DynamO Workshop Molecular models

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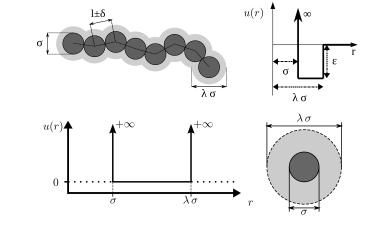
Section Outline

Modeling molecules

Thermodynamic perturbation theory

Building a multi-molecule configuration

Modeling molecules



- Bonds are enforced through an infinitely deep square-well potential.
- ► Currently within DynamO, there is built-in support for linear chains and rings.

Configuration file

dynamod command:

```
dynamod --packing-mode 2 --i1 50
```

configuration file:

```
<Interactions>
<Interaction Type="SquareBond" Diameter="0.90000000000000013"
   Lambda="1.2222222222222223"
   Name="Bonds" Elasticity="1">
        <IDPairRange Type="Chains"
        Start="0" End="15" Interval="16"/>
</Interaction>
...
</Interactions>
...
```

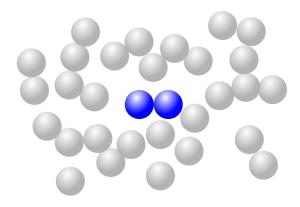
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Thermodynamic perturbation theory



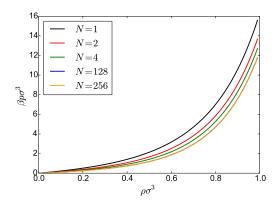
Free energy of a system of molecules:

$$\begin{split} F^{\mathrm{res}}(c_p,\beta,V,N) &= F^{\mathrm{res}}_{\mathrm{ref}}(Nc_p,\beta,V) + F^{\mathrm{bond}} \\ F^{\mathrm{bond}}(c_p,\beta,V,N) &\approx -N_p N_{\mathrm{bonds}} k_B T \ln y_{\mathrm{ref}}(I;Nc_p,\beta) \end{split}$$

where $N_{\rm bond}$ is the number of bonds per molecule.

MNB & LL DynamO Workshop 23/01/2015

Thermodynamic perturbation theory



$$Z = eta p/c_p = N + NZ_{
m ref} - (N-1) \left[1 +
ho rac{\partial \ln y_{
m ref}}{\partial
ho}
ight]$$

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Building multi-molecule configuration file

Create a single chain configuration:

```
dynamod --pack-mode 2 --i1 16 --f1 1 --thermostat 100
```

Arrange the configuration in a low density lattice:

```
dynamod --pack-mode 3 --density 0.001 \
--i1 0 --xcell 3 --ycell 3 --zcell 3 \
--s1 config.out.xml.bz2
```

Fix bonding constraints

Change bonding entry from

to

Eliminate CaptureMap tag

Compress the configuration

Compress the configuration to the desired density

```
dynarun --engine 3 --growth-rate 1 --target-density 0.1 \
  config.out.xml.bz2
```

Zero momentum and rescale velocities

Run simulation

```
dynarun --engine 1 -c 1000000 config.out.xml.bz2
```