DeePN2: A deep learning-based non-Newtonian hydrodynamic model

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Introduction

A deep learning-based non-Newtonian hydrodynamic model, which retains a multi-scaled nature by mapping the polymer configurations into a set of symmetry-preserving macro-scale features. Our goals are:

- Faithfully inherit the micro-scale interactions beyond empirical models.
- Automatically capture specific molecular structural mechanics without human intervention.
- Strictly preserve the frame indifference constraints.
- Only utilize discrete samples for model training.

DeePN² Model

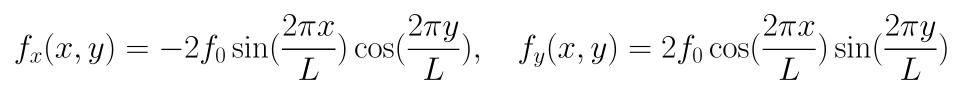
• **DeePN²:** A machine-learning-based model of non-Newtonian fluids. $\rho \frac{\mathrm{d}\mathbf{u}}{\mathrm{d}\mathbf{u}} = -\nabla p + \nabla \cdot (\boldsymbol{\tau}_{\mathrm{s}} + \boldsymbol{\tau}_{\mathrm{p}}), \quad \nabla \cdot \mathbf{u} = 0,$

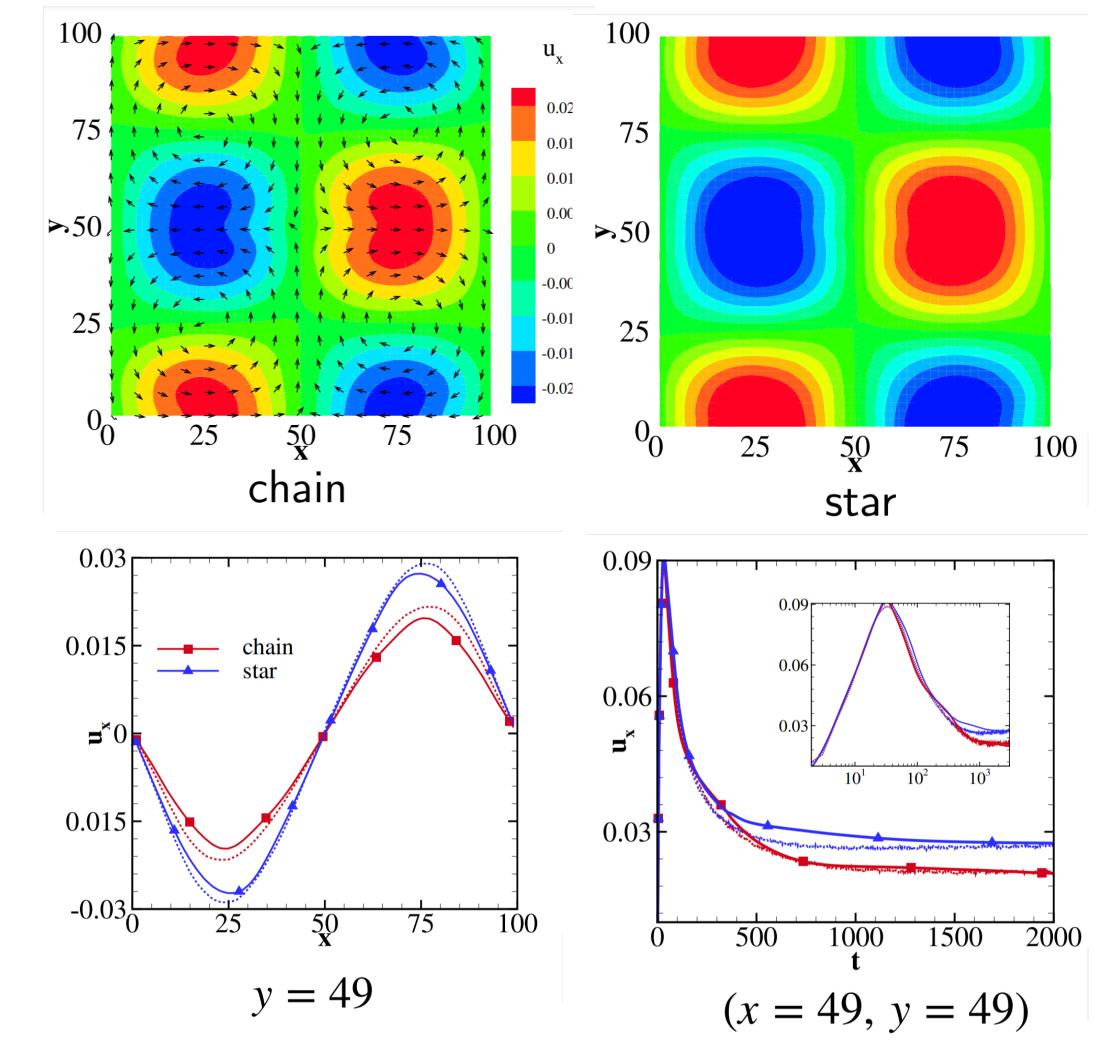
$$\boldsymbol{\tau}_{\mathrm{p}} = \mathbf{G}(\mathbf{c}), \quad \frac{\mathcal{D}\mathbf{c}_{i}}{\mathcal{D}t} = \mathbf{H}_{i}(\mathbf{c}),$$
$$\frac{\mathcal{D}\mathbf{c}_{i}}{\mathcal{D}t} = \frac{\mathrm{d}\mathbf{c}_{i}}{\mathrm{d}t} - \boldsymbol{\kappa} : \mathcal{E}_{i}, \quad \mathbf{H}_{i}(\mathbf{c}) = \frac{k_{B}T}{\gamma}\mathbf{H}_{1,i}(\mathbf{c}) - \frac{1}{\gamma}\mathbf{H}_{2,i}(\mathbf{c})$$

G, $H_{1,i}$, $H_{2,i}$, b_i , \mathcal{E}_i are represented by DNNs that rigorously **preserve** rotational symmetry and are jointly learned by:

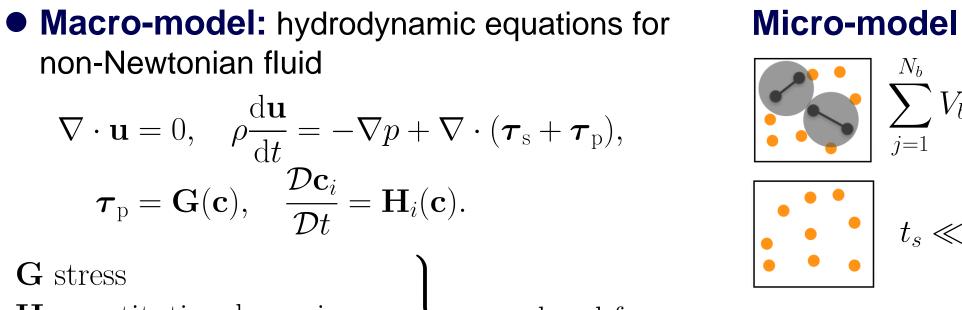
Numerical Example: 2D Green-Taylor flow

Vortex generated by:





Motivations



 \mathbf{H}_i constitutive dynamics \rightarrow unclosed form \leftarrow $\frac{\mathcal{D}\mathbf{c}_i}{\mathcal{D}t}$ objective tensor derivative

• Physical constraint: frame-indifference condition.

$$\mathbf{G}(\widetilde{\mathbf{c}}) = \mathbf{Q}\mathbf{G}(\mathbf{c})\mathbf{Q}^T, \quad \frac{\widetilde{\mathcal{D}\mathbf{c}_i}}{\mathcal{D}t} = \mathbf{Q}\frac{\mathcal{D}\mathbf{c}_i}{\mathcal{D}t}\mathbf{Q}^T$$

• Limits of existing empirical models: □ Molecular fidelity: heuristic form of **G(c) \Box** Empirical formulation: non-unique choices of $\frac{Dc}{dr}$ Generalization ability: heterogeneous molecule structural micro-mechanics

• Machine-learning difficulties: **D** Retain physical interpretation **D** Require error-prone time-series samples □ Preserve symmetry constraints

 $\mathbf{c}(t+dt)$ $\tilde{\mathbf{c}}(t+dt)$ $\mathbf{Q}(t+dt)$

 $t_s \ll t_c$

 $t_s \not\ll t_c$

$$egin{aligned} \mathbf{H}_{1,i}(\mathbf{c}) &= \left\langle \sum_{j,k=1}^{N-1} A_{jk}
abla_{\mathbf{r}_j} \cdot
abla_{\mathbf{r}_k} \mathbf{b}_i
ight
angle \ \mathbf{H}_{2,i}(\mathbf{c}) &= \left\langle \sum_{j=1}^{N-1} \sum_{k=1}^{N_b} A_{jk}
abla_{\mathbf{r}_k} V(\mathbf{r}) \cdot
abla_{\mathbf{r}_j} \mathbf{b}_i
ight
angle \ \mathcal{E}_i(\mathbf{c}) &= \left\langle \sum_{j=1}^{N-1} \mathbf{r}_j \otimes
abla_{\mathbf{r}_j} \otimes \mathbf{b}_i
ight
angle \ \mathbf{G}(\mathbf{c}) &= \left\langle \sum_{k=1}^{N_b} \mathbf{r}_k \otimes
abla_{\mathbf{r}_k} \otimes V_b(r)
ight
angle \end{aligned}$$

Remark: DeePN² achieves the following innovations • Does Not:

Learn constitutive dynamics using timeseries samples.

□ Rely on empirical choices of the constitutive closures.

□ Seek the direct approximation of the high-dimensional PDF.

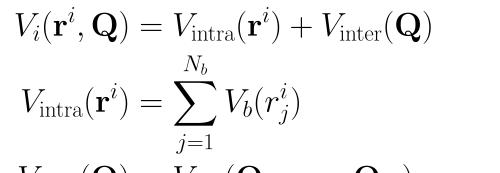
• Does:

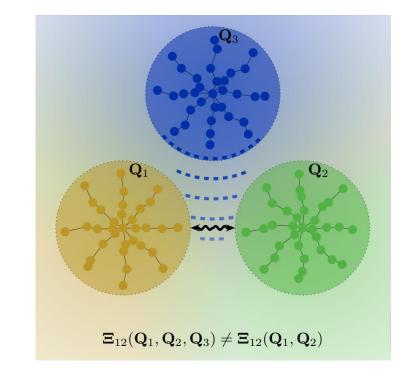
D Be reliable: systematically pass the micro-scale heterogenous molecular structures and interactions.

D Retain physical interpretation: provide a generalized form of the objective tensor derivative.

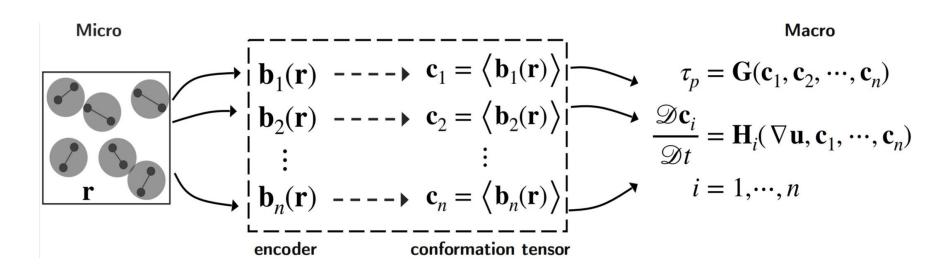
Future Work

Integrate the inter-molecule interactions into the DeePN² model

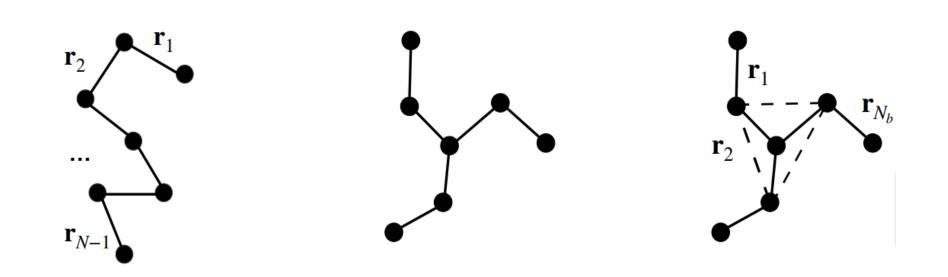




Model Construction from Micro Description



• Main idea 1: seek explicit micro-macro encoders to best approximate the macro-scale features.



r is the bond which is rotational-symmetric, r^* is the length and the distance of bonds which is rotational-invariant. Then we can construct the symmetry-preserving micro-macro encoder by

$$\mathbf{c}_i = \langle \mathbf{b}_i(\mathbf{r}) \rangle, \quad \mathbf{b}_i = \mathbf{f}_i \mathbf{f}_i^T, \quad \mathbf{f}_i = g_i(\mathbf{r}^*) \sum_{j=1}^{N-1} w_{ij} \mathbf{r}_j, \qquad 1 \le i \le n,$$

where w_{ij} and g_i will be represented by deep neural networks.

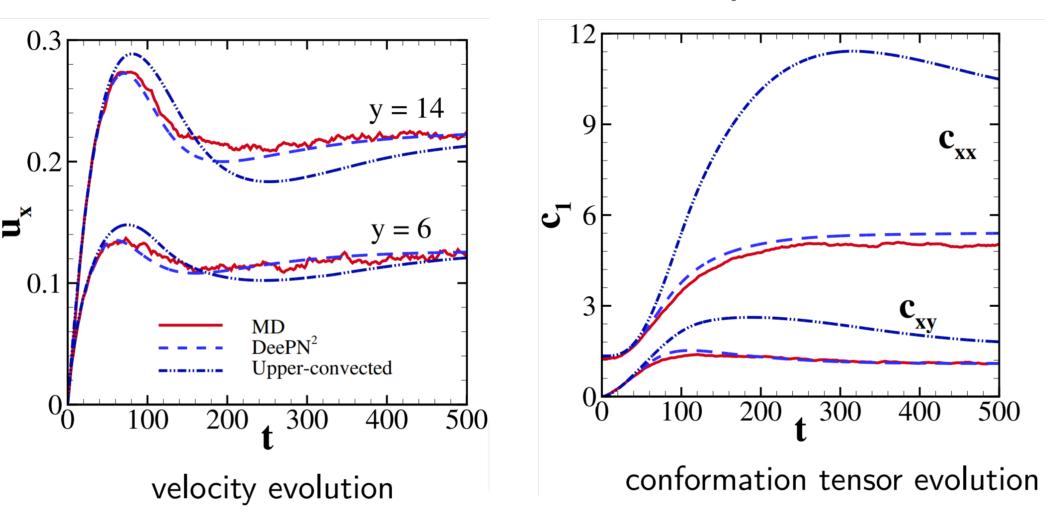
Remark:

- \Box { c_i } are not standard moments to approximate the polymer configuration density.
- \Box {**b**_{*i*}} are jointly learned for the best approximation of **G**(·) and **H**(·).

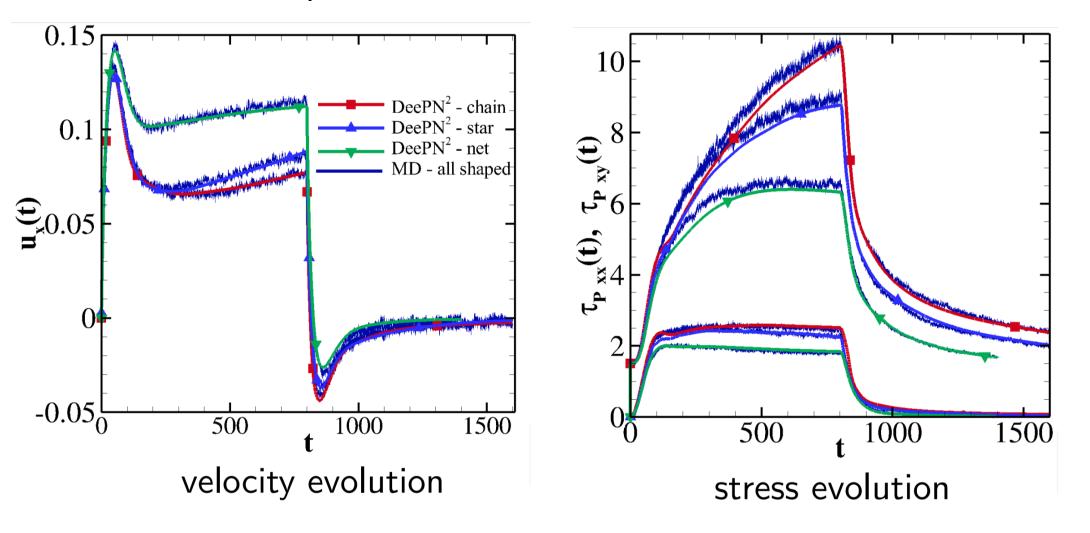
Carter Constraints: strictly preserve the rotational frameindifference symmetries.

Numerical Example: Reverse Poiseuille Flow

• **Dumbbell:** Empirical form of objective tensor derivative is limited.



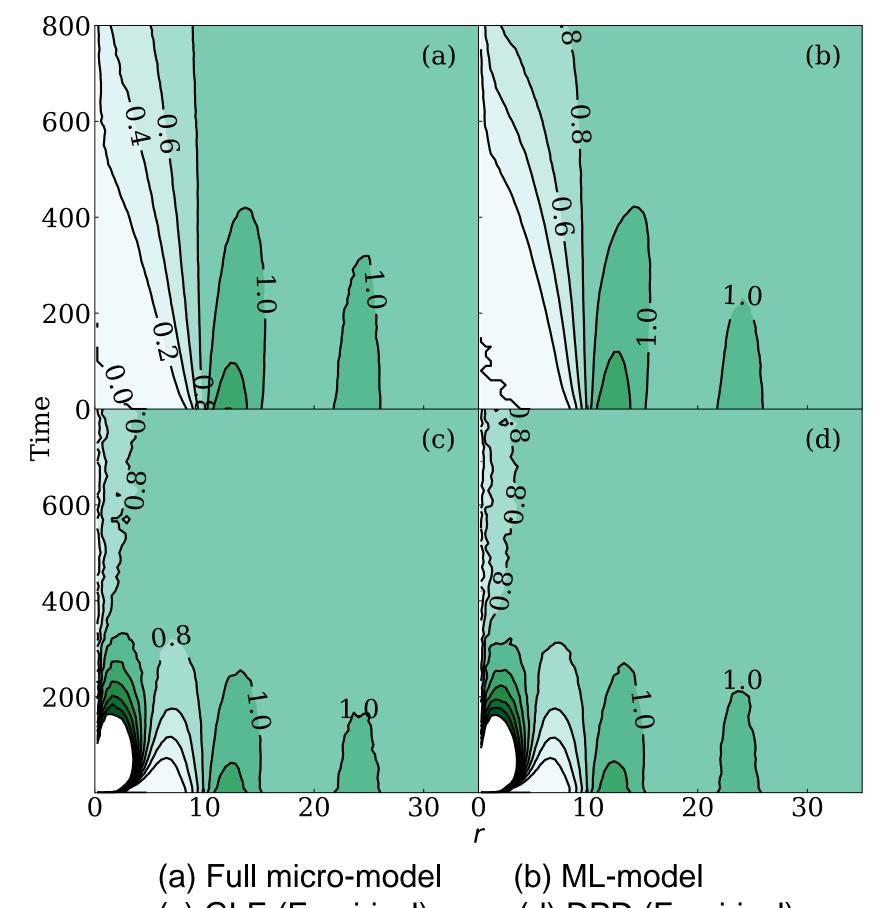
• Multi-bead molecules: the polymer suspensions exhibit different flow responses due to the different molecule structures.



$V_{\text{inter}}(\mathbf{Q}) = V_{\text{CG}}(\mathbf{Q}_1, \cdots, \mathbf{Q}_N)$

r - intramolecular coordinate **Q** - center of mass of polymer molecules

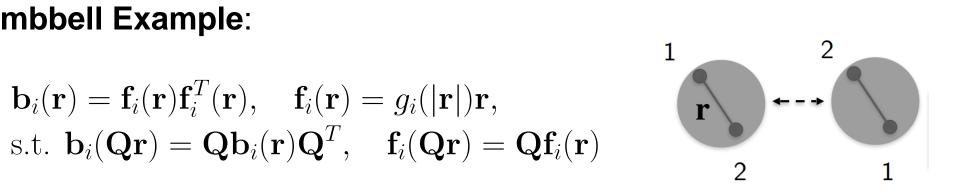
The intermolecular interaction is many-body and further effect the hydro dynamics (van-Hove function) on the collective scale.



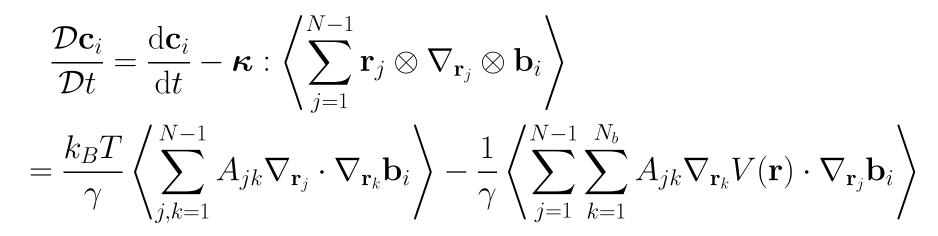
 \Box { c_i } strictly preserve symmetry constraints

 $\mathbf{b}_i(\mathbf{r}) = \mathbf{f}_i(\mathbf{r})\mathbf{f}_i^T(\mathbf{r}), \quad \mathbf{f}_i(\mathbf{r}) = g_i(|\mathbf{r}|)\mathbf{r},$

Dumbbell Example:



• Main idea 2: discrete-sample-based learning with molecular fidelity. The explicit form of micro-macro correspondence enables us to derive their evolution from the Fokker-Planck equation

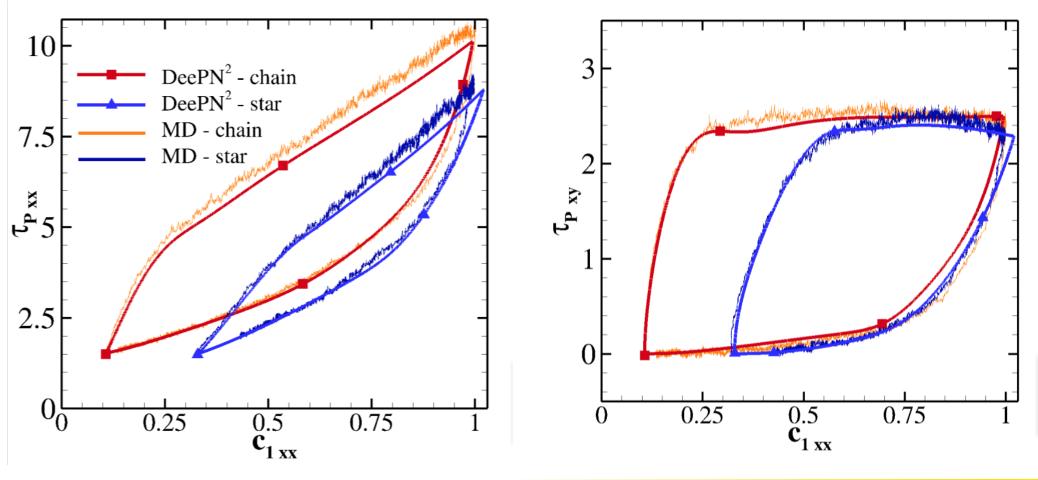


Remark:

A - generalized Rouse matrix encodes various molecule structures **U** V - potential function encodes micro-scale intramolecular interactions

□ The LHS provides a generalized objective tensor derivative with clear micro-scale physical interpretation.

Hysteresis loop: represents the strong memory effect that can not be captured by empirical (Hookean, FENE-P, etc.) models.



(d) DPD (Empirical) (c) GLE (Empirical)

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References

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