RUNNER v.1

User manual

Program description

RUNNER is a program that performs a global search for the optimal structure of atomic clusters and the structure of their low-lying isomers, during which the cluster energy is evaluated using the quantum chemical (DFT or *ab initio*) or classical interatomic potential / force field calculations.

RUNNER carries out the global minimum search using the specialized server programs for energy evaluation and local optimization of the cluster geometry. Currently, the following programs can be servers for calculating energy and local optimization:

- Gaussian 03/09/16
- NWChem v. 6/7
- GLOBUS v.1

The global search method implemented in the program is the evolutionary (genetic) algorithm combined with a taboo search. The detailed description of global optimization algorithm implemented in the program is given in the section *Global optimization algorithm implemented in RUNNER*.

Obtaining and installing the program

The program, control script and examples of input files can be downloaded from the resource https://github.com/skignatov/runner-release

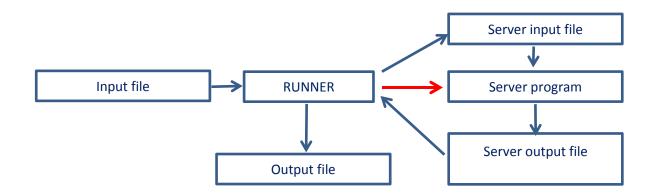
To get started, it is enough to create a separate directory on your computer, place the program, the control script (if necessary, see the section *Program operation modes*) and data files required for the selected operation mode (see the section *Files required for running*). In addition, the selected server program must also be correctly installed on the computer being used (or computers playing the role of remote computing nodes) (obtained and installed separately). The RUNNER start commands are described in the section *Running a program*.

Operating modes of the program

The program can work in several modes:

1. *Direct call mode* (Windows). In this mode, the program reads the main input data file, sequentially generates initial data files for the server program, and starts the server program on the computer being used to calculate with the generated file. After the end of

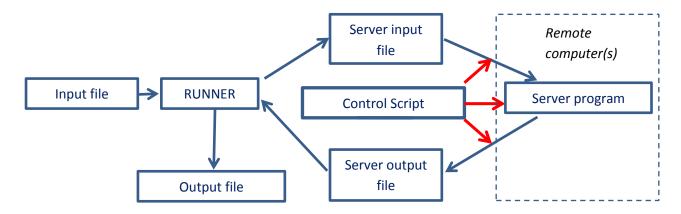
the calculation, the output file of the server program is analyzed, and information about the structure and energy of the cluster is extracted from it:



The direct call mode implements only sequential calculation of the structures under study. It is recommended to be used only for those calculations in which the calculation of energy takes a short time (for example, calculation based on empirical potentials and force fields by the GLOBUS program, quantum chemical calculation by the semiempirical method). This mode has been tested only in Windows environment.

2. SSH mode (LINUX). In this mode, the program reads the input data file, generates source data files for the server program (several files in parallel, the number of which is controlled by the MaxJobs command). The generated files are placed in the ./inp directory, which is automatically created in the current directory. A concurrently running control script (CS) runner_ssh_ <version> .sh reads files in the ./inp directory and starts server calculations on one or several remote computers (nodes) that are connected to the central node via the ssh protocol. The central computer can also act as a remote site, i.e. tasks can be run on it as well. Upon successful launch of this file, the CS moves it from Jinp to the Jout directory, while creating a copy in the Jsub directory. Files in this directory ./sub are named $\langle file \rangle$. $\langle ext \rangle$ X, where X is the number of the node on which the calculation is started. The control system cyclically checks the status of calculations at remote nodes and upon completion of the calculation (successful or emergency) downloads the results file to the ./out directory, deleting the corresponding file from ./sub. RUNNER, detecting a new output file in the ./out directory, parses it, extracting optimized structures and energies, and moves the input and output files from ./out to the ./arc directory. Depending on the reason for completing this calculation, RUNNER generates a new input file to continue the unfinished calculation or proceeds to generate a new point for global search. The generated files have the format of the name geo-XXXXXX-YY. <ext>, where XXXXXX is the number of the global search point, YY is the number of the continuation of the unfinished calculation (YY = 00 for a new file). The list of nodes on which server programs can be launched and calculations can be performed is contained in the nodes.txt file, which must be present in the startup directory. The CS can be started and stopped manually regardless of RUNNER (before or after it), or it can be started, restarted (in case of an emergency shutdown of the CS) and stopped by the program automatically (see the ShellControl command). SSH mode is recommended for tasks with long-term energy calculation (DFT, ab initio) on

workstations that are not combined into a cluster. Since quantum chemical calculations are performed in parallel at remote nodes, an increase in the number of available nodes increases the overall performance of GO.



- 3. BATCH mode. (LINUX on supercomputers or clusters). In this mode, the operation is completely similar to the SSH mode, but the control script runner batch <version> .sh starts tasks on a supercomputer (SC) or cluster using the task scheduler - slurm or pbs. The control script for pbs is different from the slurm script and is called runner_batch_pbs_<version> .sh. Unlike SSH mode, BATCH mode does not require a nodes.txt file, but two other files are required: (1) file workdirs.txt, which contains a list and description of the working directories where the results of individual tasks will appear, and (2) files of type batch.template which are forms for generating scripts for batch launching of computational tasks, formed according to the rules of the SC used. Files batch.template can be different and/or have different names for each working directory. Their respective names are specified in workdirs.txt for each working directory. When working in this mode, one or several directories are created in the program launch directory (their names are described in workdirs.txt, usually wd0, wd1, wd2 ...) so that tasks for calculating individual global search points are launched in each directory in parallel. These directories also contain the output files of this calculation and all temporary files of the server program. The CS analyzes the files in the working directories and, upon completion of the calculation, moves them to the ./out directory. After that, a new task is launched in this working directory, the input file of which is taken from the ./inp directory. This mode of operation is the main one and the most convenient when calculating on the SC. An increase in the number of working directories leads to an acceleration of the calculation only within certain limits, since a large number of simultaneously running tasks can lead to the fact that some of them will be queued or will be blocked by the scheduler due to exceeding the task limit for one user. Often, 8-12 working directories are optimal for a single system.
- 4. CRON mode, the batch mode with periodic start of control script. (LINUX on supercomputers). On some SCs, the administrator prohibits continuous operation of the control script on a central node. To work on such computers, the CRON mode is used, which is completely similar to the BATCH mode, however, the control unit and the RUNNER program do not work continuously, but are started periodically by the *cron* command with a period of several minutes. The *cron* setting is performed by the control

script itself at the initial startup. This mode is less stable than the BATCH mode and it is recommended to use it only in case of strict restrictions on the operation of the control script.

Files required for running

Hereinafter, global optimization of the Mg_{10} cluster is considered as an example. All examples use mg10 as the file name. During operation, this name should be replaced with the file name corresponding to a specific task.

File names and other parameters specified in <...> can vary at the request of the user. Files without such brackets must have names that exactly match those indicated in the table.

All data files must be in the startup directory. Programs can be located in directories to which the search path is set.

1. Direct call mode

File	Description
runner.exe (Windows)	Program RUNNER
runner.x (LINUX)	
<mg10>.inp</mg10>	The main source data file. Describes the system to be
	optimized and optimization parameters
< <i>mg.gj0</i> >	Server program data file template

2. SSH mode

File	Description
runner.x	Program RUNNER
runner_ssh_ <version>.sh</version>	Control script
<mg10>.inp</mg10>	The main source data file. Describes the system to be
	optimized and optimization parameters
< <i>mg.gj0</i> >	Server program data file template
nodes.txt	A file describing the ssh nodes on which the server
	programs will run.

3. Режим ВАТСН

File	Description
runner.x	Program RUNNER
runner_batch_ <version>.sh</version>	Control script (scheduler <i>slurm</i>). If <i>pbs</i> is used as a
	scheduler the script name is
	runner_batch_pbs<_version>.sh
< mg10 > .inp	The main source data file. Describes the system to be
	optimized and optimization parameters
< <i>mg.gj0</i> >	Server program data file template
workdirs.txt	A file describing the directories in which individual
	tasks will be run. For each directory, its name is
	indicated, whether it is active (whether it is participating
	in the work at this time) and the name of the form of the

task launch script.

<batch.template> Form (template) of the task launch script. There can be

several such files, they can have different names. These files are assigned to separate working directories in the

workdirs.txt file

4. *Режим CRON*

All files are completely identical to BATCH mode, except for the control script, which in this case is called *runner_batch_cron* <_*version*> .*sh* (for *slurm* scheduler).

Data files required for RUNNER operation

1. The main file of the input data of RUNNER <mg10>.inp

The main input data file for RUNNER is an ASCII text file. Keywords are located on lines, the first nonblank character of which is #. Blank lines and lines without a leading # are ignored. Symbol! means the beginning of a comment, it can be located anywhere on the line. All information on this line to the right of! is considered a comment.

The order of keywords, both on a line and on different lines, can be any. Uppercase and lowercase letters are the same (except for chemical symbols in the *Stoichiometry* command)

Basic commands and keywords (values in brackets < ... > are examples):

Keyword	Meaning	Default value
Stoichiometry=< <i>Mg10</i> >	Sets the stoichiometry of the cluster to be	There is no
	optimized. The symbols of a chemical element	default, the
	should be written as given in the periodic table.	command must
	Uppercase and lowercase letters are different,	always be present
	so it is wrong to write mg10 or MG10, only	
	Mg10 is correct.	
Nstruct=<100>	The number of global search cycles, i.e. the	100
	number of generated and optimized structures.	
	Only those structures are considered optimized	
	if the optimization has successfully reached the	
	stopping criteria. The total number of	
	optimizations, some of which did not reach the	
	end or ended abnormally, may exceed Nstruct.	
MaxCyc=<10000>	The number of work cycles performed by the	100000
	RUNNER program while browsing the	
	working directories. Prevents infinite looping if	
	the number of optimized structures cannot	
	reach <i>Nstruct</i> for some reason. The maximum	
	running time of the program is	
	MaxCyc*IdleTime	
IdleTime=<120>	Waiting time at the end of each working cycle,	300
Idle=<120>	in seconds	
Server= <g09></g09>	A type of server program that performs local	g09
	optimization.	
FilPat=< <i>mg.gj0</i> >	The name of the template file of the server	No default value

	I	1
	input file. The %GEO% label should be placed	
	at the geometry placement. If the server is	
	Gaussian, the $\%chk=$ parameter must be	
	present.	
Seed=(<12345>,<67890>)	Seed for initializing the random number	Internal defaults
	generator. If there is no <i>Seed</i> command, the	of the random
	generator's internal defaults will be used. Used	number generator
	to repeat the results of a previous calculation.	
MaxPool=<30>	The maximum pool size. When it is exceeded,	50
	the found local minima, the energy of which	
	exceeds the values of all structures in the pool,	
	cannot participate in the generation of new	
	structures.	
ActivePool=<0.5>	The active part of the pool, participating in the	0.7
	generation of new structures.	
MinPool=<5>	The minimum pool size at which the main GD	10
	phase begins	
MaxRestarts=<20>	The maximum number of restarts of the local	10
	optimization (LO) of the server program, i.e.	
	launches of the LO that were interrupted due to	
	an excess of the number of cycles, internal	
	errors, etc. Introduced to prevent endless server	
	work when repeating the same error. If the	
	number of restarts is exceeded, the structure is	
	placed in the ./hlp directory. This structure is	
	ignored by <i>Nstruct</i> .	
NoSaveArchive	Disable saving the results of the server	Missing (results
1 (0 2 4) 01 21 3111 (0	program in the archive. In normal operation	are saved in the
	mode, each time the LO is completed by the	./arc directory)
	server program, the input and output files of	
	this optimization (for example, *.gjf and *.log)	
	are sent to the directory ./arc (archive of	
	optimization results). If the files are very large	
	and disk space is limited, the NoSaveArchive	
	command cancels the save, saving disk space.	
GenOnly	In Serial mode, do not run a quantum chemical	Missing (start QC
Conomy	calculation, only generate input files	calculations)
BondLimits=(<1.>,<4>)	Limits of interatomic distances in Å at which	1 Å – 4 Å
Donathinus=(\1./,\4/)	atoms are considered to be bonded. Used when	111 711
	generating new structures by generation	
Sphere=7.	operators. The radius of the region in Å in which the	7 Å
Splicic—7.	atoms are generated by the operator RAND	/ A
MaxTry=10000	The maximum number of attempts to generate	5000
1v1ax 11 y = 10000	a cluster in which the link lengths satisfy	3000
	BondLimits	
Init-(>DAND\ >DND1\)		DAND DND1
Init=(<rand>,<rnd1>)</rnd1></rand>	Operators used to generate the initial	RAND, RND1
Init=(<rand=30,rnd1=< td=""><td>population in the Initial Phase. The numbers</td><td></td></rand=30,rnd1=<>	population in the Initial Phase. The numbers	
70)	after = indicate the likelihood of using the	
	given operator. If the probabilities are not	
	specified, they are considered equal (i.e. for	

	two aparators 500% and 500%) For the list of	1
	two operators - 50% and 50%). For the list of	
	available operators, see the section	
	Optimization Algorithm.	
Operators=(<pccr=50,< td=""><td>Operators used to generate the initial</td><td>PCCR=50,</td></pccr=50,<>	Operators used to generate the initial	PCCR=50,
RAND=10,RND1=10,TP	population in the Main Production Phase. The	RAND=10,
CR=20,IMOV=10>)	numbers after = show the probability of using	RND1=10,
	this operator (in%, automatically normalized to	TPCR=20,
	100). If the probabilities are not specified, they	IMOV=10
	are considered equal for all operators for which	
	the probabilities are not specified. For a list of	
	available operators, see the <i>Optimization</i>	
	Algorithm section.	
Continue		Missins
Continue	Continue the previously completed	Missing
	optimization (i.e., an optimization that has	(start new
	reached the Nstruct limit). To continue, the	optimization)
	files *.base_, *.pool_, *.sbas_, *.info_ of the	
	previous optimization will be used, they must	
	be located in the startup directory. With the	
	Continue command, the Nstruct parameter is	
	treated as an additional number of structures to	
	be examined during the ongoing optimization.	
ShellControl=	Management of starting, restarting and	If any keyword
(StartOnBeg,StartIfDown,	stopping the control script (CS).	(or the entire
StopOnFinal ,StopOnExit)	StartOnBeg – start CS in the beginning.	command) is
,	StartIfDown – restart SC if it is stopped/failed.	missing, its action
	StopOnFinal – stop SC after optimization.	is not taken.
	StopOnExit – stop SC on the program exit.	
	200 p 200 m program onto	

In this and subsequent sections, the tags and keywords shown in red in the examples must be present in the files. Other keywords and commands are optional.

Sample file <Mg10>.inp:

```
# Stoichiometry=Mg10 Nstruct=300
# Server=g09 FilPat=mg.gj0 Idle=60 NoAutoStart
!# SEED=(674127515,48622252)

# MaxPool=50 ActivePool=0.7 MinPool=10 MaxRestarts=10 !NoSaveArchive
# BondLIMIts=(1.,4.) Sphere=6. MaxTry=5000

# Init=(RND1=10,RAND=10)
# Operators=(PCCR=50,RAND=10,RND1=10,TPCR=20,IMOV=10)
```

2. *Template for server input file <mg.gj0>*

The file template for creating the input file of the server program is a slightly modified regular file of this program, in which the place where the Cartesian coordinates are located is marked with the %GEO% label. In addition, the %chk=TTT keyword (TTT is any text) must be present

in the files for the Gaussian program. During operation, the TTT text will be automatically replaced with the name of the temporary file *geo-XXXXXY-YY.chk*, where XXXXXX and YY correspond to the structure number and restart number during its optimization.

Sample file:

3. File nodes.txt

The file is an ASCII text file in which each line describes the node where the server program is running. Blank lines and lines starting with the # character are ignored. The part of the line after the # is treated as a comment.

Each line has the format:

IP	The IP address of the node at which the node is accessible via <i>ssh</i>
Active	Whether the node will be used in the current optimization (0 or 1)
User	Username for ssh connection
Nproc	The number of cores that can be used in the calculation on a given node
Mem	The amount of memory in MB that can be used on this node
Prog	Server program start command
Dir	Directory in which the calculation will be carried out

Sample file:

" .		**	3.7		-			
# ip	Active	e User	МЪ	roc Mem	Prog	Dir		
45.3.23.189	1	user1	4	16000MB	g16	/home/user1/Documents/gwork	#	0
45.3.23.190	1	user1	4	16000MB	g16	/home/user1/Documents/gwork	#	1
45.3.23.135	1	student	4	16000MB	g16	/srv1/gwork	#	2
45.3.23.158	1	student	4	16000MB	g16	/home/student/Documents/gwork	#	3
45.3.23.157	1	student	4	12000MB	g09	/home/student/Documents/gwork	#	4
45.3.23.153	1	student	8	6000MB	g09	/home/student/Documents/gwork	#	5
45.3.23.167	1	student	4	6000MB	g16	/srv1/gwork	#	6
45.3.23.165	1	student	4	6000MB	g09	/home/student/Documents/gwork	#	7
45.3.23.131	1	student	4	6000MB	g09	/home/student/Documents/gwork	#	8
45.3.23.134	1	student	4	6000MB	g09	/home/student/Documents/gwork	#	9
45.3.23.143	0	student	4	6000MB	g09	/home/student/Documents/gwork	#	10
45.3.23.144	1	student	4	6000MB	g09	/home/student/Documents/gwork	#	11

4. Файл workdirs.txt

The file is an ASCII text file, in which each line describes the directory in which the job will be launched using the BATCH method using *slurm* or *pbs*. Blank lines and lines starting with the # character are ignored. The part of the line after the # is treated as a comment.

Each line has the format:

<Dirname > <Active> <No.> <Script>

Dirname	Directory name	
Active	Whether the node will be used in the current optimization (0 or 1)	
No.	The number of the directory. Does not affect the work, used as a comment.	
Script	BATCH script template used to run jobs in this directory. Based on this template,	
	the CS creates the final batch-script in the given directory. The template must	
	contain the tags %%% and @@@ which are replaced by CS with, respectively,	
	the unique name of the task and the input file of the server program.	

Sample file:

#Dirname	Active?	No.	Script
wd0	1	0	batch.run0
wd1	1	1	batch.run0
wd2	1	1	batch.run0
wd3	1	3	batch.run0
wd4	1	4	batch.run0
wd5	1	5	batch.run0
wd6	1	6	batch.run0
wd7	1	7	batch.run0
wd8	1	8	batch.run0
wd9	1	9	batch.run0
wd10	1	10	batch.run_skx

5. File <batch.template>

The template file for creating a script for launching tasks in the BATCH and CRON modes is a slightly modified regular script of this program, in which the place where the name of the task is located is marked with a %%% mark, and the place where the name of the input file of the server program is placed is marked with @ @@. In the course of work, the text %%% will be automatically replaced with the task name in the form SXXXXXX, where S is the letter specified by the user in the *jletter* parameter in the control script file (see the section *Starting the program*), and XXXXX corresponds to the structure number during its optimization. Using different *jletter*="<S>" parameters for different systems to be optimized allows you to distinguish between the concurrent jobs displayed by the *squeue* and *qstat* commands.

Sample file (for slurm scheduler):

Sample file (for pbs scheduler):

Starting the program

1. Direct call mode

Run command (Windows):

```
runner.exe mg10.inp
```

2. SSH mode

The command to run the program in the background:

```
./runner.x mg10.inp & disown
```

Control script (CS) is launched automatically or manually before or after the program starts, depending on the *ShellControl* parameter (see the section *Input file keywords*). When starting manually, the command to start CS is:

```
./runner ssh.sh >res.log
```

At startup, the RS asks for passwords for all remote nodes, which must be entered by the user in response to requests:

```
>Password for node 1, user student: absde123
>Password for node 2, user user1: jhkk12h
...
```

After entering the passwords, the control system can be switched to the background mode:

```
Ctrl-Z
bg
disown
```

During the operation of the CS, the *res.log* file will contain updated information about its operation.

WARNING! Some versions of CS do not ask for passwords! They should be indicated inside the script body.

3. BATCH mode

In this mode, it is recommended to start control script first, which then automatically starts the program. Sample run command is (here, $_e30$ is the script version which can vary):

```
./runner_batch_e30.sh >res.log &
```

Before starting the CS, it should be edited using any editor (*mc*, *vi* ...), setting several mandatory options described at the beginning of the *runner_batch.sh* file:

```
#### Define your defaults here: ########
# These are important parameters! Check it carefully!
user="student" ← indicate you login (username)
InputFile=mg7.inp
                   ← indicate the runner input file
iletter="v"
                   ← indicate the latter to distinguish the jobs
curdir=$PWD
idle time=300
                    ← optional: sleep time, s
                    ← optional: maximum number of SC wakeups (cycles)
maxcyc=1000000
#for PBS only (prefix added by qsub to JobNo)
compname=".master1.cyberia.tsu.ru" ← for pbs: suffix of job number
                             if it is indicated in the qstat command
```

4. CRON mode

Launching the program (as well as configuring the CS before launching) is completely similar to the BATCH mode. Run command is

./runner batch cron.sh >res.log &

Global optimization algorithm implemented in RUNNER

The global optimization algorithm implemented in RUNNER is based on an evolutionary (genetic) algorithm combined with a taboo search. The genetic algorithm is based on the application of generating operators (generators) to the initial and updatable pool of structures to create descendant structures. The "best offspring" selection method is to displace the pool structures by optimized structures with higher fitness (lower total energy).

There are 10 types of generators implemented in the program:

RAND	Random generation of atoms in a region of a size set by the <i>Sphere</i> command
RND1	Generation of atoms by condensation: a new atom, randomly generated in a sphere
	of radius Sphere, moves to the center of the structure until the distance to the
	nearest atom is within the limits specified in the <i>BondLimits</i> command
PCCR	Plane-Cut Crossover: Two randomly selected structures from the pool are cut at a
	random plane so that the ratio of the number of atoms in both parts of the two
	structures is equal, and the first parts of the two structures are swapped.
TPCR	Two-Point Crossover: The Cartesian coordinates of the two structures are arranged
	in one-dimensional vectors and divided into two parts at a random position, which
	is the same for both structures. Then the first parts of the two vectors are swapped.
IMOV	Internal Move: A randomly selected atom of a randomly selected pool structure
	moves a random distance towards the center of mass of the structure.
AMOV	Angular Move: A randomly selected atom of a randomly selected pool structure
	rotates by a random angle around the center of mass of the structure.
TWST	Twist operator: A structure randomly selected from the pool is cut by a random
	plane into two parts and both parts are rotated relative to each other by a random
	angle around an axis passing through the center of mass and perpendicular to the
	plane of the cut.
HINT	Hint operator: the new structure is selected based on the "hint", i.e. structure
	provided by the user. In the case of RUNNER, a file with hint structures is placed in
	the ./add directory at any time the program is running. After generating the input
	files, the hint files are moved to the ./usd directory. The file formats are (1)
	geometry optimized server program output files; (2) text files in which hint
	structures are in NXYZ format with @geo origin marks, the end of coordinates is an
	empty line.

Global optimization begins with the formation phase of an initial set of structures ("initial population"). This phase is called the "Initial Phase". In the course of it, a set of structures is accumulated, on the basis of which "child structures" will be generated further. The initial population is formed, as a rule, by two operators RAND and RND1 ("random generation" and "random condensation"). The preference for one of the two operators can be changed using the Init command, specifying the operators to use and the likelihood of their use. In addition, hint

structures (HINT operator) can be used for the initial phase. The initial phase ends when the minimum population ("pool") of structures capable of forming "descendants" is formed. The number of such structures is determined by the *MinPool* command (10 by default). The structures included in the pool are sorted according to their absolute energy increase and are stored in the *.pool_ file.

At the end of the initial phase, the Main Production Phase begins. In its course, at each working cycle of the algorithm, new structures - "descendants" are formed by applying a given set of generators to one or more randomly selected structures from the "active pool". An active pool is a part of a common pool that unites the most beneficial structures (structures with lower energies). The share of the most profitable structures from the total pool size is set by the *ActivePool* keyword (by default 0.7, i.e. 70% of all pool structures). A decrease in this number can lead to an acceleration of optimization if the GM is close in structure to the structures already found. At the same time, this can lead to a loss of "biodiversity" and failure in the search for a GM of unusual structure. Excessive growth of *ActivePool* can result in slow optimization speed.

In the case when, in the course of optimization, structures appear that are similar in structure and energy to the structures included in the pool, only the one with a lower energy remains in the pool. The similarity of structures is determined by an algorithm based on the analysis of interatomic distances sorted in increasing order in both structures. Structures are considered similar if all such distances in the two structures differ by less than 5%. If, during the optimization process, the number of different optimized structures exceeds the maximum size of the pool (it is set by the *MaxPool* keyword), the new structure takes its place in the order of sorted energies in ascending order, and structures whose numbers exceed the size of the pool leave it.

Leaving a structure out of the pool does not mean that it is not used for further work. When generating a new structure, the program checks the similarity of the new structure with all structures for which the energy has ever been calculated in the current run. These structures are stored in the *.base_ file. This file contains not only the optimized structures, but also all the structures found in the intermediate local optimization loops executed by the server program. Before starting the energy calculation (local optimization) of a new structure, it is checked for similarity with all structures stored in *.base_ and the calculation starts only if no similar structures are found. Otherwise, the new structure is discarded and regenerated. In addition, the structure is discarded if its bond lengths exceed the limits defined by the BondLimits keyword. The number of generation attempts is determined by the MaxTry keyword (by default 5000). If during the specified number of attempts this generator could not create a new structure, the generation is performed by the RAND statement. In addition to the *.base_ file, RUNNER creates its shortened version, the * .sbas_ file. It contains only the initial (generated) and final (optimized or under-optimized) structures of each calculation; intermediate optimization cycles are omitted. The *.sbas_ file is easier to parse than *.base_. In *.base_ and *.sbas_ files, initial structures are marked with *gen, final (in this file) - with *end, optimized - with *opt. In addition, when calculating the frequencies, the structure is marked with *freq, and if there are no imaginary frequencies, *lm. If the structure has n imaginary frequencies, it is marked with *nneg $= \langle n \rangle$.

The generated structures are written to the input data file of the server program, the shape of which is set by the *FilPat* keyword. The new input data file is placed in the ./inp directory. From this directory, the control script (or the RUNNER program itself in the case of direct call mode) sends the input file to the server program and moves it to the ./out directory. Upon completion of the calculation by the server, the control script places the results file in the same directory. Finding the results file in the ./out directory, RUNNER analyzes the results, extracts the structures of each optimization step and their energies from it, and writes them to the *.base_ and * .sbas_ files. The pool and its *.pool_ file are simultaneously updated, and the input and output files from the ./out directory are moved to the ./arc directory (archive of all performed optimizations). Saving this data in the ./arc directory can be canceled with the NoSaveArchive directive. This saves disk space, but if optimization errors occur, it will be impossible to determine their cause.

The RUNNER program terminates in several cases:

- 1. Local optimization of all structures, the number of which is set by the keyword *Nstruct*, is completed;
- 2. The number of program work cycles specified by the command MaxCyc is exceeded;
- 3. File runner_stop or runner_prog_stop appeared in the program directory.

The control script terminates in several cases:

- 1. The number of the control script work cycles specified within the script body by the parameter *maxcyc* (all characters are lower case!) is exceeded;
- 2. File runner_stop или runner_shell_stop appeared in the program directory.

The presence of *runner_stop* files simultaneously stops the program and the CS, the presence of *runner_prog_stop* or *runner_shell_stop* - only the program or only the CS. The content of all these files is irrelevant, they can be empty. If these files are present in the startup directory, the operation of the program or the operating system is impossible. To get started, these files must be deleted.

If the work of the program or the control system is terminated by mistake before the completion of the global optimization (GO), the restart is carried out by a command similar to the usual start. In this case, RUNNER automatically detects its previous state and continues GO. The CS acts in a similar way. A prerequisite for a successful restart is to keep intact all files in the directories <code>./inp, ./out, ./sub</code> as well as the *.info_ file in the startup directory. It is recommended to save copies of the *.info_, *.out and res.log files before restarting. The *.info_ file stores information about the current state of the optimization being carried out; its manual editing is not recommended.

If it is necessary to continue the completed GO (GO, in which *Nstruct* structures are examined), you must use the *Continue* directive in the *.inp file. In this case, the current value of *Nstruct* determines the additional number of structures to be examined.

Author, licenses and reviews

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The program is experimental, distributed as is, under the BSD2 license.

Questions about working with the program, messages about possible inaccuracies and errors, feedback and wishes should be sent to skignatov@gmail.com