

Tracking the Liquid-Vapour interface at the Molecular Scale

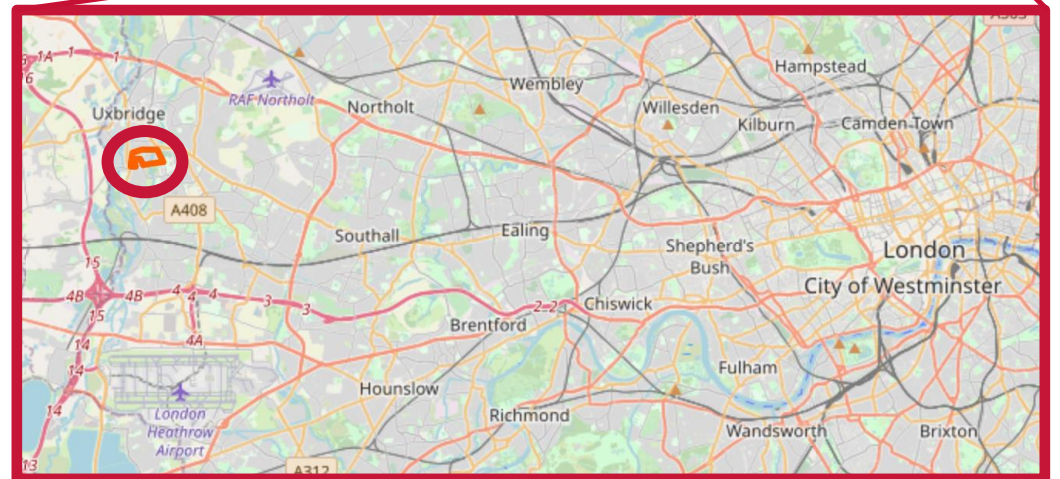
Edward Smith

19th April 2022

Brunel University London

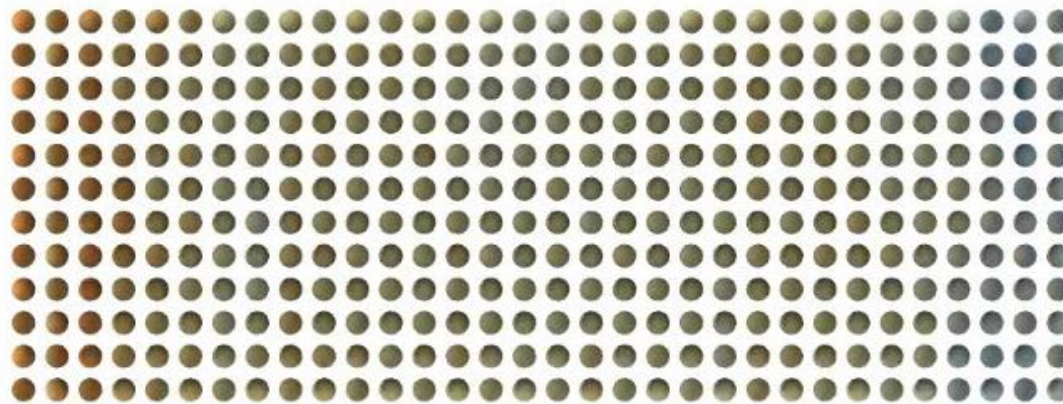


A Research-Focused Engineering University



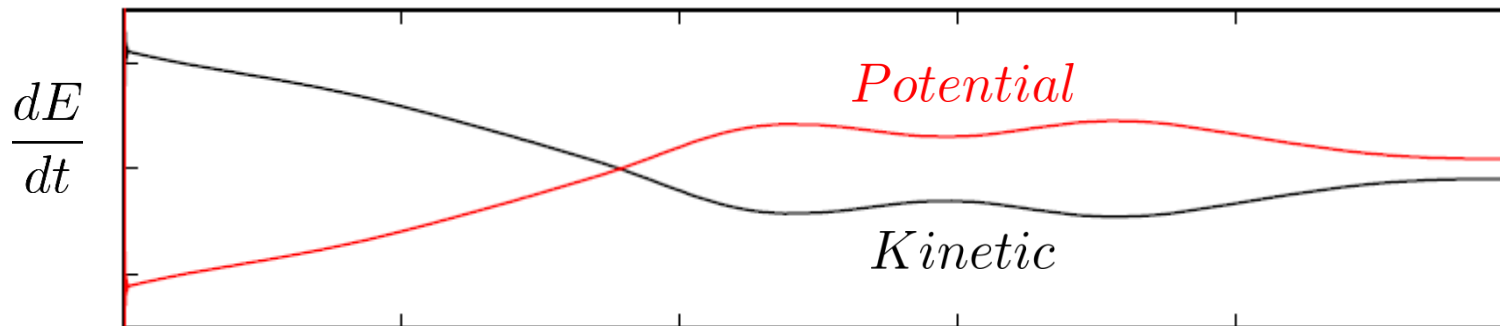
- An introduction to Molecular Dynamics (MD)
- The moving contact line
- The liquid-vapour interface - a Lagrangian reference frame
- Some applications

Molecular Dynamics



Molecular Dynamics

- Solving just Newton's law
 - Energy is automatically conserved \rightarrow total = kinetic + potential



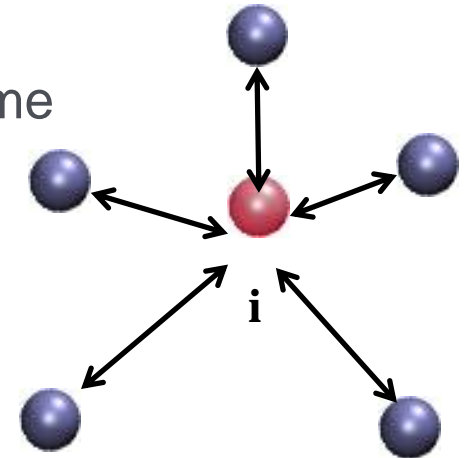
- Pressure, viscosity, heat flux and surface tension do not need to be specified and, are in fact, all outputs of the simulation
- Phase change (evaporation, condensation) occur with no additional models needed
- Solid constructed with molecular roughness
- Can model complex molecules, water, polymers, biomolecules

Molecular Dynamics

Discrete molecules in continuous space

- Molecular position evolves continuously in time
- Acceleration \rightarrow Velocity \rightarrow Position

$$\ddot{\mathbf{r}}_i \rightarrow \dot{\mathbf{r}}_i \rightarrow \mathbf{r}_i(t)$$



Acceleration obtained from forces

- Governed by Newton's law for an N-body system
- Pairwise electrostatics interactions from quantum mechanics

$$m_i \ddot{\mathbf{r}}_i = \mathbf{F}_i = \sum_{i \neq j}^N \mathbf{f}_{ij}$$

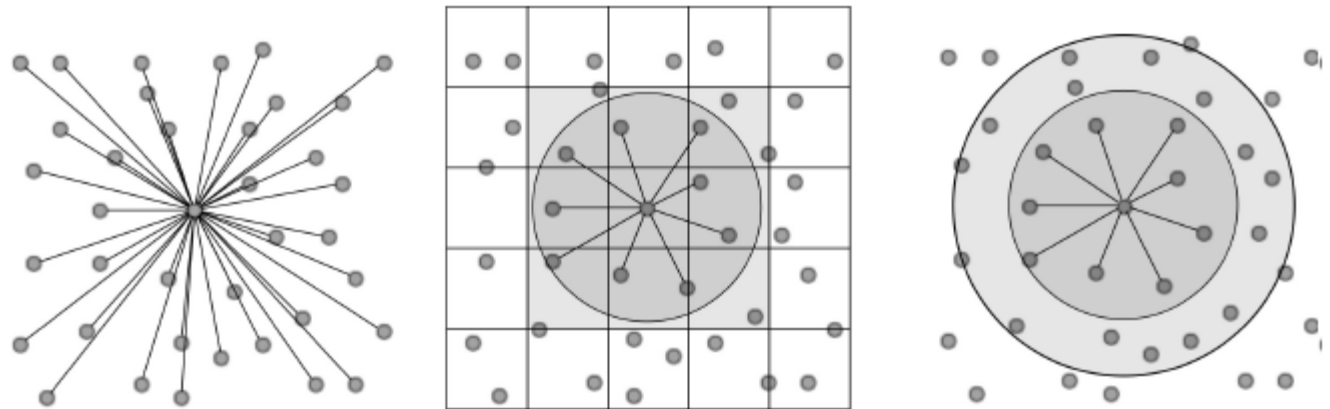
$$\Phi_{ij} = 4\epsilon \left[\left(\frac{\sigma}{r_{ij}} \right)^{12} - \left(\frac{\sigma}{r_{ij}} \right)^6 \right]$$

MD Computing

- Force Calculation

- All pairs simulation uses local cell and neighbour lists to reduce the N^2 calculation to order N

$$F_i = \sum_{j \neq i}^N f_{ij}$$



- Move particles (leapfrog in time)

$$m_i \frac{dv_i}{dt} \approx m_i \frac{v_i(t + \Delta t/2) - v_i(t - \Delta t/2)}{\Delta t} = F_i$$

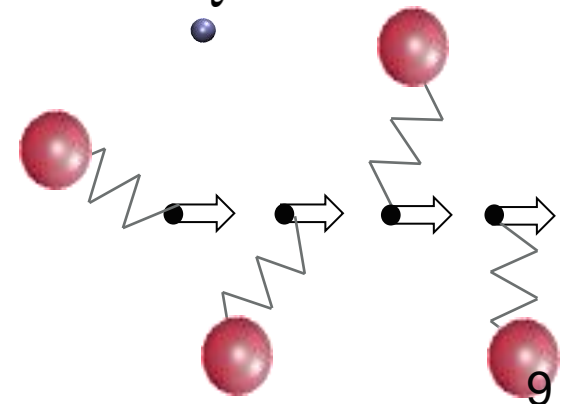
$$\frac{dr_i}{dt} \approx \frac{r_i(t + \Delta t) - r_i(t)}{\Delta t}$$

Non-Equilibrium Molecular Dynamics

- Non-Equilibrium Molecular Dynamics (NEMD) is the study of cases beyond thermodynamic equilibrium, with:
 - Temperature gradients
 - Flow of fluid (e.g. Couette or Poiseuille flow)
- Essentially fluid dynamics - temperature gradients and flows
 - Thermostats (e.g. Nosé Hoover) remove heat from system
 - Solids of molecules with (an)harmonic springs linking them to tether site
 - Sliding walls by moving molecules
 - Many techniques for inducing flows...

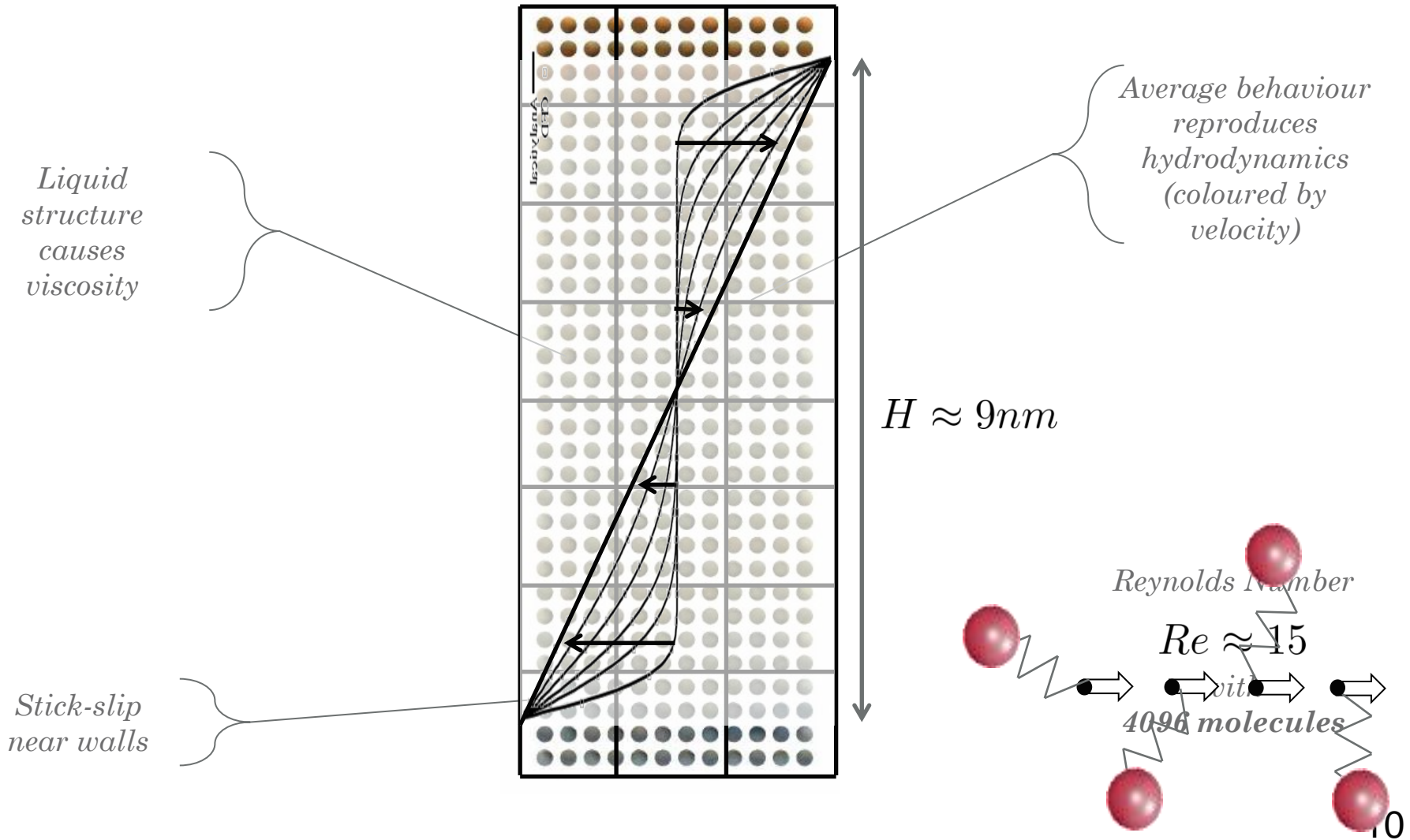
$$m_i \ddot{\mathbf{r}}_i = \mathbf{F}_i + \mathbf{F}_i^{teth} - \psi m_i \mathbf{c}_i$$

$$\dot{\psi} = \frac{1}{Q} [T - 3T_{target}]$$

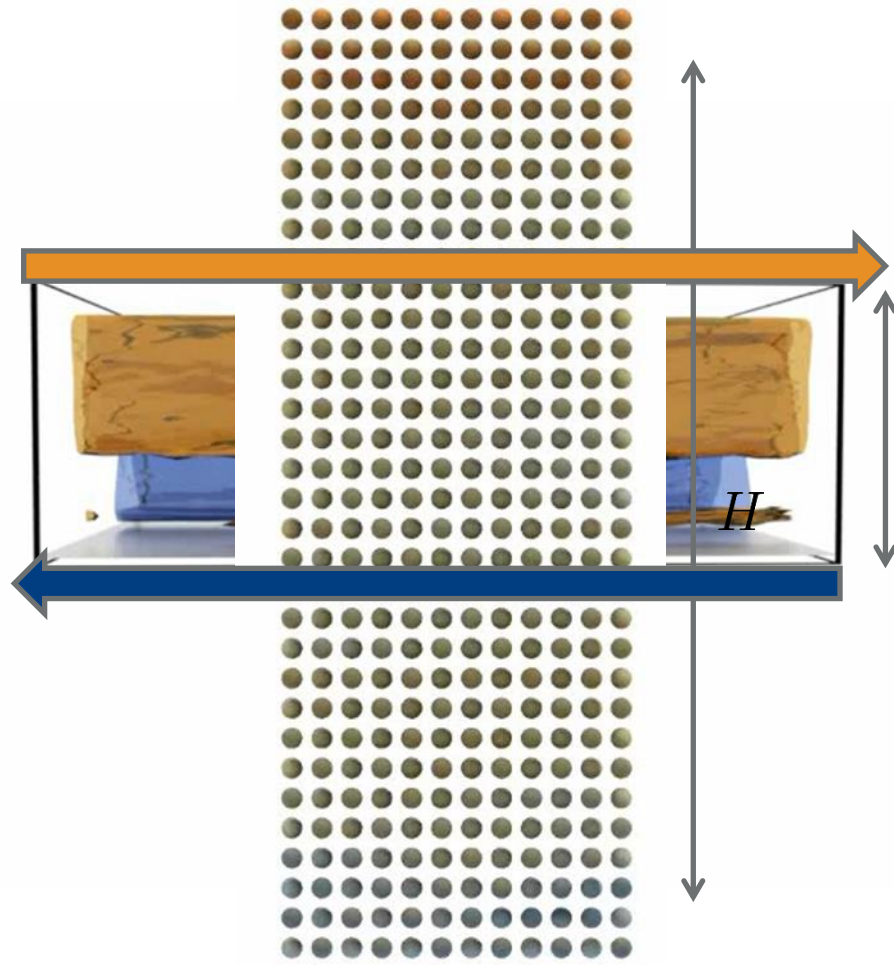




Wall Bounded Flow



Wall Bounded Flow

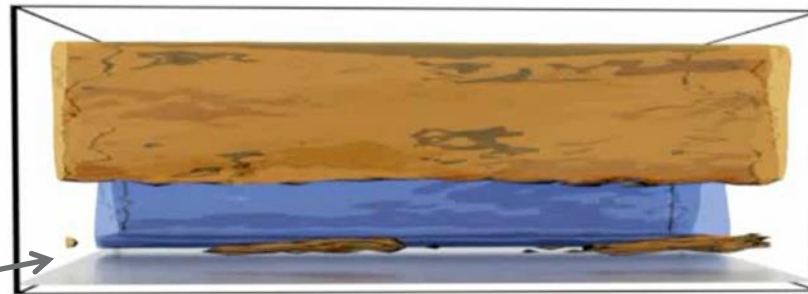


Reynolds Number
 $Re \approx 400$
with
300 million
molecules

Wall Bounded Flow

*Minimal channel Couette
flow*

$$L \approx 523nm$$



$$H \approx 190nm$$



$$W \approx 359nm$$

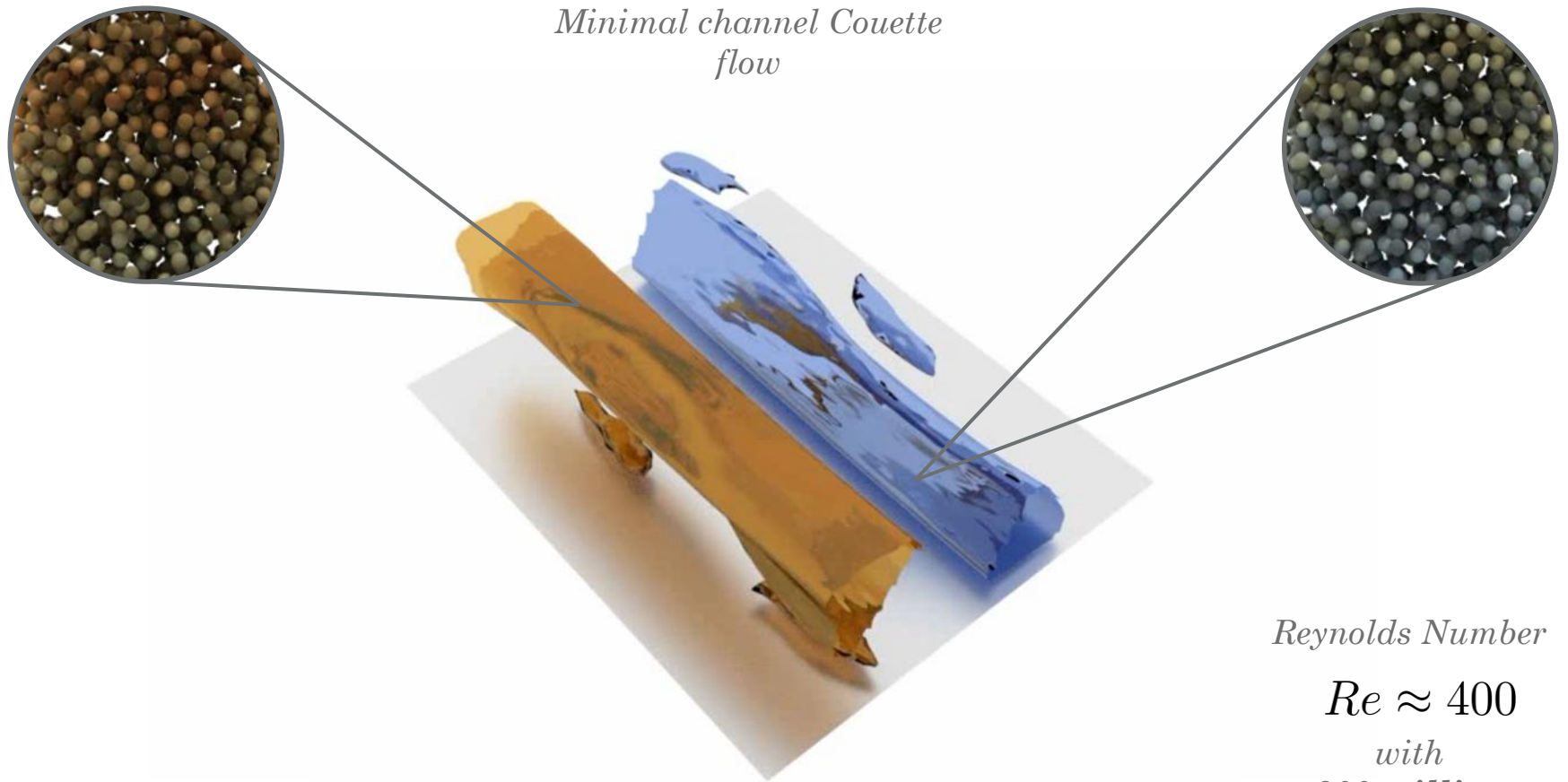


Reynolds Number

$$Re \approx 400$$

*with
300 million
molecules*

Molecular Dynamics - Turbulence



*Minimal channel Couette
flow*

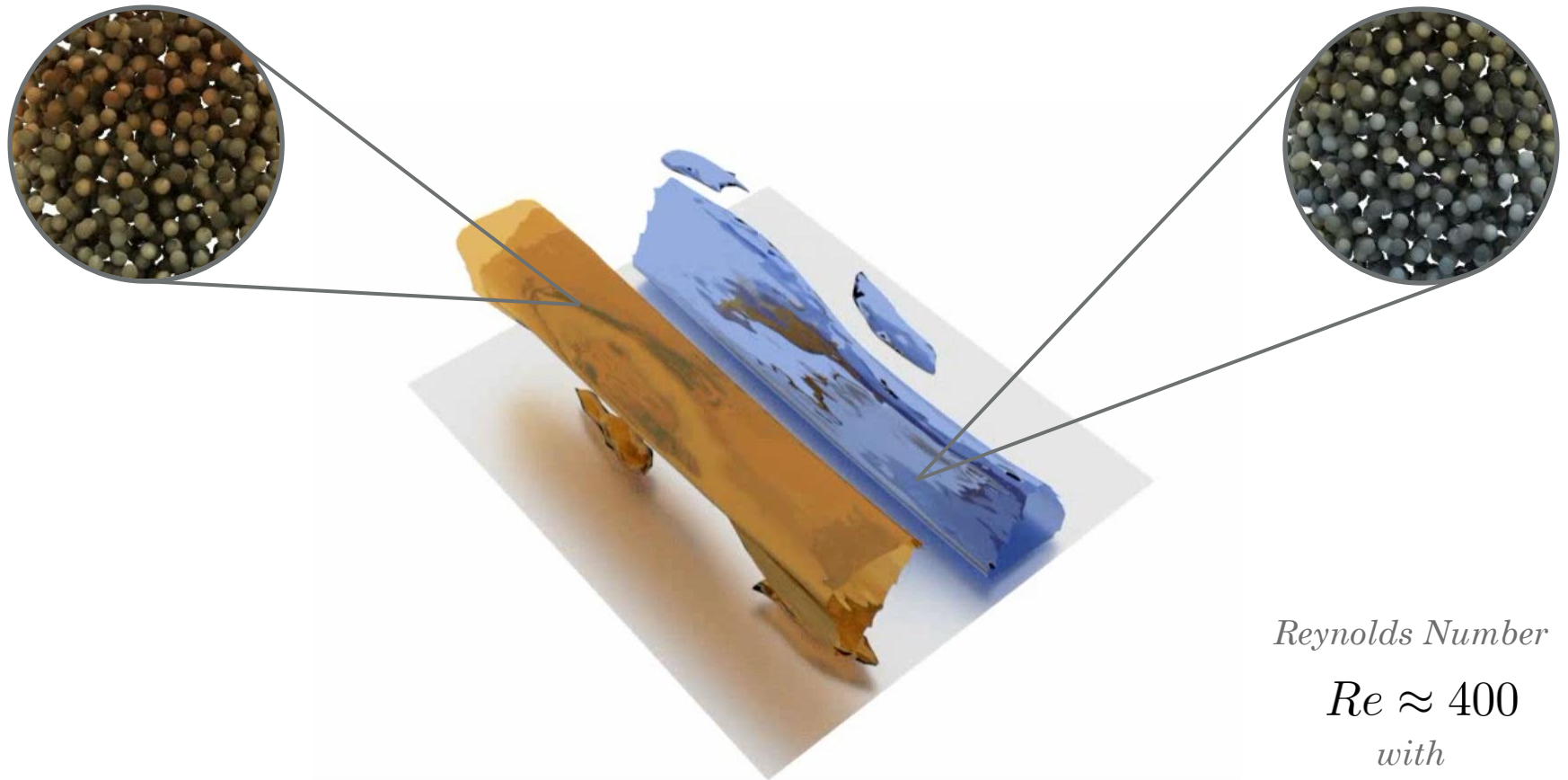
Reynolds Number

$Re \approx 400$

*with
300 million
molecules*

*Isosurfaces of turbulent kinetic
energy coloured by velocity*

Molecular Dynamics - Turbulence



Isosurfaces of turbulent kinetic energy coloured by velocity

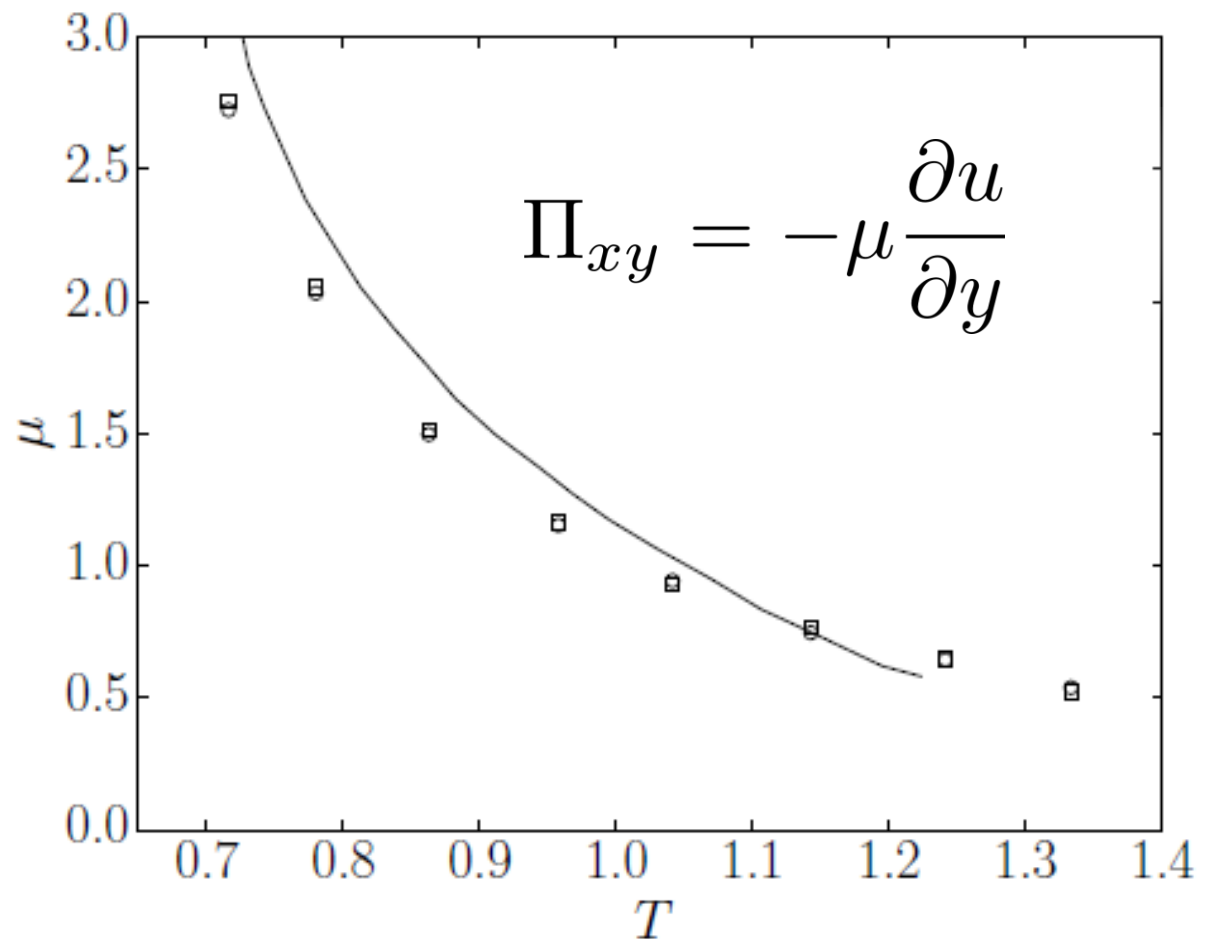
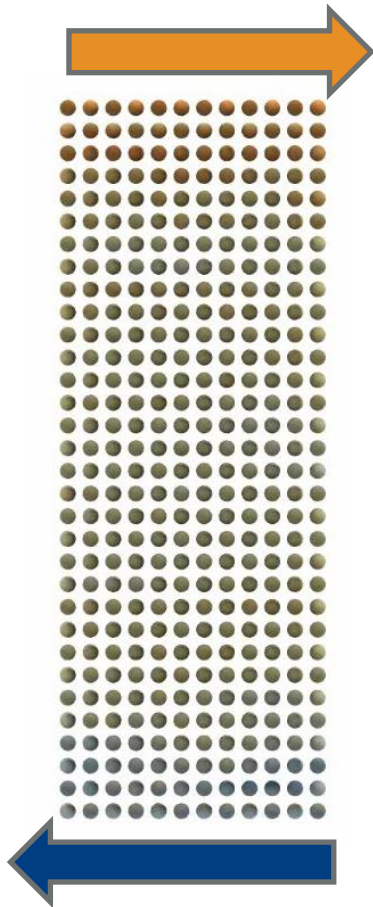
Reynolds Number

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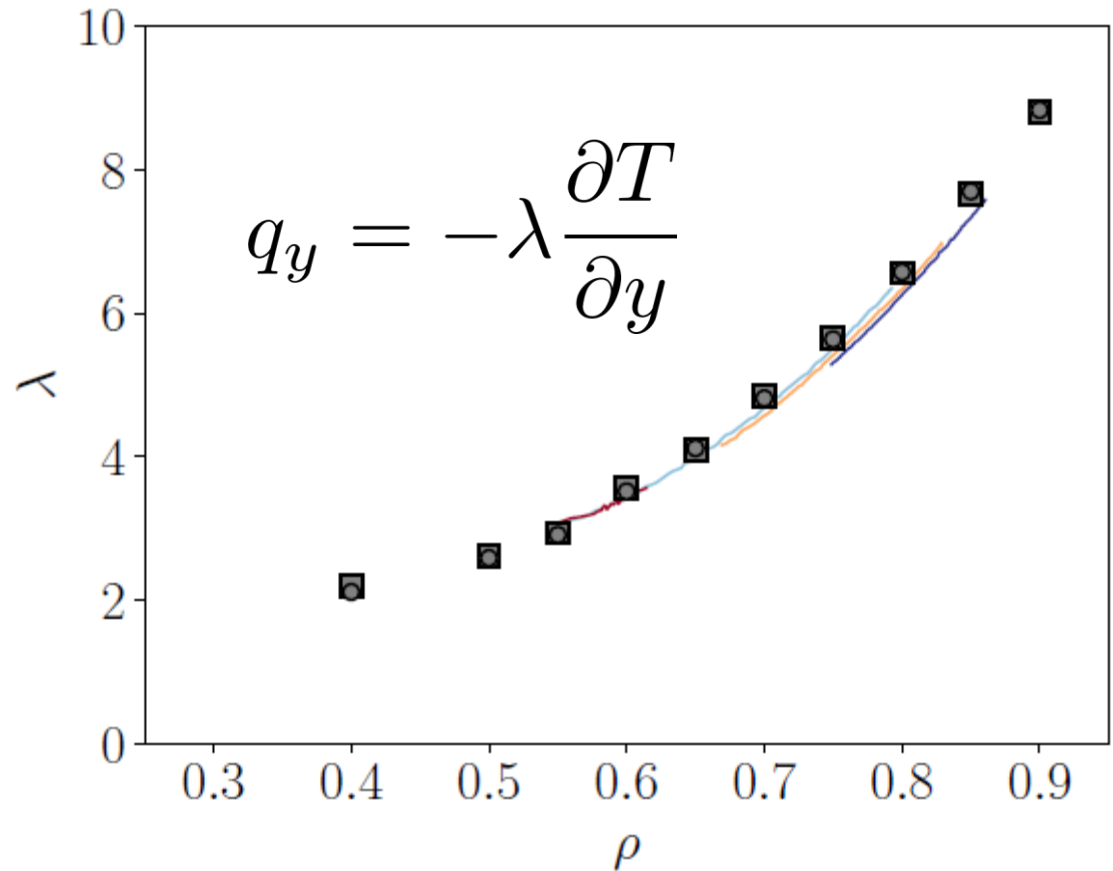
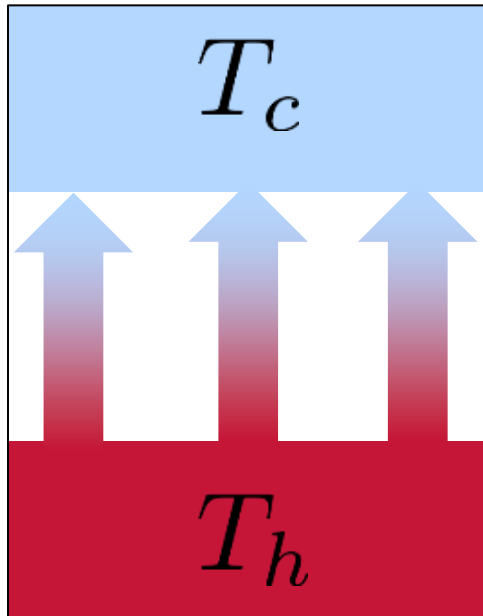
Viscosity

- Good agreement with experiments



Heat Conduction

- Good agreement with experiments

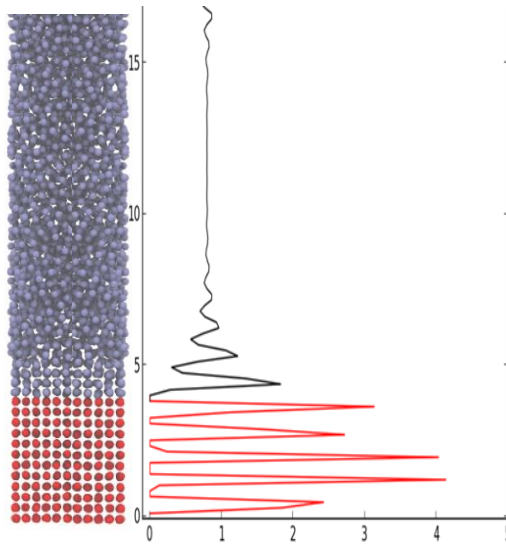




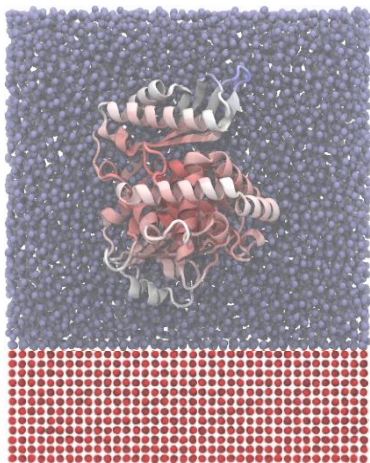
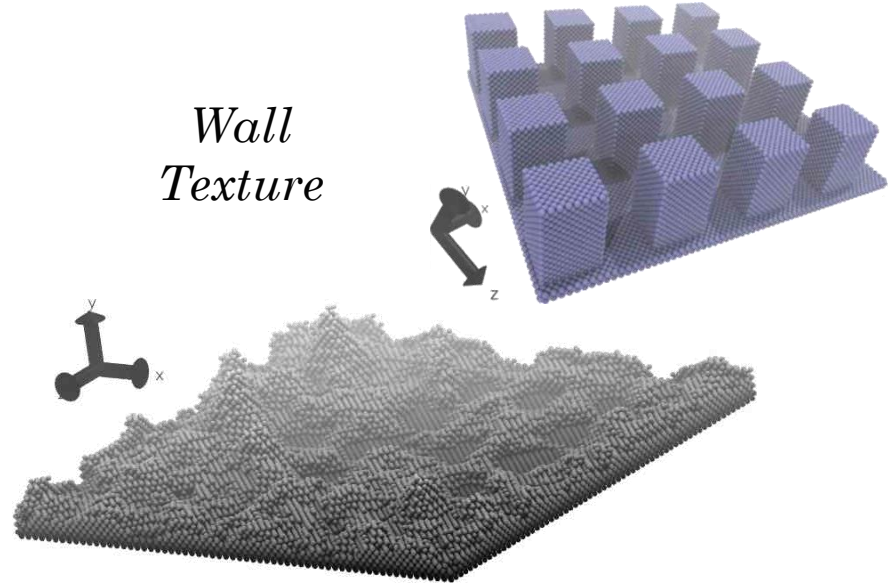
Molecular Dynamics – Complex Walls and Fluids

Liquid structure causes viscosity

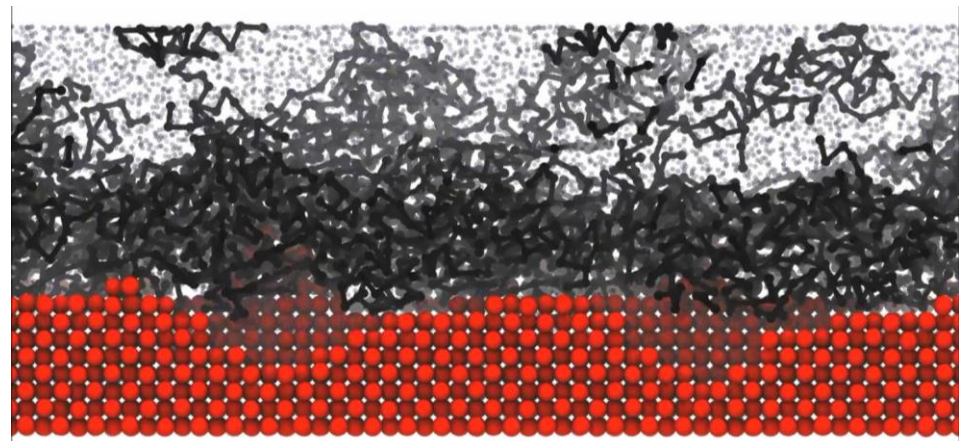
Stick-slip near walls



Wall Texture

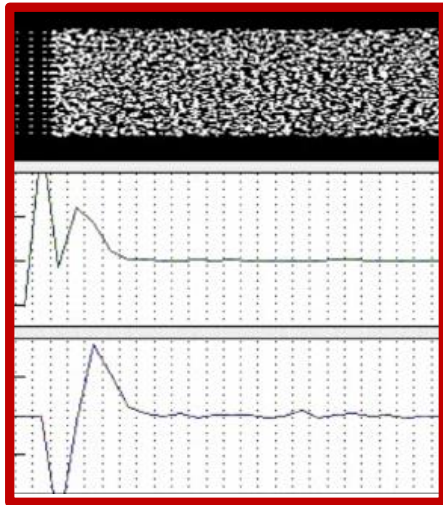


Molecules of arbitrary complexity

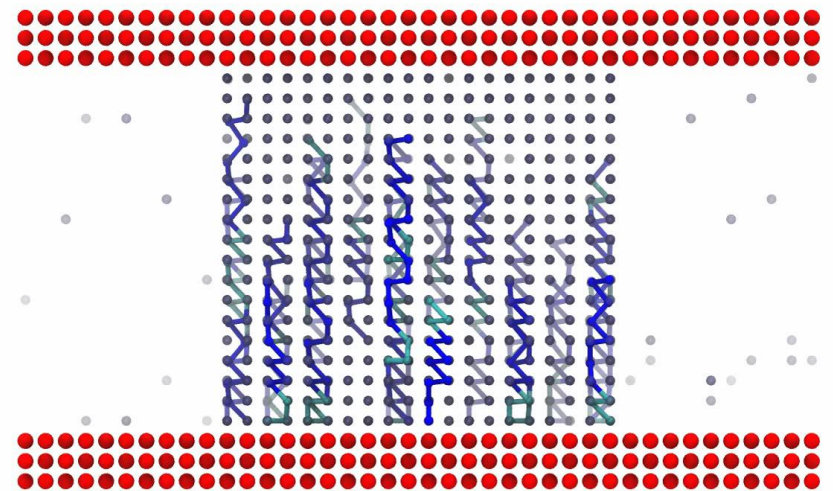


Molecular Dynamics – Shocks and Multi-Phase

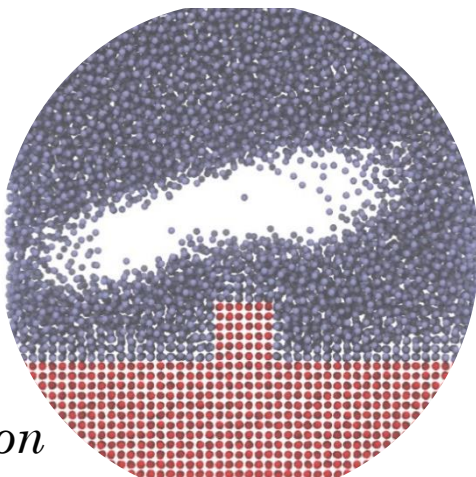
Shockwave



Droplet Formation

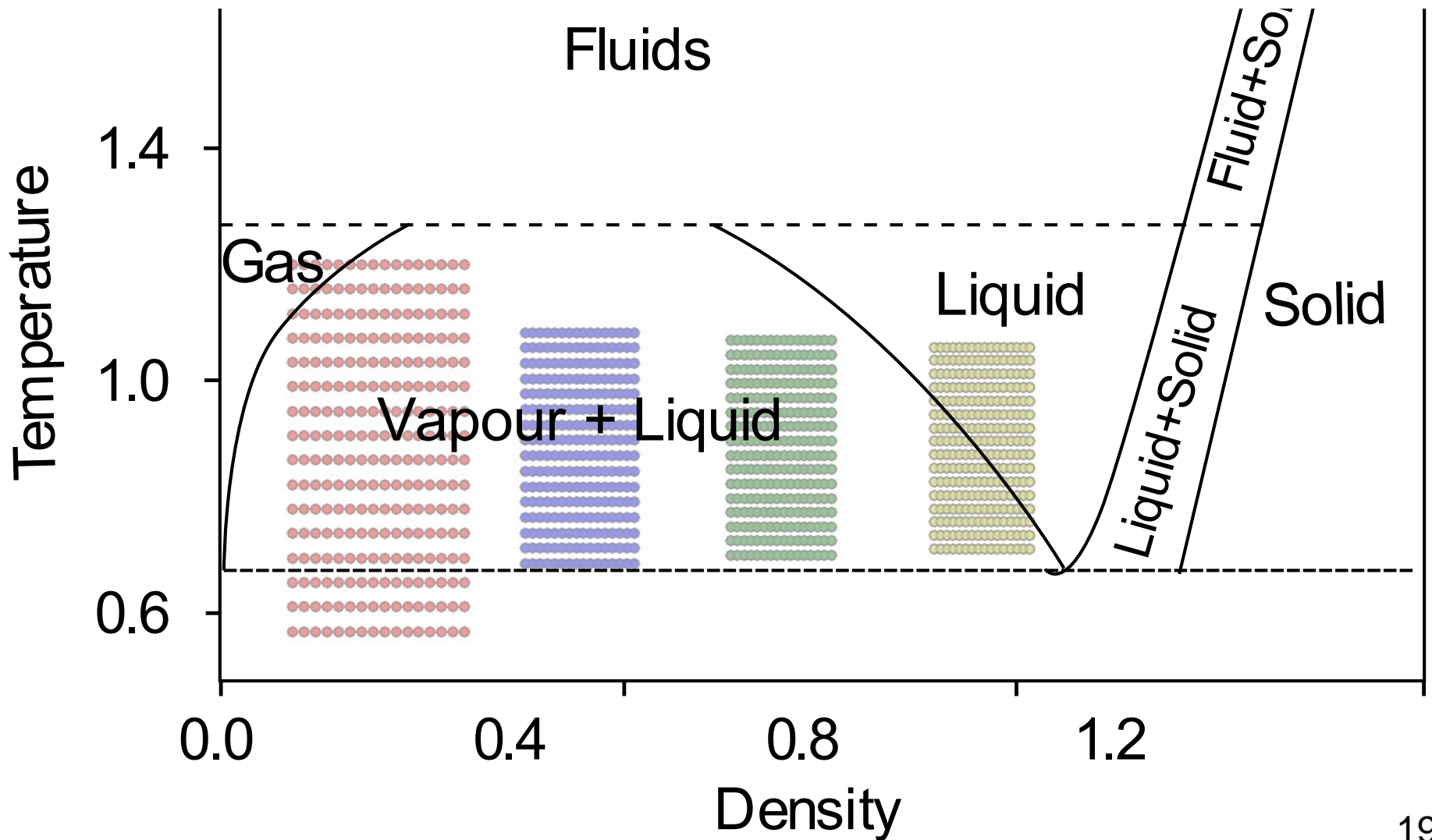


Nucleation



Contact line

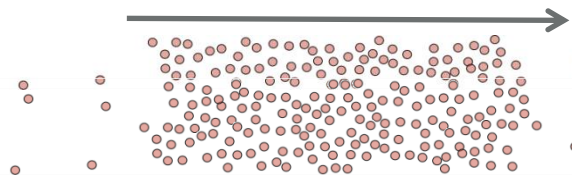
Phase Change



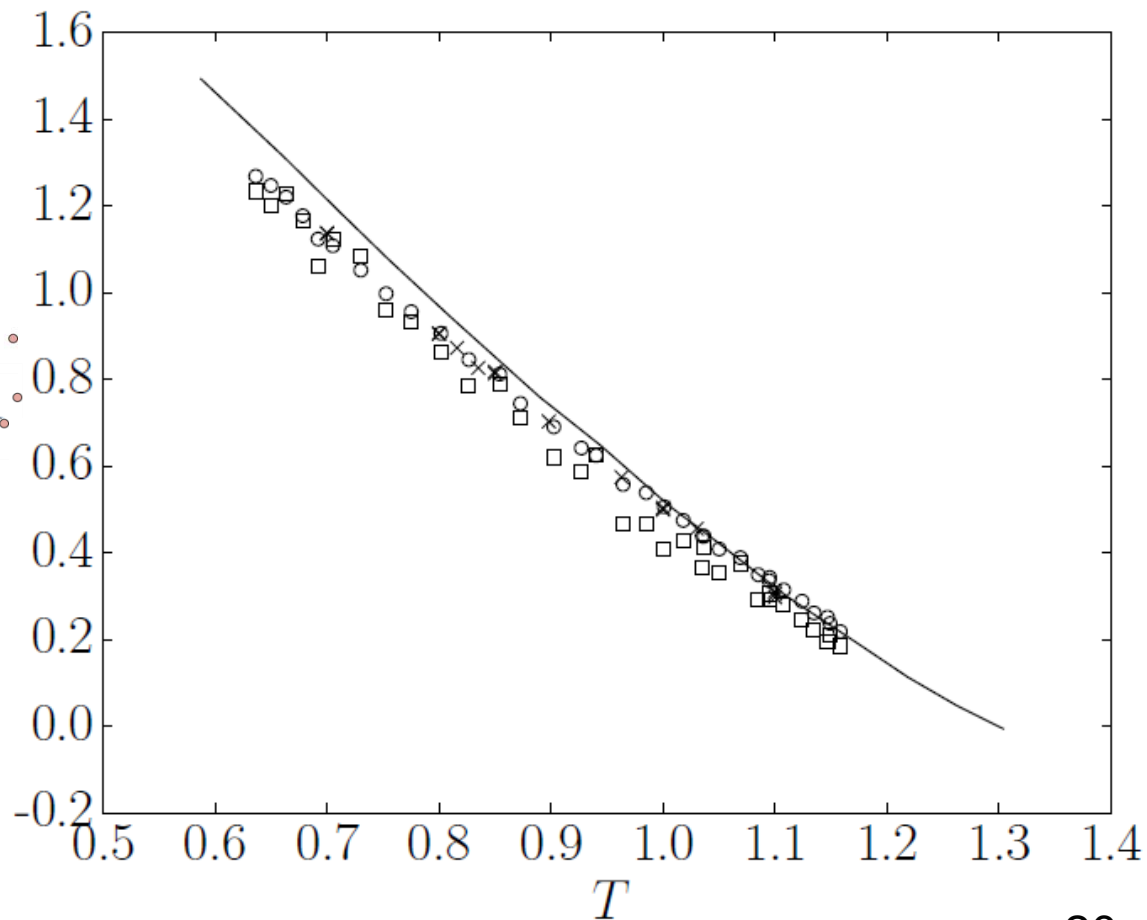
Results for Surface Tension

- Good agreement with experiments

$$\gamma = \int_{-\infty}^x [\Pi_N - \Pi_T] dx$$



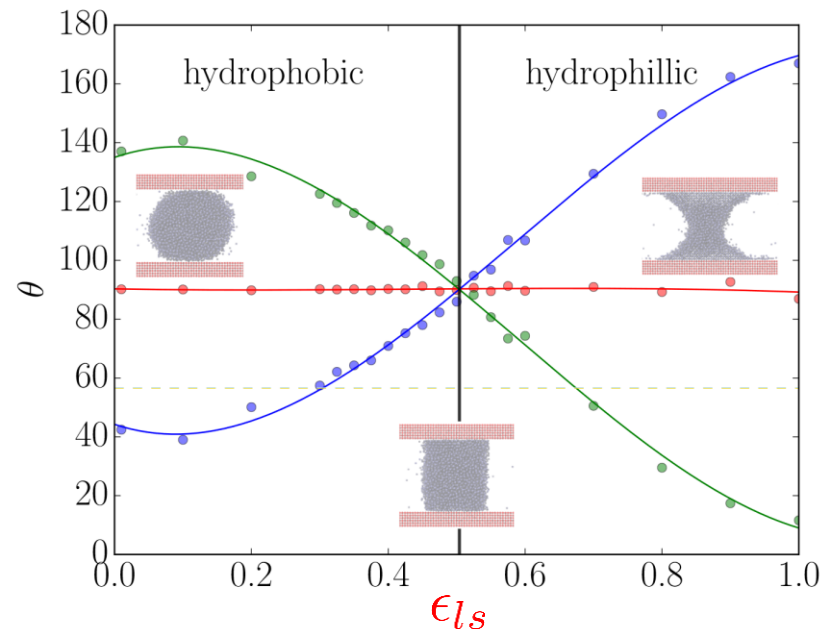
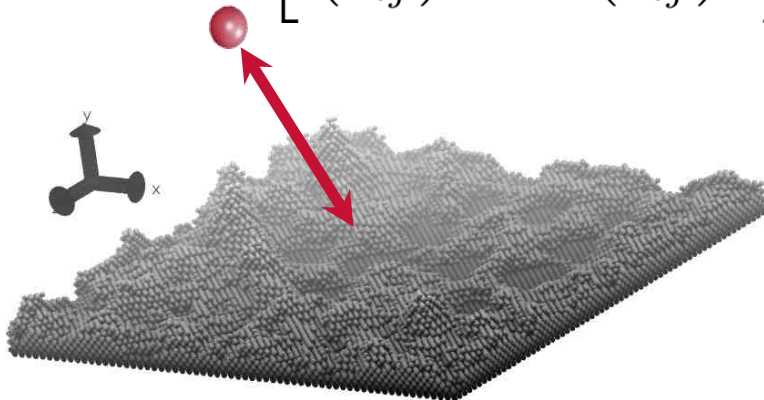
Integrate
over
Liquid
Vapour
interface(s)



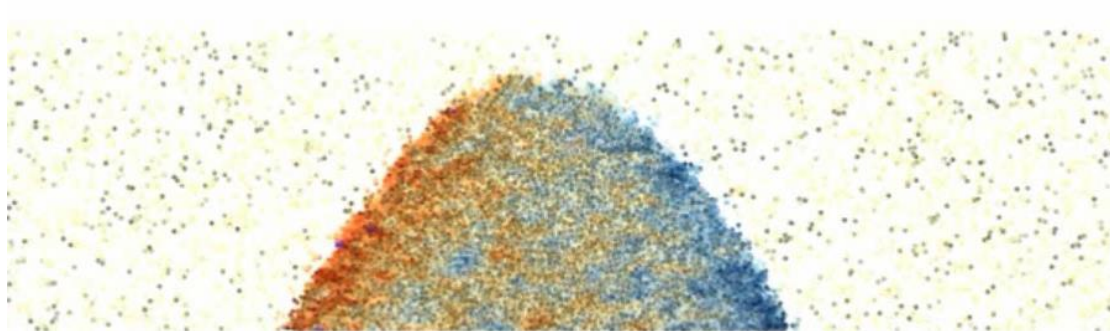
Surface Wettability

- Depends entirely on the “wetting” interaction between surface and fluid, tuned using:
 - Mixing rules, e.g. Lorentz-Berthelot $\epsilon_{ls} = \sqrt{\epsilon_l \epsilon_s}$
 - Bottom up simulation e.g. quantum mechanics
 - Top down e.g. to get desired contact angle

$$\phi(r_{ij}) = 4\epsilon_{ls} \left[\left(\frac{\sigma}{r_{ij}} \right)^{12} - \left(\frac{\sigma}{r_{ij}} \right)^6 \right]$$

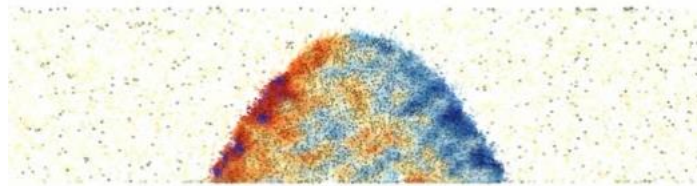


The Moving Contact Line

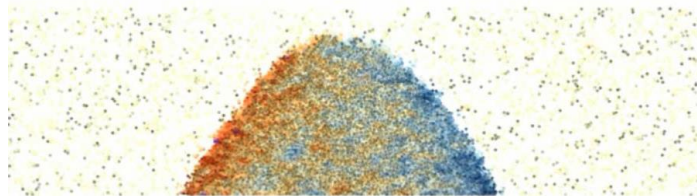


MD Simulation of Droplets

- Low Wettability



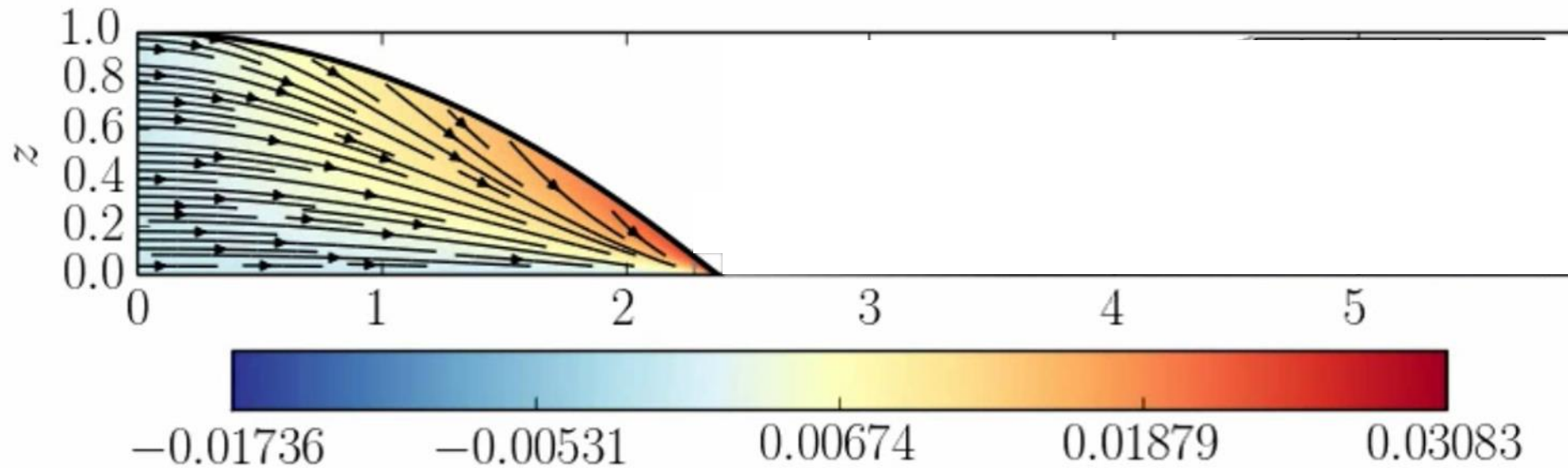
- Intermediate Wettability



- High Wettability

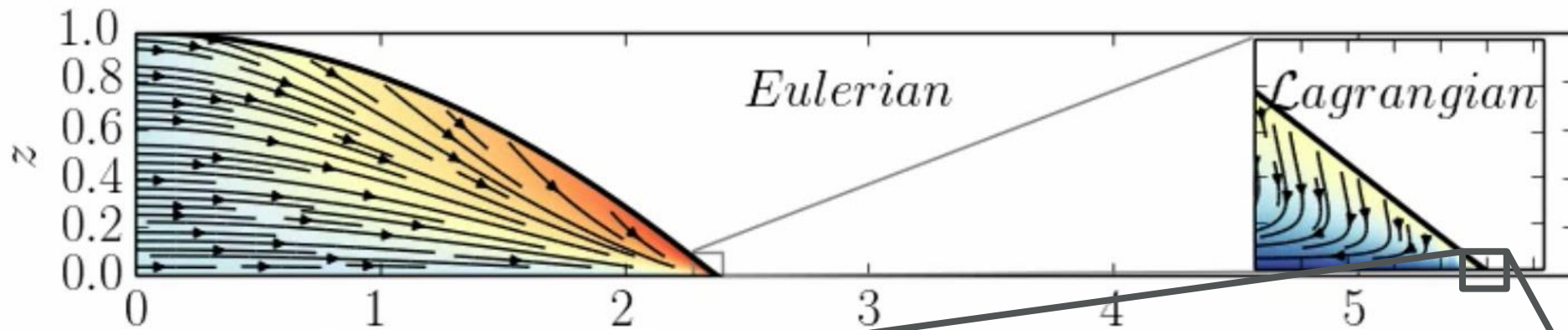


Dynamic Contact Line



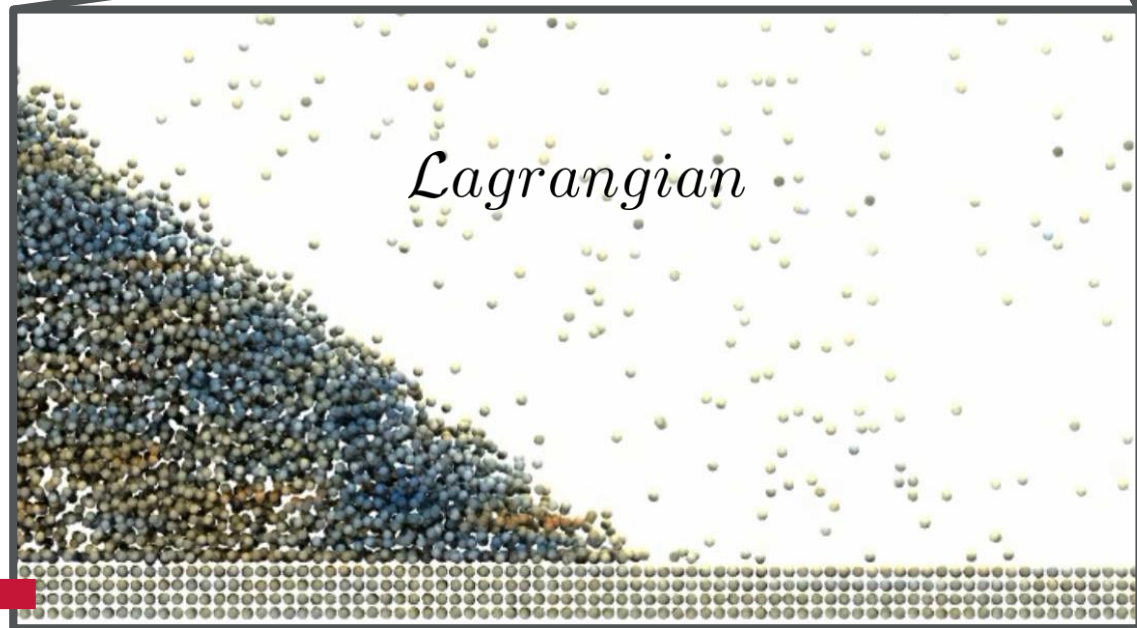


Dynamic Contact Line



- Model the moving contact line with MD
- We want contact line speed as a function of continuum contact angle

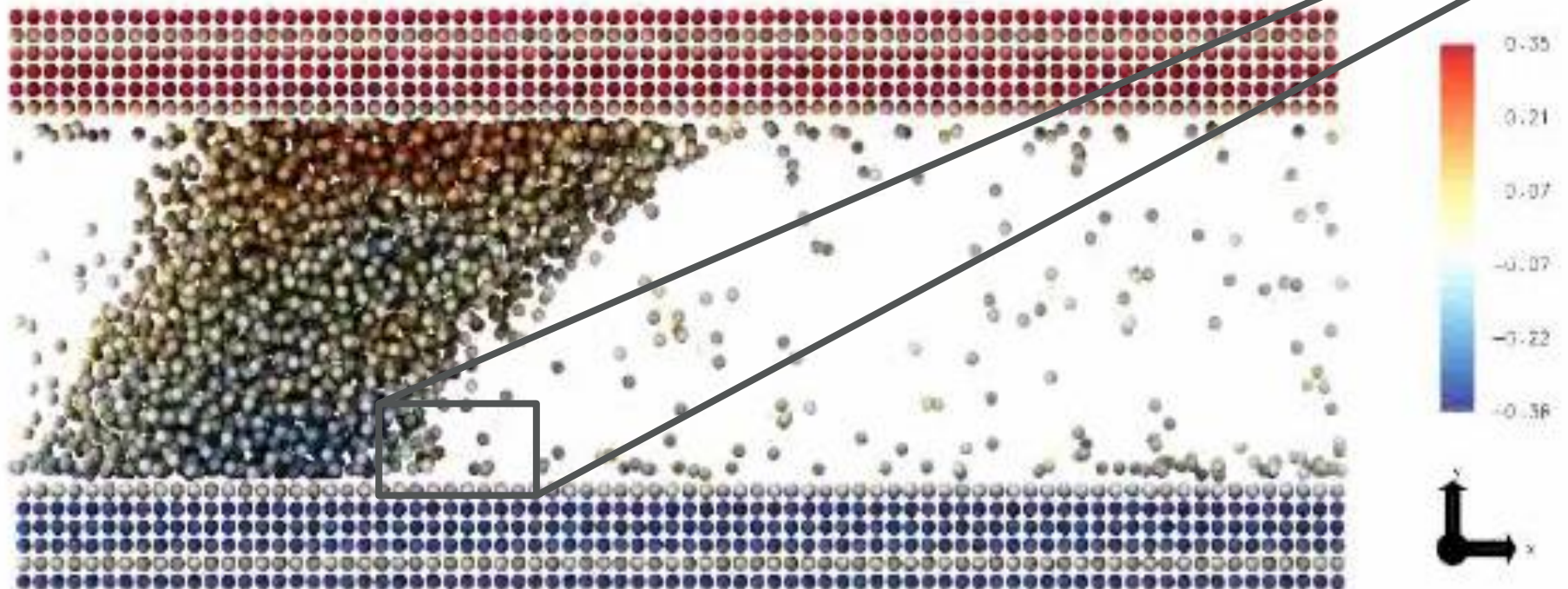
$$\frac{dx_c}{dt}$$



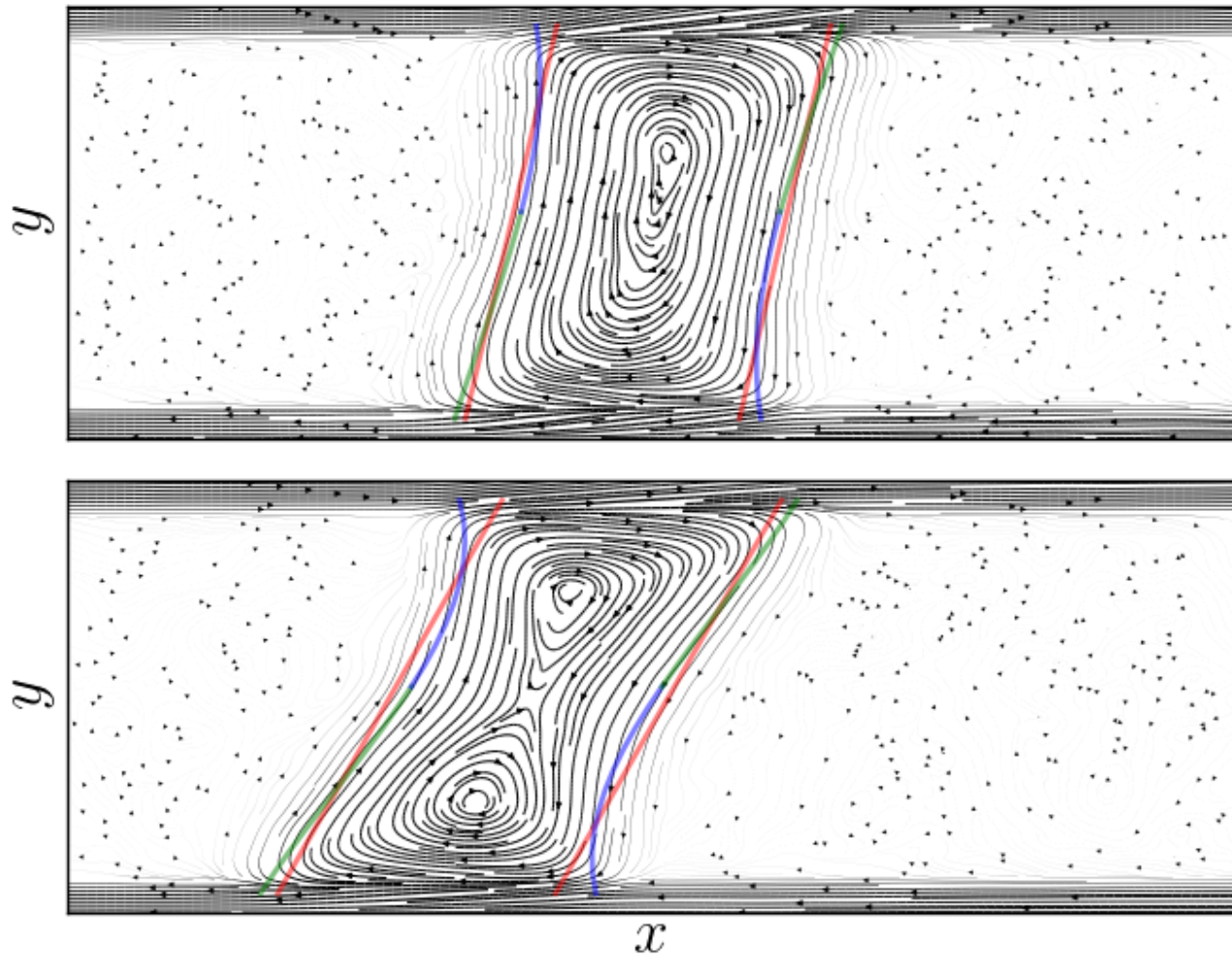


Dynamic Contact Line

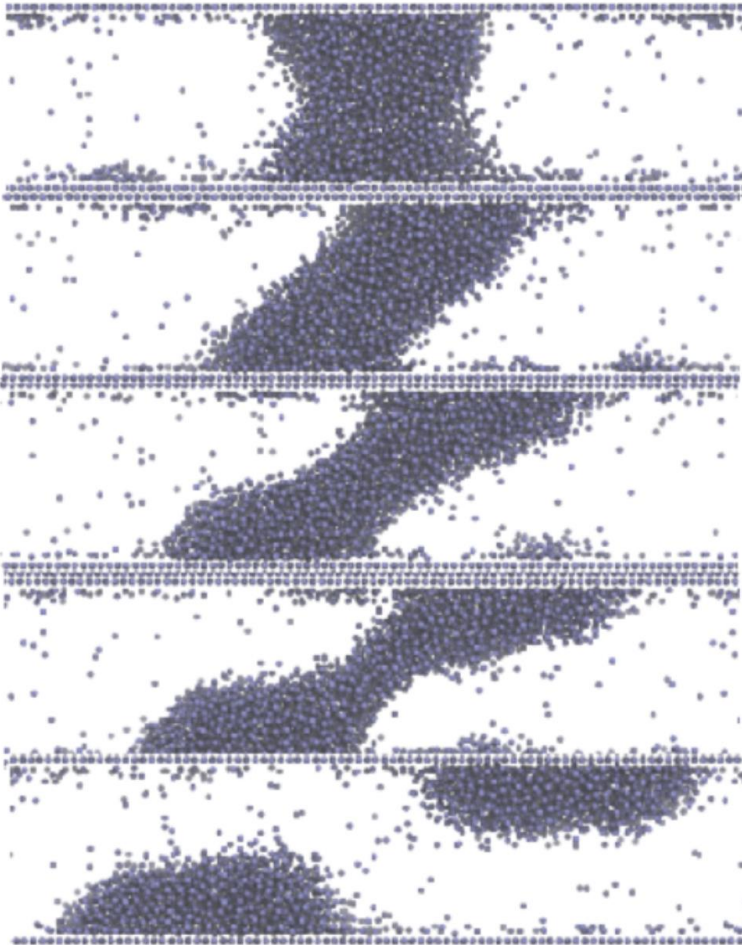
- Two fluid phases and sliding molecular walls
- Wall velocity vs contact line angle
- MD captures the sliding contact line but we need to work out how to interpret it



Flowfields



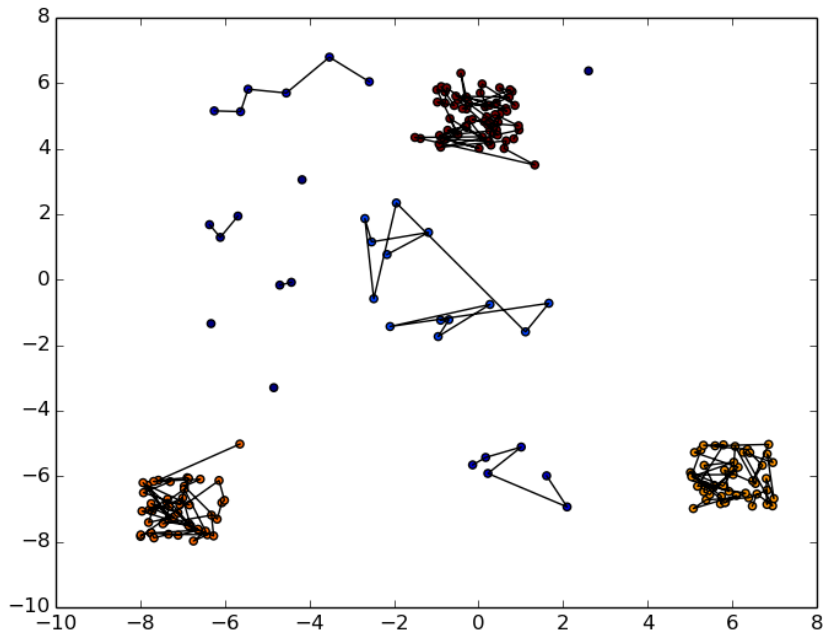
Link to Experiments



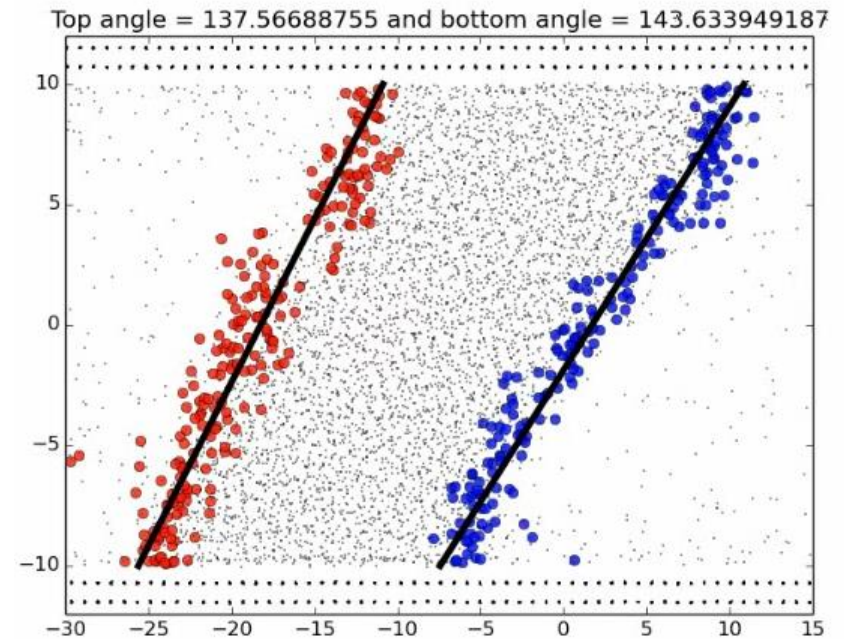
*L. Wang and T. J. McCarthy,
Langmuir, 2013, 29, 7776*

Identifying the Liquid and Surface Fitting

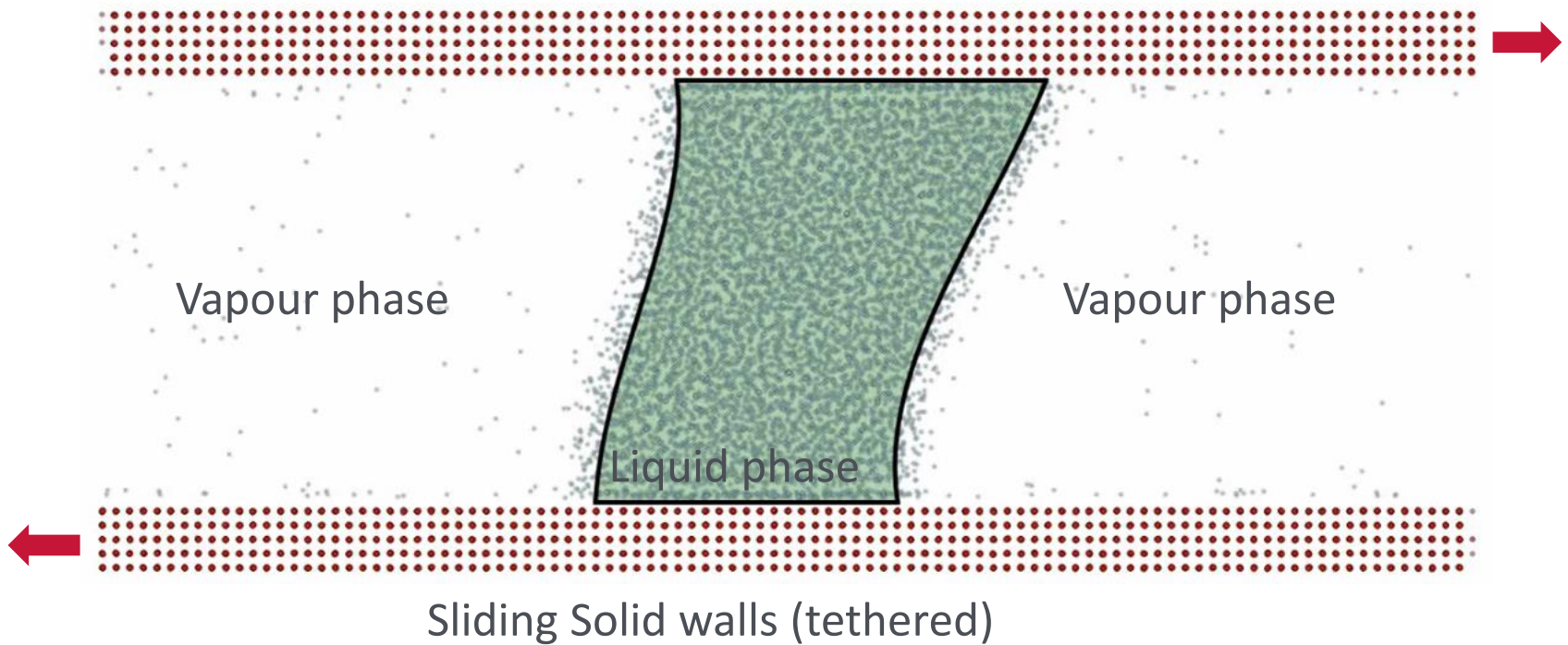
Cluster Analysis to identify the liquid droplets



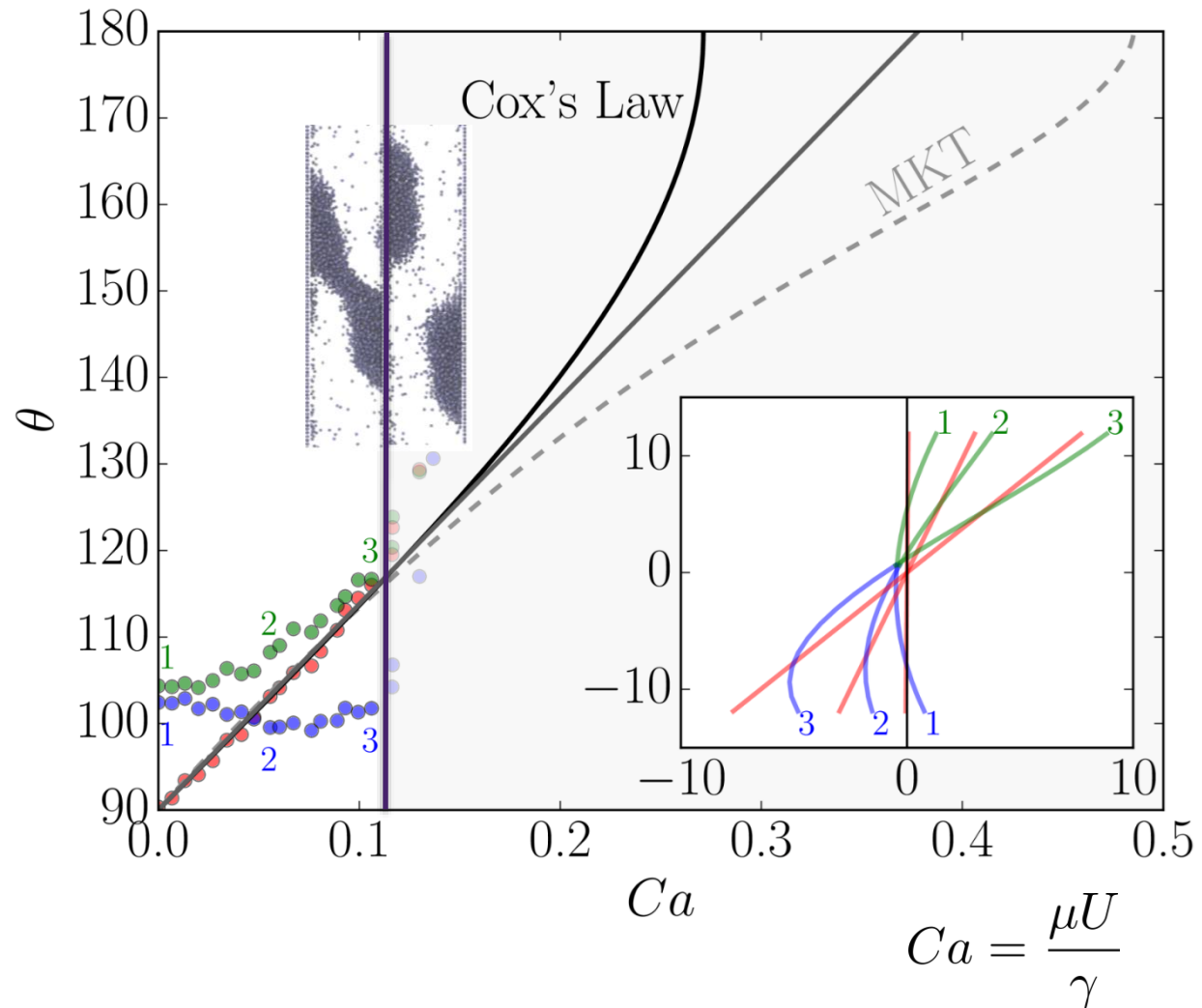
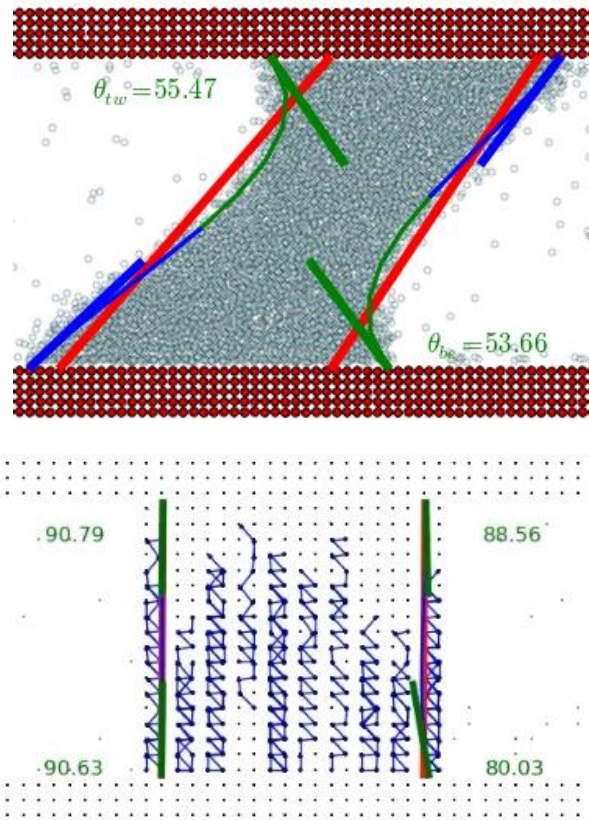
Least Square fitting to outer molecules, linear example here



A Cubic Fit to the Interface

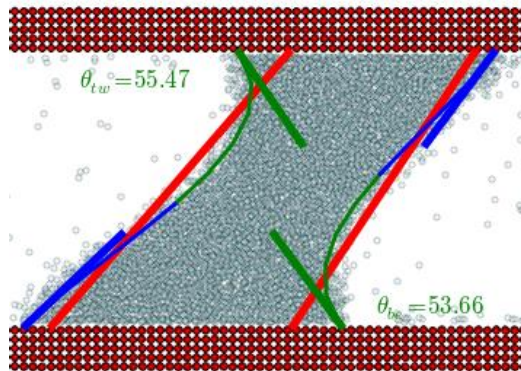


Angle vs. Velocity

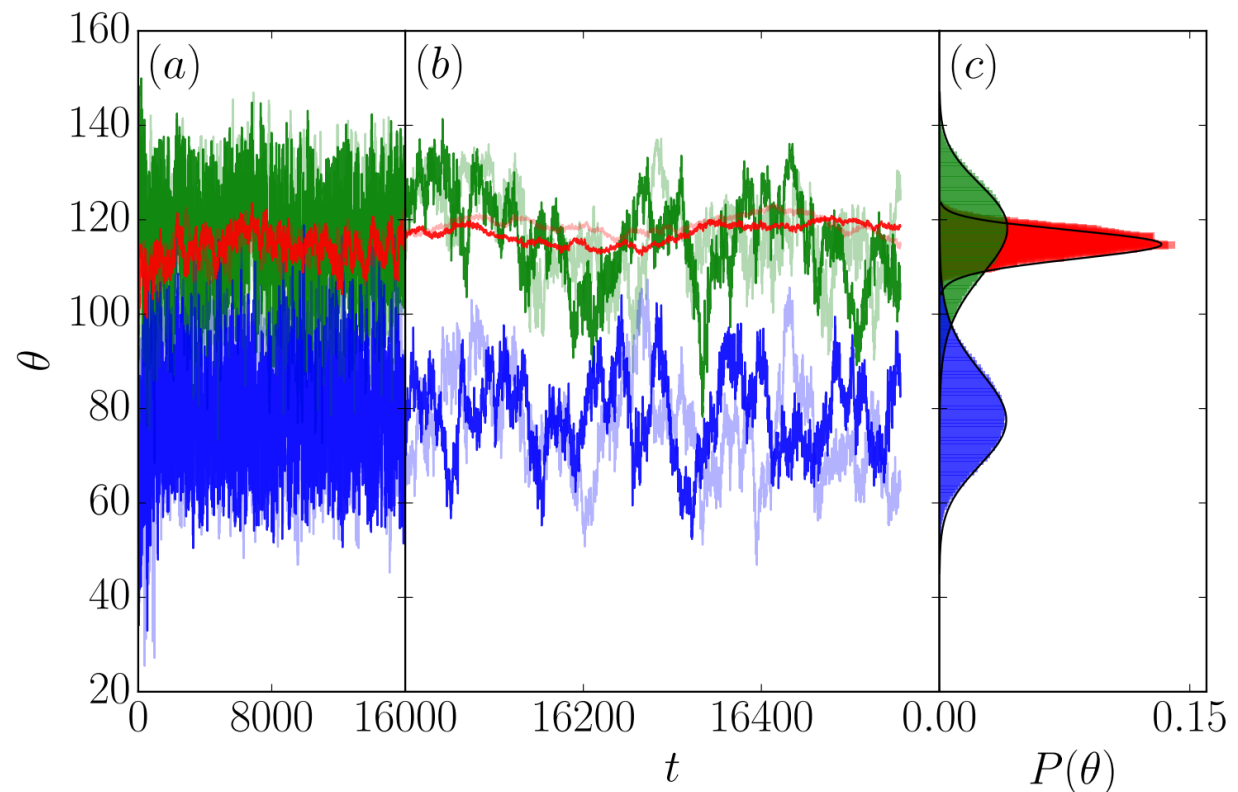


Time Evolution of Contact Angle

- Linear and Cubic Fitting angles fluctuates over time



- Steady state so we can take the probability density function of angle



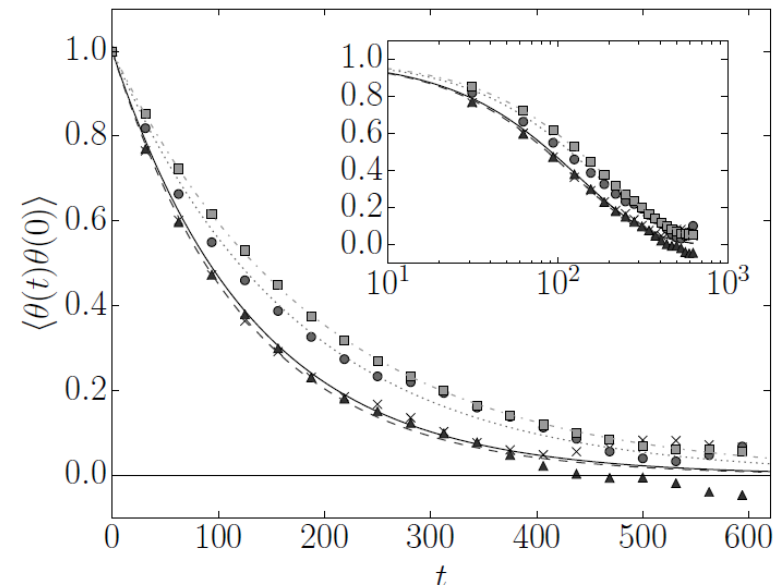
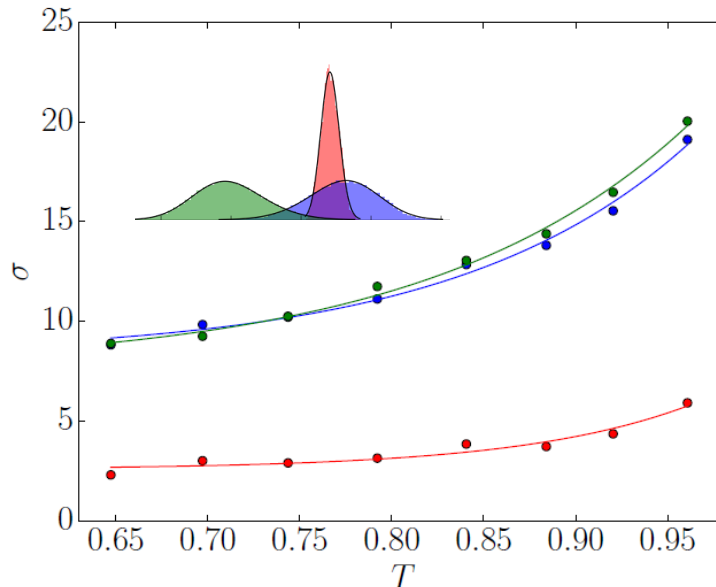
- Linear, Advancing and Receding angles*

Building this into the Continuum Model

- A Langevin Equation uses random noise to model this

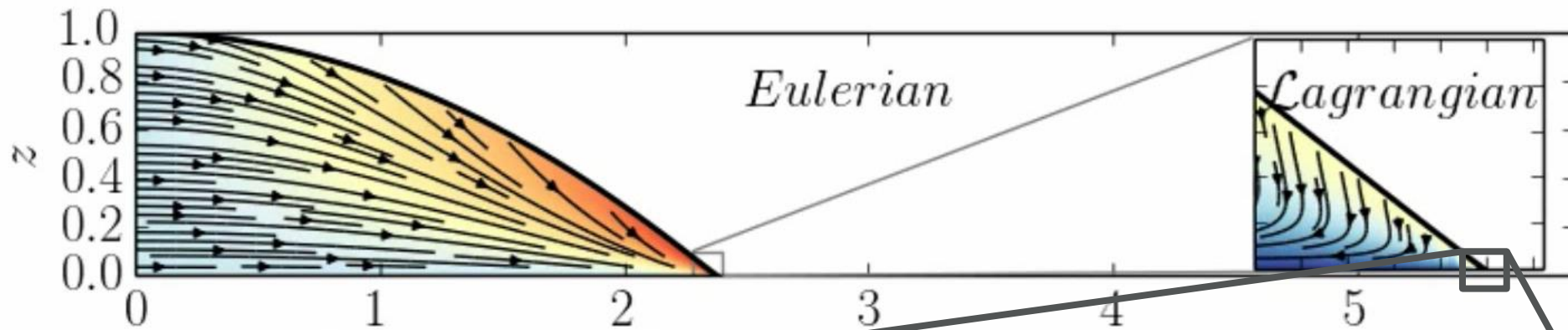
$$\dot{\theta} + \frac{k}{\Gamma} [\theta - \langle \theta \rangle] - \frac{1}{\Gamma} \xi(t) = 0 \text{ where } \langle \xi(t) \xi(t') \rangle = C \delta(t - t'),$$

- Coefficients parameterised using
 - Standard deviation – range of fluctuations
 - Autocorrelation – how quickly they decay.





Building this into the Continuum Model



- Bulk dynamics from a law like Tanners or Cox
- Add in fluctuations from the MD using the Langevin model

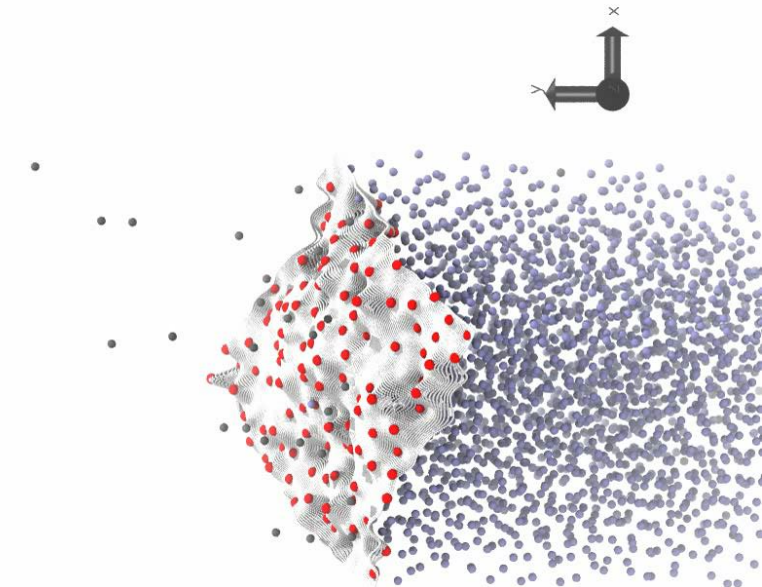
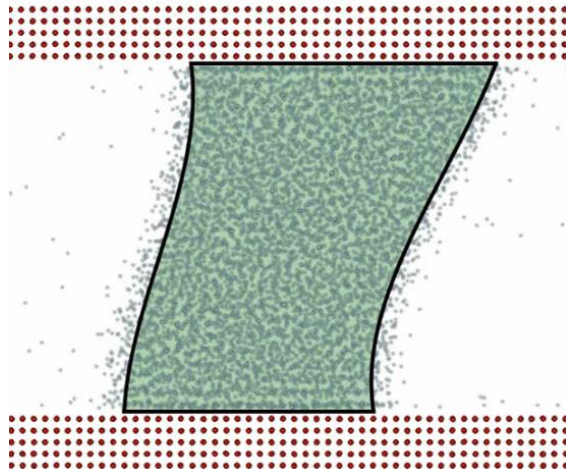
$$\frac{dx_c}{dt}$$



$$\frac{dx_c}{dt} = k(\theta_e - \langle \theta \rangle)^n$$

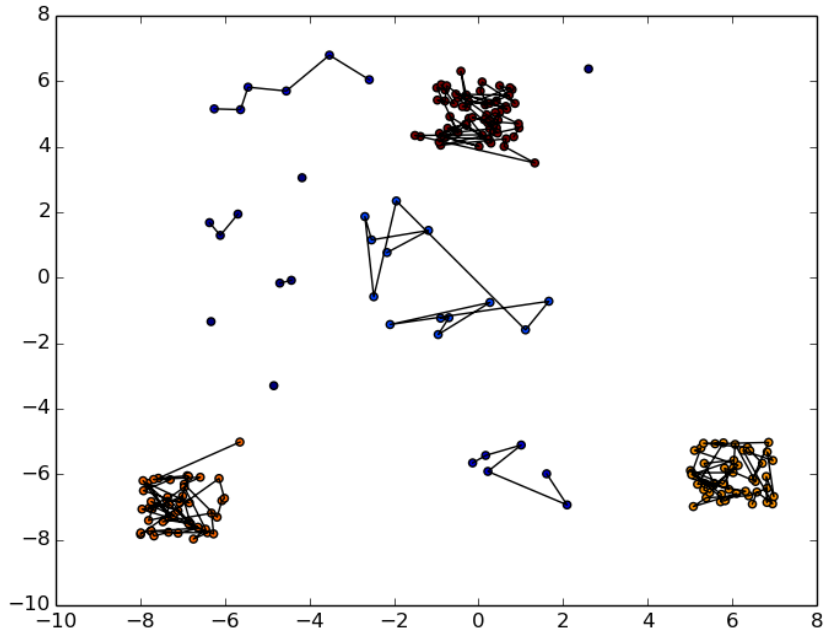
$$\dot{\theta} + \frac{k}{\Gamma} [\theta - \langle \theta \rangle] - \frac{1}{\Gamma} \xi(t) = 0$$

The Intrinsic Interface

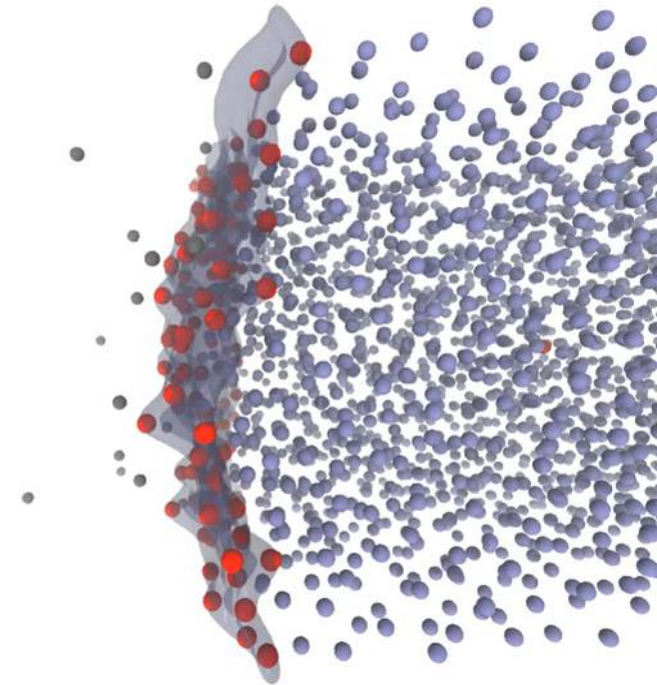


Identifying the Liquid

Cluster Analysis to identify the liquid droplets

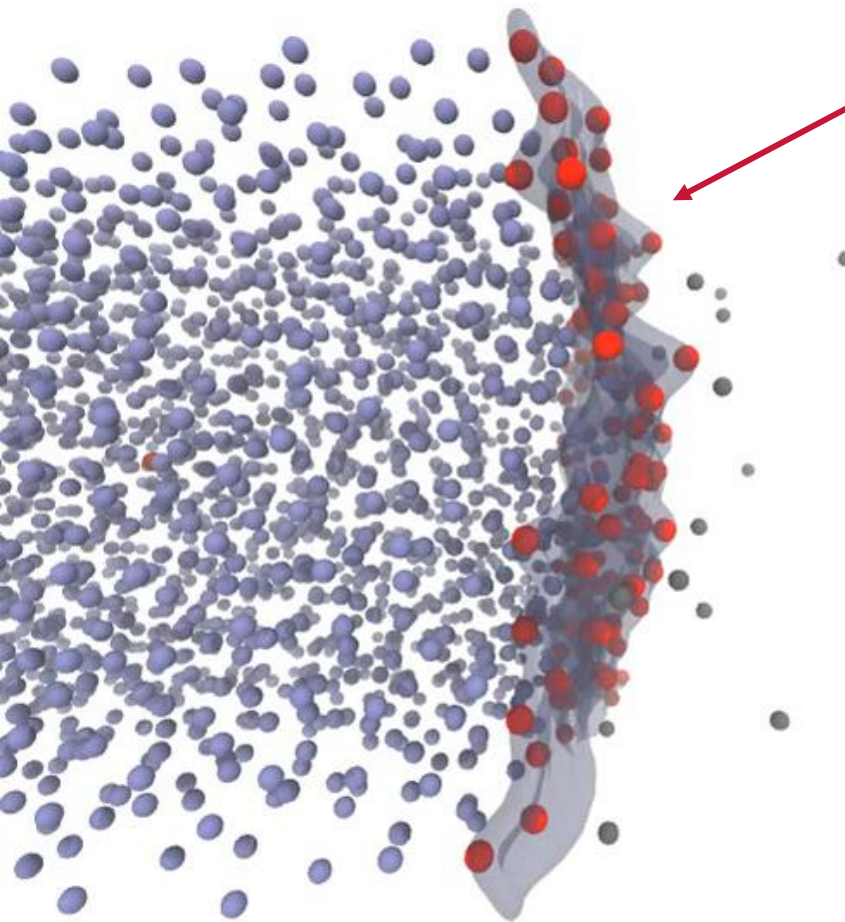


Least Square fitting to outer molecules



The Intrinsic Interface

- The Intrinsic Interface method

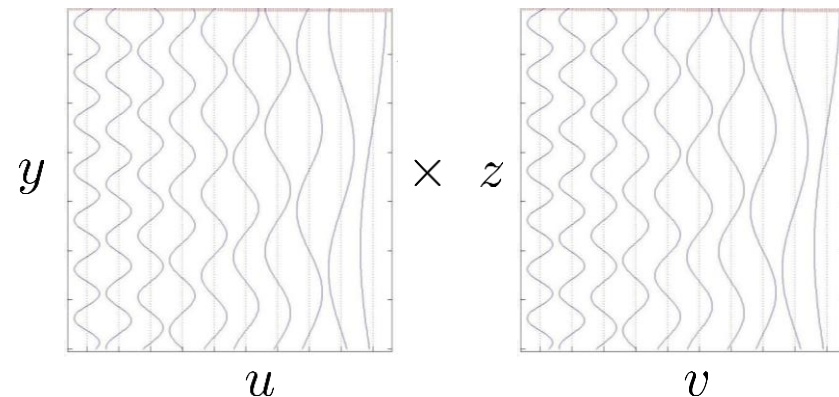


- Surface fitted by least squares

$$\xi = \sum_{\mathbf{k} < k_u} \hat{\xi}_{\mathbf{k}}(t) e^{2\pi i \mathbf{k} \cdot \mathbf{r}} \approx \sum_{\mathbf{k} < k_u} \hat{\xi}_{\mathbf{k}}(t) \xi_u(y) \xi_v(z),$$

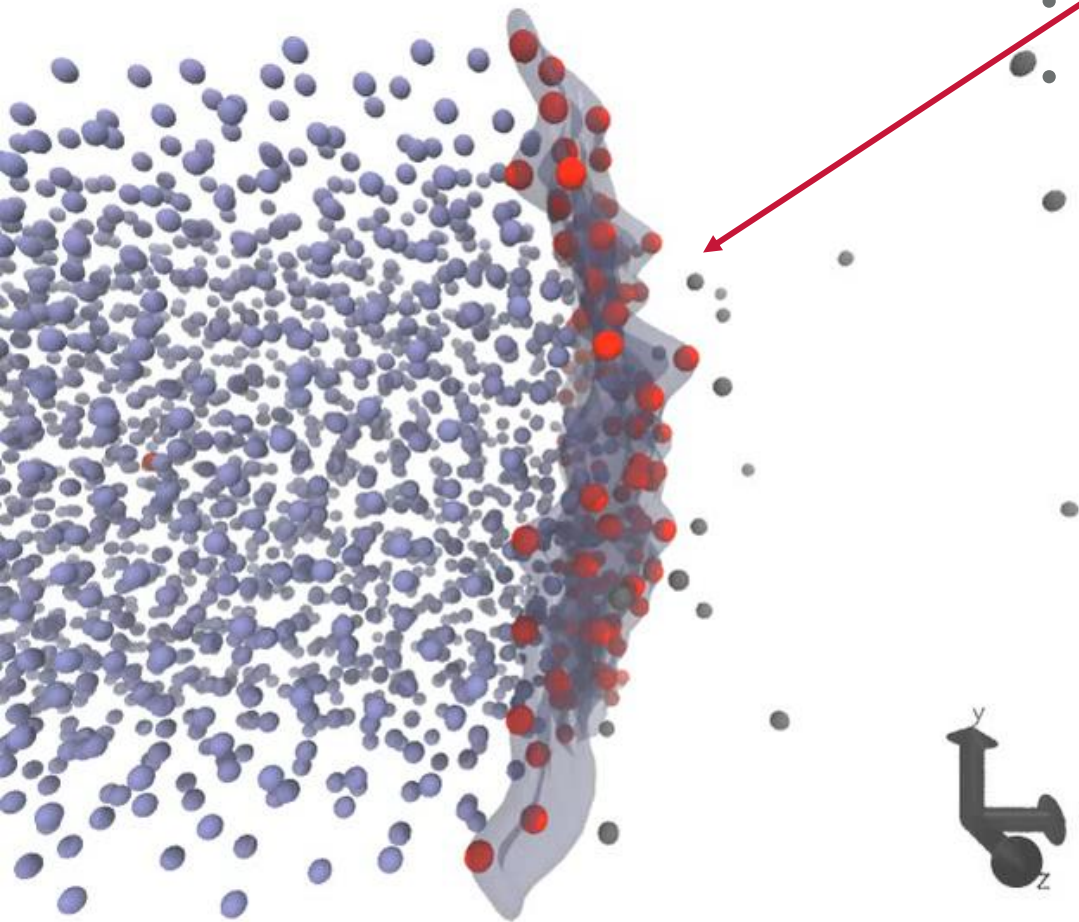
- Sum of sines and cosines down to intermolecular spacing

$$\xi_a(b) = \begin{cases} \cos(2\pi ab) & \text{if } a > 0 \\ \sin(2\pi |a|b) & \text{otherwise,} \end{cases}$$



The Intrinsic Interface

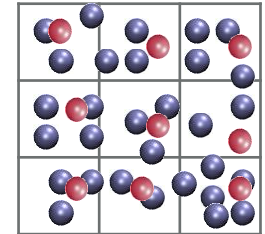
- The Intrinsic Interface method



- Surface fitted by least squares
- Sum of sines and cosines down to intermolecular spacing

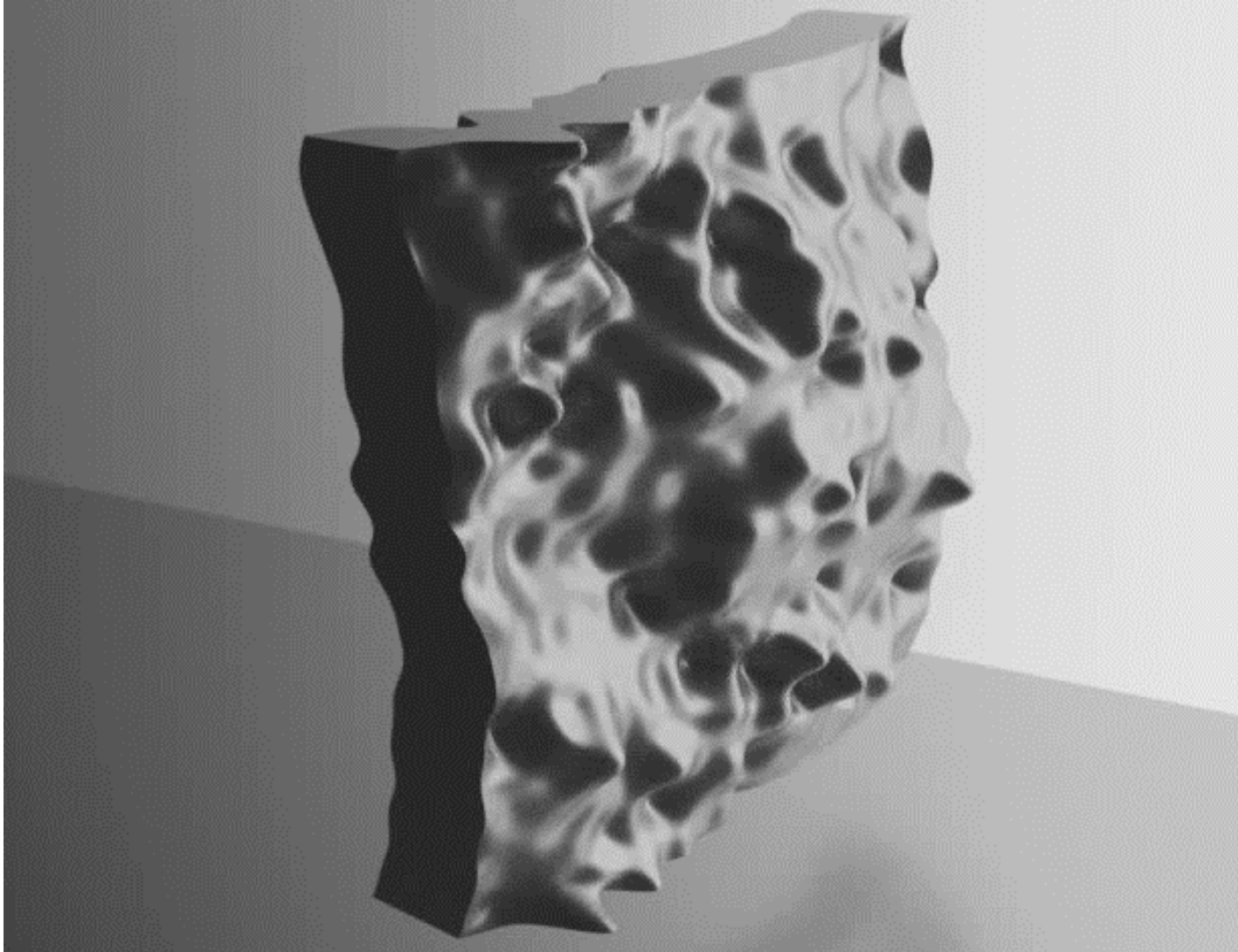
Fitting occurs in stages

- > Fit to an outer set using a grid (e.g. 3 by 3)



- > Order molecules based on distance from surface
- > Add nearest
- > Repeat until surface density reaches target value (~ 0.7)

The Intrinsic Interface

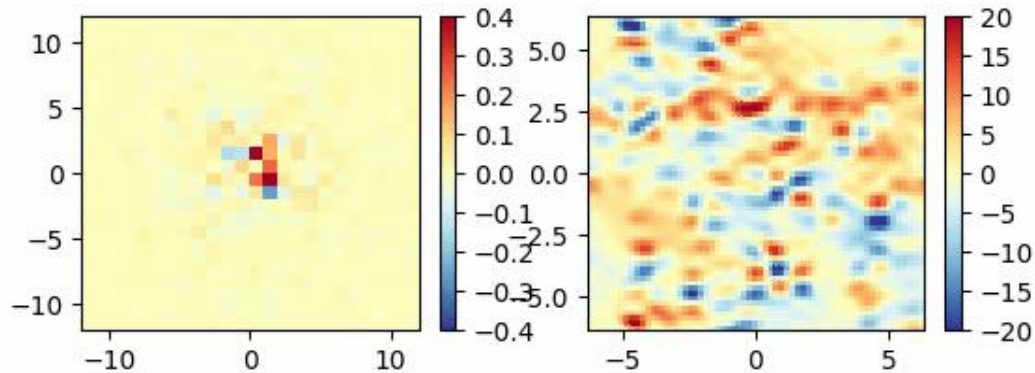


Getting Curvature

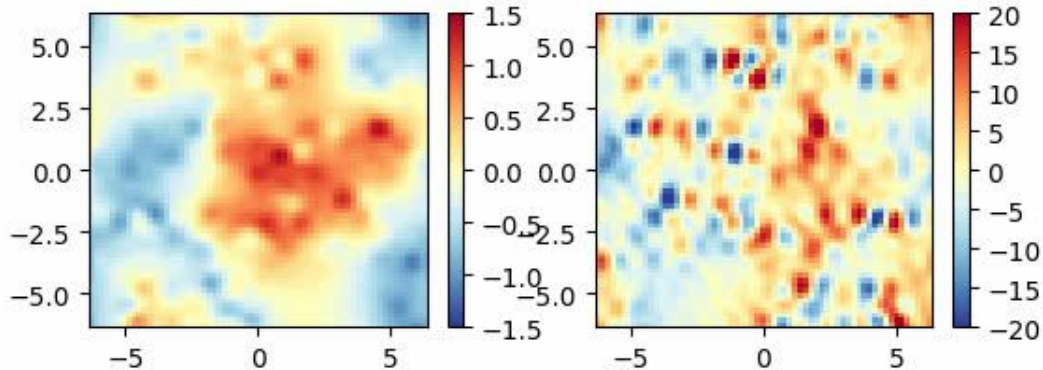
- Tensor product surface $\xi \approx \sum_{\mathbf{k} < k_u} \hat{\xi}_{\mathbf{k}}(t) \xi_u(y) \xi_v(z),$

$$\xi_a(b) = \begin{cases} \cos(2\pi ab) & \text{if } a > 0 \\ \sin(2\pi|a|b) & \text{otherwise,} \end{cases}$$

$\hat{\xi}_{\mathbf{k}}(t)$

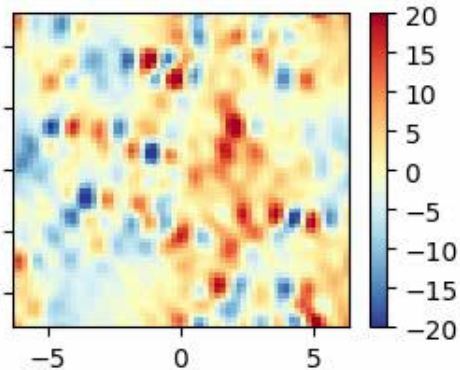
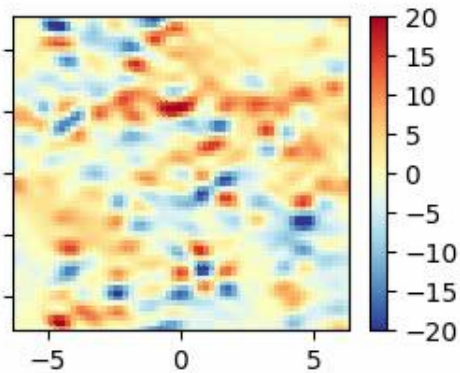


ξ



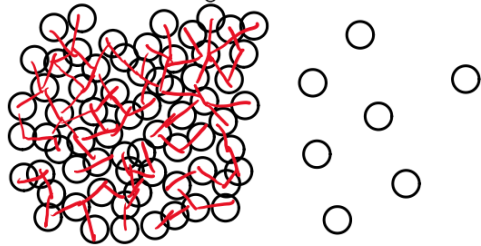
$\frac{\partial \xi}{\partial y}$

$\frac{\partial \xi}{\partial z}$

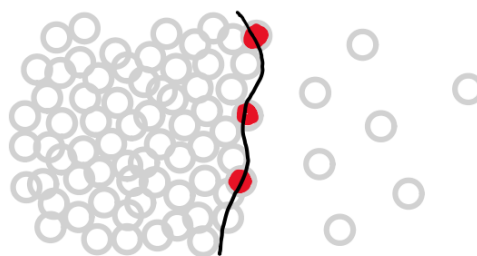


Moving Reference Frame - Overview

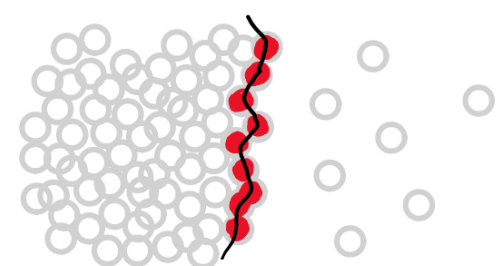
a) Cluster analysis



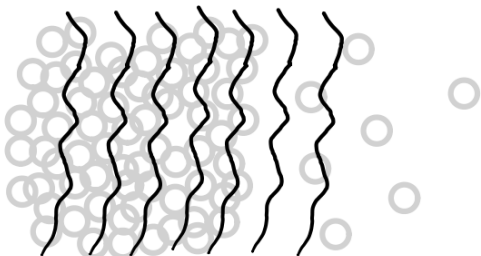
b) Fit outer molecules



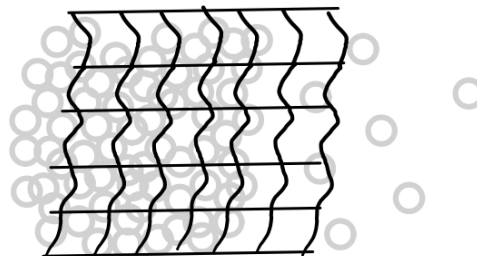
c) Refit closest molecules



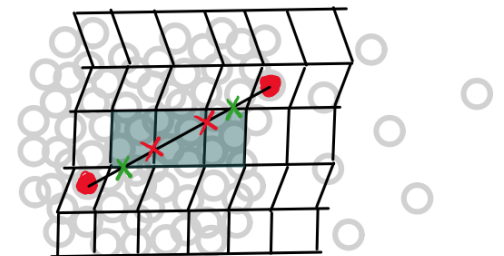
d) Intrinsic interface layers



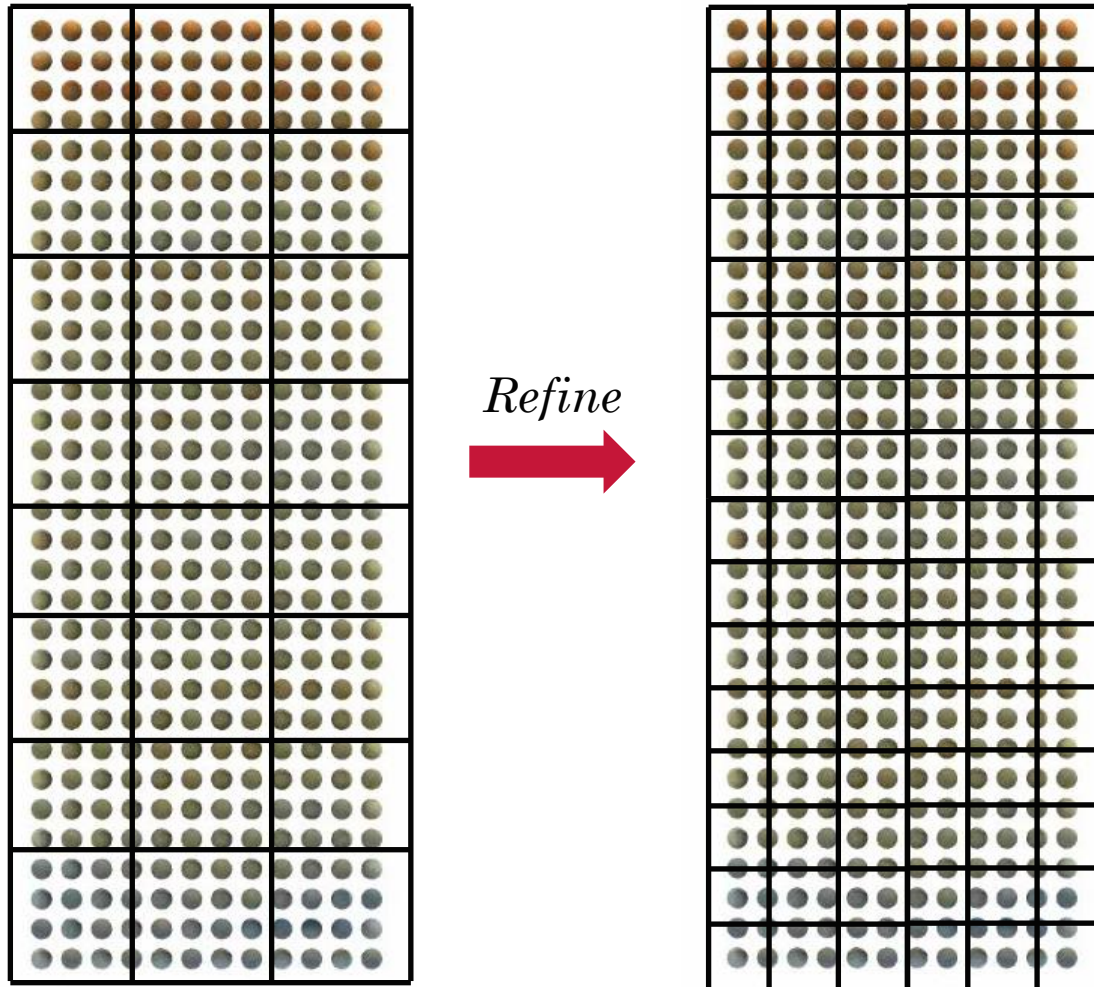
e) Create grid



f) Bilinear conversion



Molecular Dynamics - Averaging



- Density in a cell

$$\rho = \frac{1}{V} \sum_{i=1}^N \langle m_i \rangle$$

- Momentum in a cell

$$\rho \mathbf{u} = \frac{1}{V} \sum_{i=1}^N \langle m_i \mathbf{v}_i \rangle$$

- Temperature in a cell

$$T = \frac{1}{3N} \sum_{i=1}^N \langle \mathbf{v}_i^2 \rangle$$

Irving and Kirkwood (1950)



Brunel University London

TRANSPORT PROCESSES 817

777.9 cm⁻¹. The band at 809.7 cm⁻¹ has been interpreted as 2N₂, the rather large difference between calculated and observed frequency being ascribed to the proximity of the A₁ fundamental. However, this band could be an upper-stage band corresponding to 769.4 cm⁻¹. The interpretation of the faint sharp band at 838.7 cm⁻¹ as an upper-stage band, 778.4 cm⁻¹ seems more plausible than its interpretation as the R₁₂ combination band, 2104-558 cm⁻¹. The faint sharp band at 838.7 cm⁻¹ has been interpreted as an upper-stage band, 2x 407.4 cm⁻¹, rather than as the R₁₂ difference band, 1561-510 cm⁻¹, because the corresponding sum band

The Statistical Mechanical Theory of Transport Processes. IV. The Equations of Hydrodynamics

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 Gases and Certain Laboratories of Chemistry No. 1243, Pasadena, California
 (Received November 11, 1949)

The equations of hydrodynamics—continuity equation of number, equation of energy, and equation of momentum—are derived by means of the classical statistical mechanics. They are obtained for the system as a whole and not for individual molecules. In addition to the familiar terms arising from the stress tensor and heat current density are represented, respectively, as quadratures of the density and current density in the configuration space of a pair of molecules.

INTRODUCTION

THIS paper will be concerned with a derivation of the equations of hydrodynamics from the principles of the classical statistical mechanics. In particular, the separation of energy transport will be treated, and the equation of energy transport will be derived. By so doing, the stress tensor and heat current density can be expressed in terms of a kinetic part (which occurs in the kinetic theory of gases) and another term (which occurs in the kinetic theory of internal motion) involving the potential of intermolecular forces and the density and current density in and a quadrature involving the potential of intermolecular forces and the density and current density in the configuration space of a pair of molecules.

This program is amenable to a liquid, various attempts have been made to obtain a closed equation satisfied approximately by the probability distribution function in the phase space of a pair of molecules. One such attempt, known as the "superposition" assumption, has been used by Kirkwood and his co-workers. Another, a generalization of the well-known Fokker-Planck equation of stochastic theory, has been derived by Kirkwood by introducing the concepts of stress tensor and a friction constant. This latter equation has been used to obtain an expression for the heat current, and applied to problems of fluid viscosity and thermal expansion (in terms of molecular variables) for colloidal systems and nonuniform temperature, pressure, and density. The latter equation leads to equations governing the heat current and thereby an explicit expression for the coefficient of thermal expansion in an expression for the coefficient of thermal expansion. We shall assume, for purposes of simplicity, a single component, single phase fluid system consisting of molecules which interact under central forces only. It is not difficult to generalize the treatment to a multiple component or multiphase system.

* This work was supported by the U. S. ONR under Contract No. 49(52)-1-241-10. The authors are indebted to J. G. Kirkwood, J. Chem. Phys. 14, 30 (1946).

TRANSPORT PROCESSES 819

The probability distribution function (relative density of representative points in phase space) we denote by

$$\rho(\mathbf{R}_1, \dots, \mathbf{R}_N; \mathbf{p}_1, \dots, \mathbf{p}_N; t)$$

satisfying the normalization condition

$$\int \dots \int \rho(\mathbf{R}_1, \dots, \mathbf{R}_N; \mathbf{p}_1, \dots, \mathbf{p}_N; t) d\mathbf{R}_1 \dots d\mathbf{R}_N d\mathbf{p}_1 \dots d\mathbf{p}_N = 1$$

where $d\mathbf{R}_i$ stands for a volume element in the configuration space of the i th molecule, $d\mathbf{p}_i$ a volume element in the momentum space of the i th molecule. Changes in time according to the well-known Liouville equation

$$\frac{d\rho}{dt} + \sum_{i=1}^N \left[\frac{\partial \rho}{\partial \mathbf{p}_i} \cdot \frac{\partial U}{\partial \mathbf{R}_i} + \frac{\partial \rho}{\partial \mathbf{R}_i} \cdot \left(\frac{\partial U}{\partial \mathbf{p}_i} + \mathbf{V}_i \right) \right] = 0$$

where U is the potential energy of the entire system. Any dynamical variable, $\alpha(\mathbf{R}_1, \dots, \mathbf{R}_N; \mathbf{p}_1, \dots, \mathbf{p}_N; t)$, has an expectation value given at time t by

$$\langle \alpha \rangle = \int \dots \int \alpha(\mathbf{R}_1, \dots, \mathbf{R}_N; \mathbf{p}_1, \dots, \mathbf{p}_N; t) \rho(\mathbf{R}_1, \dots, \mathbf{R}_N; \mathbf{p}_1, \dots, \mathbf{p}_N; t) d\mathbf{R}_1 \dots d\mathbf{R}_N d\mathbf{p}_1 \dots d\mathbf{p}_N$$

We shall denote by $\langle \alpha \rangle_t$ the expectation value of α at time t . The product of α and f taken over phase space. Providing α does not depend on time explicitly, the rate of change of the expectation value of α is given by

$$\frac{d\langle \alpha \rangle}{dt} = \langle \dot{\alpha} \rangle = \sum_{i=1}^N \left[\left\langle \frac{\partial \alpha}{\partial \mathbf{p}_i} \cdot \frac{\partial U}{\partial \mathbf{R}_i} + \frac{\partial \alpha}{\partial \mathbf{R}_i} \cdot \left(\frac{\partial U}{\partial \mathbf{p}_i} + \mathbf{V}_i \right) \right\rangle \right]$$

By Green's theorem applied in the space of \mathbf{R}_i

$$\left\langle \frac{\partial \alpha}{\partial \mathbf{R}_i} \cdot \left(\frac{\partial U}{\partial \mathbf{p}_i} + \mathbf{V}_i \right) \right\rangle = \int \dots \int \frac{\partial \alpha}{\partial \mathbf{R}_i} \cdot \left(\frac{\partial U}{\partial \mathbf{p}_i} + \mathbf{V}_i \right) \rho d\mathbf{R}_1 \dots d\mathbf{R}_N d\mathbf{p}_1 \dots d\mathbf{p}_N$$

providing the integrated part vanishes at infinity, as $\mathbf{R}_i \rightarrow \infty$. Likewise, since \mathbf{V}_i is independent of \mathbf{p}_i , the momentum space of \mathbf{p}_i is independent of \mathbf{R}_i , and since ρ falls off rapidly as $\mathbf{p}_i \rightarrow \infty$, Green's theorem in the momentum space of \mathbf{p}_i yields

$$\left\langle \frac{\partial \alpha}{\partial \mathbf{p}_i} \cdot \frac{\partial U}{\partial \mathbf{R}_i} \right\rangle = - \int \dots \int \frac{\partial \alpha}{\partial \mathbf{R}_i} \cdot \mathbf{V}_i \rho d\mathbf{R}_1 \dots d\mathbf{R}_N d\mathbf{p}_1 \dots d\mathbf{p}_N$$

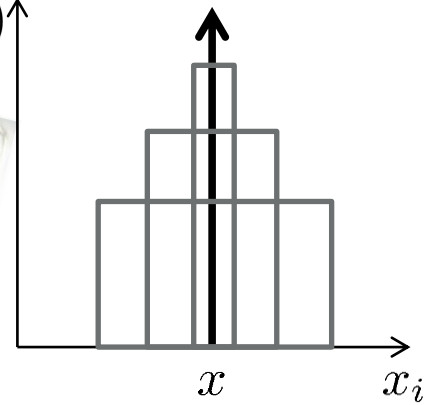
Thus, (2.6) becomes

$$\frac{d\langle \alpha \rangle}{dt} = \sum_{i=1}^N \int \dots \int \frac{\partial \alpha}{\partial \mathbf{R}_i} \cdot \mathbf{V}_i \rho d\mathbf{R}_1 \dots d\mathbf{R}_N d\mathbf{p}_1 \dots d\mathbf{p}_N$$

giving the rate of change of the expectation value of α

Consequently, $\langle \rho(\mathbf{R}_1, \dots, \mathbf{R}_N; \mathbf{p}_1, \dots, \mathbf{p}_N; t) \rangle$ is the product of this of the i th molecule to be at \mathbf{r}_i , i.e., it is the contribution (mass current density). The total momentum density

$$\delta(x - x_i) \uparrow$$



The Dirac delta infinitely high, infinitely thin peak formally equivalent to the continuum differential formulation

$$\rho(\mathbf{r}, t) = \sum_{i=1}^N \left\langle m_i \delta(\mathbf{r} - \mathbf{r}_i); f \right\rangle$$

The Control Volume Functional

- The Control volume functional is the formal integral of the Dirac delta functional in 3 dimensions (3D top hat or box car function)

$$\vartheta_i \equiv \int_{x^-}^{x^+} \int_{y^-}^{y^+} \int_{z^-}^{z^+} \delta(x_i - x) \delta(y_i - y) \delta(z_i - z) dx dy dz$$

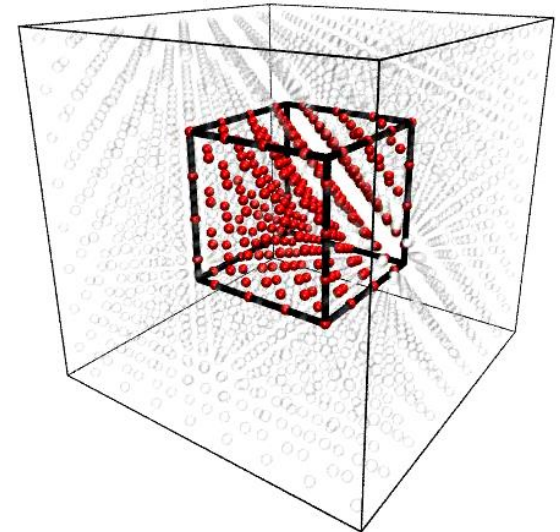
$$= [H(x^+ - x_i) - H(x^- - x_i)]$$

$$\times [H(y^+ - y_i) - H(y^- - y_i)]$$

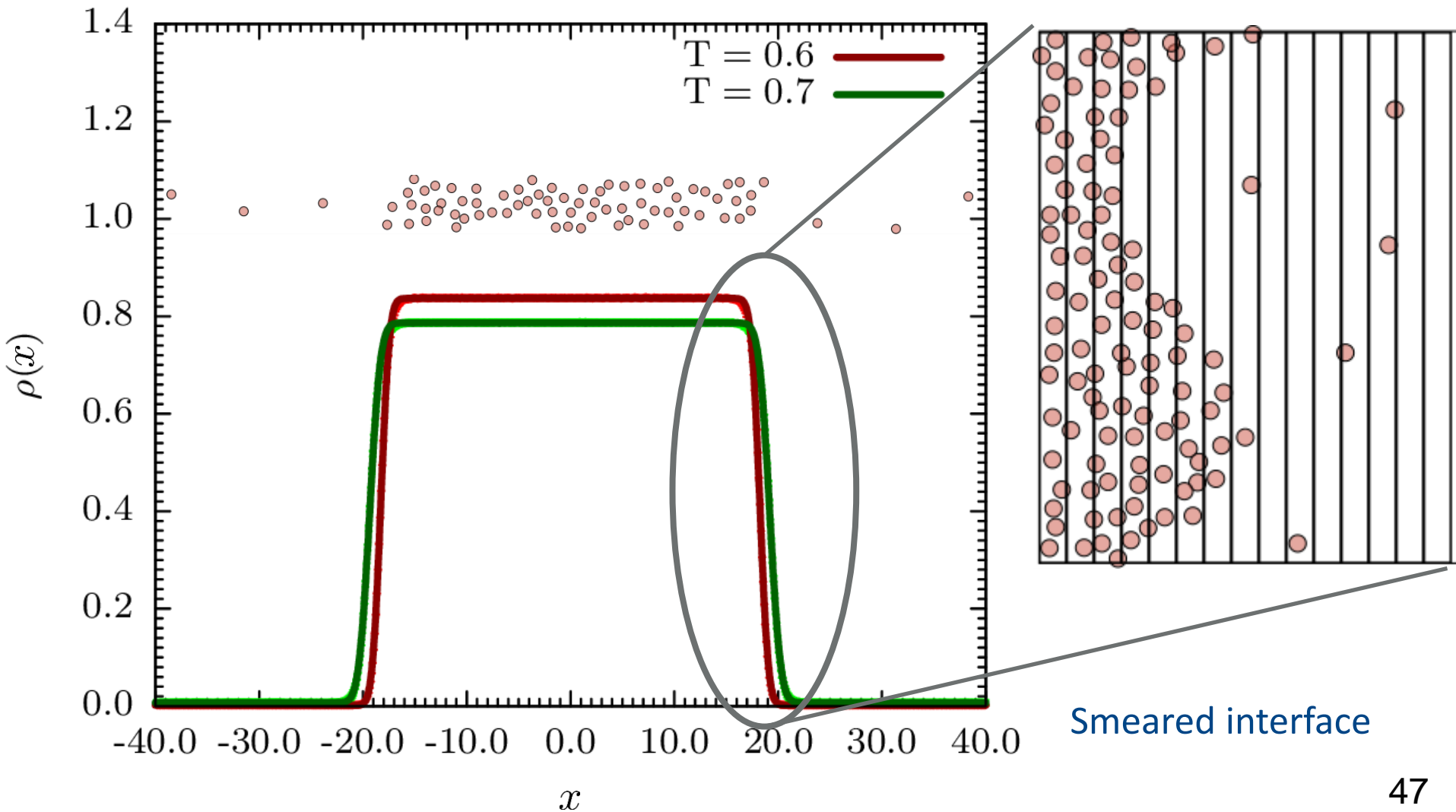
$$\times [H(z^+ - z_i) - H(z^- - z_i)]$$

- In words

$$\vartheta \equiv \begin{cases} 1 & \text{if molecule is inside volume} \\ 0 & \text{if molecule is outside volume} \end{cases}$$



Density



A more general Control Volume Functional

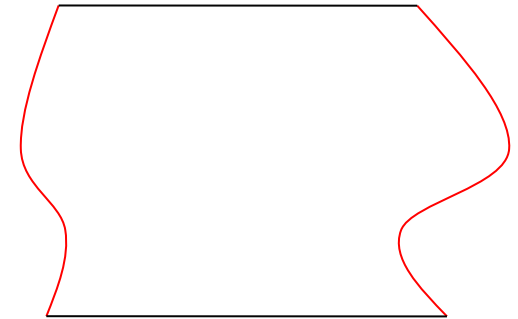
- The Control volume functional is the formal integral of the Dirac delta functional in 3 dimensions (3D top hat or box car function)

$$\vartheta_i \equiv \int_{z^-}^{z^+} \int_{y^-}^{y^+} \int_{x^- + \xi^-(y,z,t)}^{x^+ + \xi^+(y,z,t)} \delta(x - x_i) \delta(y - y_i) \delta(z - z_i) dx dy dz$$

$$= [H(x^+ + \xi^+ - x_i) - H(x^- + \xi^- - x_i)]$$

$$\times [H(y^+ - y_i) - H(y^- - y_i)]$$

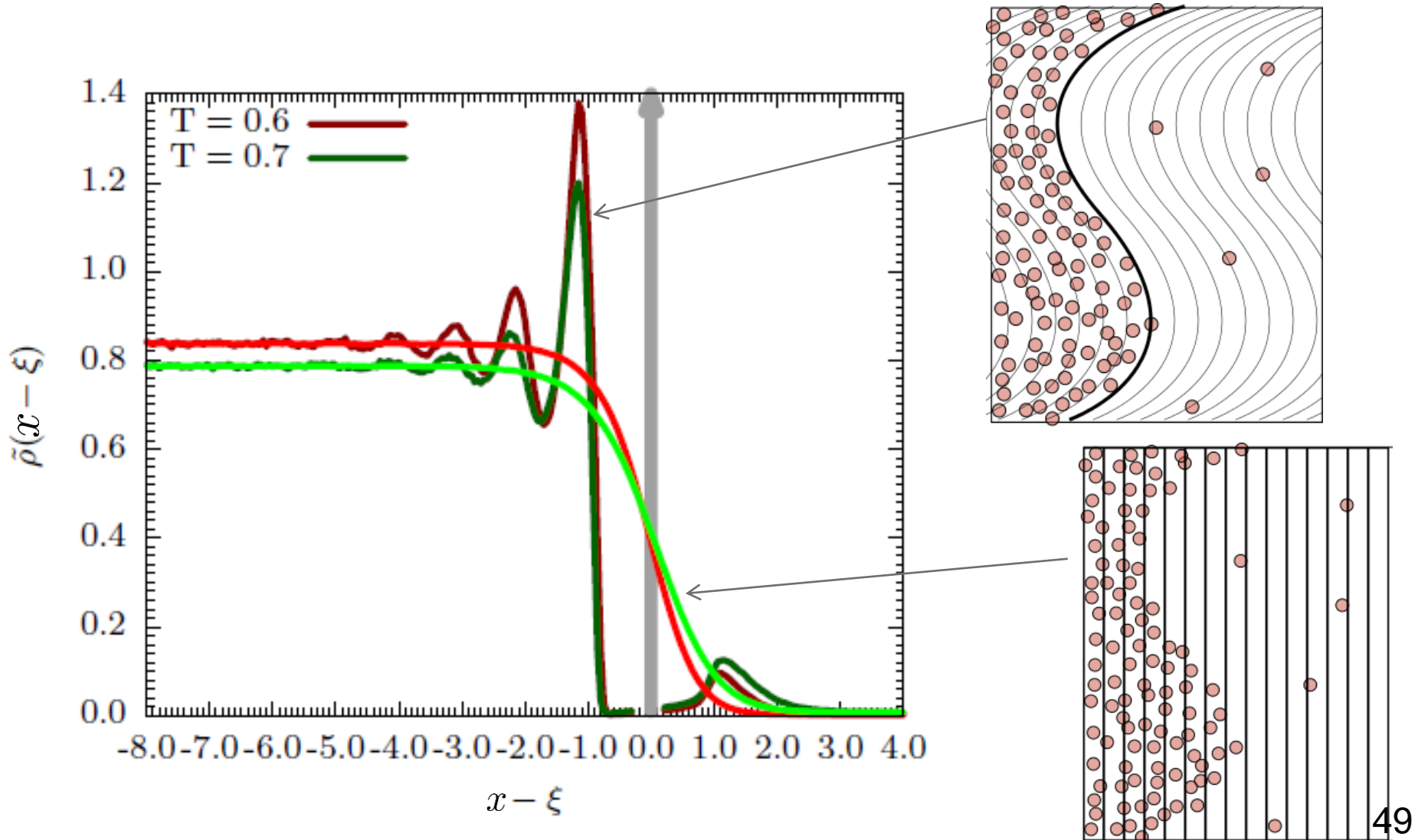
$$\times [H(z^+ - z_i) - H(z^- - z_i)]$$



- In words

$$\vartheta \equiv \begin{cases} 1 & \text{if molecule is inside the moving volume} \\ 0 & \text{if molecule is outside the moving volume} \end{cases}$$

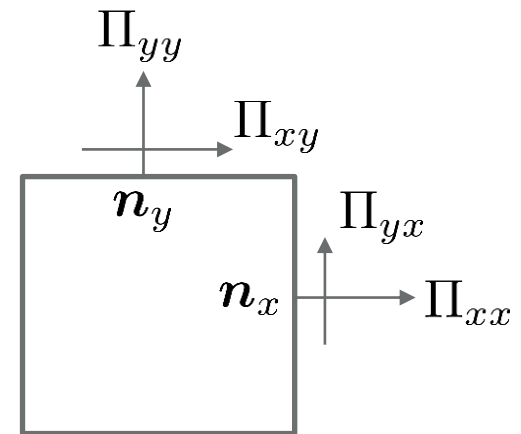
The Intrinsic Interface



Pressure (Stress)

- We need pressure to understand the surface
 - Using Cauchy's original definition
 - In two dimensions the tensor is then,

$$\mathbf{\Pi} = \begin{bmatrix} \Pi_{xx} & \Pi_{xy} \\ \Pi_{yx} & \Pi_{yy} \end{bmatrix}$$



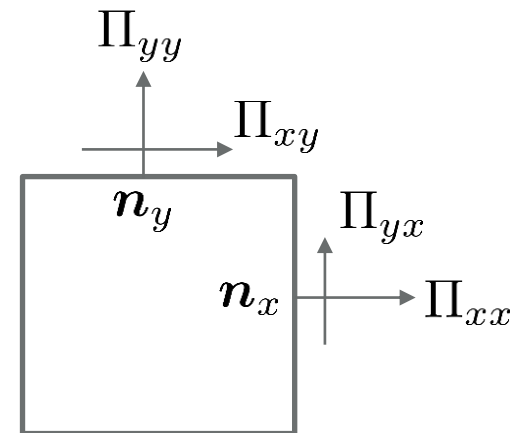
- Where pressure in molecular systems include kinetic and configurational

$$\oint_S \Pi_{xx} \cdot dS_x = \underbrace{\sum_{i=1}^N \left\langle m_i v_{xi} v_{xi} dS_{xi} \right\rangle}_{\text{Kinetic}} + \underbrace{\frac{1}{2} \sum_{i=1}^N \sum_{j \neq i}^N \left\langle f_{xij} dS_{xij} \right\rangle}_{\text{Configurational}}$$

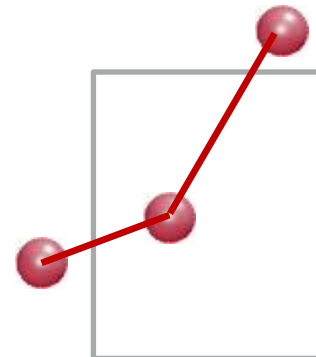
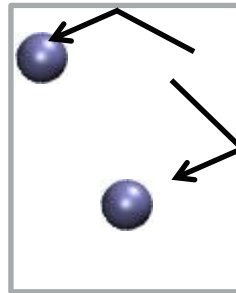
Pressure (Stress)

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$$\mathbf{\Pi} = \begin{bmatrix} \Pi_{xx} & \Pi_{xy} \\ \Pi_{yx} & \Pi_{yy} \end{bmatrix}$$



*Kinetic
theory part
Momentum due
to average of
molecules
crossing a plane
and returning*



*Configurational
part
Inter-molecular
bonds act like the
stress in a
stretched spring*

Derivative for Fluxes and Stresses

- Taking the Derivative of the CV function

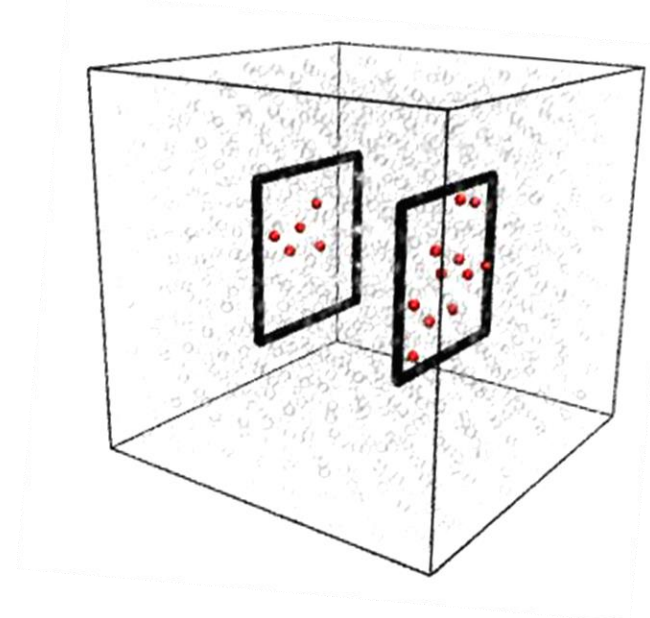
$$dS_{ix} \equiv -\frac{\partial \vartheta_i}{\partial x_i} = [\delta(x^+ - x_i) - \delta(x^- - x_i)] \\ \times [H(y^+ - y_i) - H(y^- - y_i)] \\ \times [H(z^+ - z_i) - H(z^- - z_i)]$$

- Vector form defines six surfaces

$$d\mathbf{S}_i = \mathbf{i}dS_{xi} + \mathbf{j}dS_{yi} + \mathbf{k}dS_{zi}$$

- Or in words

$$d\mathbf{S}_i \equiv \begin{cases} \infty & \text{if molecule on surface} \\ 0 & \text{otherwise} \end{cases}$$



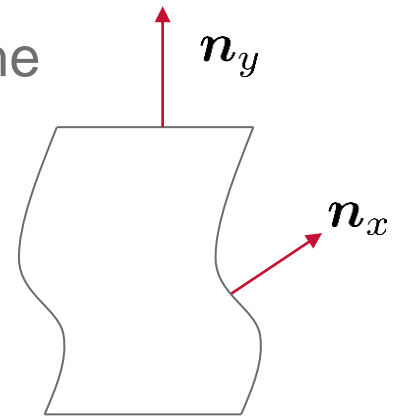
Pressure Components (normal form)

- Working through the mathematics, a form in terms of the unit vectors

$$\Pi_{\alpha}^{\text{kinetic}} = \frac{1}{\Delta S_{\alpha}} \sum_{i=1}^N m_i \dot{\mathbf{r}}_i \mathbf{r}_{i2} \cdot \frac{\mathbf{n}_{\alpha}}{|\mathbf{r}_{i2} \cdot \mathbf{n}_{\alpha}|} dS_{\alpha}$$

$$\Pi_{\alpha}^{\text{Moving surf}} = \frac{1}{\Delta S_{\alpha}} \sum_{i=1}^N m_i \dot{\mathbf{r}}_i \vartheta_t$$

$$\Pi_{\alpha}^{\text{config}} = \frac{1}{2\Delta S_{\alpha}} \sum_{i,j} \mathbf{f}_{ij} \mathbf{r}_{ij} \cdot \frac{\mathbf{n}_{\alpha}}{|\mathbf{r}_{ij} \cdot \mathbf{n}_{\alpha}|} dS_{\alpha}$$



$$\mathbf{n}_{\alpha} = \frac{\nabla_{\alpha}(\xi - z_{\alpha})}{\|\nabla_{\alpha}(\xi - z_{\alpha})\|}$$

- Calculating pressure on a surface:
 - Obtain a crossing on a surface (a moving molecule or a force)
 - Dot this with the unit vector to the surface

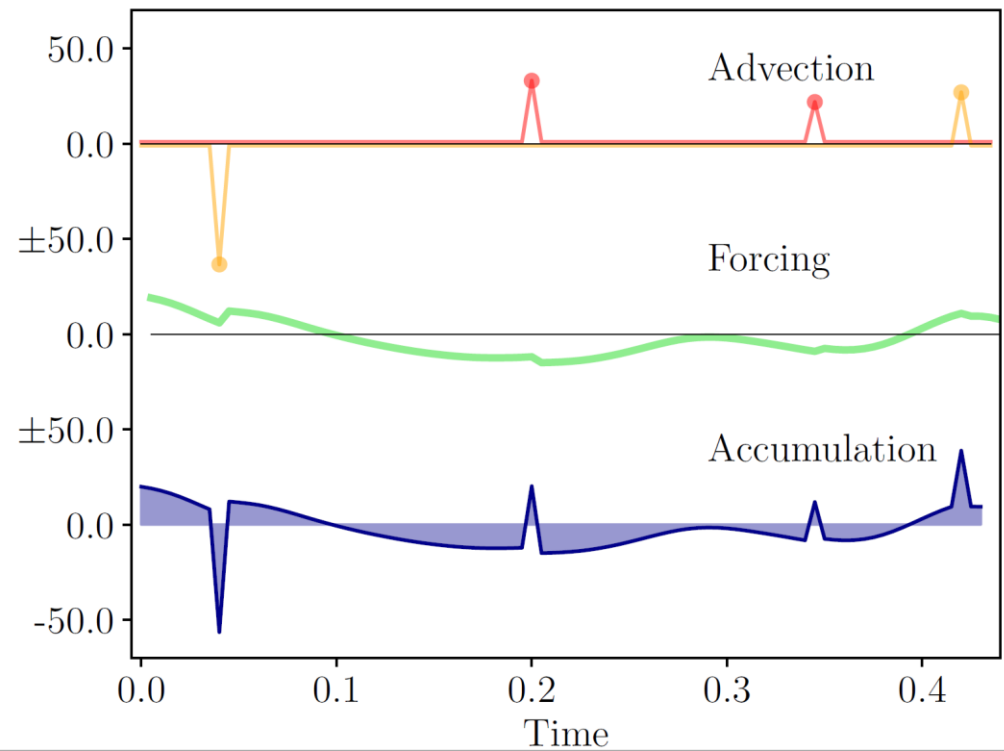
CV Balance Per Bilinear Cube

$$\text{Advection} = \frac{1}{\Delta S_z} \sum_{i=1}^N m_i \dot{\mathbf{r}}_i \frac{\mathbf{r}_{12} \cdot \mathbf{n}_z}{|\mathbf{r}_{12} \cdot \mathbf{n}_z|} dS^+ + \frac{1}{\Delta S_z} \sum_{i=1}^N m_i \dot{\mathbf{r}}_i \vartheta_t$$

$$\text{Forcing} = \frac{1}{2\Delta S_z} \sum_{i,j} \mathbf{f}_{ij} \frac{\mathbf{r}_{ij} \cdot \mathbf{n}_z}{|\mathbf{r}_{ij} \cdot \mathbf{n}_z|} dS^+$$

$$\text{Accumulation} = \frac{d}{dt} \sum_{i=1}^N m_i \dot{\mathbf{r}}_i \vartheta_i$$

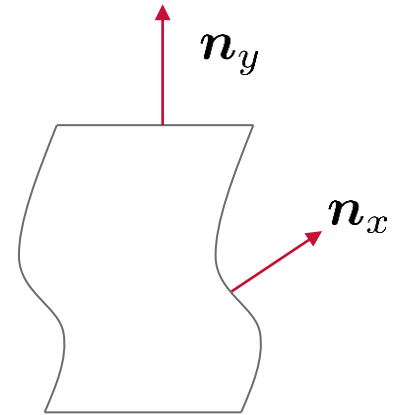
$$\text{Accumulation} = \text{Advection} + \text{Forcing}$$



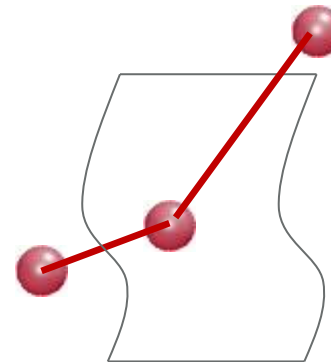
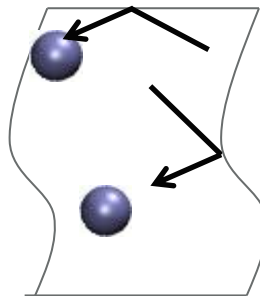
Pressure (Stress)

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 - Using Cauchy's original definition
 - In two dimensions the tensor is,

$$\mathbf{\Pi} = \begin{bmatrix} \Pi_{xx} & \Pi_{xy} \\ \Pi_{yx} & \Pi_{yy} \end{bmatrix} = \begin{bmatrix} \Pi_N & 0 \\ 0 & \Pi_T \end{bmatrix}$$

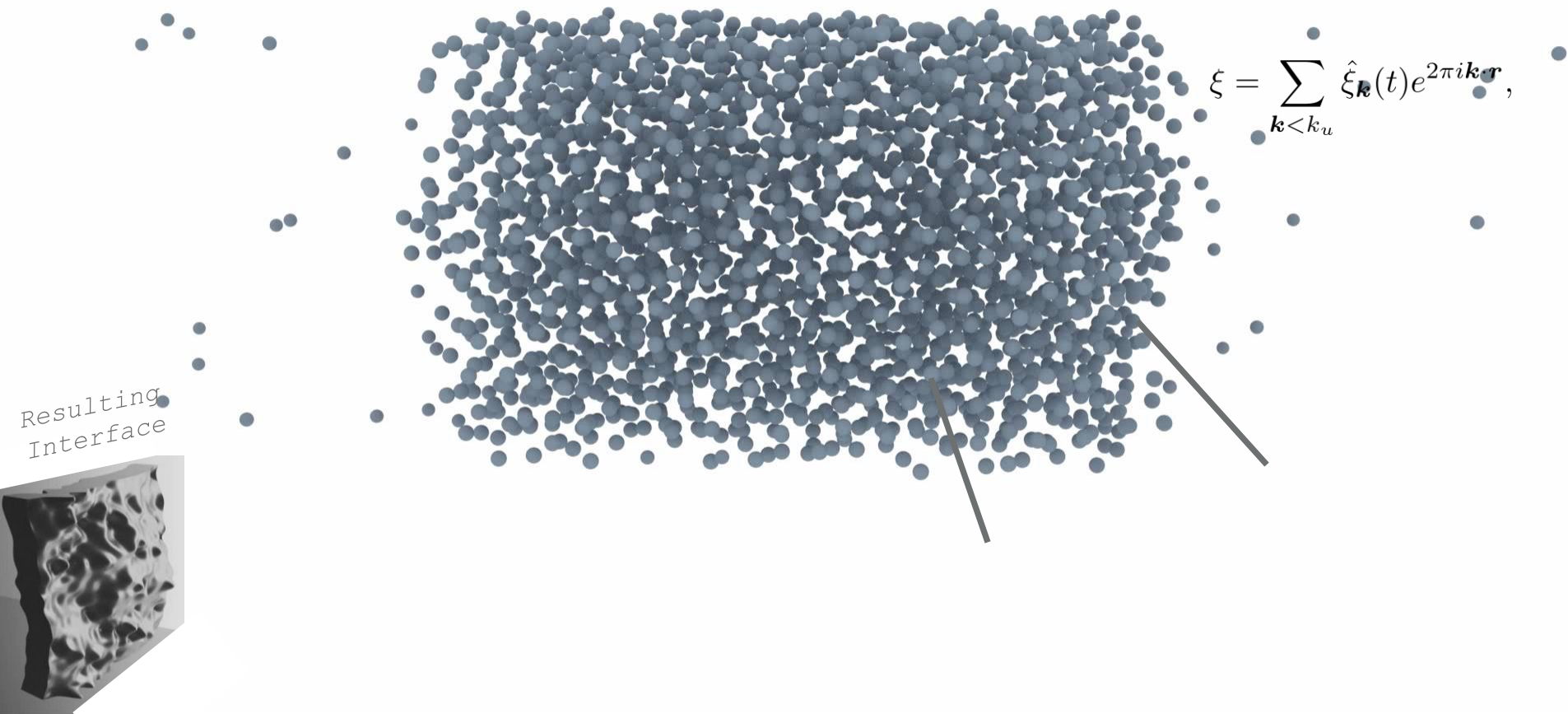


*Kinetic
theory part
Momentum due
to average of
molecules
crossing a plane
and returning*



*Configurational
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Inter-molecular
bonds act like the
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stretched spring*

Liquid-Vapour interface

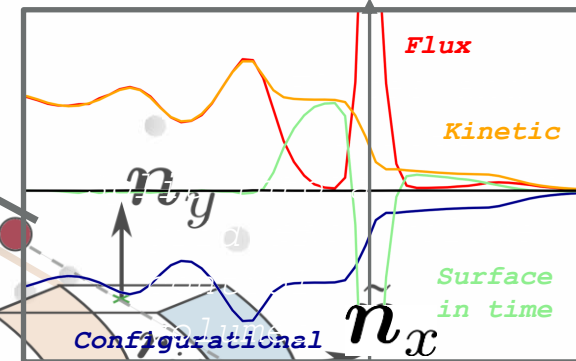




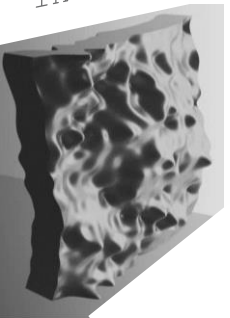
Liquid-Vapour interface

Time evolving
Fourier
surface
● fitted

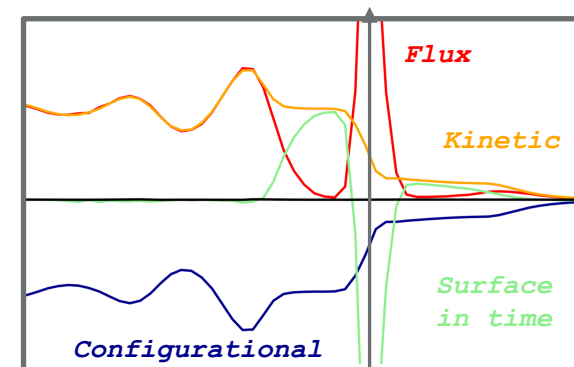
$$\xi = \sum_{\mathbf{k} < k_u} \hat{\xi}_{\mathbf{k}}(t) e^{2\pi i \mathbf{k} \cdot \mathbf{r}},$$



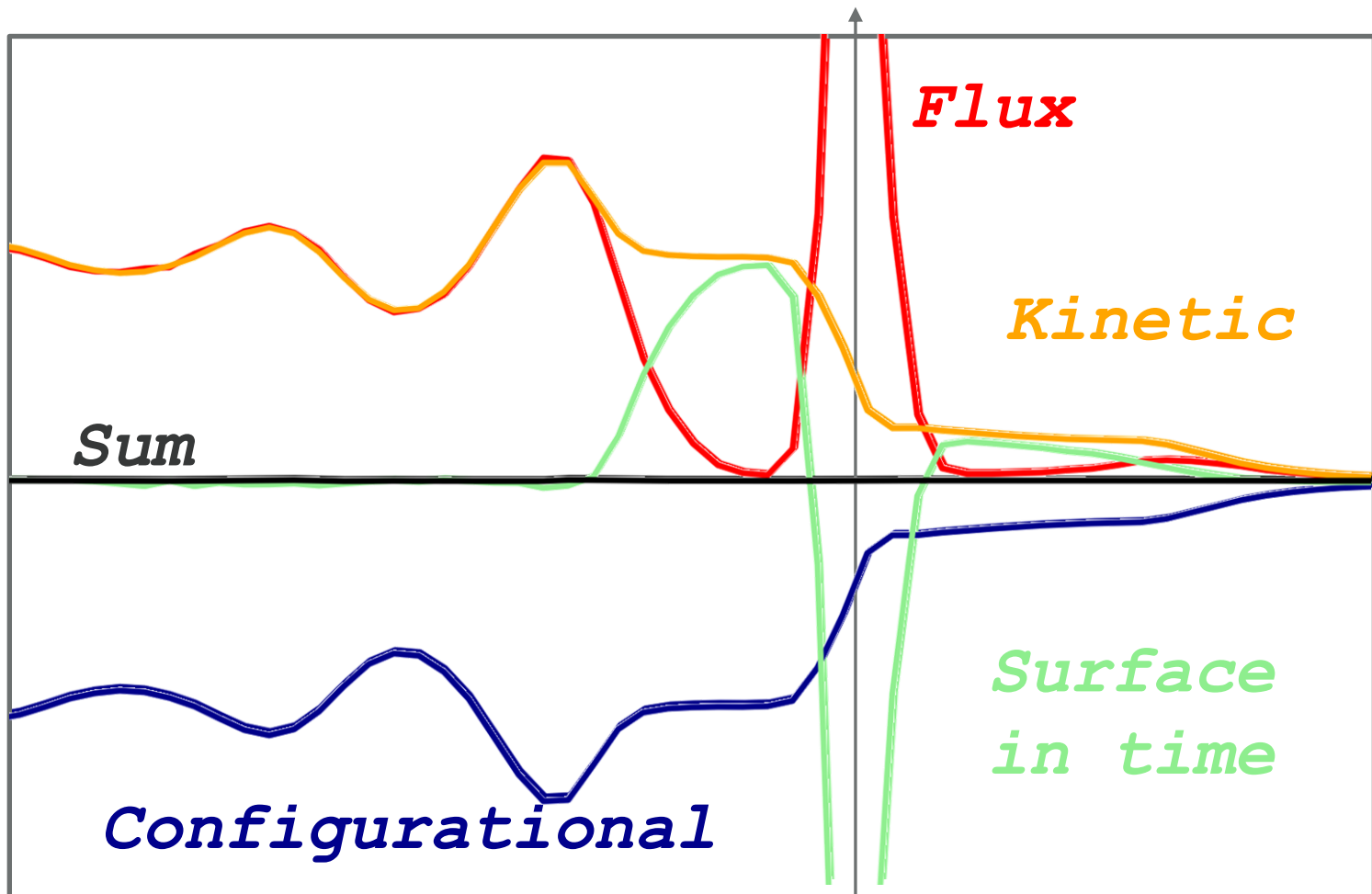
Resulting
Interface



Normal Pressure

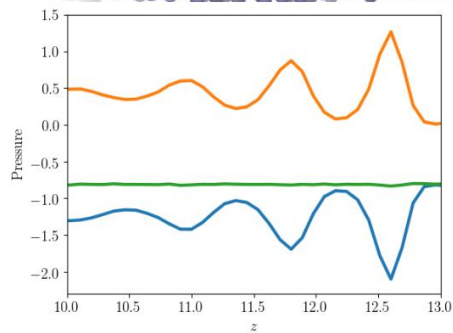
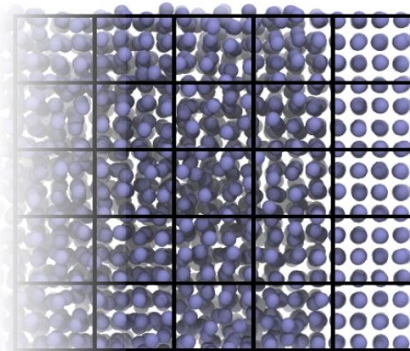


Normal Pressure

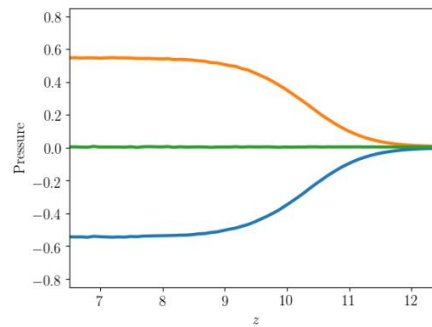
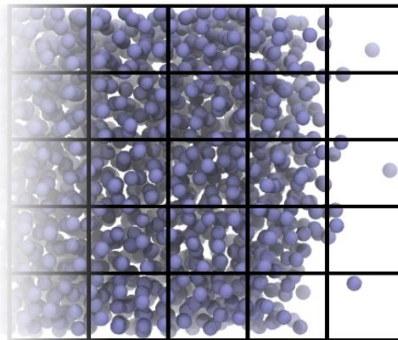


Normal Pressure is Flat (equilibrium)

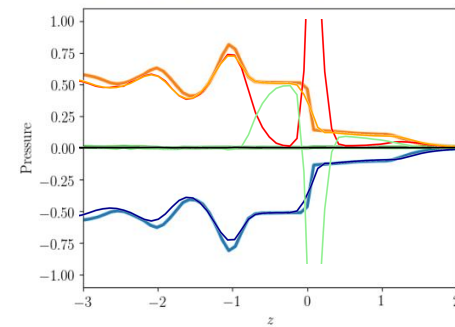
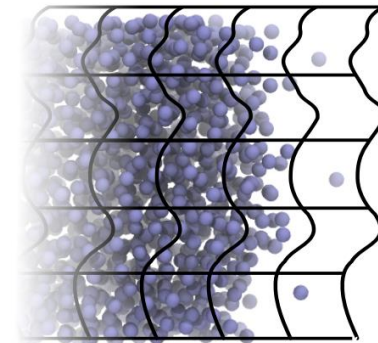
Solid-liquid
Eulerian
reference



Liquid-vapour
Eulerian
reference



Liquid-vapour
Lagrangian
reference



Other Pressure Definitions

- Only surface pressures satisfy this balance correctly

- Volume Average (VA) Pressure ●

$$\bar{\Pi}^{VA} = \frac{1}{\Delta V} \left[\sum_{i=1}^N \frac{\mathbf{p}_i \mathbf{p}_i}{m_i} \vartheta_i + \frac{1}{2} \sum_{i,j} \mathbf{f}_{ij} \mathbf{r}_{ij} \int_0^1 \vartheta_\lambda d\lambda \right]$$

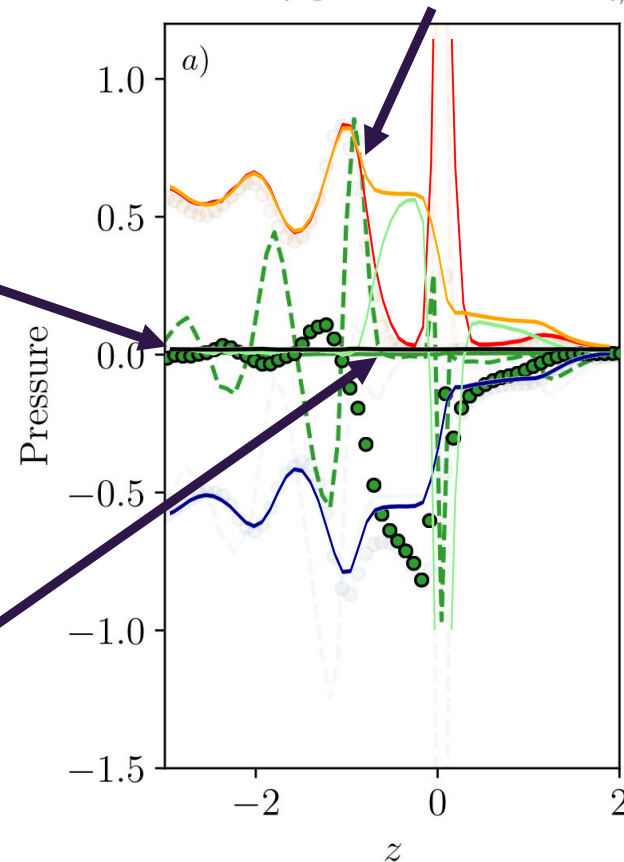
- Surface Pressure (Generalised Method of Planes) —

$$\int_{t_1}^{t_2} \rho \mathbf{u} u_z + \bar{\Pi}_z^{\text{SF}} dt = \frac{1}{\Delta S_z} \sum_{i=1}^N m_i \dot{\mathbf{r}}_i \frac{\mathbf{r}_{12} \cdot \mathbf{n}_z}{|\mathbf{r}_{12} \cdot \mathbf{n}_z|} dS^+ + \frac{1}{\Delta S_z} \sum_{i=1}^N m_i \dot{\mathbf{r}}_i \vartheta_t$$

$$\bar{\Pi}_z^{\text{SF}c} = \frac{1}{2\Delta S_z} \sum_{i,j} \mathbf{f}_{ij} \frac{\mathbf{r}_{ij} \cdot \mathbf{n}_z}{|\mathbf{r}_{ij} \cdot \mathbf{n}_z|} dS^+$$

- Irving Kirkwood 1 (IK1) or Virial Pressure

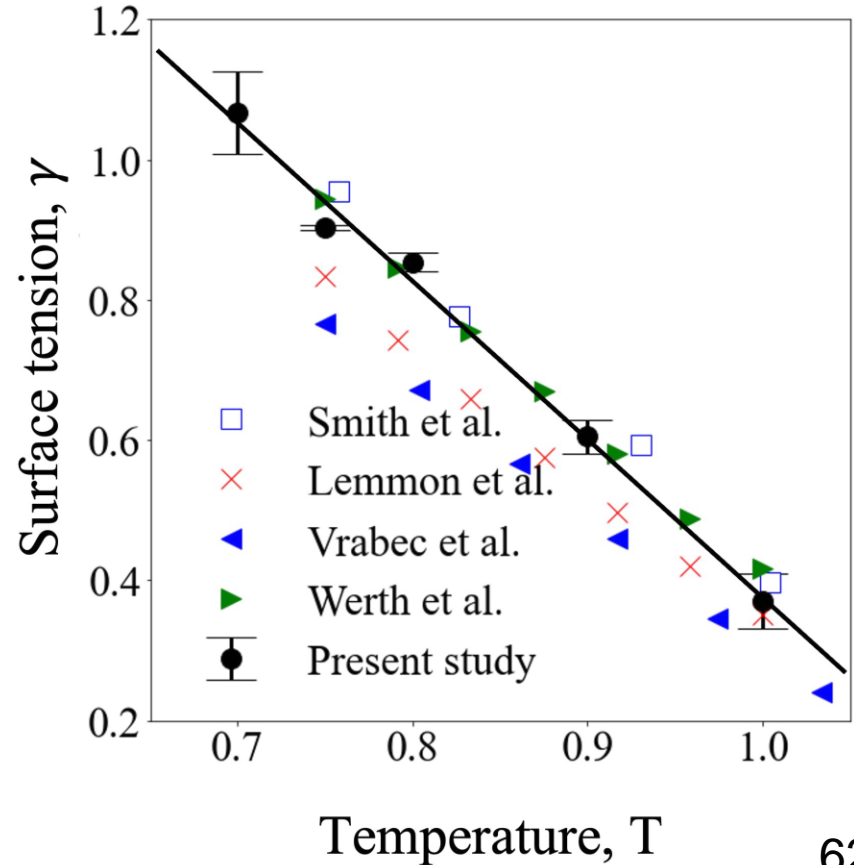
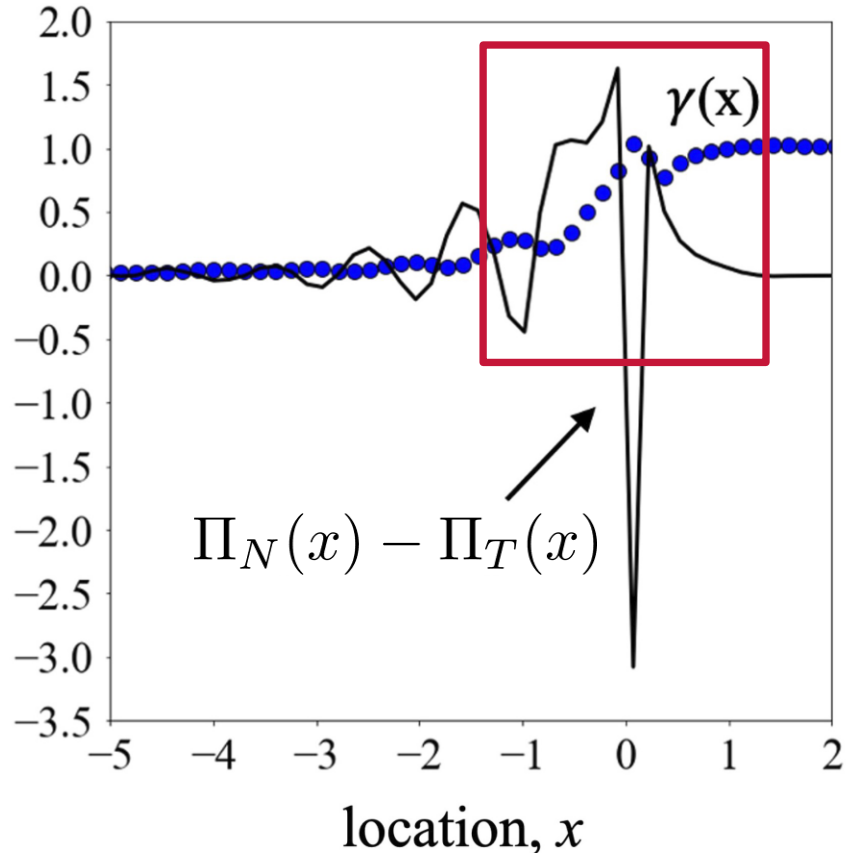
$$\bar{\Pi}^{IK1}(\mathbf{r}, t) = \sum_{i=1}^N \frac{\mathbf{p}_i \mathbf{p}_i}{m_i} \delta(\mathbf{r} - \mathbf{r}_i) + \frac{1}{2} \sum_{i,j} \mathbf{f}_{ij} \mathbf{r}_{ij} \delta(\mathbf{r} - \mathbf{r}_i).$$



Tangential Pressure and Surface Tension

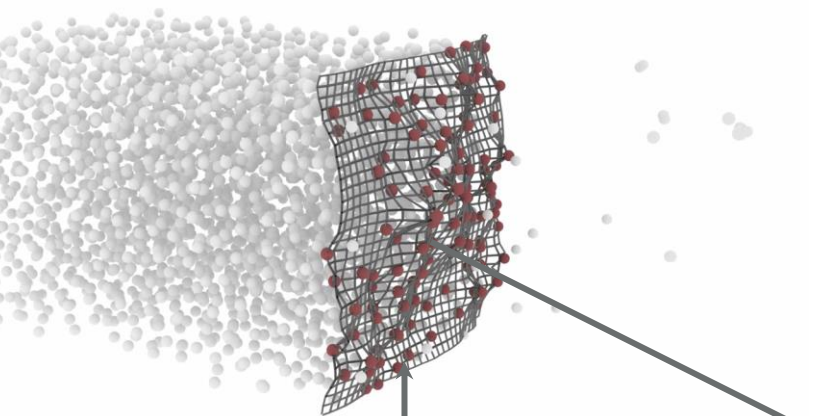
- Contribution to surface tension localised to very thin region at interface

$$\gamma = \int_{-2^{\frac{1}{6}}}^{2^{\frac{1}{6}}} [\Pi_N - \Pi_T] dx$$

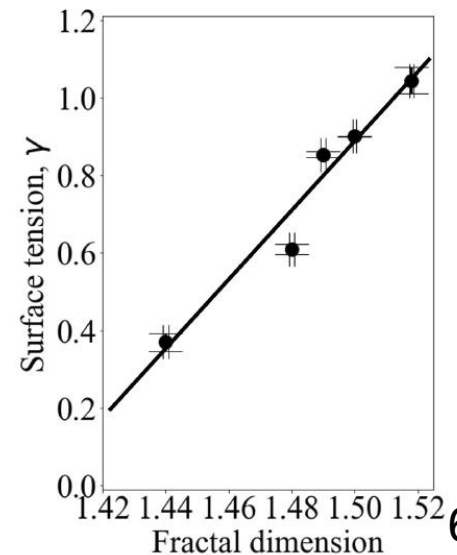
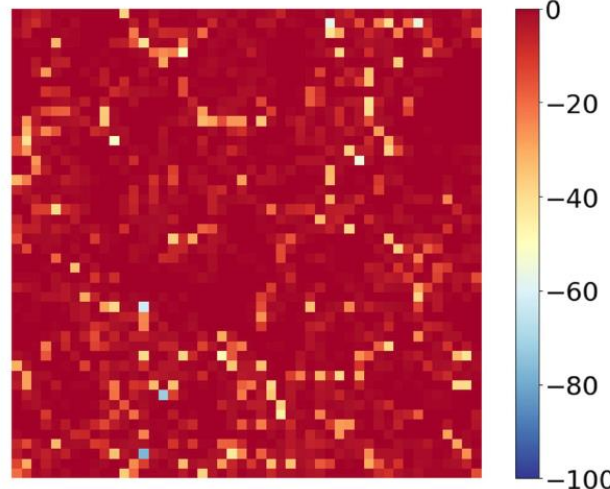
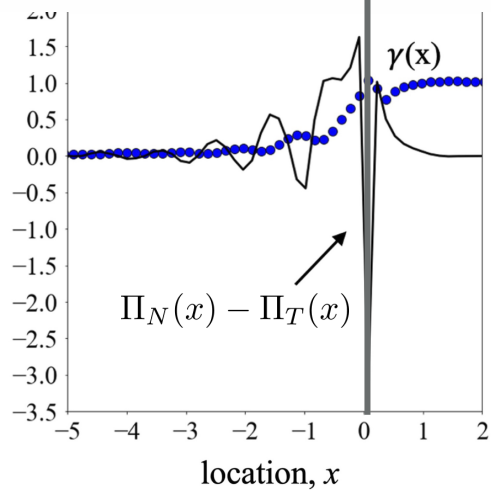


Stress Networks

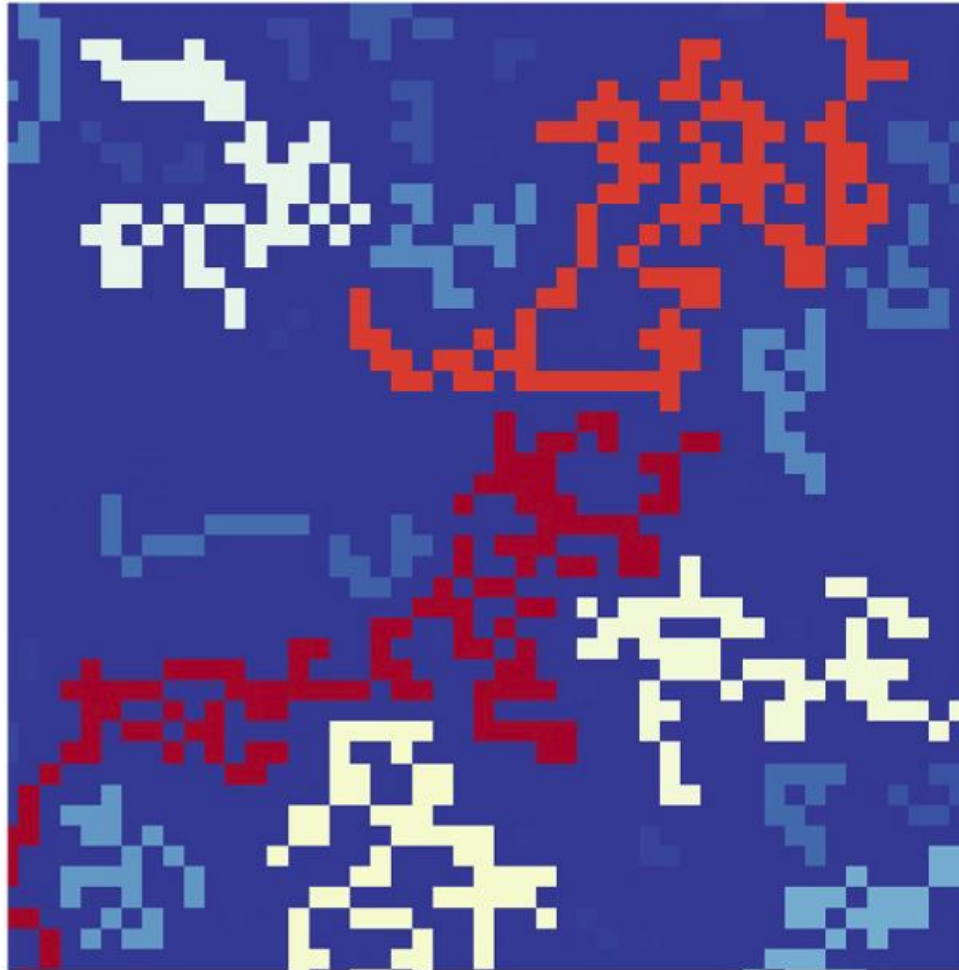
- Taking the instant Tangential Pressures on the intrinsic interface itself



Work by
**Muhammad
Rizwanur Rahman**
at
Imperial College



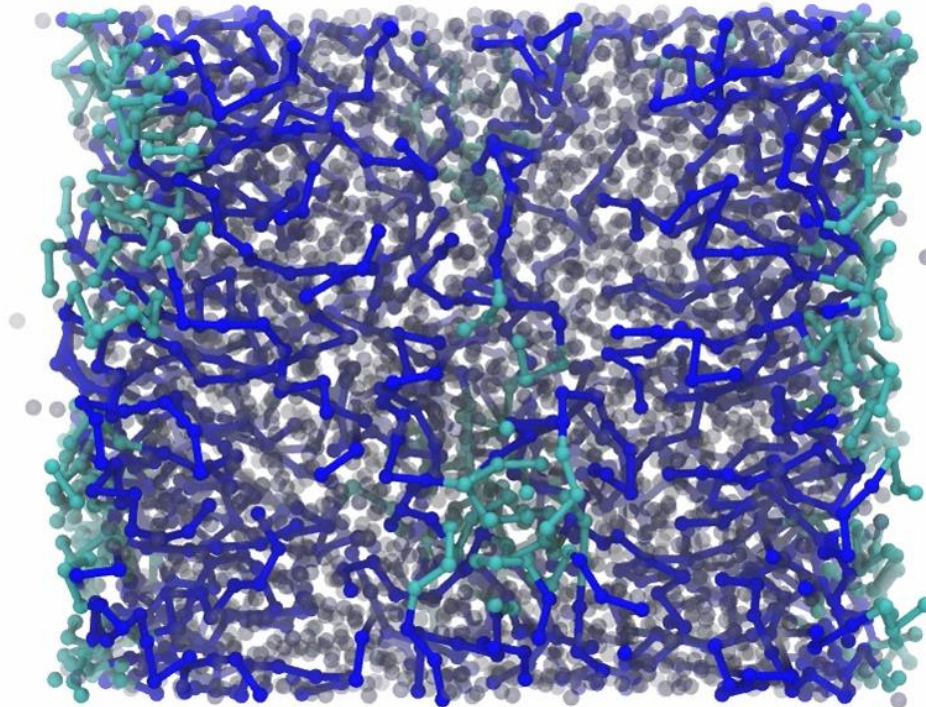
Percolating Cluster



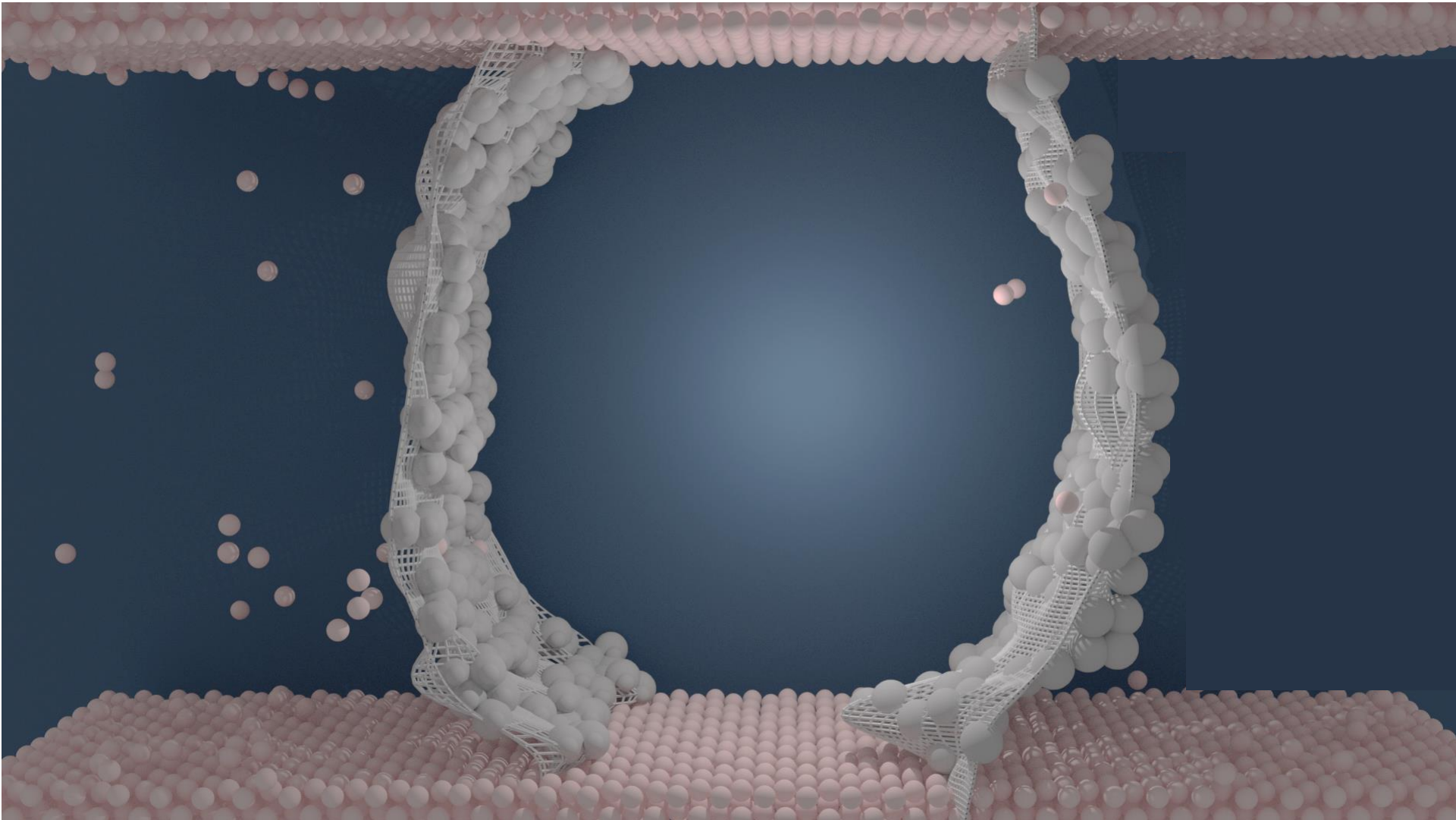
Applications

Marangoni flows

- Formation of concentrations gradients linked to surface

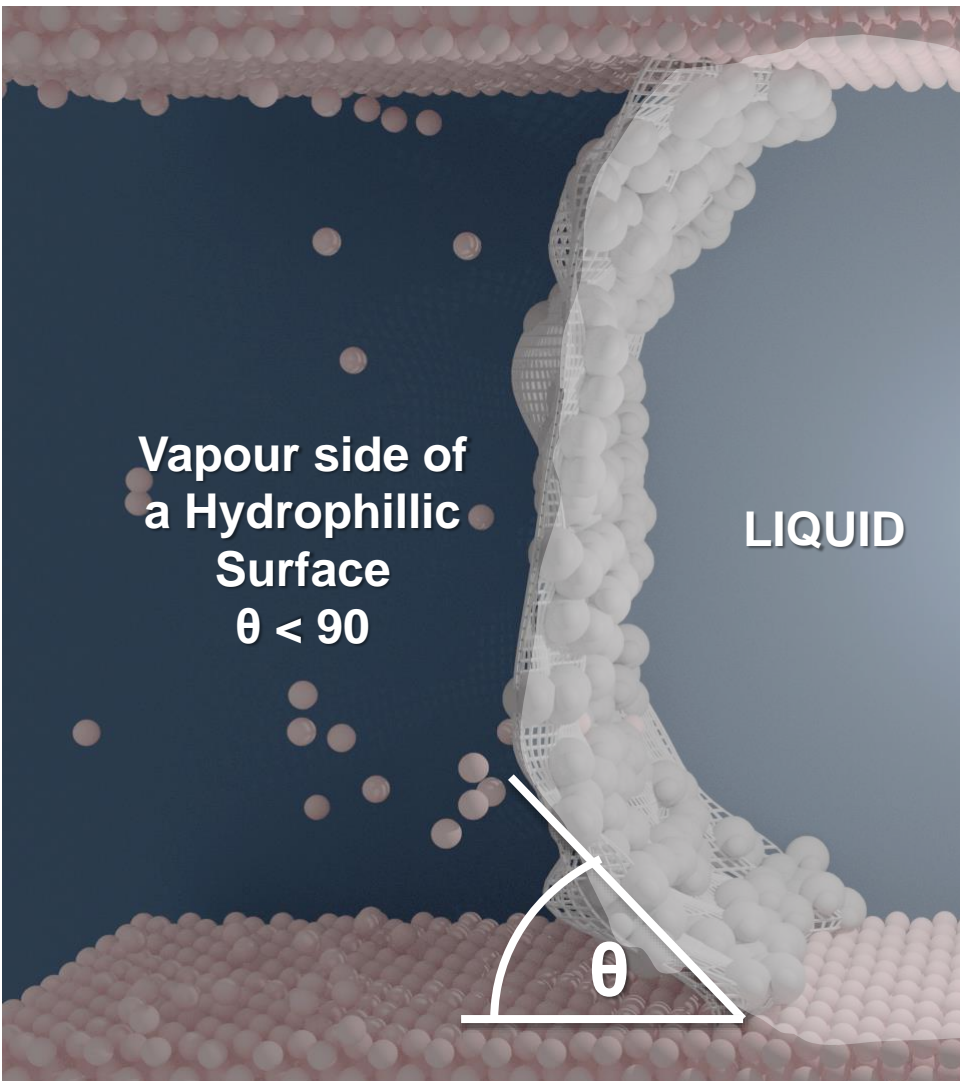


The Contact Line

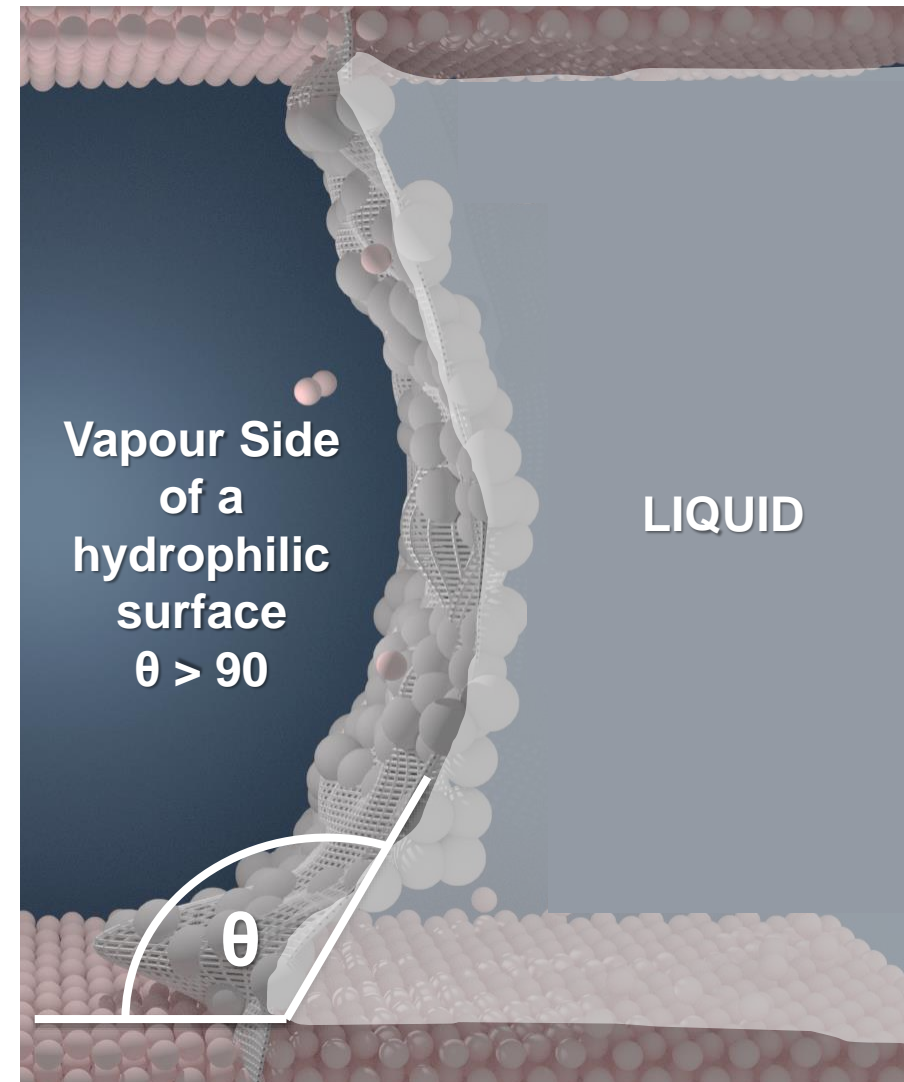




The Static Contact Line

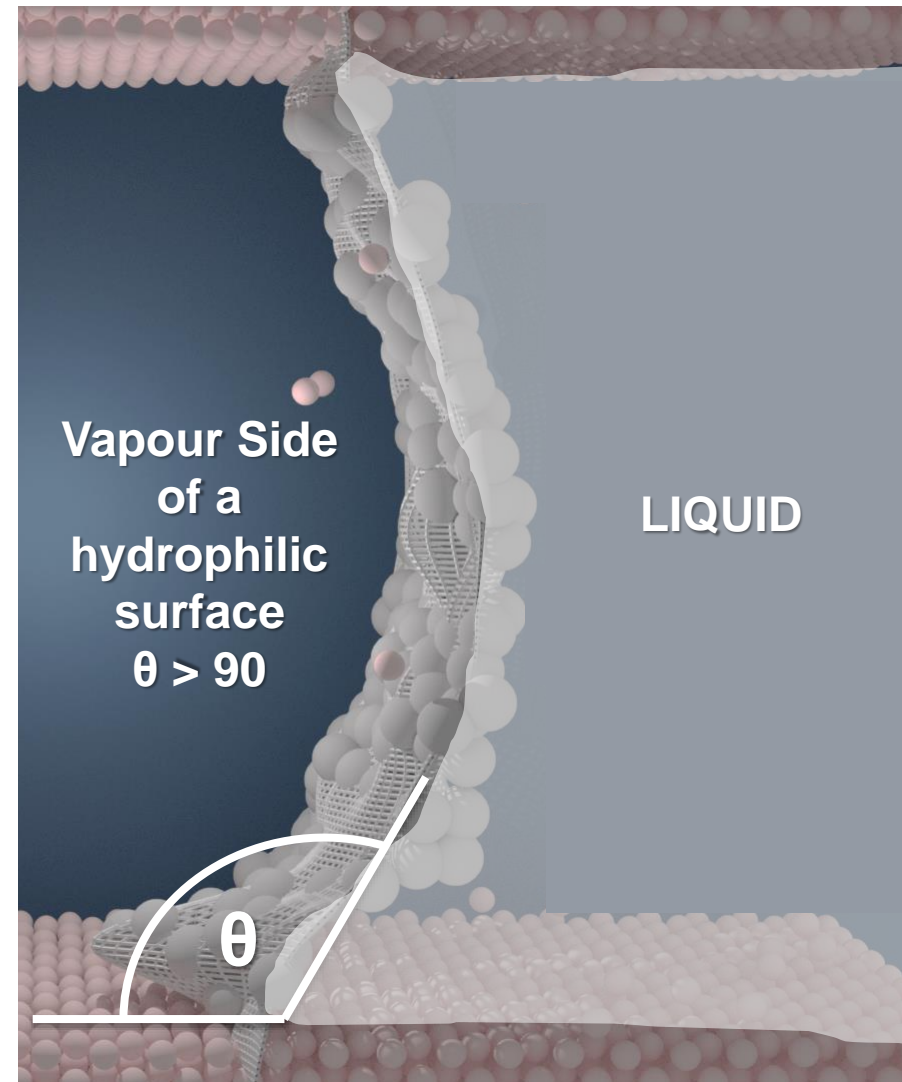
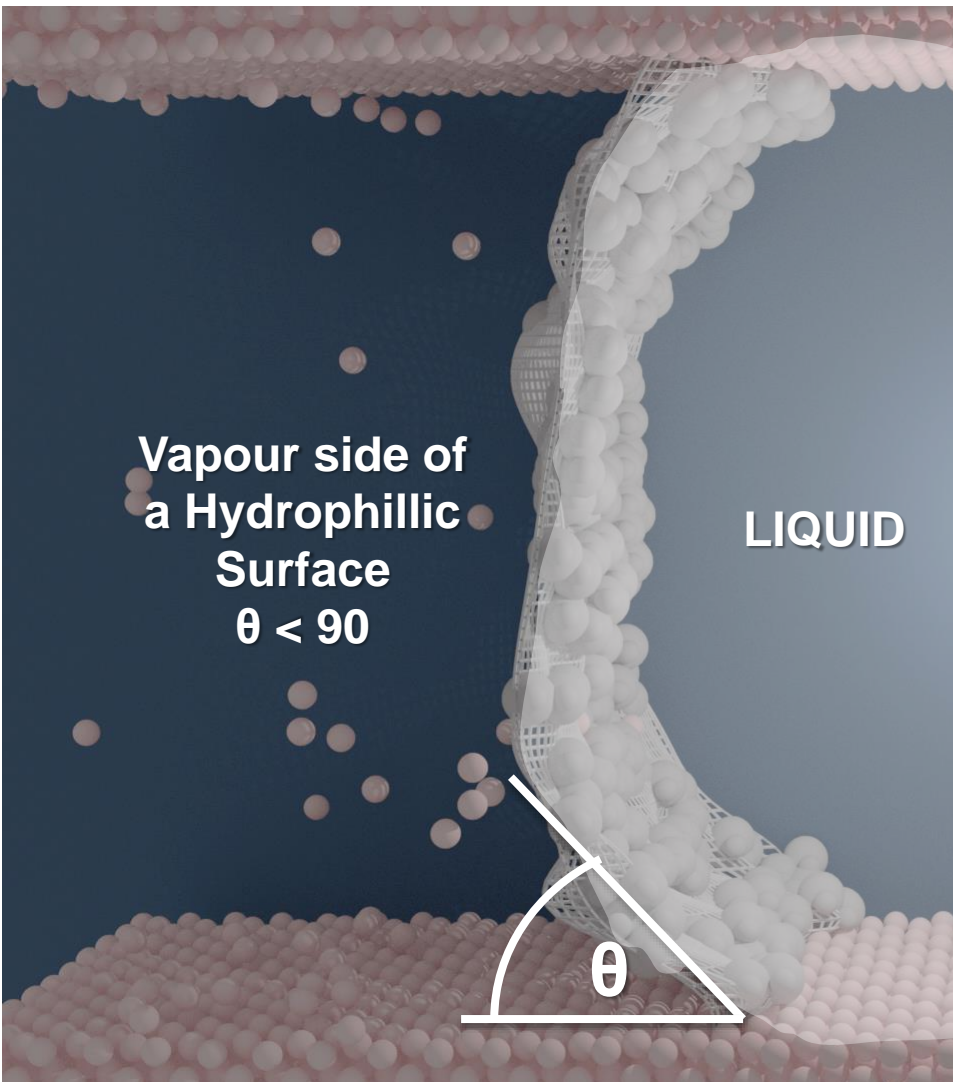


The Static Contact Line

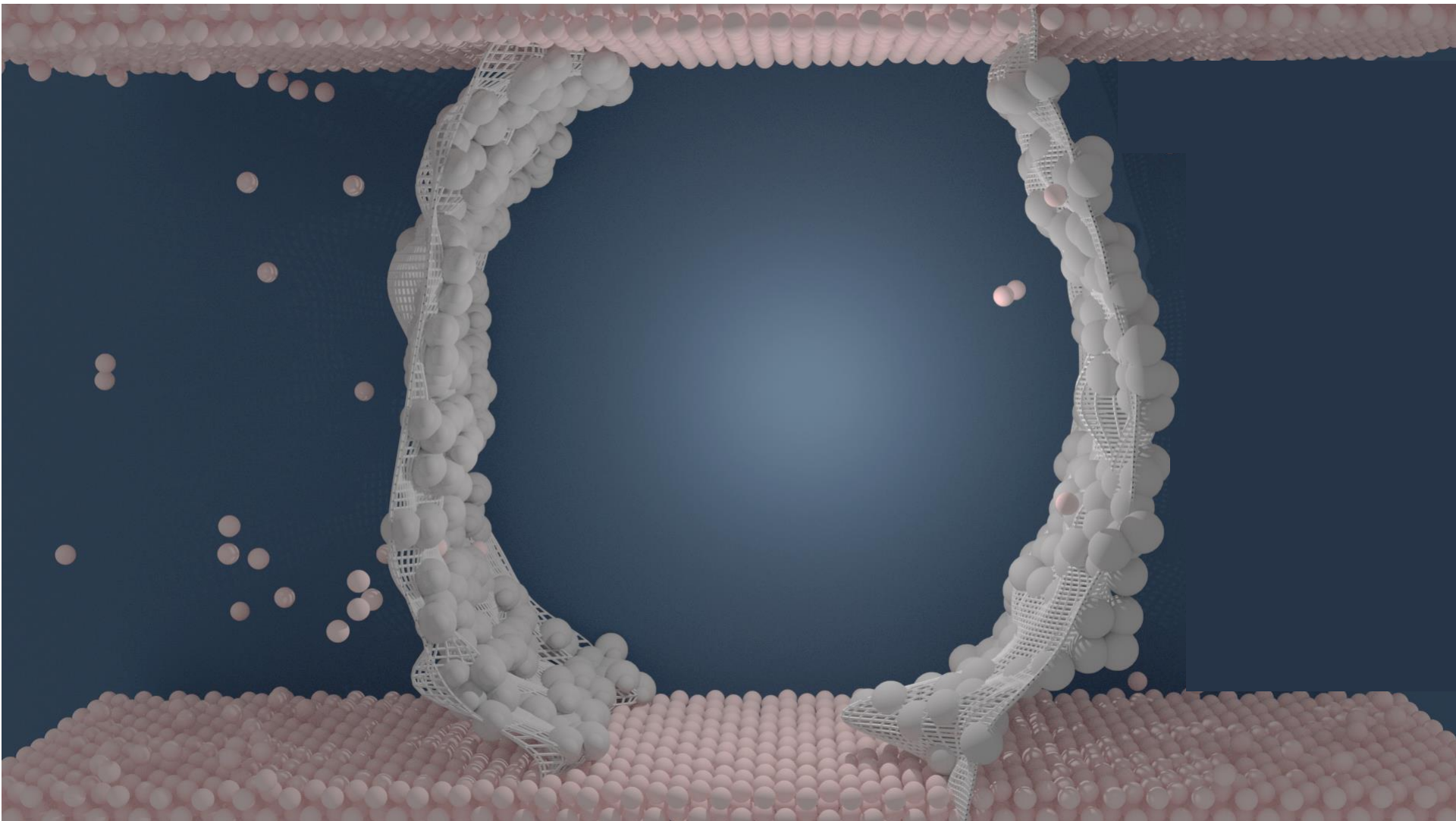




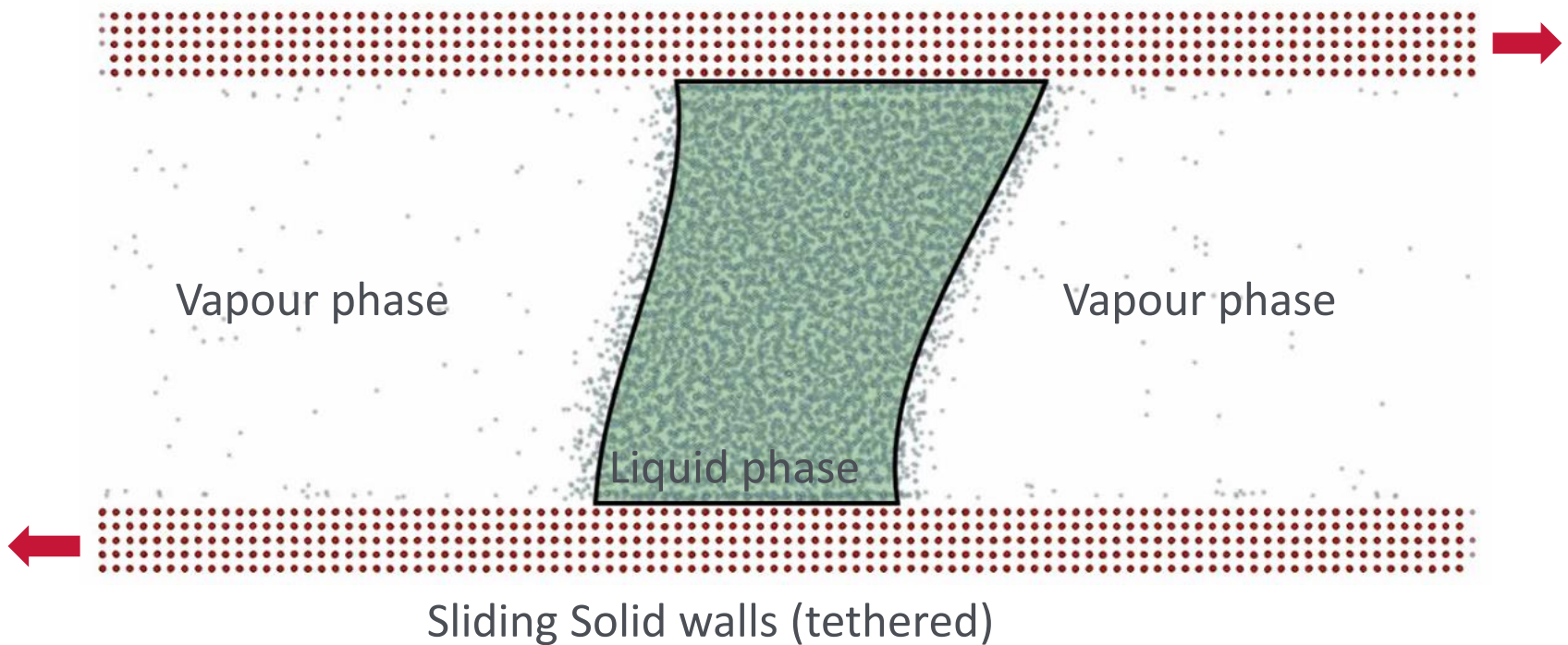
The Static Contact Line



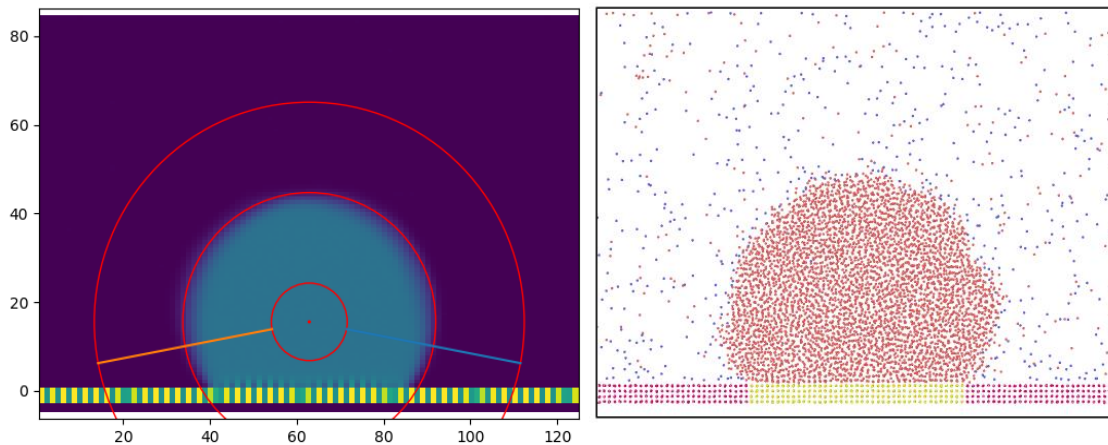
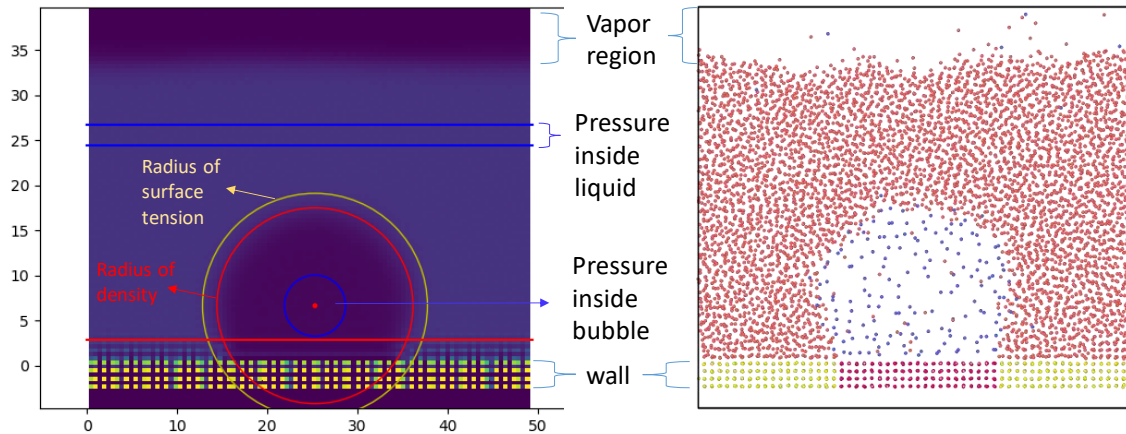
The Static Contact Line



The Dynamic Contact Line

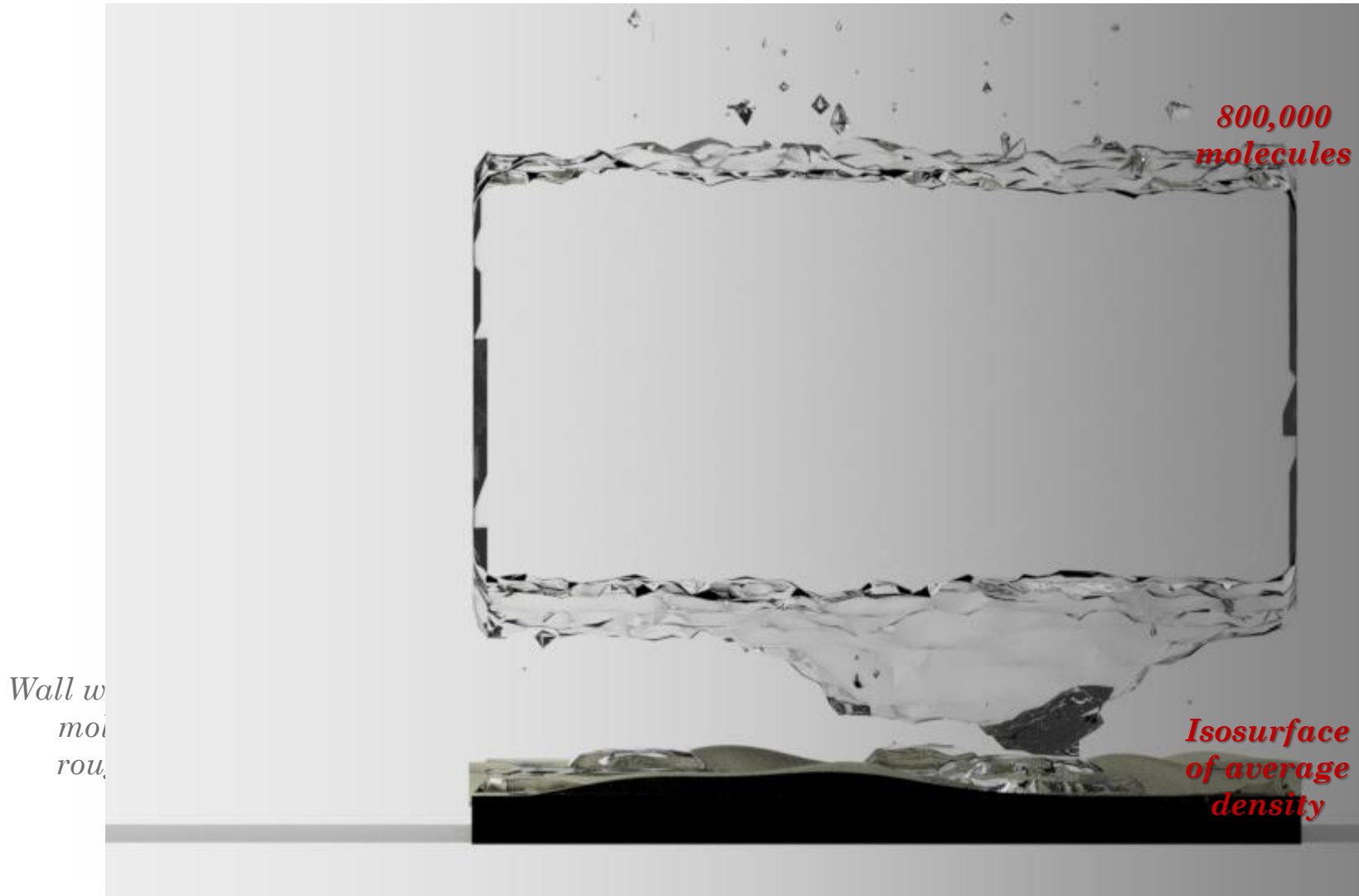


Droplet vs. Bubbles

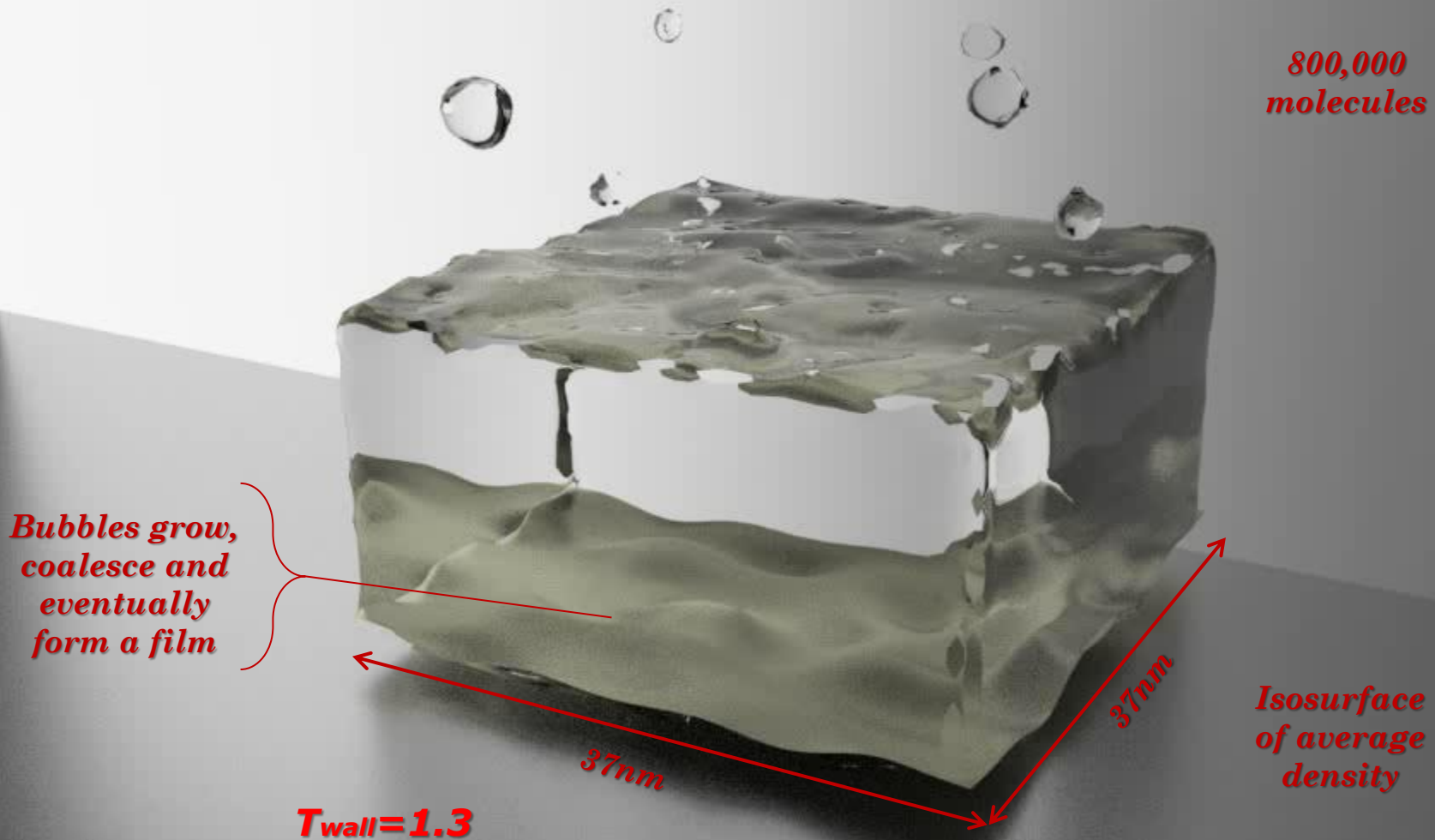


Work by
Wen Jun (Sonic)
at Imperial

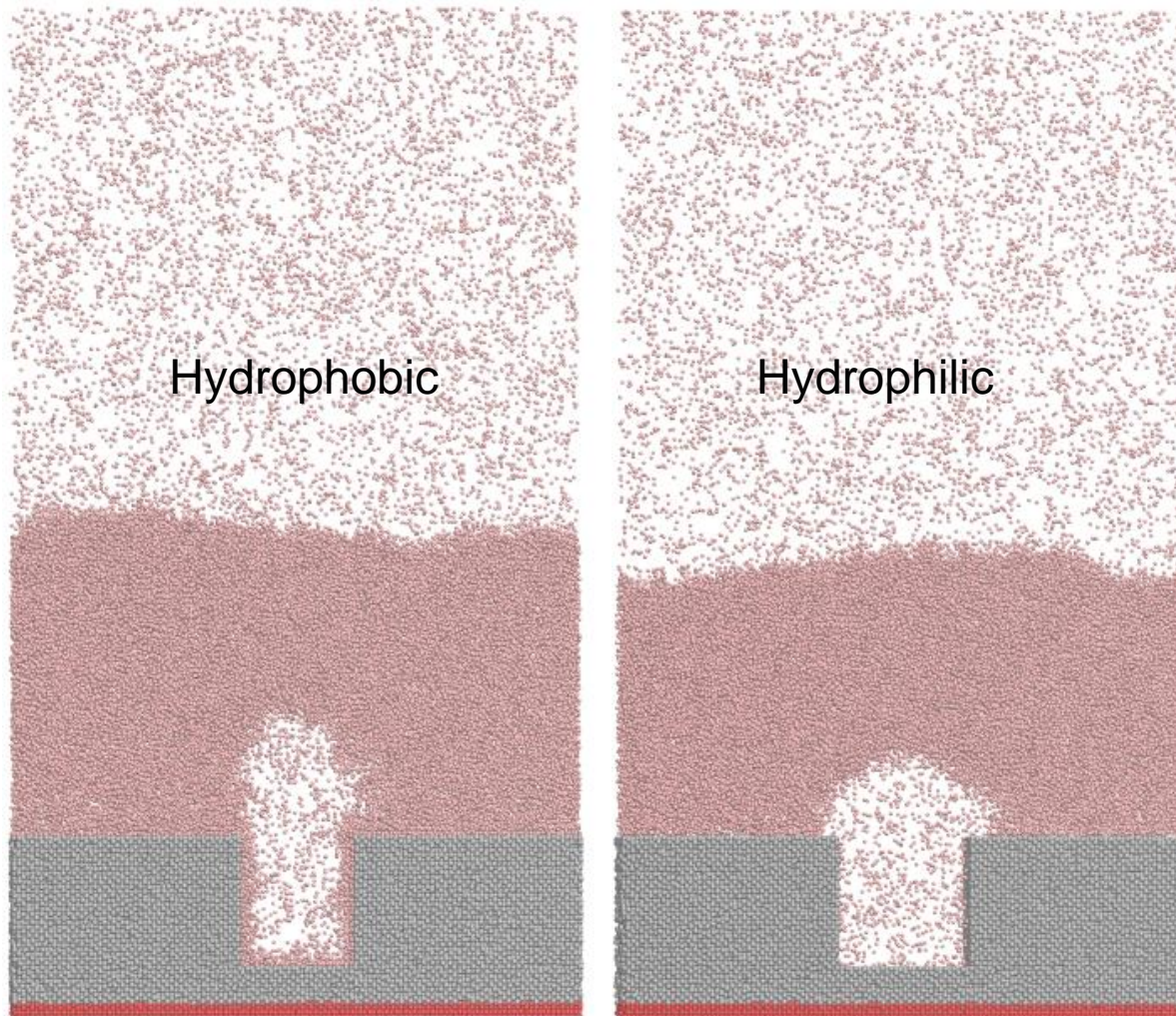
Molecular Dynamics - Nucleation



Isosurface of Density



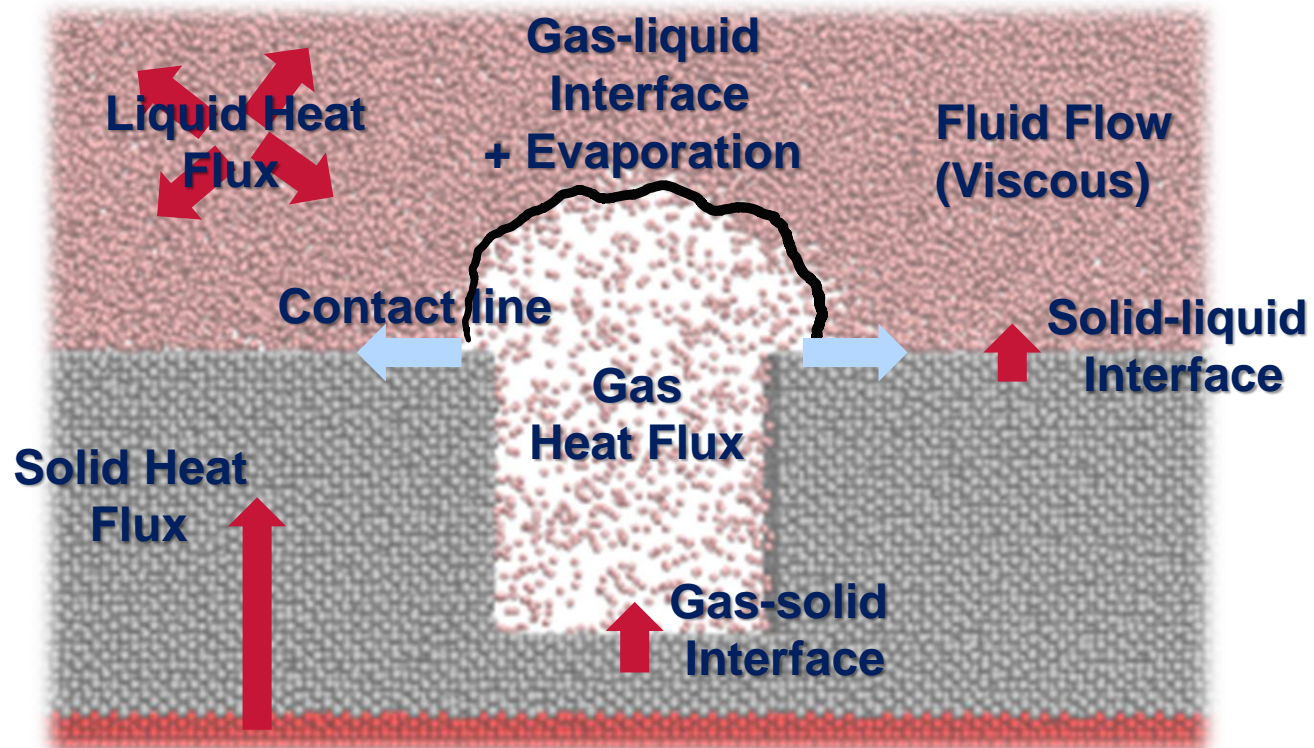
Wetting and Nucleation



Work by
Alessio Lavino
at Imperial



Factors in Bubble Nucleation



Coupled Simulation of Boiling

- Density at a point

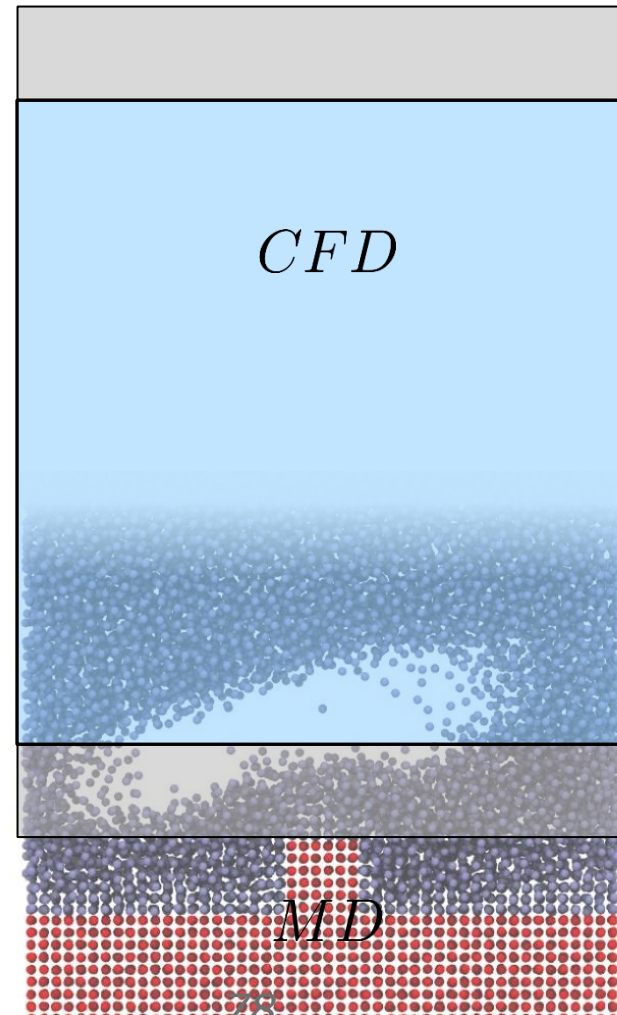
$$\rho(\mathbf{r}, t)$$



Build MD detail
into CFD

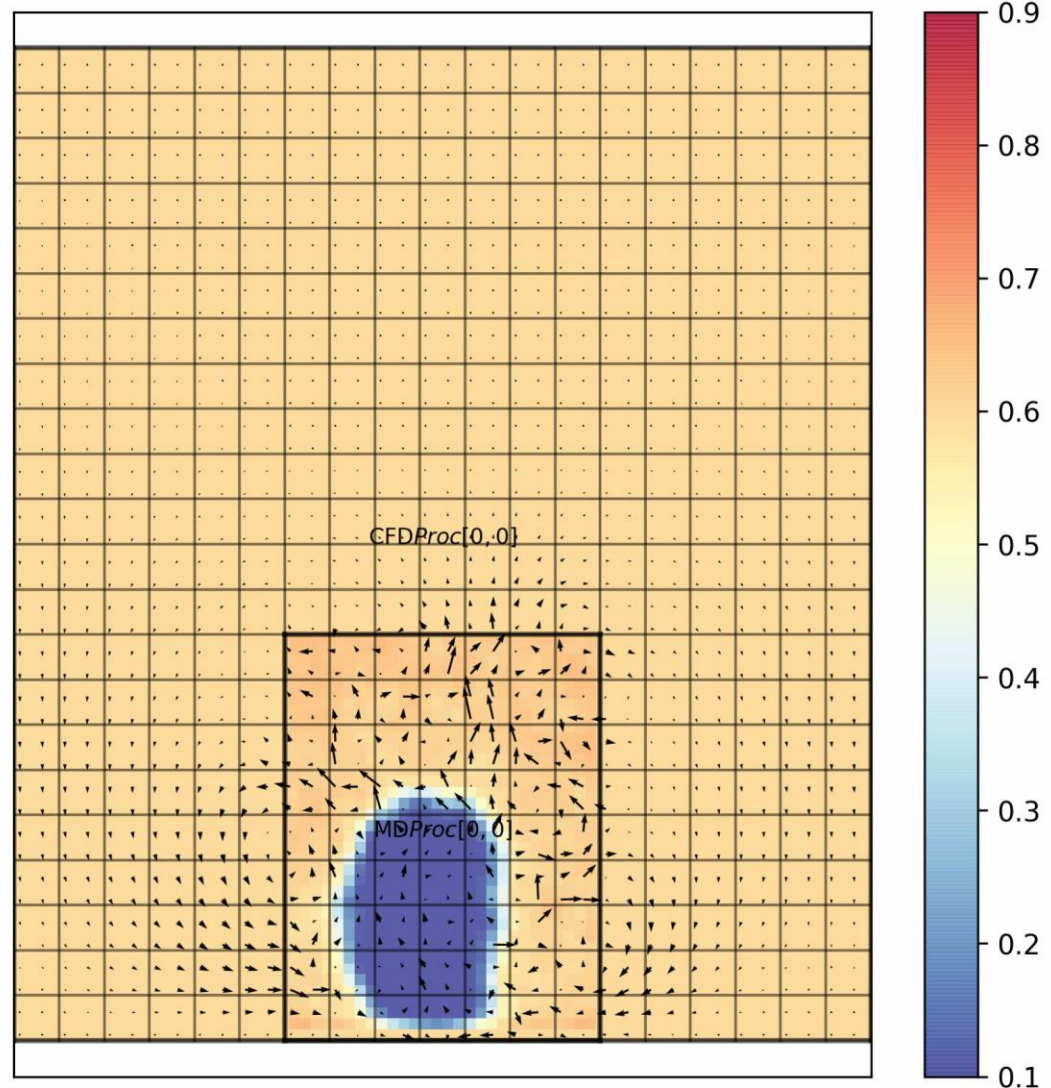
- Discrete molecules

$$m_i \text{ for all } i \text{ in } N$$



Coupled Simulation of Boiling

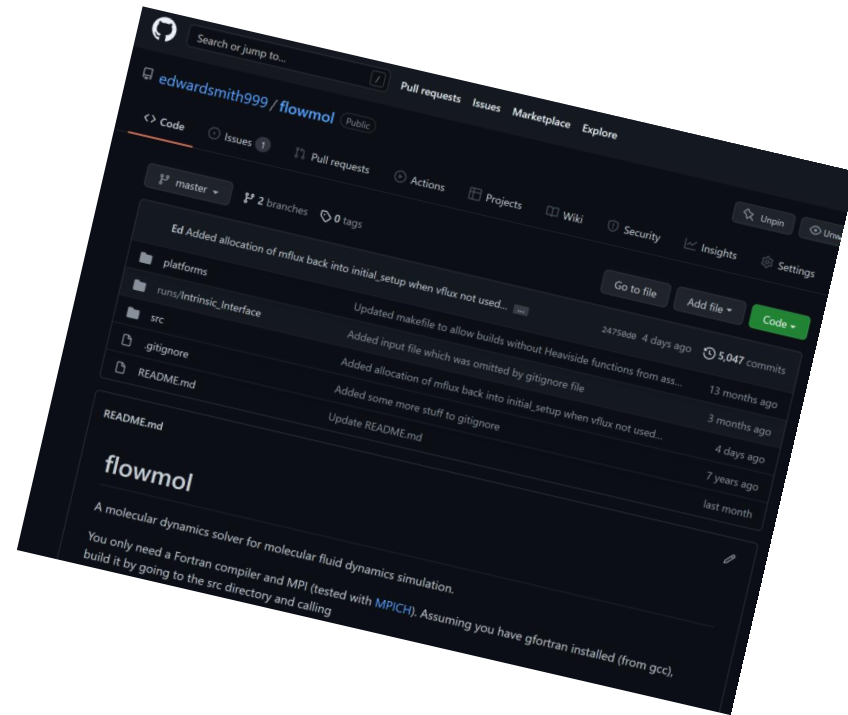
- Bubble nucleation occurs naturally in MD
- Density, velocity and temperature passed as boundary conditions



Application to Experimental Results

Open-source code

- Designed for particles
- Can be used for cells/pixels
- Surface fitting can be made as refined as necessary



- An introduction to Molecular Dynamics (MD)
- The moving contact line
- The liquid-vapour interface - a Lagrangian reference frame
- Some applications

Further Talks and Collaboration

1- Talks from Brunel

- (a) **Thermofluids** (Fluid Mech/Heat Transfer/Thermodynamics/Energy)
- (b) Manufacturing (automation/sustainability/design/materials)
- (c) Biosystems (LOC/Biomaterials)

2- Talks from York:

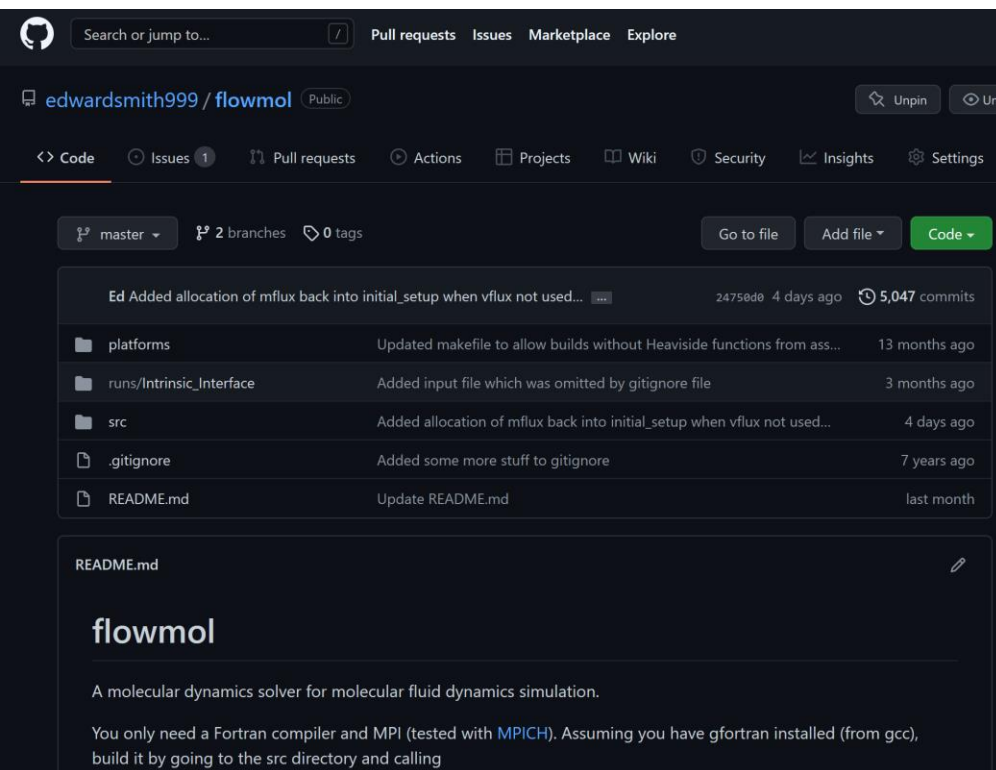
- (a) Microfluidics with or without a bio twist;
- (b) Automation with space applications;
- (c) Thermal systems with applications to vehicles; or for solar systems;
- (d) Materials (Bio, polymeric, or composites);
- (e) Droplet-surface interactions and applications in various industries.

3- UKRI (the UK funding Body) offers the UK-Canada Globalink doctoral exchange scheme, a 12 week research visit for PhD students (up to £15k)

Deadline 26th April 2022

Questions

Thank you, any questions?



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master 2 branches 0 tags

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Ed Added allocation of mflux back into initial_setup when vflux not used... 2475ed0 4 days ago 5,047 commits

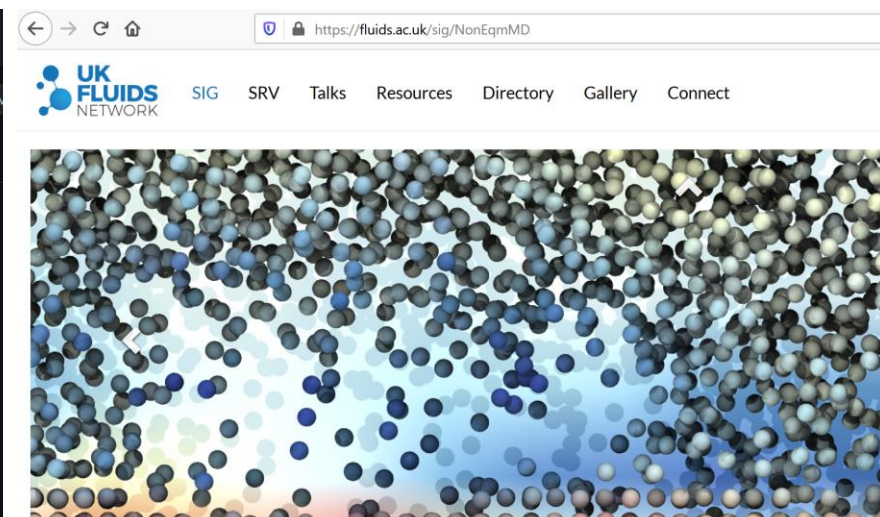
- platforms Updated makefile to allow builds without Heaviside functions from ass... 13 months ago
- runs/Intrinsic_Interface Added input file which was omitted by gitignore file 3 months ago
- src Added allocation of mflux back into initial_setup when vflux not used... 4 days ago
- .gitignore Added some more stuff to gitignore 7 years ago
- README.md Update README.md last month

README.md

flowmol

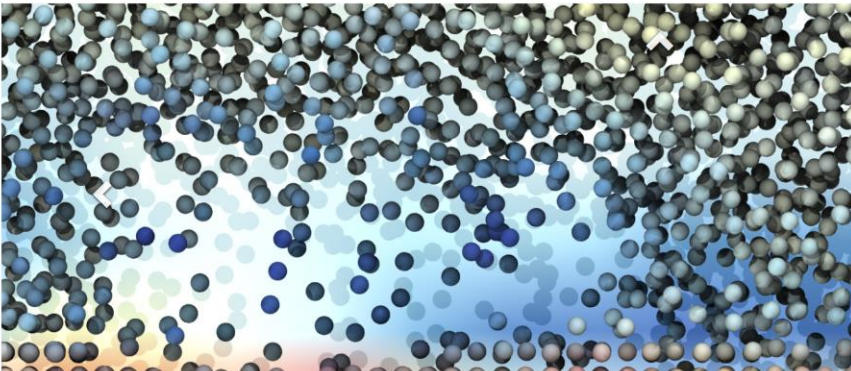
A molecular dynamics solver for molecular fluid dynamics simulation.

You only need a Fortran compiler and MPI (tested with [MPICH](#)). Assuming you have gfortran installed (from gcc), build it by going to the src directory and calling



https://fluids.ac.uk/sig/NonEqmMD

UK FLUIDS NETWORK SIG SRV Talks Resources Directory Gallery Connect



Non-equilibrium Molecular Dynamics (NEW)

Description

Non-equilibrium molecular dynamics (NEMD) is the study of fundamental fluid flow using molecular simulation - 'non-equilibrium' because the system is driven away from thermodynamic equilibrium by wall motion, temperature and pressure gradients or contains an interface, e.g. a liquid-vapour coexistence.

Please see <https://edwardsmith.co.uk>

Pressure (Stress)

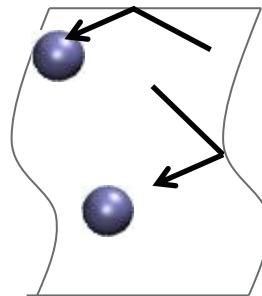
- Working through the mathematics

$$\Pi_{\alpha}^{\text{kinetic}} = \frac{1}{\Delta S_{\alpha}} \sum_{i=1}^N m_i \dot{\mathbf{r}}_i \mathbf{r}_{12} \cdot \frac{\mathbf{n}_{\alpha}}{|\mathbf{r}_{12} \cdot \mathbf{n}_{\alpha}|} dS_{\alpha}$$

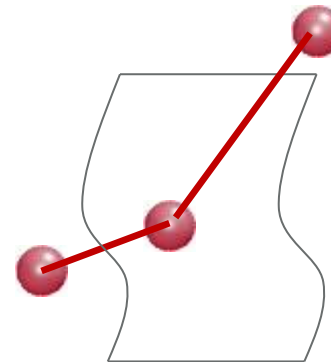
$$\Pi_{\alpha}^{\text{Moving surf}} = \frac{1}{\Delta S_{\alpha}} \sum_{i=1}^N m_i \dot{\mathbf{r}}_i \vartheta_t$$

$$\Pi_{\alpha}^{\text{config}} = \frac{1}{2\Delta S_{\alpha}} \sum_{i,j} \mathbf{f}_{ij} \mathbf{r}_{ij} \cdot \frac{\mathbf{n}_{\alpha}}{|\mathbf{r}_{ij} \cdot \mathbf{n}_{\alpha}|} dS_{\alpha}$$

*Kinetic
theory part
Momentum due
to average of
molecules
crossing a plane
and returning*



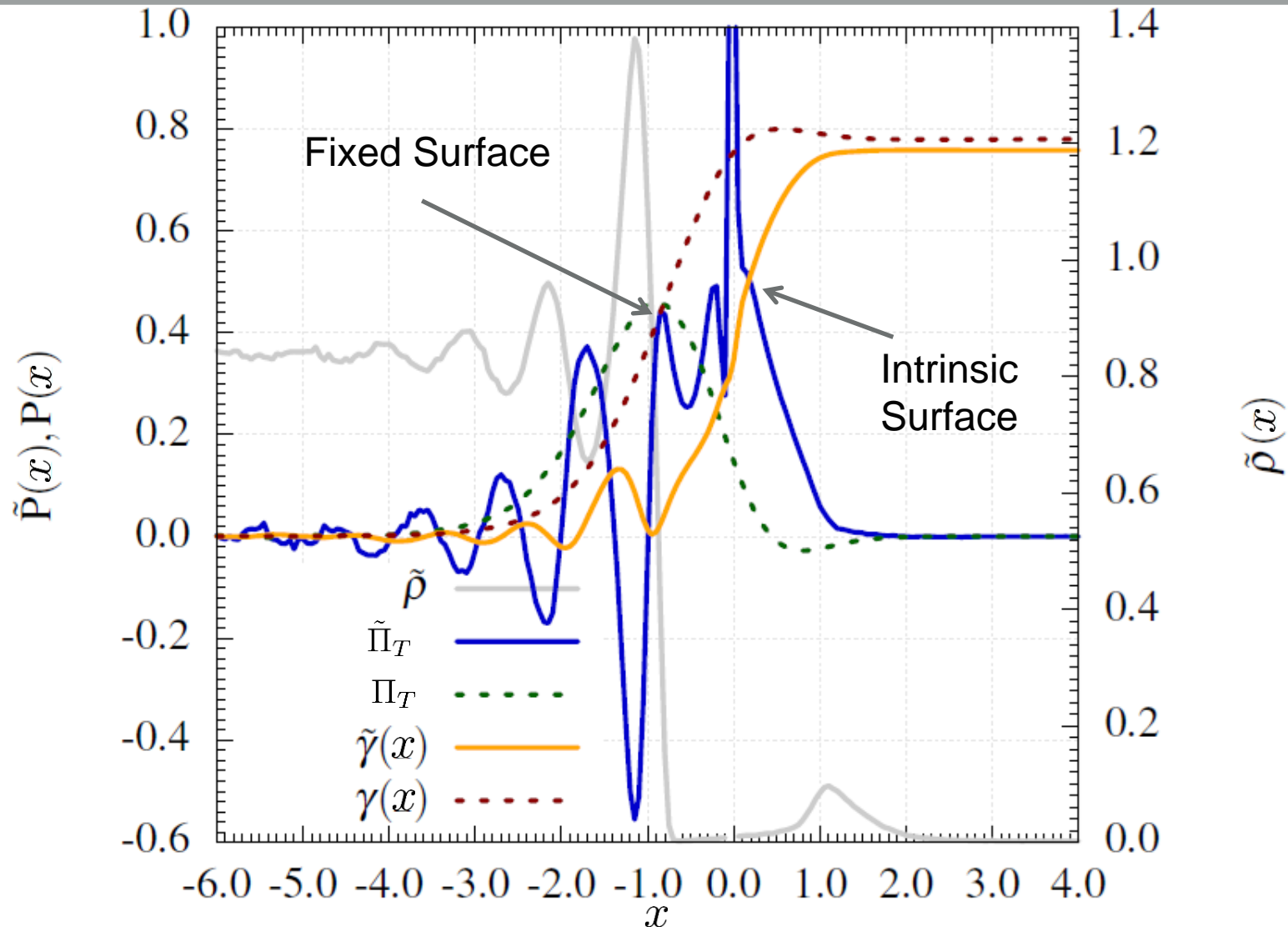
$$\dot{\mathbf{r}}_i = m_i \mathbf{v}_i + \mathbf{u}$$



*Configurational
part
Inter-molecular
bonds act like the
stress in a
stretched spring*



Tangential Pressure



Results for Surface Tension

