

# **E(n) Equivariant Normalizing Flows**

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- ▶ **Background**
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► **Introduction**

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# Introduction

This paper focus on symmetries of the **n-dimensional Euclidean group**, referred to as  $E(n)$ .

Propose a generative model equivariant to Euclidean symmetries:

## **$E(n)$ Equivariant Normalizing Flows (E-NFs)**

**Motivation:** To generate better drug candidate molecules to speed drug discovery

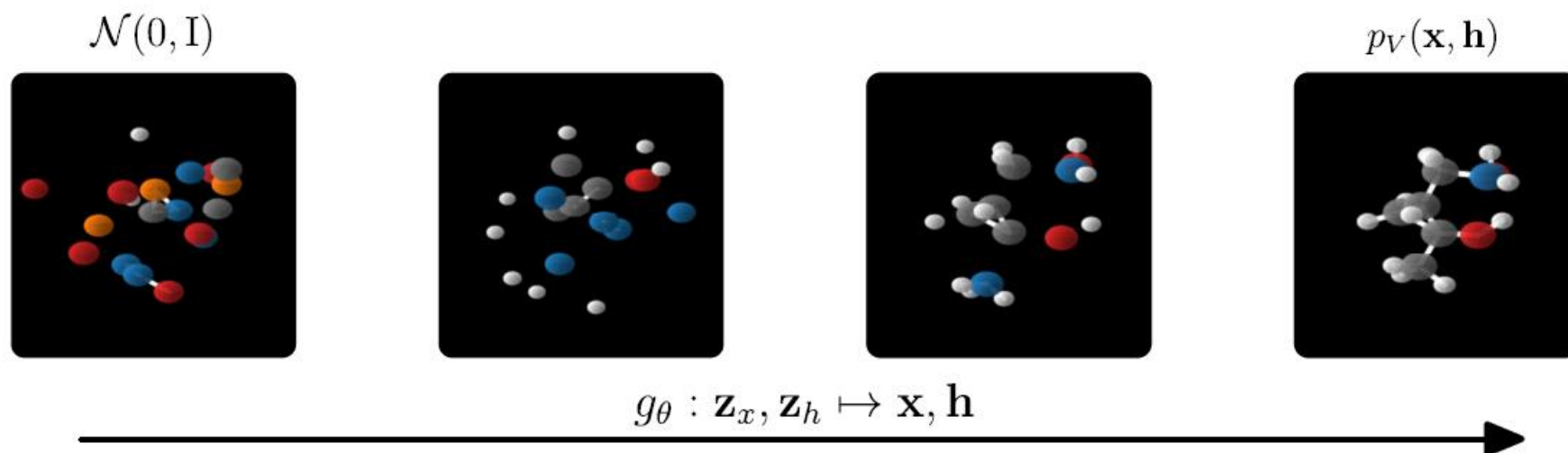


Figure 1: Overview of our method in the sampling direction. An equivariant invertible function  $g_\theta$  has learned to map samples from a Gaussian distribution to molecules in 3D, described by  $\mathbf{x}, \mathbf{h}$ .

► Introduction

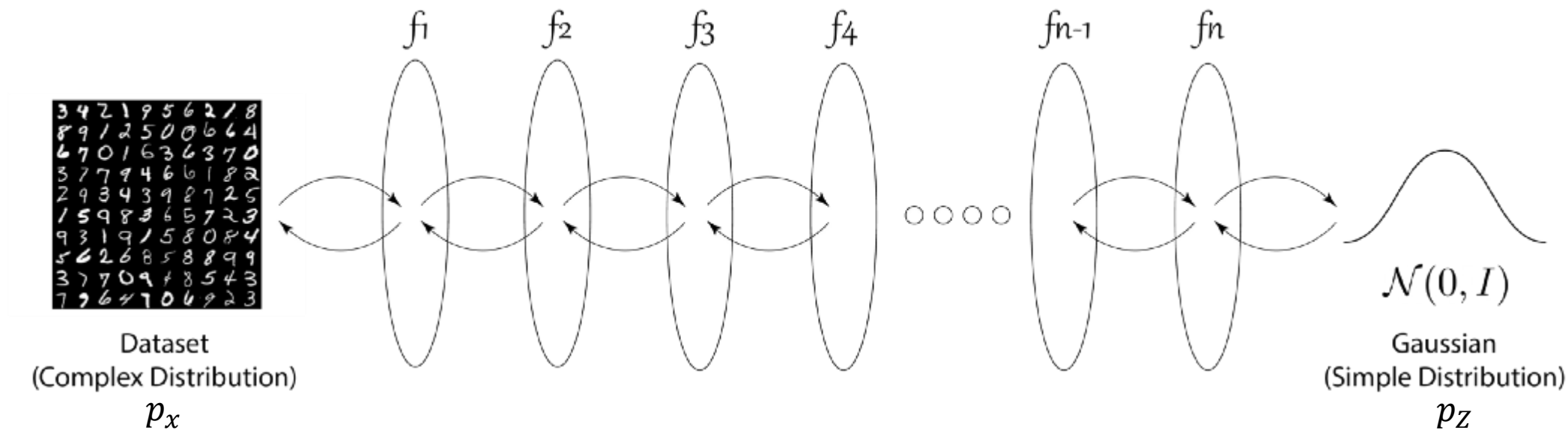
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## Background | E(n) Equivariant Normalizing Flows



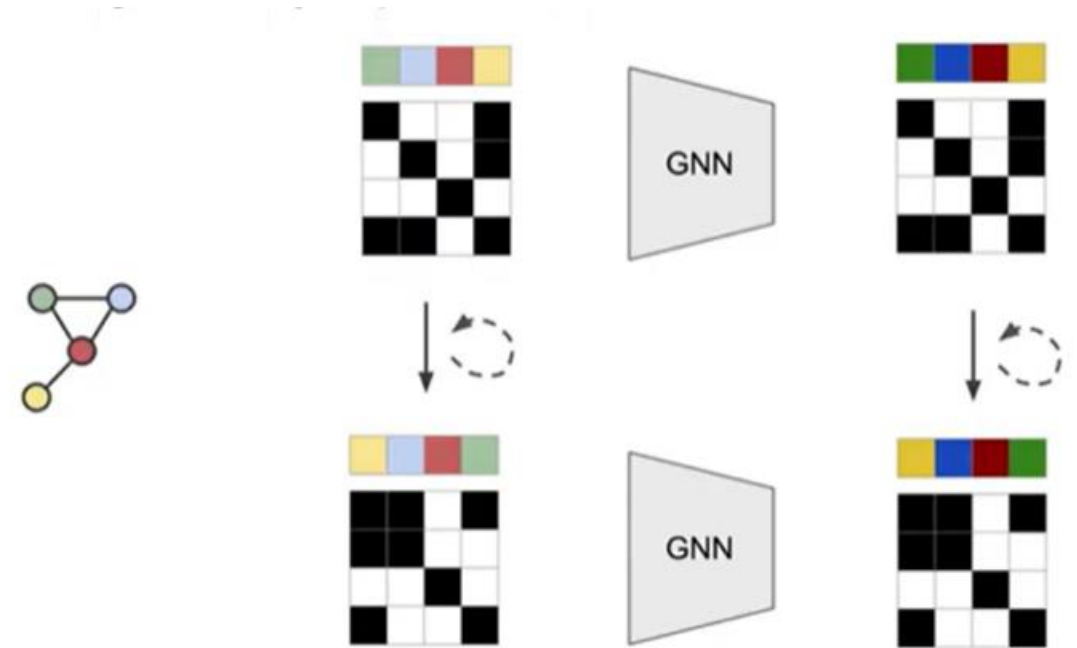
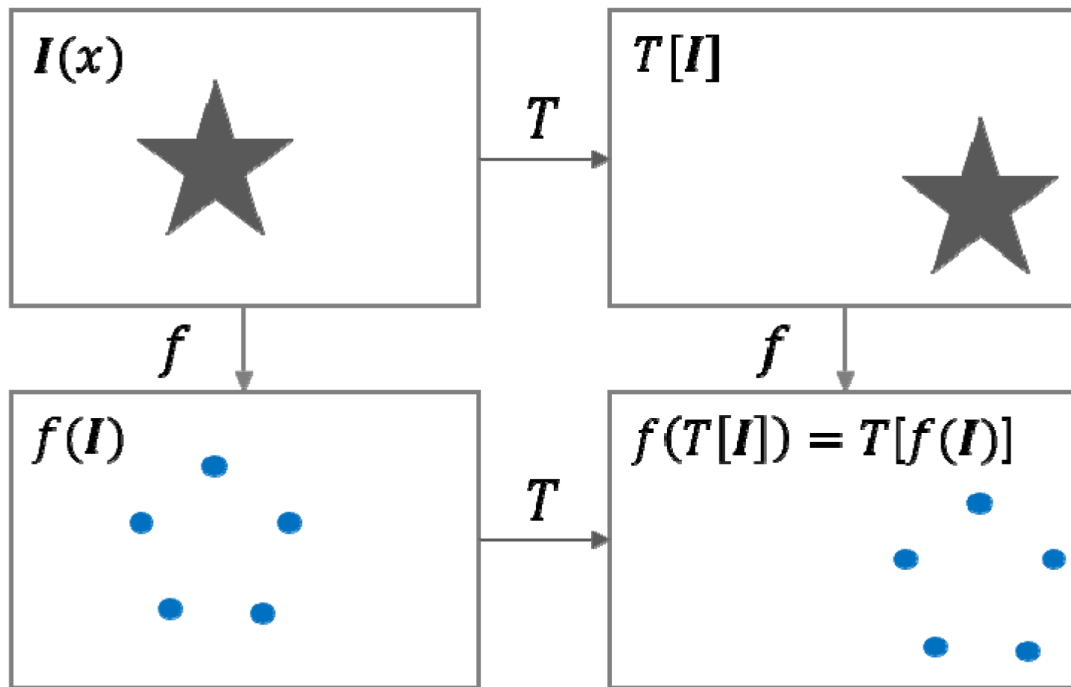
- A complex distribution  $p_x$
- A simple base distribution  $p_z$  (such as a Gaussian distribution)
- A **learnable invertible transformation  $f$**  :

$$\begin{aligned} x &= g(z) \\ z &= g^{-1}(x) = f(x) \end{aligned} \quad \rightarrow \quad \begin{aligned} p_x(x) &= p_z(z) |\det J_f(x)| \\ \log p_x(x) &= \log p_z(z) + \log |\det J_f(x)| \end{aligned}$$

## Background | E(n) Equivariant Normalizing Flows

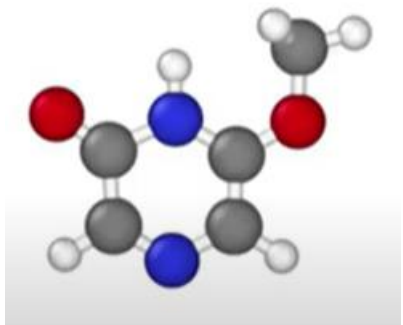
**Equivariance of a function  $f$ :** Transforming its input results in an equivalent transformation of its output.

$$f(T[I]) = T[f(I)]$$



**GNN is built to be equivalent to permutations:**  
Although we permute the indexes of nodes, it still represent the same graph.

## Background | E(n) Equivariant Normalizing Flows



### E(n) Equivariant data such as molecules in 3D:

A molecular graph  $G = (V, E)$  with nodes  $v_i \in V$  and edges  $e_{ij}$ .

Each node  $v_i$  is associated with a **position vector**  $x_i$  and **node features**  $h_i$ .

Features  $h$  have the property that they are invariant to E(n) transformations, while they do affect  $x$ .

In other words, **rotations and translations of  $x$  do not influence  $h$** .

### E(n) Equivariant Graph Neural Networks (EGNNs) (Satorras et al. ICML2021)



EGNNs is **equivariant to Euclidean symmetries**

(rotation, translation, reflection and displacement et al) .

However, this model is only able to discriminate features on nodes, and cannot generate new graphs.



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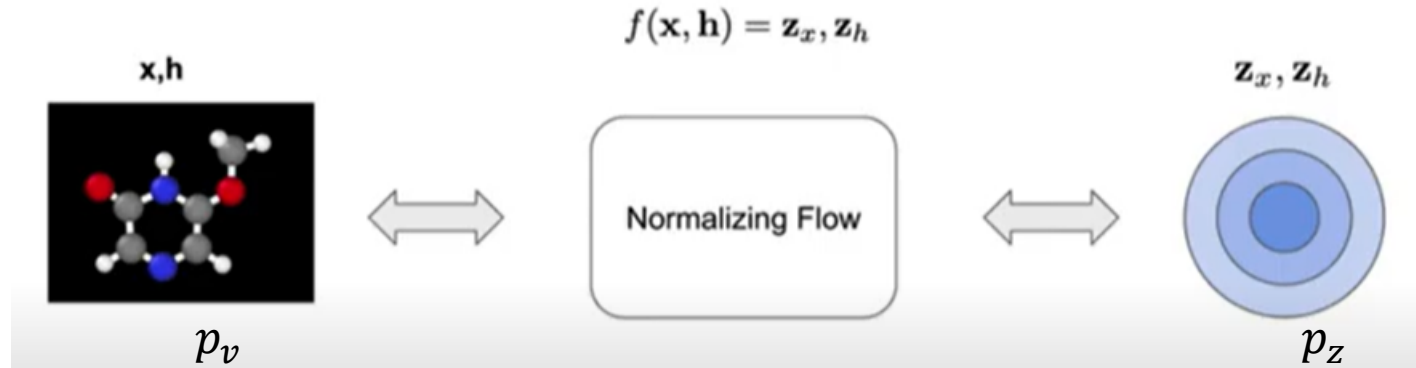
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# E(n) Equivariant Normalizing Flows

A probabilistic model for data with Euclidean symmetry.



$$V = (x, h)$$

- position vector  $x \in R^{M*n}$  embedded in a n-dimensional space
- node features  $h \in R^{M*nf}$

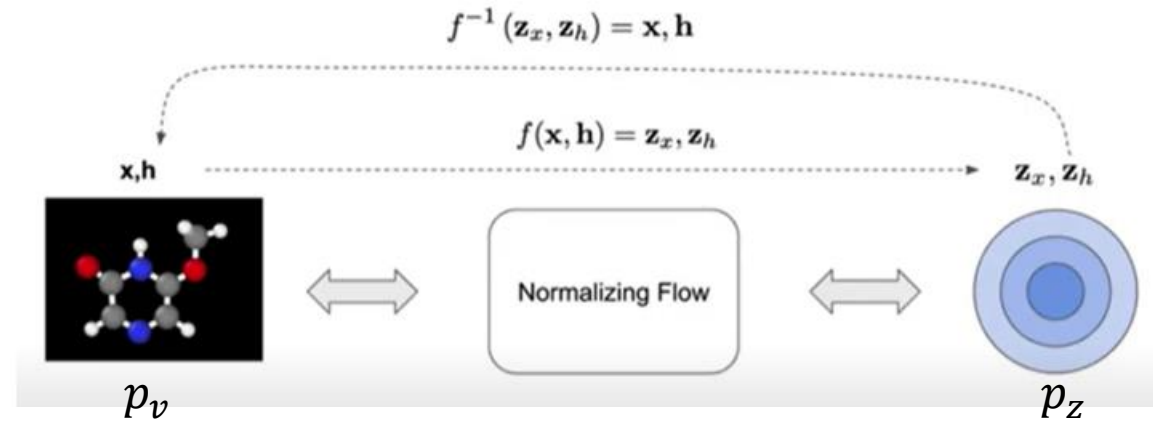
$p_z$  : A simple invariant distribution (such as a Gaussian)

$$f_\theta(x, h) = z_x, z_h \quad g_\theta(z_x, z_h) = x, h$$

$$f_\theta = g_\theta^{-1}$$

$$p_V(\mathcal{V}) = p_V(\mathbf{x}, \mathbf{h}) = p_Z(f_\theta(\mathbf{x}, \mathbf{h})) |\det J_f| = p_Z(\mathbf{z}_x, \mathbf{z}_h) |\det J_f|, \quad (6)$$

# E(n) Equivariant Normalizing Flows



$$p_V(\mathcal{V}) = p_V(\mathbf{x}, \mathbf{h}) = p_Z(f_\theta(\mathbf{x}, \mathbf{h})) |\det J_f| = p_Z(\mathbf{z}_x, \mathbf{z}_h) |\det J_f|, \quad (6)$$

- We require  $f$  to be invertible.
- We require  $f$  to be equivariant.

The solution to an ODE defined as:

$$\mathbf{z}_x, \mathbf{z}_h = f(\mathbf{x}, \mathbf{h}) = [\mathbf{x}(0), \mathbf{h}(0)] + \int_0^1 \phi(\mathbf{x}(t), \mathbf{h}(t)) dt. \quad (7)$$

Where  $x(0) = x, h(0) = h; x(1) = z_x, h(1) = z_h$

# E(n) Equivariant Normalizing Flows

The solution to an ODE defined as:

$$\mathbf{z}_x, \mathbf{z}_h = f(\mathbf{x}, \mathbf{h}) = [\mathbf{x}(0), \mathbf{h}(0)] + \int_0^1 \underbrace{\phi(\mathbf{x}(t), \mathbf{h}(t))}_{\downarrow} dt. \quad (7)$$

EGNN, but not original EGNN

- The original EGNN from (Satorras et al., ICML 2021) is unstable when utilized in an ODE because the coordinate update from Equation 5 would easily explode

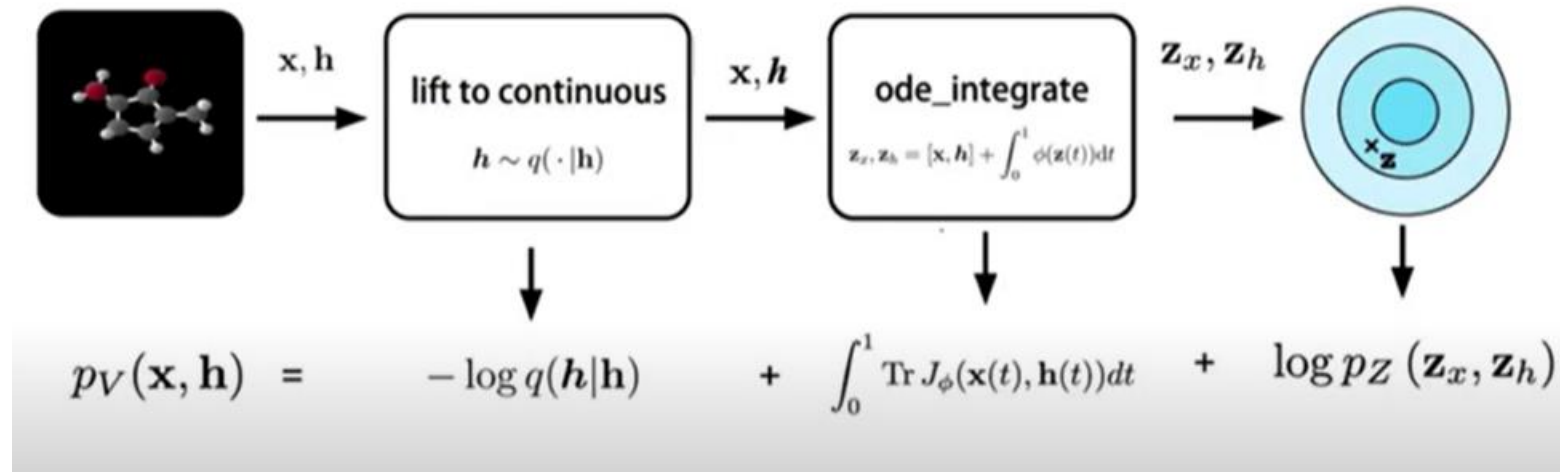
$$\mathbf{m}_{ij} = \phi_e \left( \mathbf{h}_i^l, \mathbf{h}_j^l, \|\mathbf{x}_i^l - \mathbf{x}_j^l\|^2 \right) \quad \text{and} \quad \mathbf{m}_i = \sum_{j \neq i} e_{ij} \mathbf{m}_{ij}, \quad (4)$$

$$\mathbf{x}_i^{l+1} = \mathbf{x}_i^l + \sum_{j \neq i} (\mathbf{x}_i^l - \mathbf{x}_j^l) \phi_x(\mathbf{m}_{ij}) \quad \text{and} \quad \mathbf{h}_i^{l+1} = \phi_h(\mathbf{h}_i^l, \mathbf{m}_i). \quad (5)$$

- An extension of the original EGNN from :

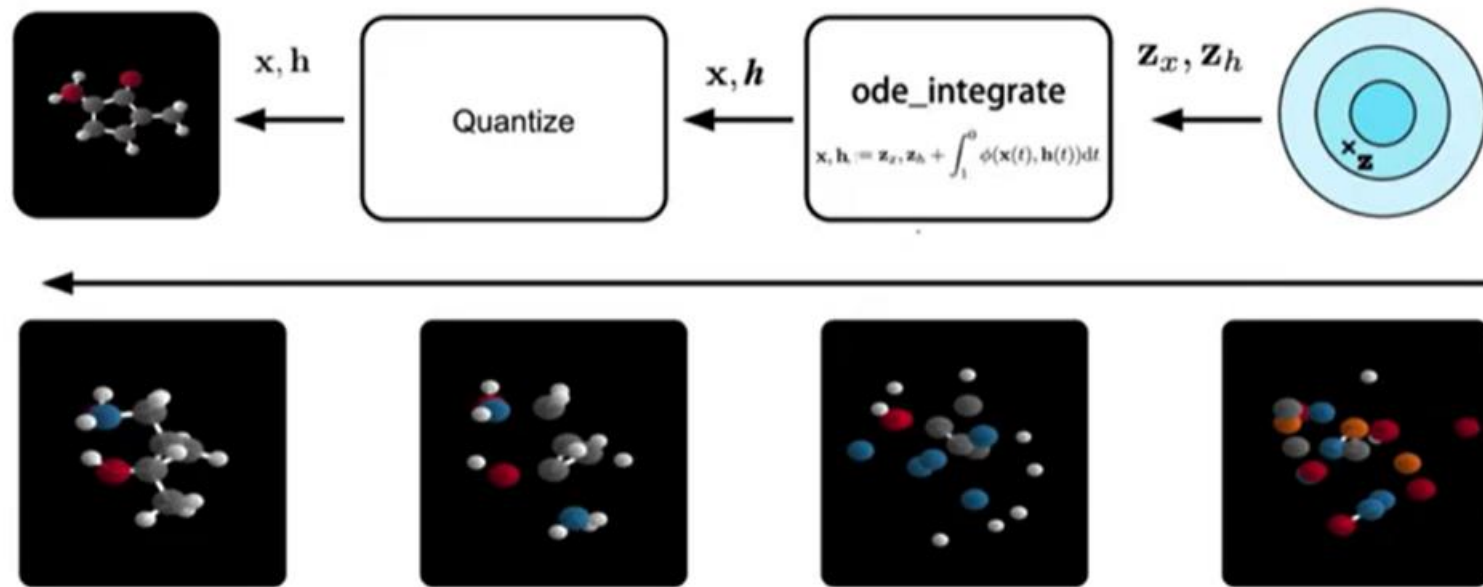
$$\mathbf{x}_i^{l+1} = \mathbf{x}_i^l + \sum_{j \neq i} \frac{(\mathbf{x}_i^l - \mathbf{x}_j^l)}{\|\mathbf{x}_i^l - \mathbf{x}_j^l\| + C} \phi_x(\mathbf{m}_{ij}) \quad (9)$$

## E(n) Equivariant Normalizing Flows | Training process



- Sample a point  $(x, h)$  from the dataset
- The discrete  $h$  is lifted to continuous  $h$
- The variables  $x, h$  are transformed by an ODE to  $z_x, z_h$
- Get the loglikelihood  $\log p_Z(z_x, z_h)$  on a simple invariant distribution (such as a Gaussian)

# E(n) Equivariant Normalizing Flows | Generation



- Sample a point  $(z_x, z_h)$  from the simple distribution
- The variables  $z_x, z_h$  are transformed by an ODE to  $x, h$
- Quantize

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# Experiments

- We compare to the state-of-the-art  $E(n)$  equivariant flows:
  - "Simple Dynamics"
  - "Kernel Dynamics" presented
- We also compare to non-equivariant variants of our method:
  - Graph Normalizing Flow (GNF)
  - GNF with attention (GNF-att)
  - GNF with attention and data augmentation (GNF-att-aug)



## Experiments | DW4 & LJ13

**DW4 and LJ13** have been synthetically generated by sampling from their respective energy functions using Markov Chain Monte Carlo (MCMC).

Table 1: Negative Log Likelihood comparison on the test partition over different methods on DW4 and LJ13 datasets for different amount of training samples averaged over 3 runs.

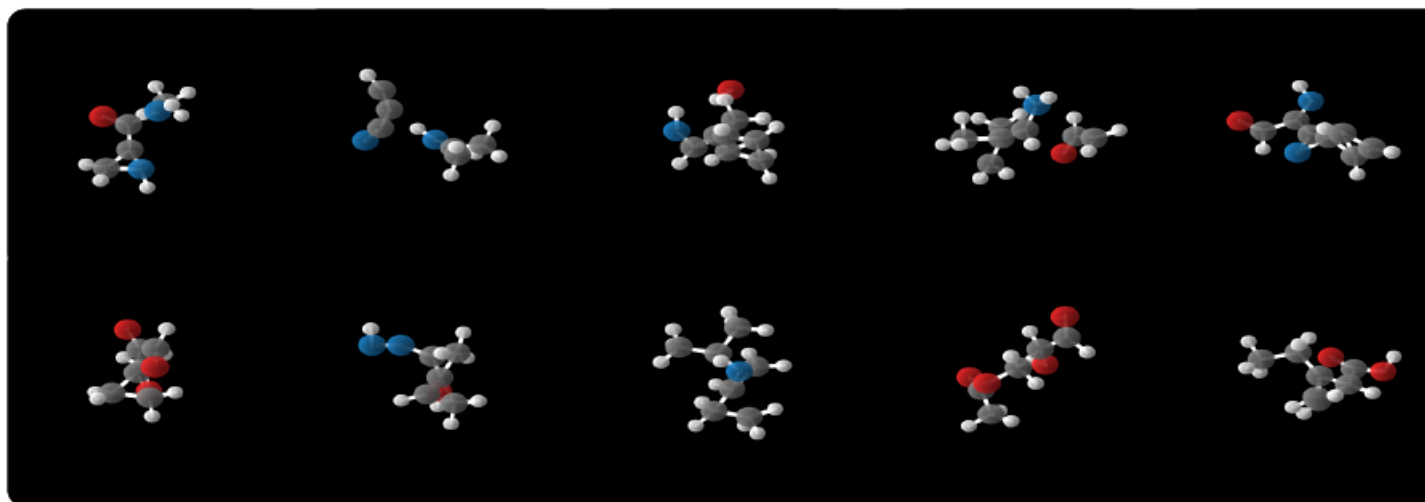
# Samples	DW4				LJ13			
	$10^2$	$10^3$	$10^4$	$10^5$	10	$10^2$	$10^3$	$10^4$
GNF	11.93	11.31	10.38	7.95	43.56	42.84	37.17	36.49
GNF-att	11.65	11.13	9.34	7.83	43.32	36.22	33.84	32.65
GNF-att-aug	8.81	8.31	7.90	7.61	41.09	31.50	30.74	30.93
Simple dynamics	9.58	9.51	9.53	9.47	33.67	33.10	32.79	32.99
Kernel dynamics	8.74	8.67	8.42	8.26	35.03	31.49	31.17	31.25
E-NF	<b>8.31</b>	<b>8.15</b>	<b>7.69</b>	<b>7.48</b>	<b>33.12</b>	<b>30.99</b>	<b>30.56</b>	<b>30.41</b>

# Experiments | QM9

QM9 (Ramakrishnan et al., 2014) is a molecular dataset in machine learning as a chemical property prediction benchmark

Table 2: Neg. log-likelihood  $-\log p_V(\mathbf{x}, \mathbf{h}, M)$ , atom stability and mol stability for the QM9 dataset.

# Metrics	NLL	Atom stability	Mol stable
<b>GNF-attention</b>	-28.2	72%	0.3%
<b>GNF-attention-augmentation</b>	-29.3	75%	0.5%
<b>E-NF (ours)</b>	<b>-59.7</b>	<b>85%</b>	<b>4.9%</b>
Data	-	99%	95.2%

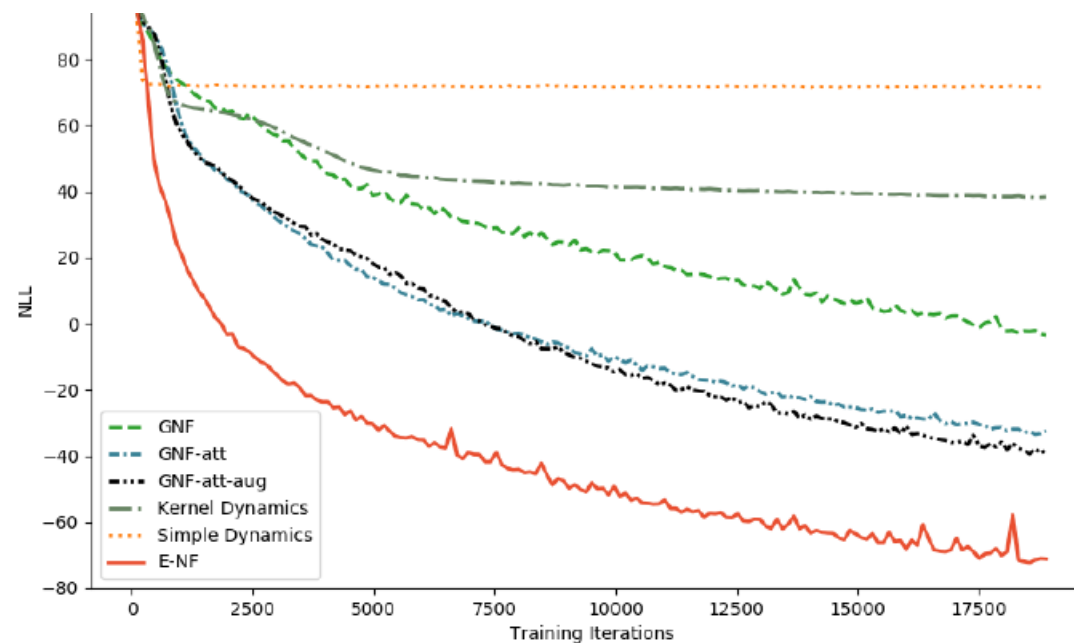


The top row contains random samples, the bottom row also contains samples but selected to be stable.

# Experiments | QM9 Positional

We introduce **QM9 Positional** as a subset of QM9 that only considers positional information and does not encode node features

# Metrics	NLL	JS(rel. dist)
Simple dynamics	73.0	.086
Kernel dynamics	38.6	.029
GNF	-00.9	.011
GNF-att	-26.6	.007
GNF-att-aug	-33.5	.006
<b>E-NF (ours)</b>	<b>-70.2</b>	.006



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# Conclusion

- The ODE type of flow makes the training expensive and slow since the same forward operation has to be done multiple times.
- E(n) Equivariant Normalizing Flows (E-NFs) are continuous-time normalizing flows that utilize an EGNN with improved stability as parametrization.
- This method may accelerate drug discovery by providing better drug candidate molecules.

**Thanks!**