



COMPUTATIONAL MATERIALS DISCOVERY LABORATORY

#### **Tutorial 8: Introduction to USPEX**

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#### **Useful links:**

 References, where the method was exhaustively described: <a href="http://uspex.stonybrook.edu/uspex.html">http://uspex.stonybrook.edu/uspex.html</a>

#### USPEX manual:

HTML: <a href="http://han.ess.sunysb.edu/uspex\_manual/">http://han.ess.sunysb.edu/uspex\_manual/</a>

PDF: <a href="http://han.ess.sunysb.edu/uspex manual/uspex manual.pdf">http://han.ess.sunysb.edu/uspex manual/uspex manual.pdf</a>

USPEX utilities: <a href="http://han.ess.sunysb.edu/">http://han.ess.sunysb.edu/</a>



#### How to start your calculations

- 1. Prepare INPUT.txt file.
- 2. Prepare files in Specific/ directory (for VASP: INCAR 1,...,INCAR N, POTCAR A, POTCAR B, etc.)
- 3. If necessary, prepare additional files, which are described in the manual.
- 4. Start calculation: USPEX -r

#### Connection information:

- ssh USPEXBNL@qsh.ess.sunysb.edu Password: \*\*\*\*\*\* (please enter the password correctly, otherwise we will be blocked)
- cd ~/workshop
- mkdir <your userid on nano.bnl.gov>
- cd <your userid on nano.bnl.gov>
- USPEX -h



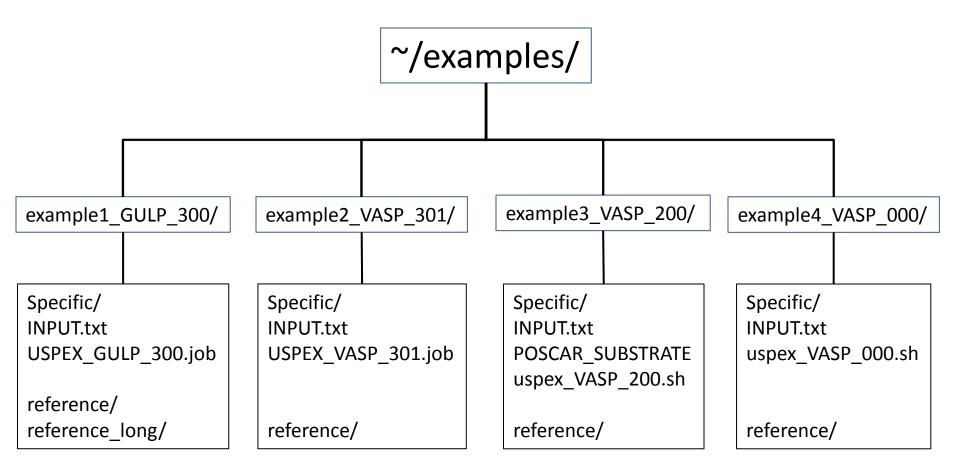


#### Examples for today:

- Bulk Mg<sub>4</sub>Al<sub>8</sub>O<sub>16</sub> (fixed composition, calculationType=300, GULP)
- Bulk Si-C (variable composition, calculationType=301, VASP)
- O<sub>2</sub> @ Si (100) (fixed composition, calculationType=200, VASP)
- C<sub>13</sub> nanoparticles (fixed composition, calculationType=000, VASP)



#### Examples file system





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MgAl<sub>2</sub>O<sub>4</sub> (28 atoms/cell) at 100 GPa pressure. Variable-cell calculation using Buckingham potentials, GULP code. This test has direct bearing on the physics of the Earth's interior!



```
TYPE OF RUN AND SYSTEM
           : calculationMethod (USPEX, VCNEB, META)
            : calculationType (dimension: 0-3; molecule: 0/1;
            : optType_(1=enthalpy, 2=volume, 3=hardness, 4=str
    % symmetries
    2-230
    % endSymmetries
11
12
    % atomType
    Mg Al O
14
    % EndAtomType
15
    % numSpecies
17
    4 8 16
18
    % EndNumSpecies
19
20
    % valences
    2 3 2
    % endValences
```

```
▷ variable optType
```

Meaning: This variable allows you to specify the qua Possible values (characters):

- enthalpy (or "1") to find the stable phases
- volume (or "2") volume minimization (to fir
- hardness (or "3") hardness maximization (to
- struc\_order (or "4") maximization of the degr structure)
- aver\_dist (or "5") maximization of average s ation

> variable calculationType

Meaning: Specifies type of calculation, i.e., whether the structure of a bulk crystal, nanoparticle, or surface is to be predicted. This variable consists of three indices: dimensionality, molecularity, and compositional variability:

dimensionality:

```
"3" — bulk crystals
"2" — surfaces, "-2"

    2D-crystals

"1" — polymers
"0" — nanoparticles
```

· molecularity:

```
"0" — non-molecular
      molecular calculations
```

variability of chemical composition in the calculation:

```
"0" — fixed composition

    variable composition
```

Default: 300 Format:

301 : calculationType

Note: If calculationType=310, i.e., a prediction for a molecular crystal is to be performed, then USPEX expects you to provide files MOL\_1, MOL\_2, ... with molecular geometries for all types of molecules, and these molecules will be placed in the newly generated structures as whole objects. Available options: 300, 301, 310, 000, 200, 201, -200 (and not yet released: 110,

- diel\_sus (or "6") maximization of the static dielectric susceptibility (only for VASP and GULP)
- gap (or "7") maximization of the band gap (only for VASP)
- diel\_gap (or "8") maximization of electrical energy storage capacity (only for VASP)
- mag\_moment (or "9") maximization of the magnetization (only for VASP)
- quasientropy (or "10") maximization of structural quasientropy





#### **INPUT.txt**:

Possible space groups for crystals

```
10 Appendix: List of 230 space groups
                : calculationMethod (USPEX, VCMEB
                 : calculationType (dimension: 0-3; molecule: 0/1; varcomp: 0/1)
                : optType (1=enthalpy, 2=volume, 3=hardness, 4=struc order, 5=aver dist)
      % symmetries
      2-230
                                                                                            Meaning: Describes the identities of each type of atom.
      % endSymmetries
                                                                                            Default: none, must specify explicitly
11
12
      % atomType
                                                                                            If you prefer to use the atomic numbers from Mendeleev's Periodic Table of the Elements
      Mg Al O
                                                                                                % atomType
      % EndAtomTvpe
                                                                                                12 14 8
                                                                                                % EndAtomType
15
                                                                                            Or, if you prefer to use atomic names, specify
      % numSpecies
                                                                                                % atomTvpe
                                                                                                Mg Si O
      4 8 16
                                                                                               % EndAtomType
                                                                                            You can alternatively specify the full names of the elements, for example:
      % EndNumSpecies
19
                                                                                                Magnesium Silicon Oxygen
20
      % valences
```

Number of atoms of each type

Valences of each type of atom



% endValences

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```
    variable numGenerations

                                                                                      Meaning: Maximum number of generations allowed for the simulation. The simulation
25
                                                                                      can terminate earlier, when the same best structure remains unchanged for stopCrit
                                                                                      generations.

    populationSize (how many individuals per generation)

                                                                                     Default: 100
             : initialPopSize (how many individuals in the first generation)
             : numGenerations (now many generations shall be calculated)
                                                                                  ▷ variable stopCrit
             : stopCrit (max number of generations with the same
                                                                                  Meaning: The simulation is stopped if the best structure did not change for stopCrit
             : reoptOld (should the old structures be reoptimized
                                                                                  generations, or when numGenerations have expired — whichever happens first.
             : bestFrac 4What fraction of current generation shall
                                                                                  Default: total number of atoms for fixed-composition runs, maximum number of atoms
33
                                                                                  maxAt for variable-composition runs.
35
                   VARIATION OPERATORS
                                                                                  > variable bestFrac
                                                                                  Meaning: Fraction of the current generation that shall be used to produce the next
                                                                                  generation.
     0.50 : fracGene (fraction of generation produced by hered)
                                                                                  Default: 0.7
           : fracRand (fraction of generation produced randomly
           : fracPerm (fraction of the generation produced by permutations)
           : fracAtomsMut (fraction of the generation produced by softmutation)
41
                       CONSTRAINTS
     % IonDistances
                                                               ▷ variable IonDistances
     0.6 0.6 0.6
                                                               Meaning: Sets the minimum inter-atomic distance matrix between different atom types.
     0.0 0.6 0.6
     0.0 0.0 0.6
                                                               Default: half of the covalent radii sum (in Å) for a corresponding atom pair.
     % EndDistances
```



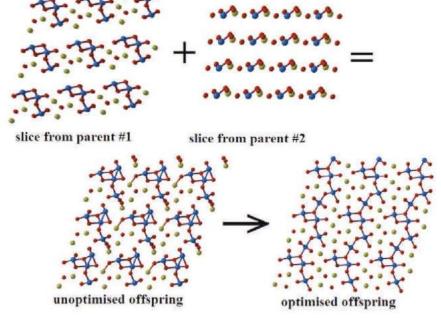


Figure 1. Heredity operator: slices of two parent structures, and the offspring structure before and after local optimization.

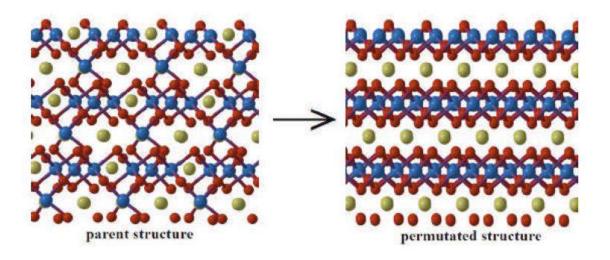
Picture from "Oganov A.R., Ma Y., Lyakhov A.O., Valle M., Gatti C. (2010). Evolutionary crystal structure prediction as a method for the discovery of minerals and materials. *Rev. Mineral. Geochem.* **71**, 271-298"



#### ▷ variable fracPerm

Meaning: Percentage of structures obtained by permutation; 0.1 means 10%, etc.

Default: 0.1 if there is more than one type of atom/molecule; 0 otherwise.



Pictures from "Oganov A.R., Ma Y., Lyakhov A.O., Valle M., Gatti C. (2010). Evolutionary crystal structure prediction as a method for the discovery of minerals and materials. *Rev. Mineral. Geochem.* **71**, 271-298"



```
: fracGene (fraction of generation produced by heredity)
     : fracRand (fraction of generation produced randomly from space groups specified by user)
0.05 : fracPerm (fraction of the generation produced by permutations)
0.25 : fracAtomsMut (fraction of the generation produced by softmutation)
   > variable fracAtomsMut
   Meaning: Specifies the percentage of structures obtained by softmutation or coormuta-
   tion.
   Default: 0.1
```

Picture from "Lyakhov A.O., Oganov A.R., Stokes H.T., Zhu Q. (2013). New developments in evolutionary structure prediction algorithm USPEX. *Comp. Phys. Comm.* **184**, 1172-1182"



## Bulk Mg<sub>4</sub>Al<sub>8</sub>O<sub>16</sub> (example1\_GULP\_300) In results1/OUTPUT.txt you will notice:

> variable fracLatMut

Meaning: Percentage of structures obtained from lattice mutations; 0.1 means 10%, etc. Default: 0 for fixed cell prediction: 0.1 otherwise. ---- VARIATION OPERATORS -The fittest 60 percent of the population used to produce next generation fraction of generation produced by heredity 0.50 fraction of generation produced by random 0.20 fraction of generation produced by softmutation 0.25 fraction of generation produced by permutation 0.05 0.10 fraction of generation produced by latmutation fraction of generation produced by rotmutation 0.00 fraction of generation produced by transmutation 0.00 ----- Variation operators applied ------3 structures produced by heredity 1 structures produced by random optimised 1 structures produced by softmutation parent structure 0 structures produced by permutation 0 structures produced by latmutation 1 structures kept as best from the previous generation O structures imported from the other USPEX Calculations 0 Seeds structures are added from Seeds/POSCARS



```
variable abinitioCode
                                                                                     Meaning: Defines the code used for every optimization step.
          DETAILS OF AB INITIO CALCULATIONS
                                                                                     Default: 1 for every optimization step (VASP)
                                                                                     Format:
     % abinitioCode
                                                                                         % abinitioCode
                                                                                         3 2 2 1 1
55
     3 3 3
                                                                                         % ENDabinit
56
     % ENDabinit
                                                                                     Note: Numbers indicate the code used at each step of structure relaxation: 1 — VASP, 2
57
                                                                                     SIESTA, 3 — GULP, 4 — LAMMPS, 5 — Neural Networks code, 6 — DMACRYS, 7 — CP2K.
                                                                                        Quantum Espresso, 9 — ASE, 10 — ATK, 11 — CASTEP.
58
     % KresolStart
59
     0.16 0.14 0.12
     % Kresolend
60
                                                                                     ▷ variable KresolStart
61
                                                                                     Meaning: Specifies the reciprocal-space resolution for k-points generation (units: 2\pi \mathring{A}^{-1}).
62
     % commandExecutable
                                                                                     Default: from 0.2 to 0.08 linearly
63
     gulp < input > output
     % EndExecutable
64
65
               : numParallelCalcs (how many parallel calculations shall be performed)
66
               : whichCluster (0: no-job-script, 1: local submission, 2: remote submission)
               : ExternalPressure (GPa)
68
     100
69
               : pickUpYN (if pickUpYN~=0, then a previous calculation will be picked up)
70
71
               : pickUpGen (at which generation shall the previous calculation be picked up)
               : pickUpFolder (number of the results folder to be used)
```



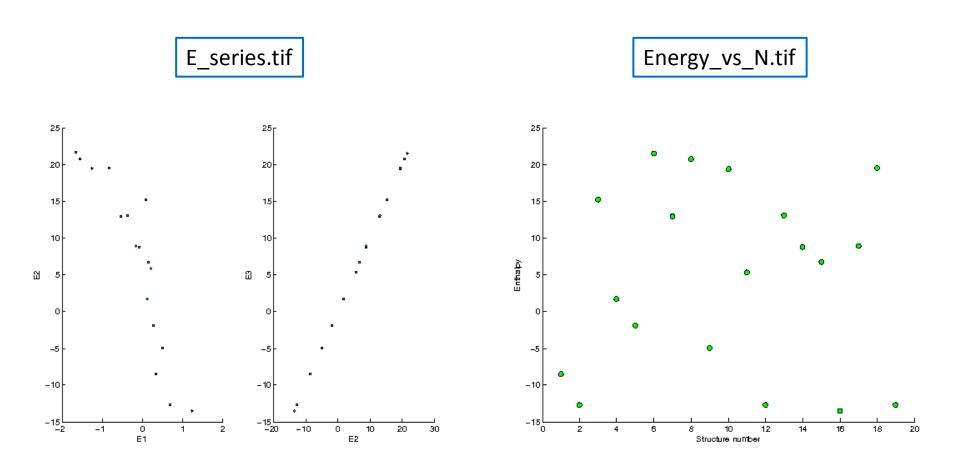
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#### Run the example:

- cp -r ~/examples/example1\_GULP\_300/.
- cd example1\_GULP\_300/
- bsub < USPEX\_GULP\_300.job</li>

Monitor **log** file, check if **USPEX\_IS\_DONE** file appeared. Then cd to **results1/** to view results.

## Bulk $Mg_4Al_8O_{16}$ (example1\_GULP\_300) Output files:

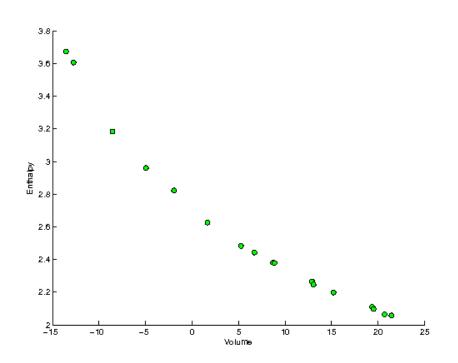


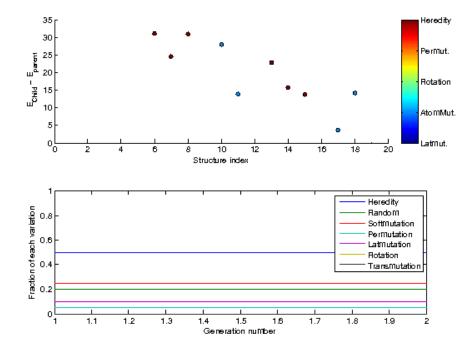




Energy\_vs\_Volume.tif

Variation-Operators.tif

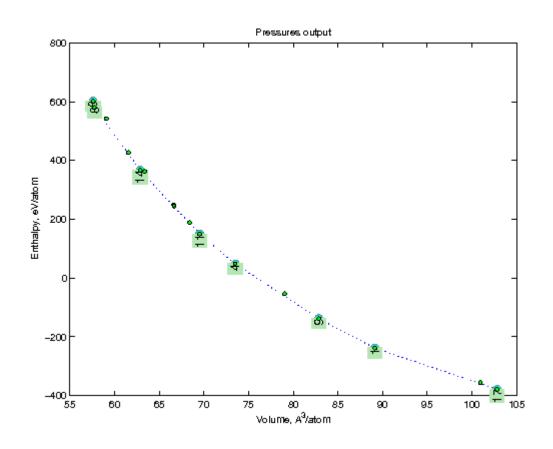






## Bulk $Mg_4Al_8O_{16}$ (example1\_GULP\_300) Output files:

Pressures\_output.png





## Bulk $Mg_4Al_8O_{16}$ (example1\_GULP\_300) Output files:

log - contains console output of USPEX
results1/generation\*/ folders - contain results in \*.mat files for the generations
results1/OUTPUT.txt - contains detailed information on each generation
results1/Parameters.txt = INPUT.txt
Individuals - summary file where you can find main results:

Gen								Enthalpy (eV)		_	Fitness	I	KPO	INT	S SYM	M	Q_entr	A_orde	r S_order
1	1	Random	[	4	8	16	]	-238.872	89.138	10.601	-238.872	[	1	1	1]	1	0.207	0.887	1.782
1	2	Random	[	4	8	16	]	-356.267	100.939	9.362	-356.267	[	1	1	1]	1	0.244	0.847	1.547
1	3	Random	[	4	8	16	1	427.272	61.546	15.354	427.272	[	1	1	1]	8	0.227	0.868	1.645
1	4	Random	[	4	8	16	]	47.898	73.509	12.855	47.898	[	1	1	1]	1	0.228	0.831	1.590
1	5	Random	[	4	8	16	]	-53.740	79.057	11.953	-53.740	[	1	1	1]	1	0.221	0.807	1.577
2	6 I	Heredity	[	4	8	16	]	601.445	57.609	16.403	601.445	[	1	1	1]	1	0.226	0.774	1.504
2	7 1	Heredity	[	4	8	16	]	362.394	63.374	14.911	362.394	[	1	1	1]	1	0.226	0.801	1.541
2	8 I	Heredity	[	4	8	16	]	580.464	57.730	16.368	580.464	[	1	1	1]	1	0.207	0.812	1.642

**BESTIndividuals** – gives this information for the best structures from each generation gatheredPOSCARS – concatenated POSCAR files gatheredPOSCARS\_order – same as gatheredPOSCARS, but with order parameter BESTgatheredPOSCARS – the same data for the best structure in each generation BESTgatheredPOSCARS\_order – same as BESTgatheredPOSCARS, but with order parameter

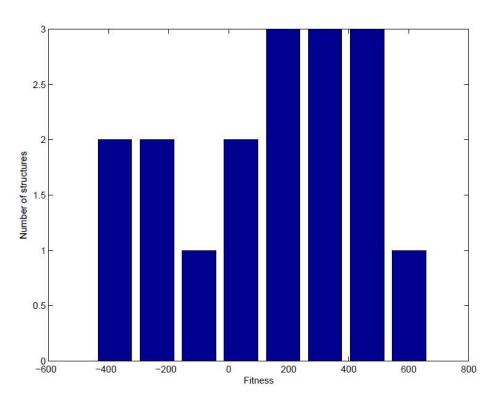




#### Output files:

goodStructures - like Individuals, but ordered by descending of stability
goodStructures\_POSCARS - concatenated POSCAR files ordered by descending of stability
enthalpies\_complete.dat - gives the enthalpies for all structures in each stage of relaxation
non\_optimized\_structures - gives all structures produced before relaxation
origin - shows which structures originated from which parents and through which
variation operators

fitnessStatistics.pdf:





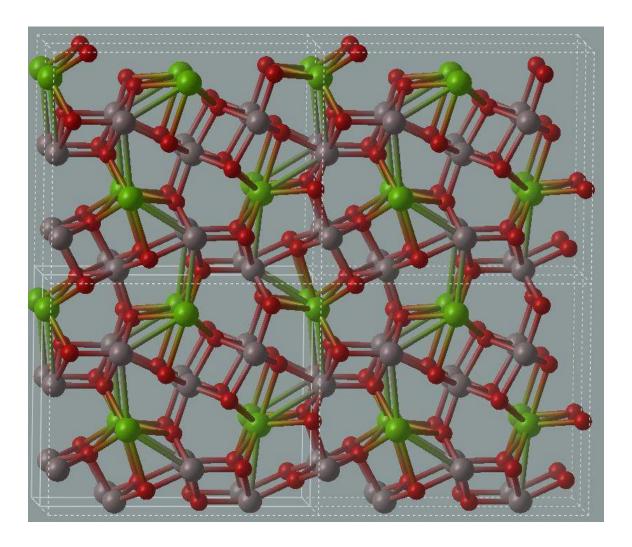
#### View gatheredPOSCARS:

- cd results1/
- python ~/examples/split\_POSCARS.py

This will produce **POSCAR\_files/** folder with splitted POSCAR files, which you can download on your computer and visualize using VESTA, VMD, etc.



### View gatheredPOSCARS (from reference\_long/):





#### Examples for today:

- Bulk Mg<sub>4</sub>Al<sub>8</sub>O<sub>16</sub> (fixed composition, calculationType=300, GULP)
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- O<sub>2</sub> @ Si (100) (fixed composition, calculationType=200, VASP)
- C<sub>13</sub> nanoparticles (fixed composition, calculationType=000, VASP)



## Bulk Si-C (example2\_VASP\_301) INPUT.txt:

```
> variable calculationType
       PARAMETERS EVOLUTIONARY ALGORITHM
                                                                                                    Meaning: Specifies type of calculation, i.e., whether the structure of a bulk crystal,
                                                                                                    nanoparticle, or surface is to be predicted. This variable consists of three indices: dimen-
 2
                                                                                                    sionality, molecularity, and compositional variability:
       USPEX : calculationMethod (USPEX WCNEB, META)
                : calculationType 4dimension: 0-3; molecule: 0/1; va

    dimensionality:

                : optType (1=enthalpy, 2=volume, 3=hardness, 4=stru
                                                                                                                   — surfaces. "-2"

    2D-crystals

       % symmetries
                                                                                                              "1" — polymers
 8
       2-230
                                                                                                              "0" — nanoparticles
       % endSymmetries
                                                                                                       · molecularity:
10
                                                                                                              "0" — non-molecular
       % atomType
                                                                                                              "1" — molecular calculations
12
       Si C

    variability of chemical composition in the calculation:

13
       % EndAtomType
                                                                                                              "0" — fixed composition
14

    variable composition

15
       % numSpecies
16
      1 0
                                                                                                    Default: 300
                                                                                                    Format:
17
       0 1
                                                                     Notes: For variable-composition calculations, you have to specify the compositional building
       % EndNumSpecies
                                                                     blocks as follows:
                                                                                                                                                      molecular crystal is to be performed,
19
                                                                          % numSpecies
                                                                                                                                                       .. with molecular geometries for all
       % valences
                                                                          203
                                                                                                                                                      n the newly generated structures as
                                                                          0 1 1
                                                                                                                                                      201, -200 (and not yet released: 110,
                                                                          % EndNumSpecies
       % endValences
                                                                     This means that the first building block has formula A<sub>2</sub>C<sub>3</sub> and the second building block has
                                                                     formula BC, where A, B and C are described in the block atomType. All structures will then
                                                                     have the formula xA_2C_3 + yBC with x, y = (0,1,2,...) — or A_{2x}B_yC_{3x+y}. If you want to do
                                                                     prediction of all possible compositions in the A-B-C system, you should specify:
```



% numSpecies 1 0 0 0 1 0 0 0 1

% EndNumSpecies

### Bulk Si-C (example2\_VASP\_301)

**INPUT.txt**:

```
> variable minAt
29
30
                                                                                             Meaning: Minimum number of atoms in the unit cell for the first generation.
                                                                                            Default: No default
32
          : bestFrac
    0.6
33
                                                                                            > variable maxAt
34
    0.40 : fracGene (fraction of generation produced by heredity)
                                                                                            Meaning: Maximum number of atoms in the unit cell for the first generation.
    0.20 : fracRand (fraction of generation produced randomly from space groups)
                                                                                            Default: No default
36
    0.20 : fracAtomsMut (fraction of the generation produced by softmutation)
37
    0.20
         : fracTrans
                                                                        Only for variable-composition calculations!
39
          : minVectorLength ( minimal length of any lattice vector)
    IonDistances
                                                                 > variable minVectorLength
    0.0 0.8
                                                                  Meaning: Sets the minimum length of a cell parameter of a newly generated structure.
    % EndDistances
                                                                  Default: 1.8 \times \text{covalent diameter of the largest atom.}
    abinitioCode (which code from CommandExecutable shall be used for calculation? |
    1 1
    ENDabinit
49
                                                                  1 for VASP
50
    % KresolStart
    0.15 0.12
    % Kresolend
55
          : numParallelCalcs (how many parallel calculations shall be performed)
          : whichCluster (0: no-job-script, 1: local submission, 2: remote submission)
    0.010 : toleranceFing (tolerance for identical structures)
58
          : pickUpYN (if pickUpYN~=0 , then a previous calculation will be picked
59
           : pickUpGen (at which generation shall the previous calculation be picked
60
          : pickUpFolder (number of the results folder to be used. If = 0 , then
61
62
    % commandExecutable
    vasp > output
    % EndExecutable
```



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### Bulk Si-C (example2\_VASP\_301)

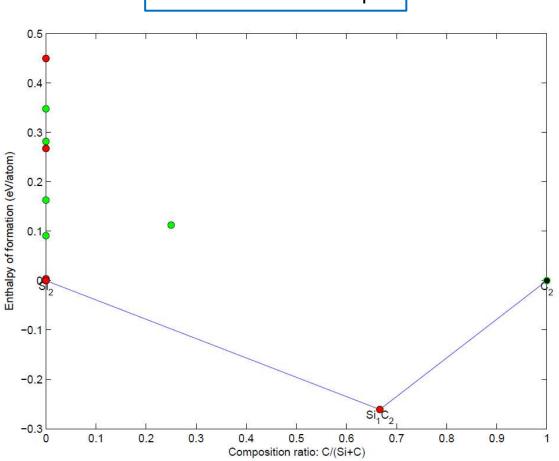
#### Run the example:

- cd ~/workshop/<your directory>
- cp -r ~/examples/example2\_VASP\_301/.
- cd example2\_VASP\_301/
- bsub < USPEX\_VASP\_301.job</li>

Monitor **log** file, check if **USPEX\_IS\_DONE** file appeared. Then cd to **results1/** to view results.

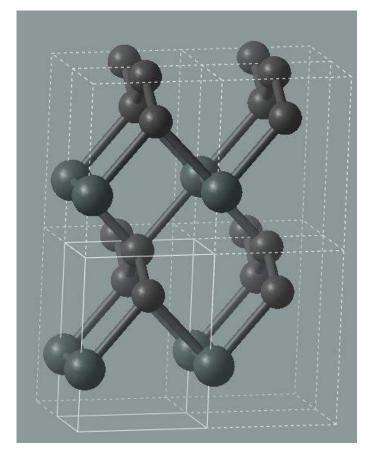
# Bulk Si-C (example2\_VASP\_301) Output files:

extended Convex Hull.pdf



## Bulk Si-C (example2\_VASP\_301) Output files:

convex\_hull – gives all thermodynamically stable compositions, and their enthalpies (per atom)
extended\_convex\_hull, extended\_convex\_hull\_POSCARS – all unique low-energy
compositions and structures



Your structures may be different!

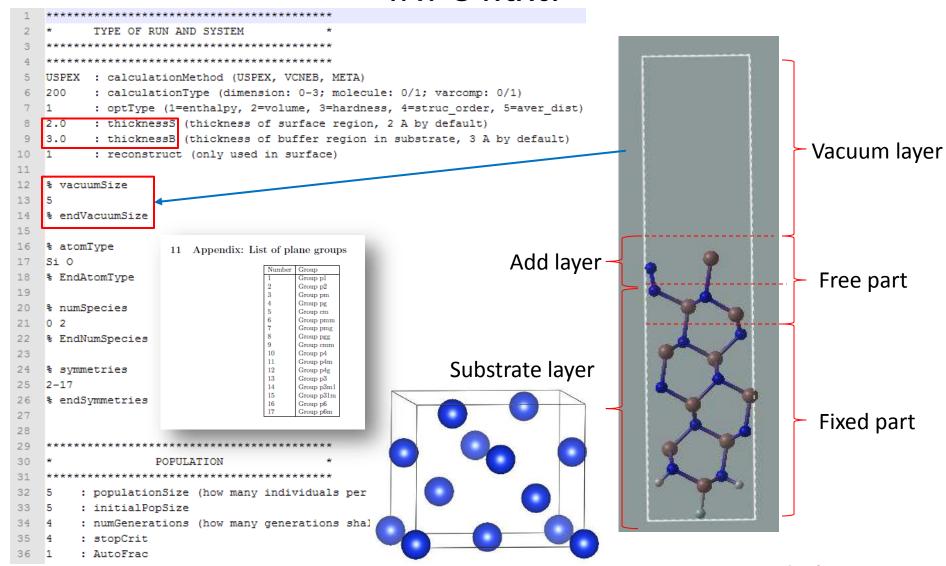




#### Examples for today:

- Bulk Mg<sub>4</sub>Al<sub>8</sub>O<sub>16</sub> (fixed composition, calculationType=300, GULP)
- Bulk Si-C (variable composition, calculationType=301, VASP)
- O<sub>2</sub> @ Si (100) (fixed composition, calculationType=200, VASP)
- C<sub>13</sub> nanoparticles (fixed composition, calculationType=000, VASP)

## O<sub>2</sub> @ Si (100) (example3\_VASP\_200) INPUT.txt:





### O<sub>2</sub> @ Si (100) (example3\_VASP\_200)

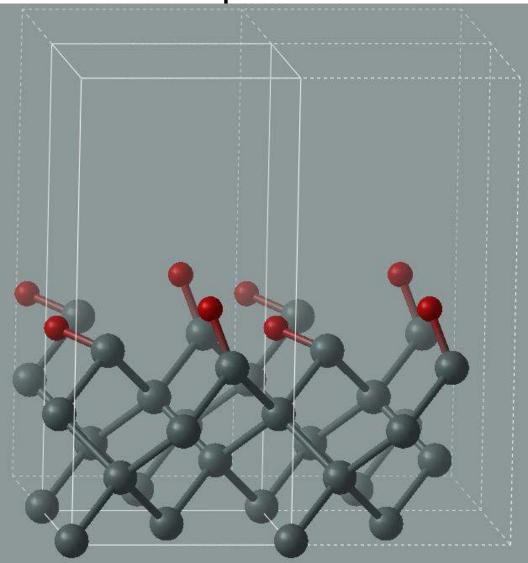
#### Run the example:

- cd ~/workshop/<your directory>
- cp -r ~/examples/example3\_VASP\_200/.
- cd example3\_VASP\_200/
- nohup ./uspex\_VASP\_200.sh > out 2>&1 &

Monitor **log** file, check if **USPEX\_IS\_DONE** file appeared. Then cd to **results1/** to view results.

## O<sub>2</sub> @ Si (100) (example3\_VASP\_200)

Output files:



Your structures may be different!



#### Examples for today:

- Bulk Mg<sub>4</sub>Al<sub>8</sub>O<sub>16</sub> (fixed composition, calculationType=300, GULP)
- Bulk Si-C (variable composition, calculationType=301, VASP)
- O<sub>2</sub> @ Si (100) (fixed composition, calculationType=200, VASP)
- C<sub>13</sub> nanoparticles (fixed composition, calculationType=000, VASP)



## C<sub>13</sub> nanoparticles (example4\_VASP\_000) INPUT.txt:

```
PARAMETERS EVOLUTIONARY ALGORITHM
    USPEX : calculationMethod (USPEX, VCNEB, META)
        : calculationType (dimension: 0-3; molecule: 0/1; varcomp: 0/1)
         : optType (optimise by: 1=enthalpy, 2=volume, 3=hardness, 4=struc order, 5=aver dist, 6=mag moment)
    % vacuumSize
    % endVacuumSize
    % What symmetry(s) have to be satisfied by the randomly created structures
    E C2 D2 C4 C3 C6 T S2 Ch1 Cv2 S4 S6 Ch3 Th Ch2 Dh2 Ch4 D3 Ch6 O D4 Cv3 D6 Td Cv4 Dd3 Cv6 Oh Dd2 Dh3 Dh4 Dh6 Oh C5 S5 S10 Cv5 Ch5 D5 Dd5 Dh5 I Ih
    % endSymmetries
    ************
   % Here come the atomic numbers of the atoms involved
   % atomType
   С
                                                                 % The following is what you know about the lattice. If you know the lattice
   % EndAtomTvpe
                                                                 % vectors, type them in as 3x3 matrix. If not, type the estimated volume.
   % numbers of species (ions/molecules/blocks) of each type
                                                                 % For variable composition - type the estimated atomic volume for each element.
   % numSpecies
                                                                 % Latticevalues (this word MUST stay here, type values below)
28
   13
   % EndNumSpecies
                                                                 % Endvalues (this word MUST stay here)
        : populationSize (how many individuals per generation)
       : numGenerations (how many generations shall be calculated)
       : stopCrit
```



### C<sub>13</sub> nanoparticles (example4\_VASP\_000)

#### Run the example:

- cd ~/workshop/<your directory>
- cp -r ~/examples/example4\_VASP\_000/.
- cd example4\_VASP\_000/
- nohup ./uspex\_VASP\_000.sh > out 2>&1 &

Monitor **log** file, check if **USPEX\_IS\_DONE** file appeared. Then cd to **results1/** to view results.

## C<sub>13</sub> nanoparticles (example4\_VASP\_000) Results:

