUMATM),
191      . 2 NLAST(NUMATM), NORBS, NELECS,NALPHA,NBETA,
192      . 3 NCLOSE,NOPEN,NDUMY,FRACT
193      C COSMO change
194      LOGICAL ISEPS, USEPS , UPDA, CFIRST
195      COMMON /ISEPS/ ISEPS, USEPS, UPDA
```

```
196      DATA CFIRST /.TRUE./
197      C end of COSMO change
198      C*****
199      C
200      C  COMPG CALCULATES (A) THE HEAT OF FORMATION OF THE SYSTEM, AND
201      C                  (B) THE GRADIENTS, IF LGRAD IS .TRUE.
202      C
203      C  ON INPUT XPARAM = ARRAY OF PARAMETERS TO BE USED IN INTERNAL COORDS
204      C          LGRAD = .TRUE. IF GRADIENTS ARE NEEDED, .FALSE. OTHERWISE
205      C          INT   = .TRUE. IF HEAT OF FORMATION IS TO BE CALCULATED
206      C          FULSCF = .TRUE. IF FULL SCF TO BE DONE, .FALSE. OTHERWISE.
207      C
208      C  ON OUTPUT ESCF = HEAT OF FORMATION.
209      C          GRAD = ARRAY OF GRADIENTS, IF LGRAD = .TRUE.
210      C
211      C*****
212      COMMON /KEYWRD/KEYWRD
213      COMMON/ANALZZ/ICALX1,ICALX2,ICALX3,ICALCN
214      CHARACTER*241 KEYWRD
215      LOGICAL DEBUG, INT, PRINT, ANALYT, LARGE, USEDICI,
216      1FORCE, TIMES, AIDER
217      DIMENSION COORD(3,NUMATM), W(N2ELEC), DEGREE(3), XPARSE(MAXPAR)
218      1,DELTAP(NMECI**2), DELTA(NMECI*MAXORB)
219      SAVE DEGREE, PRINT, DEBUG, aider,analyt,large,usedci,
220      1 force,times
221      EQUIVALENCE (W,WJ)
222      C DATA ICALCN /0/
223      logical first_call
224      data first_call /.true./
225      save first_call
226
227      !wpp
228      !上の部分について不要な部分多数あり
229      !メモ :
230      ! * gaussfgのnquantにはlink原子の数がすでに含まれている
231      ! * jqatms(i)は 原子iがQMのとき、その原子番号
232      !           Linkのとき、 -1
233      !           MMのとき、 0 を返す
234      !
235      ! 1 hartree = 627.51d0 kcal/mol
236      ! 1 bohr = 0.52917706d0 angstrom
237
238      integer io1, ierr
239      parameter (io1=70)
240
241      double precision qscale, escale, fscale
242      data qscale /18.2223d0/
243      parameter (escale=627.51d0, fscale=escale/0.52917706d0)
244
245      integer natom_gfg, nqm_gfg, nln_gfg, nqmln_gfg,
246      &          iqmln_gfg(1000), immln_gfg(1000)
247      character gh_route_gfg*400, gh_molsp_gfg*80
248
249      common /gfg/ natom_gfg, nqm_gfg, nln_gfg, nqmln_gfg,
250      &          iqmln_gfg, immln_gfg,
251      &          gh_route_gfg, gh_molsp_gfg
252
253      integer i, j, indxmm, ni_mm, jmm, jcoord
254      logical fcharg, mmflgs(natom_gfg)
255      double precision rx, ry, rz, gx, gy, gz, gmx, gmy, gmz
256      double precision ln_r, one_r, gr3
257
258      ! if(first_call) then
259      !     first_call = .false.
260      ! endif
```

```
261      !リンク原子の位置を決める
262      do i=1, nln_gfg
263          do j=1,nqm_gfg
264              if(jqatms(j)==iqm_ln_gfg(i)) then
265                  rx = crdsmm((immln_gfg(i)-1)*3+1) - coord(1,j)
266                  ry = crdsmm((immln_gfg(i)-1)*3+2) - coord(2,j)
267                  rz = crdsmm((immln_gfg(i)-1)*3+3) - coord(3,j)
268                  exit
269              endif
270          enddo
271          ln_r = 1.09d0/sqrt(rx*rx+ry*ry+rz*rz)
272          do j=1,nqm_gfg
273              if(jqatms(j)==iqm_ln_gfg(i)) then
274                  coord(1,nqm_gfg+i) = coord(1,j) + rx*ln_r
275                  coord(2,nqm_gfg+i) = coord(2,j) + ry*ln_r
276                  coord(3,nqm_gfg+i) = coord(3,j) + rz*ln_r
277                  exit
278              endif
279          enddo
280      enddo
281
282
283      ! QMと相互作用するMM原子を検索
284      fcharg = .false.
285      mmflgs = .false.
286      idxmm = 1
287      do i=1,nqmln_gfg
288          ni_mm = iqmrres(idxmm)
289          idxmm = idxmm + 1
290          do j=1,ni_mm
291              jmm = iqmrres(idxmm)
292              jcoord = (jmm-1)*3
293              mmflgs(jmm) = .true.
294              idxmm = idxmm + 1
295          enddo
296      enddo
297      if(any(mmflgs == .true.)) fcharg = .true.
298
299      ! Link原子に結合しているMM原子を省く
300      if(fcharg) then
301          do i=1, nln_gfg
302              mmflgs(immln_gfg(i)) = .false.
303          enddo
304      endif
305
306      ! gaussian input fileを作成する
307      open(io1,file='gaussian.in',status='replace',action='write')
308
309      ! gjf:header
310      write(io1,'(a)') '%nosave'
311      write(io1,'(a)') '%rwf=elec.rwf'
312      if(fcharg) then
313          write(io1,'(a)') trim(gh_route_gfg)//' force iop(6/13=1) charge'
314      else
315          write(io1,'(a)') trim(gh_route_gfg)//' force iop(6/13=1)'
316      endif
317      write(io1,'(a)')
318      write(io1,'(a)') 'QM/MM'
319      write(io1,'(a)')
320      write(io1,'(a)') gh_molsp_gfg
321
322      ! gjf:QM
323      !labels,coordは先頭から使うので、i->jqatms(i)としなくてよい
324      do i=1,nqmln_gfg
325          write(io1,'(i,3f)') labels(i),(coord(j,i),j=1,3)
```

```
326      enddo
327      write(io1,'(a)')
328
329      ! gjf:MM
330      if(fcharg) then
331          do i=1,natom_gfg
332              if(mmflgs(i)) then
333                  write(io1,'(4f)')
334                  & (crdsmm((i-1)*3+j),j=1,3), chrgs(i)/qscale
335              endif
336          enddo
337          write(io1,'(a)')
338          do i=1,natom_gfg
339              if(mmflgs(i)) then
340                  write(io1,'(4f)')
341                  & (crdsmm((i-1)*3+j),j=1,3)
342              endif
343          enddo
344          write(io1,'(a)')
345      endif
346
347      close(io1)
348
349      ! Gaussianを実行する
350
351      ierr = runqq('cat','gaussian.in >> gaussian-debug.log')
352      ierr = runqq('rm','-f elec.rwf')
353      ierr = runqq('g03b3','gaussian.in') ! output: elec.rwf
354      if(ierr== -1) then
355          write(6,*) 'FATAL ERROR IN GAUSSFG - g03 FAILS'
356          stop
357      endif
358      ierr = runqq('rm','-f gdatas.wpp')
359      ierr = runqq('./PrtGVal3.exe','elec.rwf') ! output: gdatas.wpp
360      if(ierr== -1) then
361          write(6,*) 'FATAL ERROR IN GAUSSFG - PrtGval3 FAILS'
362          stop
363      endif
364
365      ! ファイルからエネルギーと力を読み込む
366      open(io1,file='gdatas.wpp',status='old',action='read')
367
368      read(io1,'(a)')
369      ! エネルギー
370      ! read(io1,'(3f)') escf, elect, enuclr
371      ! write(*,*) '!wpp ener:', escf*escale, elect*escale, enuclr*escale
372      ! elect = 0.d0
373      ! enuclr = escf*(escale/23.060362d0)
374      ! escf = (elect+enuclr)*23.060362d0
375
376      if(escf<emin.or.emin==0.d0) emin=escf
377
378      if(icalcf/=0) return ! エネルギーだけ計算して力はupdateしない
379
380      read(io1,'(a)')
381      ! QM原子とLink原子
382      ! dxyzqmは先頭から使う
383      do i=1,nqmln_gfg
384          read(io1,'(3f)') gx, gy, gz
385          dxyzqm(1,i) = -gx*fscale
386          dxyzqm(2,i) = -gy*fscale
387          dxyzqm(3,i) = -gz*fscale
388      enddo
389      read(io1,'(a)')
390      ! MM原子
```

```

391      !dxyzclはそのままの配列
392      if(fcharg) then
393          do i=1,natom_gfg
394              if(mmflgs(i)) then
395                  read(io1,'(3f)') gx, gy, gz
396                  dxyzcl(1,i) = -gx*fscale
397                  dxyzcl(2,i) = -gy*fscale
398                  dxyzcl(3,i) = -gz*fscale
399              endif
400          enddo
401      endif
402
403      close(io1)
404
405      ! Link原子の力をQM原子,MM原子に分配する
406      do i=1, nqm_gfg
407          gx = dxyzqm(1,nqm_gfg+i)
408          gy = dxyzqm(2,nqm_gfg+i)
409          gz = dxyzqm(3,nqm_gfg+i)
410          do j=1,nqm_gfg
411              if(jqatms(j)==iqmln_gfg(i)) then
412                  rx = crdsmm((imm_ln_gfg(i)-1)*3+1) - coord(1,j)
413                  ry = crdsmm((imm_ln_gfg(i)-1)*3+2) - coord(2,j)
414                  rz = crdsmm((imm_ln_gfg(i)-1)*3+3) - coord(3,j)
415                  exit
416              endif
417          enddo
418          one_r = 1.d0/sqrt(rx*rx+ry*ry+rz*rz)
419          rx = rx*one_r
420          ry = ry*one_r
421          rz = rz*one_r
422          ln_r = 1.09d0*one_r
423          gr3 = gx*rx+gy*ry+gz*rz
424          gmx = ln_r*(gx-gr3*rx)
425          gmy = ln_r*(gy-gr3*ry)
426          gmz = ln_r*(gz-gr3*rz)
427          dxyzcl(1,imm_ln_gfg(i)) = dxyzcl(1,imm_ln_gfg(i)) + gmx
428          dxyzcl(2,imm_ln_gfg(i)) = dxyzcl(2,imm_ln_gfg(i)) + gmy
429          dxyzcl(3,imm_ln_gfg(i)) = dxyzcl(3,imm_ln_gfg(i)) + gmz
430          do j=1,nqm_gfg
431              if(jqatms(j)==iqmln_gfg(i)) then
432                  dxyzqm(1,j) = dxyzqm(1,j) + (gx-gmx)
433                  dxyzqm(2,j) = dxyzqm(2,j) + (gy-gmy)
434                  dxyzqm(3,j) = dxyzqm(3,j) + (gz-gmz)
435                  exit
436              endif
437          enddo
438          dxyzqm(1,nqm_gfg+i) = 0.0d0
439          dxyzqm(2,nqm_gfg+i) = 0.0d0
440          dxyzqm(3,nqm_gfg+i) = 0.0d0
441      enddo
442
443      ! write(*,'(a)') 'gaussfg: gradient(QM)'
444      ! do i=1,nqm_ln_gfg
445      !     write(*,'(i5,3(2x,e15.7))') jqatms(i),(dxyzqm(j,i),j=1,3)
446      ! enddo
447      ! write(*,'(a)') 'gaussfg: gradient(MM)'
448      ! do i=1,natom_gfg
449      !     write(*,'(i5,3(2x,e15.7))') i,(dxyzcl(j,i),j=1,3)
450      ! enddo
451
452      4000 RETURN
453      END

```