

FRC Forcefield Files in MedeA

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1 Introduction

As described in the previous section Forcefields for Materials Simulations, forcefields, or interatomic potentials describe interatomic interactions in classical molecular simulations. The `.frc` format in *MedeA* is used for forcefield parameter files, which contain all the necessary information for using a specific forcefield for running a classical molecular dynamics (MD) or Monte Carlo (MC) simulation. This section briefly describes the contents of *MedeA* `frc` files.

2 Getting Started

Lines starting with an exclamation point `!` are considered comments and no action is taken for such lines. The `.frc` formatted forcefield file consists of several sections, which start with a `#` character. The order with which these sections appear in the `.frc` file is irrelevant. In the beginning of each section there might be some lines starting with a `>` character, which are special comments related to that specific section.

3 Forcefield Type

The first line of an `.frc` formatted file is the identification of the forcefield:

```
!MD forcefield 1
```

Currently, the type must be `1`

4 Versions and References

Forcefield files may be augmented with the addition of new parameters or the refinement of old ones. In order to keep track of the changes and be able to easily and quickly locate them, version numbers and references

are used, placed in the first two columns of a forcefield parameter entry. Version numbers have the general form of a release number, followed by a revision number, e.g. 1.2 refers to version 1 and revision 2. For the same entry, the data of the entry with the highest version number are included and the rest are ignored, while they remain in the file to facilitate keeping track of parameter changes.

For example, an entry for the nonbond parameters might look like this:

```
#nonbond(9-6) pcff+ 200
@type r-eps
@combination sixth-power
!Ver Ref I r eps
!----
1.0 1 c0 3.8540 0.0150
2.0 19 c0 3.7500 0.0070
```

which means that new nonbond parameters have been added in version 2.0 (*ref. 19*) for the atom type “c0” since version 1.0 (*ref. 1*). The new parameters (version 2.0) will be the ones used in the simulation but the old parameters remain in the forcefield file, for tracking purposes. Why these parameters have changed and by whom may be mentioned in the corresponding reference. The references have the sole purpose of providing comments and documentation to the user and are disregarded when the forcefield file is employed by the *MedeA* simulation environment. References have the following form:

```
#reference 30
@Author D. Rigby
@Date 10-Sep-2014
```

Original published compass molecular oxygen parameters re-optimized to fit density **and** DHvap along entire saturation curve.

In order for *MedeA* to use the latest (highest) version for all parameters, in the beginning of the frc file (after the forcefield type definition), the latest (highest) version number needs to be defined:

```
#version pcff+.frc 1.0 12-August-2010
#version pcff+.frc 1.1 7-September-2010
#version pcff+.frc 2.0 15-September-2010
#version pcff+.frc 3.1 26-August-2013
```

The dates mentioned in this section facilitate tracking of changes and are not used elsewhere. Each time a new set of parameters is introduced in the frc file, the user or developer may add this version number in the beginning of the frc file, as shown above. In this example, the 3.1 version is the highest version that will be used in case of multiple entries for the same set of parameters.

5 Description

A short description of the forcefield and some information on its use may be provided in the frc file. Description sections have the following form:

```
#description
Finnis-Sinclair potential for Zr

These potentials were imported from the NIST Interatomic Potentials
Repository Project on 13 June 2015 Please see
http://www.ctcms.nist.gov/potentials/ for more details of the project,
and the web pages below for more information about these particular
potentials. NIST requests that if you find these potentials useful, and
publish results with them, that you please reference

C.A. Becker, et al., "Considerations for choosing and using force fields
and interatomic potentials in materials science and engineering,"
```

Current Opinion **in** Solid State **and** Materials Science, 17, 277–283 (2013). <http://www.ctcms.nist.gov/potentials>

as well **as** the appropriate reference **for** the potential itself:

M.I. Mendeleev **and** G.J. Ackland, "Development of an interatomic potential **for** the simulation of phase transformations

in zirconium," *Phil. Mag. Lett.*, 87, 349–359 (2007). DOI: 10.1080/09500830701191393.

The file at NIST was 'Zr_1.eam.fs'. Information **from the** file's header:

Source: Potential #1 from [M.I. Mendeleev and G.J. Ackland, *Phil. Mag. Letters* 87, 349–359 (2007).]

Contact information: mendeleev@ameslab.gov

Thursday, Nov 29, 2007 The potential was taken **from** v2_3_hcp (**in** C:\SIMULATION.MD\Zr\Results\v2_3)

Please see the following web page **for** more detail:

<http://www.ctcms.nist.gov/potentials/Zr.html>

6 Inclusion of Other FRC File(s)

There may be different “flavors” of a given forcefield, i.e. different versions that have been published over time, representing additions and extensions to a forcefield. In such cases, it is often convenient to include one frc file into another. This allows the use of parts of included forcefield(s) without the need to duplicate the data therein and facilitates handling of the forcefield. To *include* an frc file, a key word, include, preceded by a hash symbol may be employed at the beginning of the frc file, after the versions and description sections:

```
#include pcff.frc
```

7 Definition of Forcefield

In one frc file, there will be one or more defined forcefields. One table listing the interactions used in the forcefield is necessary for each of the forcefields defined.

The section of the definition of each forcefield starts with a *define* header:

```
#define cvff\_nocross\_nomorse
```

followed by a table with the list of all the sections that need to be included in this forcefield, including versions, reference numbers, function, label(s):

```
!Ver Ref Function Label
!-----
2.0 18 atom\_types cvff
1.0 1  equivalence cvff
2.0 18 auto\_equivalence cvff\_auto
1.0 1  hbond\_definition cvff
2.0 18 quadratic\_bond cvff cvff\_auto
2.0 18 quadratic\_angle cvff cvff\_auto
2.0 18 torsion\_1 cvff cvff\_auto
2.0 18 out\_of\_plane cvff cvff\_auto
```

```
1.0 1 nonbond(12-6) cvff
3.0 32 bond\_increments cvff
3.0 31 templates cvff
```

In the above example, the first column provides the version for each interaction type, the second column provides the corresponding reference, the third column provides the name of the function (see the following sections of this document), and the fourth and last column provides the label of the section(s) included. More than one label (each label must correspond to one and only one section in the frc file for a specific function) can be included in the label column. In the example above, there is only one label “cvff” for “atom.types” but there are two “cvff” and “cvff_auto” for “quadratic.angle”. The name that appears in the header of the forcefield definition table is the one that will be used by this forcefield. In the above example, that will be: cvff_nocross_nomorse.

If there are multiple forcefields defined in the frc file, reading the forcefield in, all the forcefields will be read and the default will be the one used, unless otherwise specified, for example:

```
#define clayff-dioctahedral default
```

Forcefield files may define a type for a forcefield:

```
#force_field_type mesoscale
```

The type specification is used by the *MedeA* environment to control what system a forcefield can be applied to.

7.1 Atom Types

The atom types that are utilized in the forcefield, are defined in the atom.types section. For each atom type, the version number, the reference, the type name, the mass and the number of connections are listed, and there is an extra field where a comment may optionally be introduced. This information and is used to assign parameters to the atom types.

```
#atom_types pcff+ 200 !Ver Ref Type Mass Element connection Comment !— — — — —
```

For all-atom forcefields, each atom is assigned an atom type, for which all relevant forcefield parameters need to be assigned.

For united-atom forcefields, a united atom type is assigned to one of the atoms present in the system (e.g. for a CH₃ group, a CH₃ atom type is assigned to the carbon atom of this group) while the hydrogens, that are “grouped” together into the united atom particle, are assigned a “UnitedH” atom type and are disregarded from the simulation.

7.2 Equivalences

Similar atom types may have equivalent behavior, as far as certain interactions are concerned. To use the minimum number of parameters that need to be defined in the parameter file and avoid duplication an equivalence table can be employed in the frc file.

In the equivalences table, there is one column for each different type of interaction (e.g. nonbond, bond, angle, torsion, charges etc) of a given forcefield. For each atom type for which some of the interactions may be equivalenced with those of another atom type, the name of that atom type is defined. That is done for every different interaction term, as shown in the following example:

```
#equivalence pcff+ 200
! Equivalences
! -----
!Ver Ref Type NonB Bond Angle Torsion OOP
!-----
1.0 1 c0 c0 c c c c
```

In this example, for the atom_type “c0” unique nonbond parameters will be used, i.e. the ones introduced specifically for this atom type in the corresponding section of the forcefield file. For the bond, angle, torsion and out-of-plane terms, the parameters that will be used for “c0” are those that are defined for the atom type “c”; therefore there is no need to define any of the interaction parameters for the bond, angle, torsion and out-of-plane terms for “c0”.

7.3 Scaling

Certain forcefields apply scaling factors to interactions between atoms which also interact in some other way. Usually, non-bond interactions between atoms which are bonded directly or to a common third atom are not included in the forcefield energy.

The scaling table permits to explicitly and globally specify the scaling factors for 1,2-, 1,3- and 1,4-interactions for both electrostatic and Van der Waals energies:

#scaling		Martini				Coulomb	VdW
!Ver	Ref	I	J	K	L		
1.0	1	*	*			0.0	0.0
1.0	1	*	*	*		1.0	1.0
1.0	1	*	*	*	*	1.0	1.0

In this example the 1,2-interactions are excluded from the forcefield energy, while the 1,3- and 1,4-interactions are fully included.

7.4 Nonbond (van der Waals) Interactions

Nonbond interactions are describing the interactions between nonbonded atoms (inter- or intra-molecular interactions).

6-9 Lennard-Jones

The “6-9” Lennard-Jones nonbond interaction is:

$$E_{ij} = \varepsilon_{ij} \left[2 \left(\frac{r_{ij,min}}{r_{ij}} \right)^9 - 3 \left(\frac{r_{ij,min}}{r_{ij}} \right)^6 \right] \quad (1)$$

where i and j are the two non-bonded atoms (or united atoms) that are interacting, ε_{ij} is the potential well depth of the interaction and $r_{ij,min}$ is the interatomic distance at which the energy is minimized (i.e. where the interatomic force is zero).

Equation (1) can be equivalently written as:

$$E_{ij} = \frac{A_{ij}}{r_{ij}^9} - \frac{B_{ij}}{r_{ij}^6} \quad (2)$$

where A_{ij} and B_{ij} are the interaction parameters of atoms i and j and r_{ij} is the interatomic distance.

One can easily convert one of the above formulae (1), (2) into the other:

$$\begin{aligned} A_{ij} &= 2\varepsilon_{ij}r_{ij,min}^9 & B_{ij} &= 3\varepsilon_{ij}r_{ij,min}^6 \\ \varepsilon_{ij,min} &= \frac{4B_{ij}^3}{27A_{ij}^2} & r_{ij,min} &= \left(\frac{3A_{ij}}{2B_{ij}} \right)^{1/3} \end{aligned} \quad (3)$$

The *MedeA* environment can read parameters in either format, to facilitate the introduction of these parameters into the frc file in the form in which they have been derived and published by their authors. In this way, conversion errors are avoided and the values present in the frc file can easily be compared to those that are published in the literature.

The interaction parameters are introduced for ij interactions, i.e. interaction of identical atom types. Hetero-atomic interactions are defined thereafter as arithmetic, geometric or sixth-power averages, i.e. using a specific “combining” rule.

6-12 Lennard-Jones

The “6-12” Lennard-Jones nonbond interaction is:

$$E_{ij} = \varepsilon_{ij} \left[\left(\frac{r_{ij,min}}{r_{ij}} \right)^{12} - \left(\frac{r_{ij,min}}{r_{ij}} \right)^6 \right] \quad (4)$$

where i and j are the two non-bonded atoms (or united atoms) that are interacting, ε_{ij} is the potential well depth of the interaction and $r_{ij,min}$ is the interatomic distance at which the energy is minimized (i.e. where the interatomic force is zero).

Equation (4) can be equivalently written as:

$$E_{ij} = 4 \cdot \varepsilon_{ij} \left[\left(\frac{r_{ij,0}}{r_{ij}} \right)^{12} - \left(\frac{r_{ij,0}}{r_{ij}} \right)^6 \right] \quad (5)$$

where i and j are the two non-bonded atoms (or united atoms) that are interacting, ε_{ij} is the potential well depth of the interaction and $r_{ij,0}$ is the interatomic distance at which the energy is zero (often mentioned as σ).

Equation (5) can be equivalently written as:

$$E_{ij} = \frac{A_{ij}}{r_{ij}^{12}} - \frac{B_{ij}}{r_{ij}^6} \quad (6)$$

where A_{ij} and B_{ij} are the interaction parameters of atoms i and j and r_{ij} is the interatomic distance.

One can easily convert one of the above formulas (4), (5), (6) into the other:

$$\begin{aligned} A_{ij} &= \varepsilon_{ij} \cdot r_{ij,min}^{12} & B_{ij} &= 2 \cdot \varepsilon_{ij} \cdot r_{ij,min}^6 \\ A_{ij} &= 4 \cdot \varepsilon_{ij} \cdot r_{ij,0}^{12} & B_{ij} &= 4 \cdot \varepsilon_{ij} \cdot r_{ij,0}^6 \\ \varepsilon_{ij} &= \frac{B_{ij}^2}{4 \cdot A_{ij}} & r_{ij,min} &= \left(\frac{2 \cdot A_{ij}}{B_{ij}} \right)^{1/6} \end{aligned} \quad (7)$$

Again, the *MedeA* environment can employ parameters in either format (A - B or ε - r_{min} or ε - r_0), to facilitate the introduction of these parameters in the frc file, in the same way as they have been introduced and published by the forcefield authors. In this way, conversion errors are avoided and the values present in the frc file can very easily be compared to those that are published in the literature.

The interaction parameters are introduced for ij interactions, i.e. interaction of identical atom types. Hetero-atomic interactions are defined thereafter as arithmetic (8), geometric (9) or sixth-power (10) averages, i.e. using a specific “combining” rule.

The type of the potential (r -eps for use of Eq. (4a) or r_0 -eps for use of Eq. (4b) or A - B for use of Eq. (4c)) and the combining rule are specifically defined in the beginning of the nonbond 6-12 section (*arithmetic*, or *geometric* or *sixth-power*), e.g.:

```
@combination arithmetic
@type r0-eps
```

The format of the 6-12 Lennard-Jones nonbond energy term section is:

```
#nonbond(12-6) AUA
> E = 4.0\*eps(ij) [(r0(ij)\*/r(ij))\*\*12 - (r0(ij)\*/r(ij))\*\*6]
> where r0(ij)\* = 0.5\*((r0(i)\*) + (r0(j)\*))
> and eps(ij) = sqrt(eps(i) \* eps(j))
@combination arithmetic
@type r0-eps
```

```
@units Sigma Ang
@units Epsilon K
!Ver Ref I r0 eps
!-----
1.0 1 CH3-AUA 3.6072 120.15
```

All lines starting with “>” are comments. Any change in the functional form will not be reflected in the calculations in the code.

Tip: If the units are not defined in the beginning of this section, the default units will be used, i.e. kcal mol⁻¹ for ϵ_{ii} and Å for $r_{ii,min}$ and $r_{ii,0}$, kcal mol⁻¹ Å¹² for A and kcal mol⁻¹ Å⁶ for B .

Arithmetic Combining Rule

$$\epsilon_{ij} = \sqrt{\epsilon_{ii} \cdot \epsilon_{jj}} \quad r_{ij,min} = \frac{r_{ii,min} + r_{jj,min}}{2} \quad (8)$$

Geometric Combining Rule

$$\epsilon_{ij} = \sqrt{\epsilon_{ii} \cdot \epsilon_{jj}} \quad r_{ij,min} = \sqrt{r_{ii,min} \cdot r_{jj,min}} \\ A_{ij} = \sqrt{A_{ii} \cdot A_{jj}} \quad B_{ij} = \sqrt{B_{ii} \cdot B_{jj}} \quad (9)$$

Sixth-Power Combining Rule

$$\epsilon = \frac{\sqrt{\epsilon_{ii}\epsilon_{jj}} \cdot 2 \cdot r_{ii,min}^3 \cdot r_{jj,min}^3}{r_{ii,min}^6 + r_{jj,min}^6} \quad (10)$$

The type of the potential (*r-eps* for use of Eq. (1) or *A-B* for use of Eq. (2)) and the combining rule are specifically defined in the beginning of the nonbond 6-9 section (*arithmetic*, or *geometric* or *sixth-power*), e.g.:

```
@type r-eps
@combination sixth-power
```

The format of the 6-9 Lennard-Jones nonbond energy term section is:

```
#nonbond(9-6) pcff+ 200
> E = eps(ij) [2(r(ij)\*/r(ij))\*\*9 - 3(r(ij)\*/r(ij))\*\*6]
> where r(ij) = [(r(i)\*\*6 + r(j)\*\*6))/2]\*\*(1/6)
>
> eps(ij) = 2 sqrt(eps(i) \* eps(j)) \*
> r(i)^3 \* r(j)^3/[r(i)^6 + r(j)^6]
@type r-eps
@combination sixth-power
!Ver Ref I r eps
!-----
1.0 1 ar 3.8800 0.2000
1.0 13 Br 5.4135 0.07993
```

All lines starting with “>” are comments. Any change in the functional form will not be reflected in the calculations in the code.

Tip: If the units are not defined in the beginning of this section, the default units will be used, i.e. kcal mol⁻¹ for ϵ_{ii} and Å for σ_{ii} , kcal mol⁻¹ Å⁹ for A and kcal mol⁻¹ Å⁶ for B .

7.5 Buckingham

The “Buckingham” nonbond interaction is:

$$E_{ij} \cdot e^{-r_{ij}/\rho_{ij}} - \frac{C_{ij}}{r_{ij}^6} \quad (11)$$

where A , ρ and C are constants and r_{ij} is the interatomic distance. Equation (11) can be equivalently written as:

$$E_{ij} \cdot e^{-B_{ij} \cdot r_{ij}} - \frac{C_{ij}}{r_{ij}^6} \quad (12)$$

where A , B and C are constants and r_{ij} is the interatomic distance. One can easily convert one of the above formulas (11) and (12) into the other:

$$B_{ij} = \frac{1}{\rho_{ij}} \quad (13)$$

that has the ability to read parameters in either format (*A-Rho-C* or *A-B-C*), to facilitate the introduction of these parameters in the frc file, in the same way as they have been introduced and published by the forcefield authors. In this way, conversion errors are avoided and the values present in the frc file can very easily be compared to those that are published in the literature.

No combining rule is needed, as the interaction parameters (A , ρ and C) are provided in the frc file for pairs of atoms.

The type of the potential (*A-Rho-C* for use of Eq. (11) or *A-B-C* for use of Eq. (12)) is specifically defined in the beginning of the nonbond Buckingham section, e.g.:

```
@type A-Rho-C
```

The format of the Buckingham nonbond energy term section is:

```
#nonbond(exp-6) inorganic
> E = Aij\*exp(-r/Rhoij) - Cij/r^6
@type A-Rho-C
@units A eV
@units Rho Ang
@units C eV\*Ang^6
!Ver Ref I J A Rho C
!-----
1.0 1 Ag1+ O2- 962.197 0.3000 0.0
```

All lines starting with “>” are comments. Any change in the functional form will not be reflected in the calculations in the code.

Tip: If the units are not defined in the beginning of this section, the default units will be used, i.e. kcal mol⁻¹ for A_{ij} , Å for ρ_{ij} , Å⁻¹ for B_{ij} , and kcal mol⁻¹ Å⁶ for C .

7.6 Shinoda-DeVane-Klein (sdk)

Mesoscale simulations with the SPICA forcefield use different functional forms for Van der Waals interactions depending on the bead types interacting. For pairs with water a 12-4 potential is used:

$$E_{ij} = \frac{3\sqrt{3}}{2} \epsilon_{ij} \left[\left(\frac{\sigma_{ij}}{r_{ij}} \right)^{12} - \left(\frac{\sigma_{ij}}{r_{ij}} \right)^4 \right] \quad (14)$$

while for all other pairs a standard Lennard-Jones potential is applied:

$$E_{ij} = \frac{27}{4} \epsilon_{ij} \left[\left(\frac{\sigma_{ij}}{r_{ij}} \right)^9 - \left(\frac{\sigma_{ij}}{r_{ij}} \right)^6 \right] \quad (15)$$

To ease the use of this forcefield the sdk nonbond energy term section can be used in forcefield files:

```
#nonbond(sdk) SPICA
@type r0-eps
@combination explicit
> E = 27/4*eps*[(sigma/r)^9-(sigma/r)^6]
> E = 3*sqrt(3)/2*eps*[(sigma/r)^12-(sigma/r)^4] (pairs involving W beads)
@units eps kcal/mol
@units sigma Ang
!Ver Ref I J cgtype sigma eps
!---- ---
1.0 1 C2T C2T lj9_6 4.8115 0.4003
1.0 6 CM2 W lj12_4 4.1722 0.2788
```

Combining rules cannot be applied for the Shinoda-DeVane-Klein potential. Therefore, only “explicit” is allowed in the specification and all parameters need to be listed.

All lines starting with > are comments. Any change in the functional form will not be reflected in the calculations in the code. The “cgtype” parameter is directly passed on to *MedeA LAMMPS*. Possible values are: “lj9_6”, “lj12_4” or “lj12_6”.

Tip: If the units are not defined in the beginning of this section, the default units will be used, i.e. kcal mol⁻¹ for ϵ_{ij} and Å for σ_{ij} .

7.7 Electrostatic Interactions

Bond Increments

The bond increments section provides the bond increments that are used from certain forcefields to assign charges to atoms. The charge on an atom is calculated by:

$$q_i = \sum \delta_{ij} \quad (16)$$

where q_i is the charge on atom i and δ_{ij} is the bond increment between an atom and the other atom that it is bonded with. The sum runs over all atoms bonded to atom i .

The format of the bond-increments section is:

```
#bond_increments pcff
!Ver Ref I J DeltaIJ DeltaJI
!---- ---
2.1 11 Ag Ag 0.0000 0.0000
3.0 10 az oah 0.3013 -0.3013
```

where I and J are the atom types of the two atoms that are bonded and δ_{ij} and δ_{ji} are the bond increments that will be added to atoms I and J , respectively.

Tip: The charges are given in units of multiple of electron charge (i.e. 1.0 is the charge on a proton).

Charges

The charges section provides the charges that correspond to specific atom types. This is an alternative way of using bond increments, or can be used in addition to the use of bond-increments. For example, to assign charges, bond increments can be used for covalently bonded species and charges can be used for e.g. ions.

The format of the charges section is:

```
#charge pcff+
! Formal charge on ions
!Ver Ref I q
!-----
3.2 45 Br -1.0
3.2 45 ca+ 2.0001
```

where I is the atom bearing a charge q .

Tip: The charges are given in units of multiple of electron charge (i.e. 1.0 is the charge on a proton).

7.8 Bond Interactions

For bonded atoms a bond interaction term may be used, depending on the description of a forcefield. A bond is representing a chemical bond and may be rigid or flexible.

If a bond is rigid, the length of the bond is constant and equal to a pre-defined value.

If a bond is flexible, the length of the bond is allowed to fluctuate around an equilibrium value. The bond may be physically thought of as a spring that is connecting the two atoms. The length of the spring under no strain is the equilibrium length of the bond. The bond term may be described by different functional forms (depending on the forcefield) and defines how easily the bond length may be augmented or reduced or equivalently, how “stiff” the spring is.



Quadratic Bond

A quadratic bond energy term is:

$$E = k_2 \cdot (l - l_{eq})^2 \quad (17)$$

where l is the bond length, l_{eq} is the equilibrium bond length and k_2 is the force constant.

The format of the quadratic bond energy term section is:

```
#quadratic\_bond oplsa
> E = K2 \* (R - R0)^2
!Ver Ref I J R0 K2
!-----
1.1 6 C CA 1.4900 400.0000
```

The third and fourth columns of the quadratic bond section refer to atom types I and J , which correspond to the bonded atoms. The fifth and sixth columns ($R0$ and $K2$) refer to the equilibrium bond lengths and force constants of Equation (17), respectively.

All lines starting with “>” are comments and are only considered as such by . Any change in the functional form will not be reflected in the calculations in the code.

Tip: If the units are not defined in the beginning of this section, the default units will be used, i.e. kcal mol⁻¹ Å⁻² for $K2$ and Å for the equilibrium bond length.

Quartic Bond

A quartic bond energy term is:

$$E = k_2 \cdot (l - l_{eq})^2 + k_3 \cdot (l - l_{eq})^3 + k_4 \cdot (l - l_{eq})^4 \quad (18)$$

where l is the bond length, l_{eq} is the equilibrium bond length and k_2 , k_3 , k_4 are the force constants for the quadratic, cubic and quartic terms, respectively. The format of the quartic bond energy term section is:

```
#quartic\_bond pcff+ 200
> E = K2 \* (R - R0)^2 + K3 \* (R - R0)^3 + K4 \* (R - R0)^4
!Ver Ref I J R0 K2 K3 K4
!-----
1.0 3 c n\_2 1.4432 319.1593 -586.3243 961.4143
```

The third and fourth columns of the quartic bond section refer to atom types I and J , which correspond to the bonded atoms. The fifth column ($R0$) refers to the equilibrium bond lengths of the bonded atoms. The sixth, seventh and eighth columns ($K2$, $K3$ and $K4$) refer to the force constants of Equation (18), respectively.

Tip: If the units are not defined in the beginning of this section, the default units will be used, i.e. kcal mol⁻¹ Å⁻² for $K2$, kcal mol⁻¹ Å⁻³ for $K3$ and kcal mol⁻¹ Å⁻⁴ for $K4$ and Å for the equilibrium bond length.

Morse Bond

A Morse bond energy term is:

$$E_{ij} = D_b \left[1 - e^{-\alpha(l_{ij} - l_0)} \right]^2 \quad (19)$$

where D_b is the bond dissociation energy, l_{ij} is the instantaneous bond distance, l_0 is the equilibrium bond length and α is the Morse anharmonicity parameter. The format of the Morse bond energy term section is:

```
#morse\_bond cvff
> E = D \* (1 - exp(-ALPHA \* (R - R0)))^2
!Ver Ref I J R0 D ALPHA
!-----
2.3 23 no o- 1.2178 140.2486 2.0000
```

The third and fourth columns of the Morse bond section refer to atom types I and J , which correspond to the bonded atoms. The fifth column ($R0$) refers to the equilibrium bond lengths of the bonded atoms. The sixth and seventh (D and $ALPHA$) refer to the constants of Equation (19).

Tip: If the units are not defined in the beginning of this section, the default units will be used, i.e. kcal mol⁻¹ for D , Å for the equilibrium bond length $R0$ and Å⁻¹ for the Morse anharmonicity parameter $ALPHA$.

Rigid Bond

A rigid bond is a bond for which only the bonded atoms and the equilibrium bond length need to be specified. In the case of atoms bonded with a rigid bond, there is no stretching or shrinking of the bond length. The format of the rigid bond term section is:

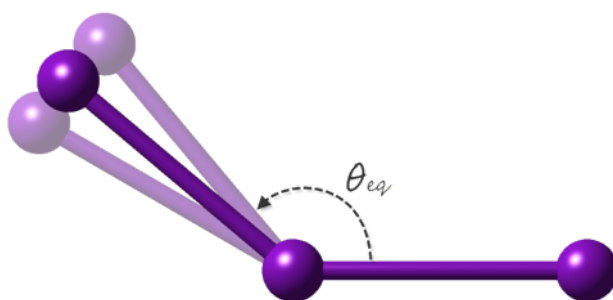
```
#rigid\_bond EH
!Ver Ref I J R0
!-----
1.1 1 C-arom-TraPPE-EH C-arom-TraPPE-EH 1.4
```

The third and fourth columns of the rigid bond section refer to atom types I and J , which correspond to the bonded atoms. The fifth column (RO) refers to the bond lengths of the bonded atoms.

Tip: If the units are not defined in the beginning of this section, the default units will be used, i.e. Å for the equilibrium bond length.

7.9 Angle-Bending Interaction

For bonded atoms an angle-bending interaction term may be used, depending on the description of a force-field. Three atoms, participating in succeeding bonds are interacting through such a term. If an angle is rigid, the angle formed by the three successively bonded atoms is constant and equal to a pre-defined value. If an angle is flexible, the value of the angle is allowed to fluctuate around an equilibrium value. The angle-bending term may be described by different functional forms (depending on the forcefield) and defines how easily the angle may be augmented or reduced.



Quadratic angle-bending potential

A quadratic angle-bending energy term section is:

$$E_{ijk} = k_2(\theta - \theta_0)^2 \quad (20)$$

where θ , θ_0 is the reference bond angle and k_2 is the quadratic force constant.

The format of the quadratic angle-bending term section is:

```
#quadratic\_angle oplsaa
> E = K2 \* (Theta - Theta0)^2
!Ver  Ref I    J    K    Theta0    K2
!-----
1.1  6   CA   C    O    120.4000  80.0000
```

The third, fourth and fifth columns of the quadratic angle-bending section refer to atom types I , J and K , which correspond to the successively bonded atoms. The sixth column ($Theta0$) refers to the equilibrium angle of the bonded atoms and the seventh column ($K2$) refers to the quadratic force constant.

Tip: If the units are not defined in the beginning of this section, the default units will be used, i.e. degrees for the equilibrium angle and kcal mol⁻¹ rad⁻² for the force constant $K2$.

Quadratic cosine angle-bending potential

A quadratic cosine angle-bending energy term section is:

$$E_{ijk} = k_2 \cdot (\cos\theta - \cos\theta_0)^2 \quad (21)$$

where θ is the bond angle, θ_0 is the reference bond angle and k_2 is the quadratic cosine force constant.

The format of the quadratic cosine angle-bending term section is:

```
#quadratic\_cosine\_angle AUA
@type K/2
@units K2 K
@units Theta0 degree
> E = K \* (cosTheta - cosTheta0)^2
!Ver Ref I      J      K      Theta0  K2
!-----
1.0  1  C-arom-AUA  O-ROR-AUA  C-aliph-AUA  112.0  69000.00
```

The third, fourth and fifth columns of the quadratic cosine angle-bending section refer to atom types I , J and K , which correspond to the successively bonded atoms. The sixth column ($Theta0$) refers to the equilibrium angle of the bonded atoms and the seventh column ($K2$) refers to the quadratic cosine force constant.

Tip: If the units are not defined in the beginning of this section, the default units will be used, i.e. degrees for the equilibrium angle and kcal mol⁻¹ for the force constant $K2$.

Quartic angle-bending potential

A quartic angle-bending energy term section is:

$$E_{ijk} = k_2(\theta - \theta_0)^2 + k_3(\theta - \theta_0)^3 + k_4(\theta - \theta_0)^4 \quad (22)$$

where θ is the bond angle, θ_0 is the reference bond angle and k_2 , k_3 and k_4 are the quartic force constants.

The format of the quartic angle-bending term section is:

```
#quartic\_angle pcoeff+ 200
> Delta = Theta - Theta0
> E = K2 \* Delta^2 + K3 \* Delta^3 + K4 \* Delta^4
!Ver Ref I      J      K      Theta0  K2      K3      K4
!-----
1.0  7  c      c      ct      112.7000  58.3500  0.0000  0.0000
```

The third, fourth and fifth columns of the quartic angle-bending section refer to atom types I , J and K , which correspond to the successively bonded atoms. The sixth column ($Theta0$) refers to the equilibrium angle of the bonded atoms and the seventh, eighth and ninth columns ($K2$, $K3$, $K4$) refer to the quartic force constants.

All lines starting with ">" are comments. Any change in the functional form will not be reflected in the calculations in the code.

Tip: If the units are not defined in the beginning of this section, the default units will be used, i.e. degrees for the equilibrium angle and kcal mol⁻¹ rad⁻², kcal mol⁻¹ rad⁻³ and kcal mol⁻¹ rad⁻⁴ for the force constants $K2$, $K3$ and $K4$, respectively.

Rigid angle-bending

A rigid angle is an angle for which only the bonded atoms and the equilibrium angle need to be specified. In the case of a rigid angle, there is no fluctuating of the angle around its equilibrium value.

The format of the rigid angle term section is:

```
# rigid\_angle trappeUA
!Ver Ref I      J      K      Theta
```

```
!-----
1.1 1 CH-arom-TrapPE-UA CH-arom-TrapPE-UA CH-arom-TrapPE-UA 120.00
```

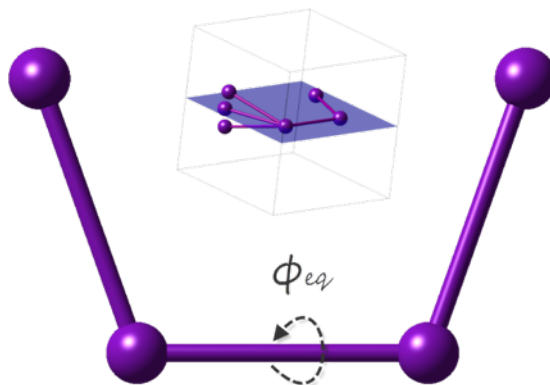
The third and fourth columns of the rigid angle section refer to atom types I and J , which correspond to the bonded atoms. The fifth column (RO) refers to the equilibrium angle of the bonded atoms.

Tip: If the units are not defined in the beginning of this section, the default units will be used i.e. degrees for the equilibrium angle.

Note: Rigid angles can be used in *MedeA-GIBBS* but not in *MedeA-LAMMPS* simulations.

7.10 Torsion Interaction

For bonded atoms a torsion interaction term may be used, depending on the description of a forcefield. Four atoms, participating in successive bonds interact through such a term. If a torsion angle is rigid, the angle formed by the four successively bonded atoms is constant and equal to a pre-defined value. If a torsion angle is flexible, the value of the angle is allowed to fluctuate around an equilibrium value. The torsion angle term may be described by different functional forms (depending on the forcefield). This term specifies how torsion angle variations affect system energies and forces.



One term (cosine) torsion

A (one term) cosine torsion energy term section is:

$$E_{ijkl} = K_{\phi} \cdot [1 + \cos(n \cdot \phi - \phi_0)] \quad (23)$$

where I , J , K and L are the atoms that are bonded to each other (in that succeeding order) forming the torsion angle ϕ , ϕ is the instantaneous torsion angle, ϕ_0 is the equilibrium torsion angle for this quadruplet of atoms, K_{ϕ} is half the energy barrier height and n is the periodicity of the torsion.

The format of the (one term) cosine torsion energy term section is:

```
#torsion\_1 cvff
> E = Kphi \* [ 1 + cos(n\*Phi - Phi0) ]
!Ver Ref I J K L Kphi n Phi0
!-----
2.3 23 \* cp no \* 10.0000 2 180.0000
1.9 17 cp cp c cp 0.6750 4 0.0000
```

The third, fourth, fifth and sixth columns of the (one term) cosine torsion section refer to atom types I , J , K and L , which correspond to the successively bonded atoms. The sixth column (K_{ϕ}) refers to the energy

constant, the seventh column (n) refers to the periodicity of the torsion and the eighth column ($Phi0$) refers to the reference torsion angle.

All lines starting with “>” are comments. Any change in the functional form will not be reflected in the calculations in the code.

Tip: Asterisks are recognized as wildcards, i.e. atoms matching any atom types, as shown in the first line in the example above. Neither or both end atom types (I and L) must be wildcards.

Tip: If the units are not defined in the beginning of this section the default units will be used, i.e. degrees for the equilibrium angle and kcal mol⁻¹ for $Kphi$.

Three term (cosine) torsion

A (three term) cosine torsion energy term section is:

$$\begin{aligned}
 E_{ijkl} = & V_1 \cdot [1 - \cos(\phi - \phi_1)] \\
 & + V_2 \cdot [1 - \cos(2 \cdot \phi - phi_2)] \\
 & + V_3 \cdot [1 - \cos(3 \cdot \phi - phi_3)]
 \end{aligned}
 \tag{24}$$

where I , J , K and L are the atoms that are bonded to each other (in that succeeding order) forming the torsion angle ϕ , ϕ is the instantaneous torsion angle, ϕ_1 , ϕ_2 , and ϕ_3 are the reference torsion angles for this quadruplet of atoms and V_1 , V_2 , V_3 , are the energy barrier heights.

The format of the (three term) cosine torsion energy term section is:

```
#torsion\_3 pcff
> E = SUM(n=1,3) { V(n) \* [ 1 + cos(n\*Phi - Phi0(n)) ] }
!Ver Ref I    J    K    L    V(1)    Phi1(0) V(2)    Phi2(0) V(3)    Phi3(0)
!-----
 3.0 10  oah  az   oah  hoa  0.2821  0.0    -0.0644  0.0    0.0752  0.0
```

The third, fourth, fifth and sixth columns of the (one term) cosine torsion section refer to atom types I , J , K and L , which correspond to the successively bonded atoms. Columns 7, 9 and 11 ($V(1)$, $V(2)$ and $V(3)$) refer to the energy constants, while columns 8, 10 and 12 ($Phi(1)$, $Phi(2)$ and $Phi(3)$) refer to the equilibrium torsion angles. The equilibrium torsion angles are usually 0 or 180 degrees.

All lines starting with “>” are comments. Any change in the functional form will not be reflected in the calculations in the code.

Tip: If the units are not defined in the beginning of this section the default units will be used, i.e. degrees for the equilibrium angle and kcal mol⁻¹ for the energy constants.

OPLS torsion

An OPLS torsion energy term section is:

$$\begin{aligned}
 E_{ijkl} = & \frac{V_1}{2} \cdot [1 + \cos(\phi + \phi_1)] \\
 & + \frac{V_2}{2} \cdot [1 - \cos(2 \cdot \phi + \phi_2)] \\
 & + \frac{V_3}{2} \cdot [1 + \cos(3 \cdot \phi + \phi_3)] \\
 & + \frac{V_4}{2} \cdot [1 + \cos(4 \cdot \phi + \phi_4)]
 \end{aligned}
 \tag{25}$$

where I, J, K and L are the atoms that are bonded to each other (in that succeeding order) forming the torsion angle ϕ , ϕ is the instantaneous torsion angle, ϕ_1, ϕ_2, ϕ_3 and ϕ_4 are the reference torsion angles for this quadruplet of atoms and V_1, V_2, V_3 and V_4 are the energy barrier heights.

The format of the OPLS torsion energy term section is:

```

#torsion\_opls oplsa
> E = SUM(n=1,4) { [V(n)/2] \* [ 1 - ((-1)^n)cos(n\*Phi + Phi0(n)) ] }
> with '1-4' interactions scaled by 0.5
@units V kcal/mol
@units Phi degree
!Ver  Ref I   J   K   L   V1      Phi0  V2      Phi0  V3      Phi0  V4      Phi0
!-----
  1.1  6   O   C   CA  CA  0.0000  0.0    2.1000  0.0    0.0000  0.0    0.0000  0.0
  
```

The third, fourth, fifth and sixth columns of the OPLS torsion section refer to atom types I, J, K and L , which correspond to the successively bonded atoms. Columns 7, 9, 11 and 13 (V_1, V_2, V_3 and V_4) refer to the energy constants, while columns 8, 10, 12 and 14 (Φ_1, Φ_2, Φ_3 and Φ_4) refer to the reference torsion angles.

All lines starting with “>” are comments. Any change in the functional form will not be reflected in the calculations in the code.

Tip: If the units are not defined in the beginning of this section the default units will be used, i.e. degrees for the equilibrium torsion angle and kcal mol⁻¹ for the energy constants.

TraPPE Torsion

A TraPPE torsion energy term section is:

$$\begin{aligned}
 E_{ijkl} = & C_0 + C_1 \cdot [1 + \cos(\phi)] + \\
 & C_2 \cdot [1 - \cos(2 \cdot \phi)] + \\
 & C_3 \cdot [1 + \cos(3 \cdot \phi)]
 \end{aligned}
 \tag{26}$$

where I, J, K and L are the atoms that are bonded to each other (in that succeeding order) forming the torsion angle ϕ , ϕ is the instantaneous torsion angle and C_0, C_1, C_2 and C_3 are the energy barrier heights.

The format of the TraPPE torsion energy term section is:

```

#torsion\_trappe trappeUA-flexible
> E = C0 + C1 \* [1 + cos(phi)] + C2 \* [1 - cos(2\*phi)] + C3 \* [1 + cos(3\*phi)]
> with '1-4' electrostatic interactions scaled by 0.5 and van der Waals ignored
@units C0 K
@units C1 K
@units C2 K
@units C3 K
@units Phi degree
!Ver  Ref I           J           K           L           C0      C1      C2      C3
!-----
  <- C3
  !-----
  <------
  
```



```
1.1 7 CH-TraPPE-UA C-TraPPE-UA CH2-TraPPE-UA CH2-TraPPE-UA 0.00 0.00 0.00
↪461.29
```

The third, fourth, fifth and sixth columns of the TraPPE torsion section refer to atom types *I*, *J*, *K* and *L*, which correspond to the successively bonded atoms. Columns 7, 8, 9 and 10 (*C1*, *C2*, *C3* and *C4*) refer to the energy constants.

All lines starting with > are comments. Any change in the functional form will not be reflected in the calculations in the code.

Tip: If the units are not defined in the beginning of this section the default units will be used, i.e. kcal mol⁻¹ for the energy constants.

AUA torsion

An AUA torsion energy term section is:

$$E_{ijkl} = \sum_{n=0}^8 A_n \cdot (\cos\phi)^n \quad (27)$$

where *I*, *J*, *K* and *L* are the atoms that are bonded to each other (in that succeeding order) forming the torsion angle ϕ , ϕ is the instantaneous torsion angle and *A*₀, *A*₁, *A*₂, *A*₃, *A*₄, *A*₅, *A*₆, *A*₇, and *A*₈ are the energy barrier heights.

The format of the AUA torsion energy term section is:

```
#torsion\_aua AUA
> E = A(n) \* (cos(phi))^n
@lj14 0.0
@el14 0.0
@units A0 K
@units A1 K
@units A2 K
@units A3 K
@units A4 K
@units A5 K
@units A6 K
@units A7 K
@units A8 K
!Ver Ref I J K L A0 A1 A2 A3
↪A4 A5 A6 A7 A8
!-----
↪-----
1.0 1 CH-aliph-AUA CHx-aliph-AUA O-OH-AUA H-OH-AUA 339.41 353.97 58.34 -751.72
↪0.0 0.0 0.0 0.0 0.0
```

The third, fourth, fifth and sixth columns of the AUA torsion section refer to atom types *I*, *J*, *K* and *L*, which correspond to the successively bonded atoms. Columns 7, 8, 9 and 10 (*A0*, *A1*, *A2*, *A3*, *A4*, *A5*, *A6*, *A7* and *A8*) refer to the energy constants.

All lines starting with > are comments. Any change in the functional form will not be reflected in the calculations in the code.

Tip: If the units are not defined in the beginning of this section the default units will be used, i.e. K for the energy constants.

Rigid Torsion

A rigid torsion angle is an angle for which only the bonded atoms and the equilibrium torsion angle need to be specified. In the case of a rigid torsion angle, there is no fluctuating of the torsion angle.

The format of the rigid torsion angle term section is:

```
#rigid\_angle AUA
!Ver Ref I J K Theta0
!-----
1.0 1 CH-arom-AUA CH-arom-AUA CH-arom-AUA 120.00
```

Tip: If the units are not defined in the beginning of this section the default units will be used, i.e. degrees for the equilibrium torsion angle.

7.11 Cross Interactions

Quadratic Bond-bond interaction

A quadratic bond-bond interaction potential, or cross-term, section is:

$$E_{bb'} = k_{bb'} \cdot (r - r_0) \cdot (r' - r'_0) \quad (28)$$

where r and r_0 are the instantaneous bond length and the equilibrium bond length of the first bond respectively, r' and r'_0 are the instantaneous bond length and the equilibrium bond length of the second bond respectively and $k_{bb'}$ is the interaction force constant.

The format of the quadratic bond-bond interaction energy term section is:

```
#bond-bond          pcff+    200
> E = K(b,b') * (R - R0) * (R' - R0')
!Ver Ref    I    J    K    K(b,b')
!-----
1.0    6    o1=   c2=   o1=   275.4350
```

The third, fourth and fifth columns of the quadratic bond-bond interaction energy term section refer to atom types I , J and K , with J being the central atom of the angle formed from the two bonds. Atoms I and J are the atom types of the first bond and atoms J and K are the atoms of the second bond. The equilibrium bond lengths r_0 and r'_0 are taken from the appropriate bond section of the parameter file.

Tip: If the units are not defined in the beginning of this section, the default units will be used, i.e. Å for the bonds and kcal mol⁻¹ Å⁻² for the $k_{bb'}$ constant.

Quadratic bond-angle interaction

A quadratic bond-angle interaction potential, or cross-term, section is:

$$E_{ba} = k_{b\theta} \cdot (r - r_0) \cdot (\theta - \theta_0) \quad (29)$$

where r and r_0 are the instantaneous bond length and equilibrium bond length respectively, θ and θ_0 are the instantaneous bond angle and equilibrium bond angle respectively, and $k_{b\theta}$ is the quadratic force constant.

The format of the quadratic bond-angle interaction energy term section is:

```
#bond-angle          pcff

> E = K * (R - R0) * (Theta - Theta0)

!Ver Ref      I      J      K      K(b,theta)  K(b',theta)
!-----
3.0 10      oah     az     oah      21.5223
3.0 10      oah     az     oas      5.5077      45.3577
```

The third, fourth and fifth columns of the quadratic bond-angle interaction energy term section refer to atom types I , J and K , with J being the central atom of the angle. If the atoms I and K are the same, only one force constant is provided in the sixth column. If the atoms I and K are different, then an additional force constant is provided in the seventh column. The values of the equilibrium bond length r_0 and equilibrium angle θ_0 are taken from the appropriate bond and angle term sections of the file.

Tip: If the units are not defined in the beginning of this section, the default units will be used, i.e. Å for the bonds, degrees for the angles and kcal mol⁻¹ Å⁻¹ rad⁻¹ for the quadratic force constant.

Angle-angle interaction

An angle-angle interaction potential, i.e. the interaction between atoms that form two angles which share a bond, is:

$$E_{\theta\theta'} = V \cdot (\theta - \theta_0) \cdot (\theta' - \theta'_0) \quad (30)$$

where V is the force constant, θ and θ_0 are the instantaneous and equilibrium values for the first bond angle and θ' and θ'_0 are the instantaneous and equilibrium values for the second bond angle.

The format of the angle-angle interaction energy term section is:

```
#angle-angle          pcff

> E = K * (Theta - Theta0) * (Theta' - Theta0')

!Ver Ref      I      J      K      L      K(theta,theta')
!-----
3.0 10      oah     az     oah     oah      11.3873
```

where I , J , K and L are the atom types of the atoms involved in the two angles (θ and θ'). Atoms I , J and K define the first angle (θ) and atoms K , J and L define the second angle (θ'), with the bond between atoms J and K being the common bond of the two angles. The values of the equilibrium angles (θ_0 and θ'_0) are obtained from the appropriate angle sections.

Tip: If the units are not defined in the beginning of this section, the default units will be used, i.e. kcal mol⁻¹ rad² for the quadratic force constant.

Bond-Torsion Interaction

End Bond-Torsion

An interaction potential between a three term (cosine) torsion and the end bonds of the torsion is:

$$E_{ebt} = (r - r_0) [V_1 \cdot \cos\phi + V_2 \cdot \cos(2\phi) + V_3 \cdot \cos(3\phi)] \quad (31)$$

where r is the instantaneous bond length, r_0 is the equilibrium bond length, $V_1/V_2/V_3$ are the force constants and ϕ is the instantaneous torsion angle.

The format of the end bond-torsion interaction energy term is:

```
#end_bond-torsion_3  pcff

> E = (R - R0) * SUM { V(n) * cos[n*phi] }

!                                     LEFT                                     -
↪  RIGHT                                     -
!                                     -----                                     -
↪  -----                                     -
!Ver Ref  I      J      K      L      F (1)      F (2)      F (3)      F (1) -
↪  F (2)      F (3)                                     -
!-----                                     -
↪ -----                                     -
1.0  1  c      c      c      c      -0.0732      0.0000      0.0000
1.0  1  c      c      c      c=     -0.6028      0.0000      0.7675      1.0356 -
↪  0.0000      0.0506
```

where I , J , K and L are the atom types involved in the torsion. V_1 , V_2 and V_3 (noted in the table as LEFT $F(1)$, $F(2)$ and $F(3)$) are the force constants for the interaction of the bond between atoms I and J with the torsion between atoms I , J , K and L , V'_1 , V'_2 and V'_3 (noted in the table as RIGHT $F(1)$, $F(2)$ and $F(3)$) are the force constants for the interaction of the bond between atoms I and J with the torsion between atoms I , J , K and L . If the bonds between atoms I and J and between K and L are identical, i.e. atom types I and L are the same and atom types J and K are also the same, then only one set of force constants should be provided, as shown in the example above. The equilibrium bond lengths are obtained from the appropriate bond section.

Tip: If the units are not defined in the beginning of this section, the default units will be used, i.e. kcal mol⁻¹ Å⁻¹ for the force constants, Å for the bond length and degrees for the torsion angle.

Middle Bond-Torsion

An interaction potential between a three term (cosine) torsion and the end bonds of the torsion is:

$$E_{mbt} = (r - r_0) [V_1 \cdot \cos\phi + V_2 \cdot \cos(2\phi) + V_3 \cdot \cos(3\phi)] \quad (32)$$

where r is the instantaneous bond length, r_0 is the equilibrium bond length, $V_1/V_2/V_3$ are the force constants and ϕ is the instantaneous torsion angle.

The format of the middle bond-torsion interaction energy term is:

```
#middle_bond-torsion_3 pcff

> E = (R - R0) *
>      { F(1) * cos(phi) + F(2) * cos(2 * phi) + F(3) * cos(3 * phi) }

!Ver Ref  I      J      K      L      F (1)      F (2)      F (3)
!-----                                     -
1.0  1  c      c      c      c      -17.7870      -7.1877      0.0000
```

where I , J , K and L are the atom types involved in the torsion, V_1/V_2 and V_3 (noted in the table as $F(1)$, $F(2)$ and $F(3)$) are the force constants for the interaction of the bond between atoms J and K with the torsion between atoms I , J , K and L . The equilibrium bond lengths are obtained from the appropriate bond section.

Tip: If the units are not defined in the beginning of this section, the default units will be used, i.e. kcal mol⁻¹ Å⁻¹ for the force constants, Å for the bond length and degrees for the torsion angle.

Angle-torsion interaction

An interaction potential between an angle and a torsion defined by the same atoms is:

$$E_{at} = (\theta - \theta_0) [V_1 \cdot \cos\phi + V_2 \cdot \cos(2\phi) + V_3 \cdot \cos(3\phi)] \quad (33)$$

where θ is the instantaneous bond angle and θ_0 is the equilibrium angle, $V_1/V_2/V_3$ are the force constants and ϕ is the instantaneous torsion angle.

The format of the angle-torsion interaction energy term is:

```
#angle-torsion_3      pcff

> E = (Theta - Theta0) *
>      { F(1) * cos(phi) + F(2) * cos(2 * phi) + F(3) * cos(3 * phi) }

!                                     LEFT
↪ RIGHT
!
↪-----
!Ver Ref  I    J    K    L    F(1)    F(2)    F(3)    F(1)
↪      F(2)    F(3)
!-----
↪-----
1.0  1  c    c    c    c    0.3886   -0.3139   0.1389
1.0  1  c    c    c    c-   16.6010   0.1267   3.1777   -0.7732
↪      2.4204   -1.5184
```

where I , J , K and L are the atom types involved in the torsion, V_1/V_2 and V_3 (noted in the table as LEFT $F(1)$, $F(2)$ and $F(3)$) are the force constants for the interaction of the bond between atoms I , J and K with the torsion between atoms I , J , K and L , V'_1/V'_2 and V'_3 (noted in the table as RIGHT $F(1)$, $F(2)$ and $F(3)$) are the force constants for the interaction of the bond between atoms J , K and L with the torsion between atoms I , J , K and L . The equilibrium angles are obtained from the appropriate angle sections.

Tip: If the units are not defined in the beginning of this section, the default units will be used, i.e. kcal mol⁻¹ rad⁻¹ for the energy constants and degrees for the angles.

7.12 Out-of-plane, Improper Torsion Interaction

Wilson out-of-plane interaction

This section supplies the parameters used for the potential when the out-of-plane coordinate is defined according to the angle between one bond from the central atom and the plane defined by the other two bonds. An out-of-plane potential is usually applied to planar groups containing an sp² central atom bonded to three other atoms. Examples are amide nitrogens, amide carbons, and the carbon atoms in a benzene ring. The out-of-plane potential acts to keep the central atom in the plane defined by the other three atoms. The functional form is:

$$E = V \cdot \chi^2 \quad (34)$$

where V is the force constant and χ is the instantaneous Wilson out-of-plane angle.

The format of the out-of-plane potential (Wilson definition) section is:

```
#wilson_out_of_plane  pcff

> E = K * (Chi - Chi0)^2

!Ver Ref  I    J    K    L    KChi    Chi0
```

!-----	-----	-----	-----	-----	-----	-----	-----
1.0	1	nr	c+	nr	nr	54.4060	0.0000

where I , J , K and L are the atom types involved in the out-of-plane term, J being the central atom, and $\text{Chi}0$ the instantaneous out-of-plane angle. This term is asymmetric with respect to the outer atoms (I and L) but is made symmetric by summing over the three different out-of-planes defined by a trigonal center.

Tip: If the units are not defined in the beginning of this section, the default units will be used, i.e. kcal mol⁻¹ rad⁻² for the force constant and degrees for the angle.

7.13 Embedded Atom Method

Introduction and functional form

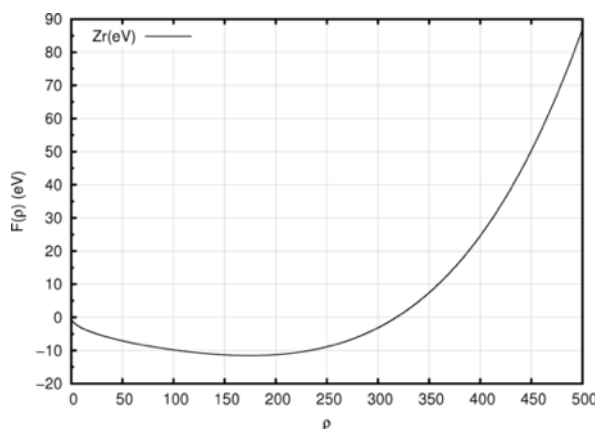
As its name implies, the embedded atom method accounts for the behavior of an atom placed in a defined electron density. The method therefore captures a significant portion of the physical reality of metallic bonding. Related to the effective medium theory of Norskov and Lang [1], the embedded atom method (EAM) was developed by Daw and Baskes [2]. The approach represents the total energy of the system as two additive terms, a pairwise sum of interactions between atoms, and a term representing the electron density of each atomic site, as shown in Equation (35) below.

$$E = \frac{1}{2} \sum_{\substack{i, j = 1 \\ j \neq i}}^N V_{ij}(r_{ij}) + \sum_{i=1}^N F_i(\rho_i) \quad (35)$$

E is the total energy of the system, i and j indicate the unique pairs of atoms within the N atoms of the system, r_{ij} is their interatomic separation, $V_{ij}(r_{ij})$ is a pairwise potential, and $F_i(\rho_i)$ is the embedding function for atom i which depends on the electron density, ρ_i , experienced by that atom:

$$\rho_i = \sum_{\substack{j = 1 \\ j \neq i}}^N \phi_j(r_{ij}) \quad (36)$$

To evaluate a given atom's embedding function, one needs to compute the electron density at the position of atom i . This is obtained by a superposition of "atomic densities", which are described by a density function, $\phi_j(r)$, as shown in Equation (36).



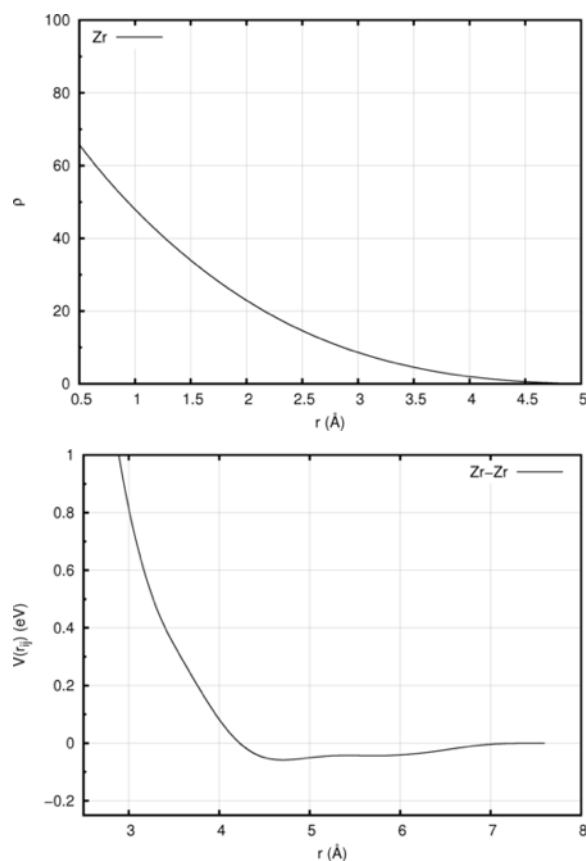
[1] Norskov, J. K., & Lang, N. D. (1980). Effective-medium theory of chemical binding: Application to chemisorption. *Physical Review B*, 21 (6), 2131.

[2] Daw, M. S., & Baskes, M. I. (1984). Embedded-atom method: Derivation and application to impurities, surfaces, and other defects in metals. *Physical Review B*, 29 (12), 6443.

A typical EAM embedding function, illustrated using the Zr EAM forcefield of Mendeleev and Ackland [3].

The embedding function, $F_i(\rho_i)$, provides an essential degree of freedom in the description of metallic bonding. If this term were linear with respect to varying density, the overall energetic description would be equivalent to a standard two body representation. However, the curvature of the embedding term with varying electron density provides an account of the effects of many body interactions. A common form of the embedding function for an EAM forcefield is shown in Figure 4. Here increasing electron density yields progressively more negative embedding energies, until a minimum value is attained beyond which increasing electron density yields less favorable system energies. Figures 5 and 6 provide views of the density function, employed to compute the electron density at a given site (Figure 5), and the interaction function (Figure 6) which is reminiscent of a typical two-body interaction function.

Computing energies and forces based on Equations (35) and (36) can be achieved rapidly as each of the terms are functions of interatomic separation and such separations and their derivatives with respect to atomic coordinates can be rapidly evaluated. In practice, to avoid restricting the form of the functions employed, and to promote calculation efficiency, numerically splined tabulated look-up tables are employed in most EAM calculations for the necessary functions. The resulting forcefield files are therefore large numerical tables. For individual elements three such tables are required, representing the pairwise function, the embedding function, and the density function.



Handling alloy systems requires provision for interaction functions describing the pairwise interaction of each element, in addition to embedding, and density functions. For an n -component alloy there will be $n(n+1)/2$ pairwise interaction functions, n -embedding functions, and, associated density functions. The determination of these functions is challenging, and is complicated by the fact that the creation of a description suitable for a single element provides little information for the behavior of that element in an alloy or compound. Consequently, EAM forcefields are typically developed for specific systems and the description of a given element cannot trivially be combined directly with the description for another element, as assumptions about the two density functions, for example, may not be compatible.

[3] Mendeleev, M. I., & Ackland, G. J. (2007). Development of an interatomic potential for the simulation of phase transformations in zirconium. *Philosophical Magazine Letters*, 87 (5), 349-359.

Despite such specificity, the merit of EAM forcefields is their ability to rapidly and accurately describe the bonding of metallic systems. EAM forcefields allow the simulation of:

- Structures - for example atomic configurations in the vicinity of grain boundaries
- Energies - for example relative polymorph energies and defect energies
- Diffusivity - for example through the use of mean squared displacements of sets of atoms in molecular dynamics trajectories
- Thermal expansivity - for example employing constant pressure simulations as a function of temperature to predict the response of a lattice to a temperature ramp
- Melting of metals and thermodynamic properties of the liquid state

EAM forms included in the *MedeA* Environment

The *MedeA* environment supports standard ‘Finnis-Sinclair’ format EAM forcefield files, with extensions to permit detailed referencing of the source of the particular EAM description and atom type assignment. Such a file contains named sections expressing atom types, any atom equivalences, the standard Finnis-Sinclair format EAM function tables, information for partial charge assignment, and template information to assign forcefield atom types based on rules concerning topology and element type. This overall format is the standard employed by all *MedeA* environment forcefields.

In addition to standard Finnis-Sinclair EAM forcefields, the environment also supports the EAM parameterization described by Zhou and co-workers [4]. Here, mixing rules have been implicitly included in the design of the forcefield, and any combination of the elements: Cu, Ag, Au, Ni, Pd, Pt, Al, Pb, Fe, Mo, Ta, W, Mg, Co, Ti, or Zr may be handled. It is likely that this generality results in diminished accuracy in some circumstances. However, when the effects of alloy formation are of interest, in the creation of layered metallic structures, for example, this description is highly effective.

Finally, the environment also supports an EAM parameterization from Bonny *et al.* [5] where the pair potential is defined in terms of analytical spline functions. In this specific functional form, the density function is defined as the product of a Thomas-Fermi screening function and a cutoff:

$$\phi(r) = S \frac{\exp(-\beta r)}{r} \cdot \frac{x^4}{1+x^4} \quad (37)$$

with:

$$x = \frac{r - r_c}{h} \quad (38)$$

where S , β , r_c , and h are parameters. The format of the density function term section is:

```
#Bonny\_atomic\_density eam
> Phi(r) = S*(exp(-beta*r)/r)*x^4/(1+x^4)
> x(r) = (r-rc)/h
@type S
@units S eV
@type rc
@units rc Ang
!Ver Ref I S beta rc h
!-----
1.0 1 Fe 27.8689586 2.0 4.0 0.25
```

All lines starting with “>” are comments and any change in the functional form represented in the comment will not be reflected in the calculations in the code. The embedding function is parameterized as:

$$F(\rho) = A\sqrt{\rho} + B\rho + C\rho^2 + D\rho^4 \quad (39)$$

[4] Zhou, X. W., Johnson, R. A., & Wadley, H. N. G. (2004). Misfit-energy-increasing dislocations in vapor-deposited CoFe/NiFe multilayers. *Physical Review B*, 69 (14), 144113.

[5] Bonny, G., Terentyev, D., Pasianot, R. C., Poncé, S., & Bakaev, A. (2011). Interatomic potential to study plasticity in stainless steels: the FeNiCr model alloy. *Modelling and simulation in materials science and engineering*, 19 (8), 085008.

lines is a good solution if one needs to edit the frc file.

The parameterization of Bonny *et al.* [6] for Fe-Ni-Cr is readily available in the environment.

7.14 Tabulated Potentials

Tabulated data can be used in lieu of an analytical expression for a potential. Tabulated potentials are useful for situations in which analytical potentials are unknown or are more computationally expensive to evaluate compared to looking up values in a table. One common application of tabulated potentials is for mapping atomistic energies and forces to the interactions between corresponding coarse-grained particles.

When defining a tabulated potential for bonds, angles, torsion, or non-bonded interactions, minimum and maximum values for the distance or angle variable must be specified. Typically, the intermediate values are specified in regular intervals, but non-uniform intervals can also be specified. The tabulated potential can then be interpolated by the compute engine in multiple ways and is also subject to standard considerations such as non-bond cutoff distances for evaluating pairwise interactions. It is possible for compute engines to treat the provided tabulated potential in the following ways:

- A lookup table that finds the nearest table entry to the desired distance or angle.
- Perform linear interpolation between the two nearest table entries for the desired distance or angle.
- Fit cubic splines to the tabulated potential.

Thus for the same tabulated potential data, it is possible for a compute engine to calculate different energies or forces for an interaction based on the style in which the tabulated potential data is interpolated. Accordingly, care must be taken so that a tabulated forcefield is interpolated in the same manner used during parameterization. If it is unclear which method of interpolation is most appropriate for a given forcefield, a sensitivity analysis can be carried out to assess the impact of interpolation choice on the performance of the forcefield.

Tabulated Bond

The format of the tabulated bond energy and force term section is:

```
#tabulated_bond mesohexane
> E = f(r)
!Ver Ref  I      J      Data
!-----
 1.0  1   B1     B2     {
N 5951 R 0.100000 12.000000
1  1.00000e-01  7.1125335e+02  2.8403548e+02
2  1.02000e-01  7.1068528e+02  2.8403542e+02
3  1.04000e-01  7.1011721e+02  2.8403545e+02
...
5951  1.20000e+01  1.2984599e+02  -4.1317500e-01
}
```

All lines starting with “>” are comments and are only considered as such. Any change in the functional form will not be reflected in the calculations in the code.

In the interaction header, the third and fourth columns refer to atom types *I* and *J*, which correspond to the bonded atoms. The fifth column, *Data*, refers to the tabular data that is contained in the braces, {}. The tabular data contains its own header information that denotes the following settings:

- The number of table entries, *N*, is a required parameter and is set to an integer value. This is the only required parameter and all subsequent parameters are optional.
- After specifying the value of *N*, the following parameters can be specified:

[6] Bonny, G., Terentyev, D., Pasianot, R. C., Poncé, S., & Bakaev, A. (2011). Interatomic potential to study plasticity in stainless steels: the FeNiCr model alloy. *Modelling and simulation in materials science and engineering*, 19 (8), 085008.

- *R*: specify two values that correspond to distance bounds between which *N* uniform distance increments exist.
- *RSQ*: specify two values that correspond to squared distance bounds between which *N* uniform squared distance increments exist.
- *FPRIME*: specify two values that correspond to the derivative of the force at the minimum and maximum distance. If *FPRIME* is not specified, these will be estimated by the first two and last two forces listed. These force derivatives are needed for spline construction routines.

After the *Data* header information, *N* lines are written that contain, from right to left, the index value ranging from 1:*N*, the bond distance value, the bond energy value, and the bond force value.

Tabulated Angle

The format of the tabulated angle energy and force term section is:

```
#tabulated_angle mesohexane
> E = f(th)
!Ver Ref      I      J      K      Data
!-----
  1.0  1      B1     B2     B1      {
N 9001 R 0.0 180.0
1  0.00000e+00  0.0000000e+00 -4.1416272e-01
2  2.00000e-02  8.2832544e-03 -4.1416272e-01
3  4.00000e-02  1.6566509e-02 -4.1416272e-01
...
9001  1.80000e+02  2.9081248e+01 -3.0471378e-01
}
```

All lines starting with “>” are comments and are only considered as such. Any change in the functional form will not be reflected in the calculations in the code.

In the interaction header, the third, fourth, and fifth columns refer to atom types *I*, *J*, and *K*, which correspond to the angle atoms. The sixth column, *Data*, refers to the tabular data that is contained in the braces, {}. The tabular data contains its own header information that denotes the following settings:

- The number of table entries, *N*, is a required parameter and is set to an integer value. This is the only required parameter and all subsequent parameters are optional.
- After specifying the value of *N*, the following parameters can be specified:
 - *R*: specify two values that correspond to angle bounds between which *N* uniform angle increments exist.
 - *RSQ*: specify two values that correspond to squared angle bounds between which *N* uniform squared angle increments exist.
 - *FPRIME*: specify two values that correspond to the derivative of the force at the minimum and maximum distance. If *FPRIME* is not specified, these will be estimated by the first two and last two forces listed. These force derivatives are needed for spline construction routines.

After the *Data* header information, *N* lines are written that contain, from right to left, the index value ranging from 1:*N*, the angle value, the angle energy value, and the angle force value.

Tabulated Torsion

The format of the tabulated torsion energy and force term section is:

```
#tabulated_torsion mesohexane
> E = f(th)
!Ver Ref      I      J      K      L      Data
!-----
```

```

1.0 1 B1 B2 B1 B1 {
N 9001 R 0.0 180.0
1 0.00000e+00 0.0000000e+00 -4.1416272e-01
2 2.00000e-02 8.2832544e-03 -4.1416272e-01
3 4.00000e-02 1.6566509e-02 -4.1416272e-01
...
9001 1.80000e+02 2.9081248e+01 -3.0471378e-01
}

```

All lines starting with “>” are comments and are only considered as such. Any change in the functional form will not be reflected in the calculations in the code.

In the interaction header, the third, fourth, fifth, and sixth columns refer to atom types *I*, *J*, *K*, and *L*, which correspond to the torsion atoms. The seventh column, *Data*, refers to the tabular data that is contained in the braces, {}. The tabular data contains its own header information that denotes the following settings:

- The number of table entries, *N*, is a required parameter and is set to an integer value. This is the only required parameter and all subsequent parameters are optional.
- After specifying the value of *N*, the following parameters can be specified:
 - *R*: specify two values that correspond to torsion angle bounds between which *N* uniform torsion angle increments exist.
 - *RSQ*: specify two values that correspond to squared torsion angle bounds between which *N* uniform squared torsion angle increments exist.
 - *FPRIME*: specify two values that correspond to the derivative of the force at the minimum and maximum distance. If *FPRIME* is not specified, these will be estimated by the first two and last two forces listed. These force derivatives are needed for spline construction routines.

After the *Data* header information, *N* lines are written that contain, from right to left, the index value ranging from 1:*N*, the torsion angle value, the torsion angle energy value, and the torsion angle force value.

Non-bonded Interactions

The format of the non-bonded interaction energy and force term section is:

```

#nonbond(table) mesohexane
> E = f(r)
@combination explicit
!Ver Ref I J Data
!---- ---
1.0 1 B1 B1 {
N 746 R 0.100000 15.00000
1 1.0000000000e-01 1.1885847280e+03 1.5814323250e+03
2 1.2000000000e-01 1.1573550360e+03 1.5412787530e+03
3 1.4000000000e-01 1.1269232530e+03 1.5016480930e+03
...
746 1.5000000000e+01 0.0000000000e+00 -4.6973142030e+00
}

```

All lines starting with “>” are comments and are only considered as such. Any change in the functional form will not be reflected in the calculations in the code.

In the interaction header, the third and fourth columns refer to atom types *I* and *J* which correspond to the non-bonded atoms. The fifth column, *Data*, refers to the tabular data that is contained in the braces, {}. The tabular data contains its own header information that denotes the following settings:

- The number of table entries, *N*, is a required parameter and is set to an integer value. This is the only required parameter and all subsequent parameters are optional.
- After specifying the value of *N*, the following parameters can be specified:

- *R*: specify two values that correspond to distance bounds between which *N* uniform distance increments exist.
- *RSQ*: specify two values that correspond to squared distance bounds between which *N* uniform squared distance increments exist.
- *FPRIME*: specify two values that correspond to the derivative of the force at the minimum and maximum distance. If *FPRIME* is not specified, these will be estimated by the first two and last two forces listed. These force derivatives are needed for spline construction routines.

After the *Data* header information, *N* lines are written that contain, from right to left, the index value ranging from 1:*N*, the distance value, the non-bonded energy value, and the non-bonded force value.

8 Forcefields with *MedeA-LAMMPS* and *MedeA-GIBBS*

Interaction		<i>MedeA LAMMPS</i>	<i>MedeA GIBBS</i>
Nonbond	LJ6-9	✓	✓
	LJ6-12	✓	✓
	Buckingham	✓	
	SDK	✓	✓
	Bond-increments	✓	✓
	Charges	✓	✓
	mie	✓	✓
	Tabulated	✓	
Bond	Quadratic	✓	✓
	Quartic	✓	✓
	Morse	✓	
	Rigid		✓
	Tabulated	✓	
Angle	Quadratic	✓	✓
	Quadratic cosine	✓	✓
	Quartic	✓	✓
	Rigid		✓
	Tabulated	✓	
Torsion	One-term cosine	✓	✓
	Three-term cosine	✓	✓
	OPLS	✓	✓
	TraPPE	✓	✓
	AUA		✓
	Rigid		✓
	Tabulated	✓	
Mesoscale	Martini	✓	✓
	SPICA	✓	✓
EAM	Finnis-Sinclair	✓	✓

Continued on next page

Table 1 – continued from previous page

Interaction		<i>MedeA LAMMPS</i>	<i>MedeA GIBBS</i>
	Zhou	✓	✓
	Bonny	✓	✓
	MEAM	✓	
3-body	Tersoff	✓	
	Stillinger-Weber	✓	
Reactive	Streitz-Mintmire	✓	
	ReaxFF	✓	
	COMB3	✓	
Machine Learned	SNAP	✓	
	NNP	✓	