

Updates and Progress

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- Paper read: 20/1280
 - EQUIBIND: Geometric Deep Learning for Drug Binding Structure Prediction
 - Hamiltonian Graph Networks with ODE Integrator
 - Graph Normalizing Flows
 - PointFlow: 3D Point Cloud Generation with Continuous Normalizing Flows [Reading]
 - A Lagrangian Approach to Information Propagation in Graph Neural Networks [Reading]

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- CNF for generation & Coding

Numerical integrators for solving ODEs

- Given a first-order ODE and initial conditions, numerical integration can be used to approximate solutions to the initial value problem

$$\mathbf{y} \equiv \mathbf{y}(t) \quad , \quad \dot{\mathbf{y}} \equiv \frac{d\mathbf{y}}{dt} = f_{\mathbf{y}}(t, \mathbf{y}) \quad , \quad \mathbf{y}(t_0) = \mathbf{y}_0 \quad (1)$$

- The family of Runge-Kutta (RK) integrators are fully differentiable and can generate trajectories via iterations of the form, $y_{n+1} = RK(t_n, \Delta t, y_n, f_{\dot{y}})$

Hamiltonian Mechanics

- In Hamiltonian mechanics, the Hamiltonian, $\mathcal{H}(q, p)$, is a function of the canonical position, q , and momentum, p , coordinates

$$\dot{\mathbf{q}} \equiv \frac{d\mathbf{q}(t)}{dt} = \frac{\partial \mathcal{H}}{\partial \mathbf{p}} \quad , \quad \dot{\mathbf{p}} \equiv \frac{d\mathbf{p}(t)}{dt} = -\frac{\partial \mathcal{H}}{\partial \mathbf{q}} \quad , \quad (\dot{\mathbf{p}}, \dot{\mathbf{q}}) \equiv \frac{d(\mathbf{p}, \mathbf{q})}{dt} = f_{\dot{\mathbf{p}}, \dot{\mathbf{q}}}(\mathbf{q}, \mathbf{p}) \quad (2)$$

Delta graph network (DeltaGN)

- Baseline, DeltaGN directly predicts changes to (\mathbf{q}, \mathbf{p})

$$(\mathbf{q}, \mathbf{p})_{n+1} = (\mathbf{q}, \mathbf{p})_n + (\Delta \mathbf{q}, \Delta \mathbf{p})_n, (\Delta \mathbf{q}, \Delta \mathbf{p})_n \leftarrow GN(\Delta t, \mathbf{q}_n, \mathbf{p}_n, c, \phi)$$

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- The \mathbf{c} are static parameters (masses, spring constants) of the system, and ϕ are the neural network parameters.
- The GN's signature matches the integrator's, and so is analogous to learning an integrator.

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- Node output of a GN to model the per-particle time derivatives, and provide the GN, together with the initial conditions and , to an RK integrator

$$(\mathbf{q}, \mathbf{p})_{n+1} = RK(\Delta t, (\mathbf{q}, \mathbf{p})_n, f_{\dot{\mathbf{q}}, \dot{\mathbf{p}}}^{\text{OGN}}) \quad , \quad f_{\dot{\mathbf{q}}, \dot{\mathbf{p}}}^{\text{OGN}} \equiv GN(\mathbf{q}, \mathbf{p}, c, \phi) = (\dot{\mathbf{p}}, \dot{\mathbf{q}}) \quad (4)$$

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- The $f_{\dot{\mathbf{q}}, \dot{\mathbf{p}}}$ is a function that the integrator can use to operate on any (\mathbf{q}, \mathbf{p}) and query more than once

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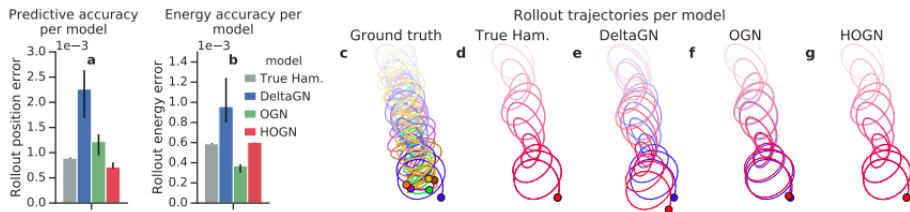
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Results

- Train and Test approach on datasets consisting of particle systems where particle j exerts a spring force on particle i , as defined by Hooke's law, $F^{ij} = -k^{ij}(\mathbf{q}^i - \mathbf{q}^j)$ limiting particles between 4 and 9 particles

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- Rollout position error: RMS position error averaged across all examples, dimensions, particles, and sequence axis.
- Rollout energy error: Energy error is calculated as the RMS of the deviation between the mean energy of a trajectory and the initial energy (normalized by the initial energy, to yield relative errors), averaged across all examples



Connection to our work

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$$V = \sum_i^{N_v} \sum_j^{N_{hops}} V(r_{ij}) = f_{PE}(\mathcal{G}(\mathcal{V}, \mathcal{E}), t) \quad (6)$$

$$U = \sum_i^{N_v} \frac{1}{2} m \dot{r}_i^2 = f_{KE}(\mathcal{G}(\mathcal{V}, \mathcal{E})_t, \mathcal{G}(\mathcal{V}, \mathcal{E})_{t-1}, t) \quad (7)$$

$$\mathcal{L} = U - V, \quad \frac{d(U + V)}{dt} = 0 \quad (8)$$

Connection to our work

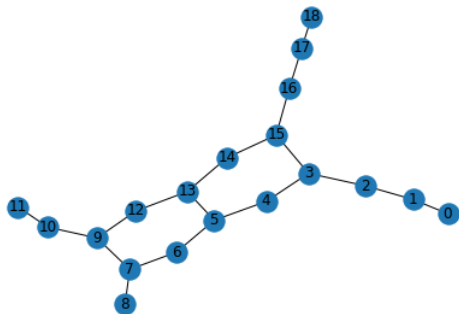
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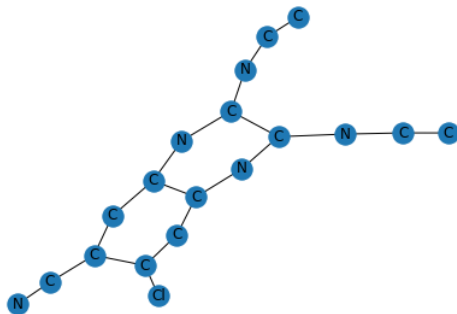
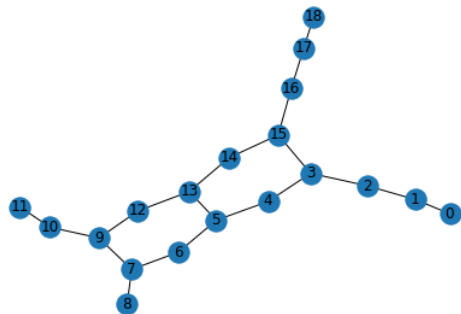
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$$v_i(t) = v_i(t-1) + g\left(\underbrace{\frac{\partial \mathcal{L}}{\partial r_i}}_{\text{Over each node}}, \underbrace{\frac{\partial \mathcal{L}}{\partial r_{ij}}}_{\text{Over each connection}}\right) \quad (9)$$

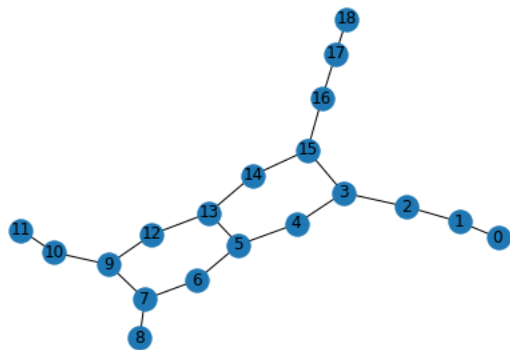
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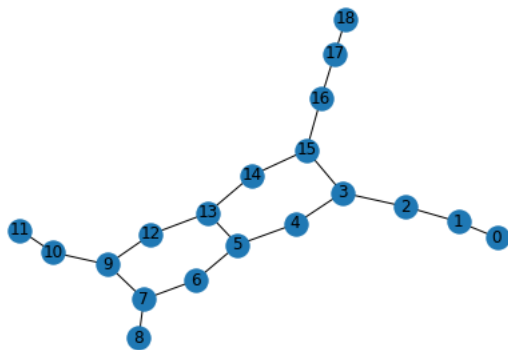


How to do?



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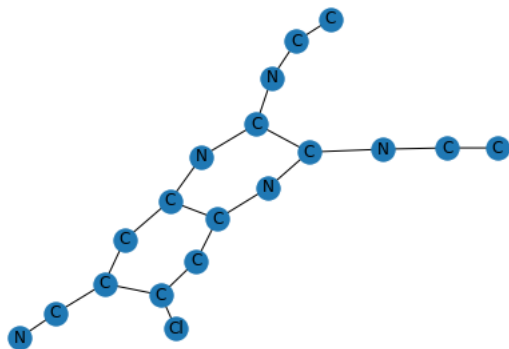
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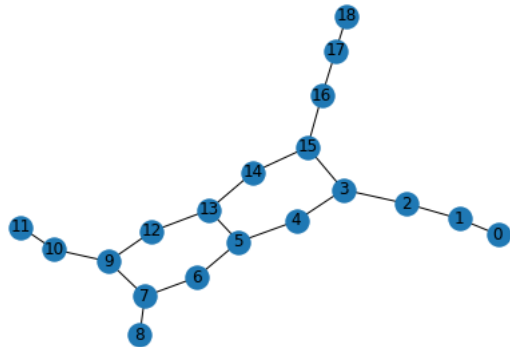
$$\log(p(z_1)) = \log(p(z_0)) - \log \left| \det \frac{\partial f}{\partial z_0} \right|$$

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- At the end get node label (type of atom) having the highest density

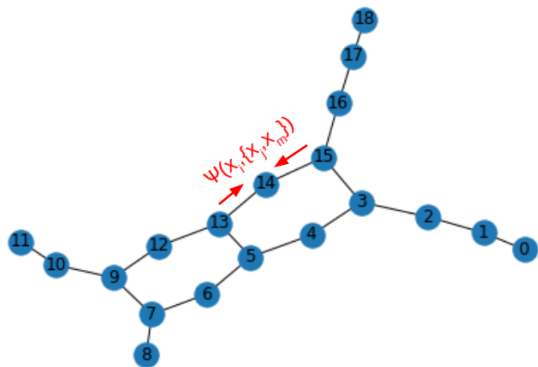
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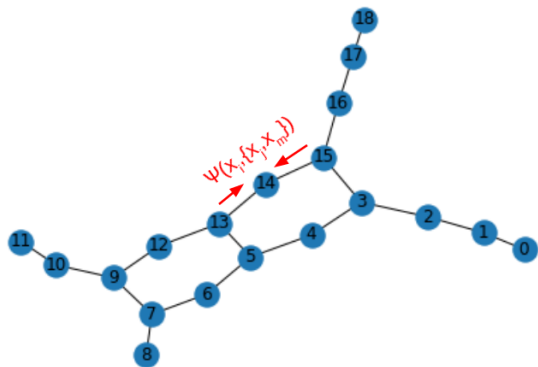
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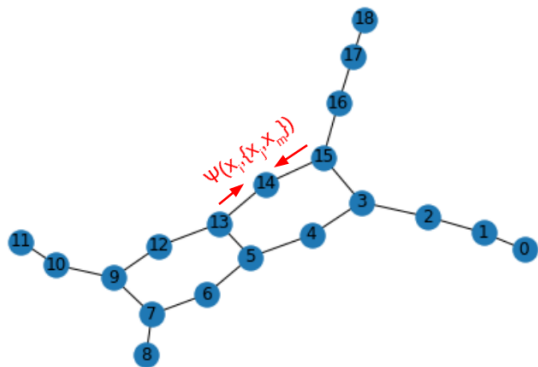


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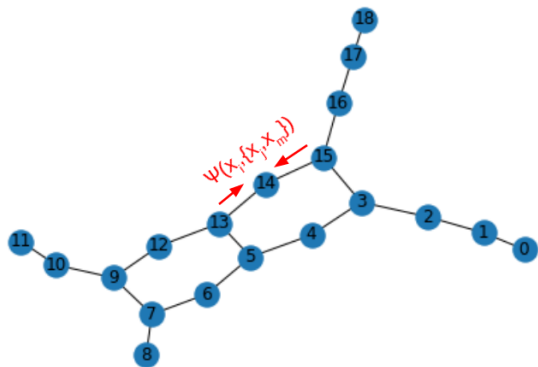
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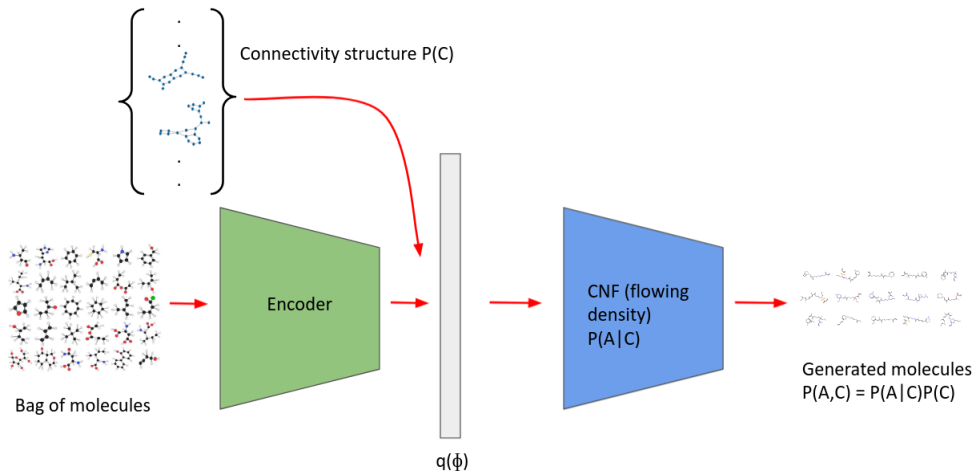
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- Recursively updated during each iteration and used as a feature input

Extension

- We can extend it as an encoder-decoder formalism as shown above



Master Equation

The drug design process can be represented as :

$$P(Prop, f, 3D, A, C) = P(Prop|f, 3D) \cdot P(f|3D, A) \cdot \underbrace{P(3D|A, C)}_{\text{Z-matrix}} \cdot \underbrace{P(A|C)P(C)}_{\text{We are here}} \quad (10)$$

THANK YOU