

# Documentation - MD\_Evaluation

---

Program to evaluate MD output files of bulk simulations

Notes:

- usable for LAMMPS simulations in metal units
- required files provided by specific LAMMPS scripts

## 1. Overview

---

Programm evaluates three different simulation types:

- Evaluation of **single run simulations** for transport properties (**single\_run**)
- **Time decomposition method (TDM)** for transport properties (**tdm**)
- Evaluation of **VLE simulations** (two phase simulations) (**vle**)

### Single run simulations

**Required Folder structure:**

- *SIM\_1* [contains all output files of a EMD simulation of one state point]
- ...

### TDM

**Required Folder structure:**

- *STATE\_1* [contains all simulations of one state point]
  - *Sim\_001*
  - *Sim\_002*
  - ...
- ...

### VLE simulations

**Required Folder structure:**

- *FOLDER\_1* [contains all simulations done for one substance]
  - *Sim\_T1* [VLE simulation at temperature T1]
  - *Sim\_T2* [VLE simulation at temperature T2]
  - ...
- ...

## 2. Input File

---

**Name:** 'INPUT.txt' (has to be in same folder as 'Master' file)

**General structure:**

- first line: '#' + 'keyword'
- second line: value assigned to keyword

**Keywords:**

---

Name	Type	Modes	Description
<b>mode</b>	string [ <i>single_run</i> , <i>tdm</i> , <i>vle</i> ]	all	defines the mode of the simulations/evaluation
<b>folder</b>	string	all	path to main folder containing all simulation data (see chapter 1)
<b>ensemble</b>	string [ <i>NVT</i> , <i>NVE</i> , <i>NpT</i> ]	<i>single_run</i> , <i>tdm</i>	ensemble to evaluate
<b>timesteps_EQU</b>	integer ( $\geq 0$ )	all	number of timesteps to ignore at the start of each simulation
<b>DO_single</b>	integer [0,1]	<i>tdm</i>	1 - evaluate single folders, 0 - single folders already evaluated
<b>DO_state</b>	integer [0,1]	<i>tdm</i>	1 - evaluate complete thermodynamic state (main folder)
<b>N_boot</b>	integer ( $\geq 0$ )	<i>tdm</i>	number of bootstrapping repetitions
<b>corr_length</b>	integer ( $\geq 0$ )	<i>single_run</i>	length (timesteps) of correlation function
<b>span_corr_fun</b>	integer ( $\geq 0$ )	<i>single_run</i>	timesteps between single correlation functions

Example *single*:

```
#mode
single_run

#folder
C:/path2simulations/sim_1

#folder
C:/path2simulations/sim_*

#ensemble
NVT

#timesteps_EQU
0

#corr_length
100000

#span_corr_fun
20000
```

Example *TDM*:

```
#folder
C:/path2simulations/state_1

#folder
C:/path2simulations/state_*

#ensemble
NVT

#timesteps_EQU
0

#DO_single
1

#DO_state
```

1

#N\_boot  
100

## Output

---

Units:

- Length: [L] = m
- Mass: [m] = g/mol
- Time: [t] = s
- Temperature: [T] = K
- Pressure: [p] = MPa
- Density: [ $\rho$ ] = g/ml
- Energy: [E] = eV
- Viscosity: [ $\eta$ ] = Pa\*s
- Diffusion Coefficient: [D] = m<sup>2</sup>/s
- Thermal Conductivity: [ $\lambda$ ] = W/(m\*K)