

Symmetry-Preserving Diffusion Models via Target Symmetrization

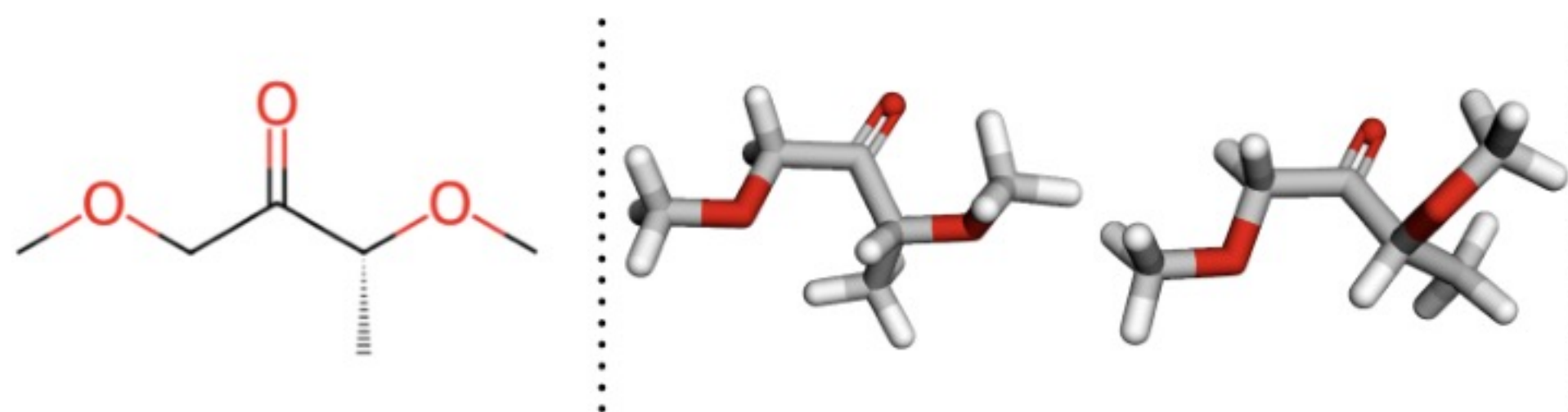
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We propose a **loss function** that enables **diffusion models** to respect **data symmetries** without **architectural constraints** or **data augmentation**, ensuring **stable, low-variance training**.

Introduction

Molecular graph

Symmetric conformer



- **Diffusion models** struggle with symmetries (e.g., rotations, reflections) since the **empirical dist.** is not necessarily **G-invariant**.

- Existing methods:

- **Equivariant denoisers** → complex to design.

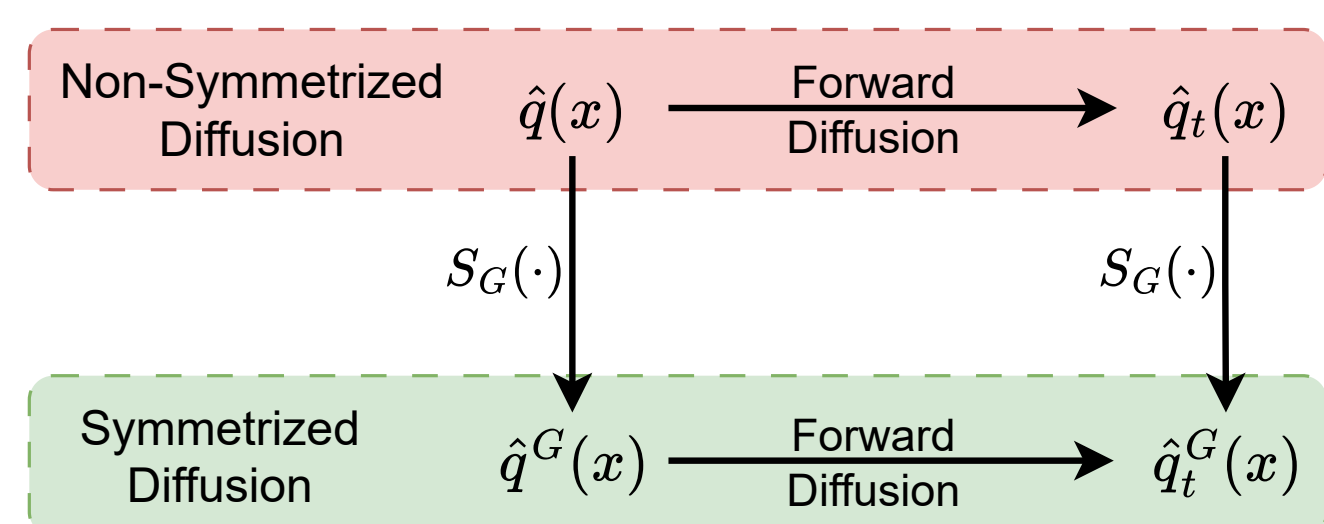
$$\mathcal{L}_t(\phi) = E_{(x_0, x_t) \sim \hat{q}_t(x_0, x_t)} [\omega(t) \|\phi(x_t, t) - x_0\|^2]$$

s.t. $\phi(g \circ x_t, t) = g \circ \phi(x_t, t) \quad \forall x_t \in \Omega \text{ and } \forall g \in G$

- **Data augmentation** to approximate G-invariant dist. → expensive.

$$\mathcal{L}_t^G(\phi) = E_{(x_0, x_t) \sim \hat{q}^G(x_0, x_t)} [\omega(t) \|\phi(x_t, t) - x_0\|^2]$$

- Our idea: **Symmetrized loss** → simple, stable, symmetry-aware.



Method

$$\bar{\mathcal{L}}_t^G(\phi) = E_{(x_0, x_t) \sim \hat{q}_t(x_0, x_t)} [\omega(t) \|\phi(x_t, t) - \phi_G^*(x_t, t)\|^2]$$

s.t. $\phi_G^*(x_t, t) = \int_{\Omega} x_0 \hat{q}_t^G(x_0 | x_t) dx_0$

- The minimizer is **provably G-equivariant**.
- **Reduced gradient variance** → stable training.

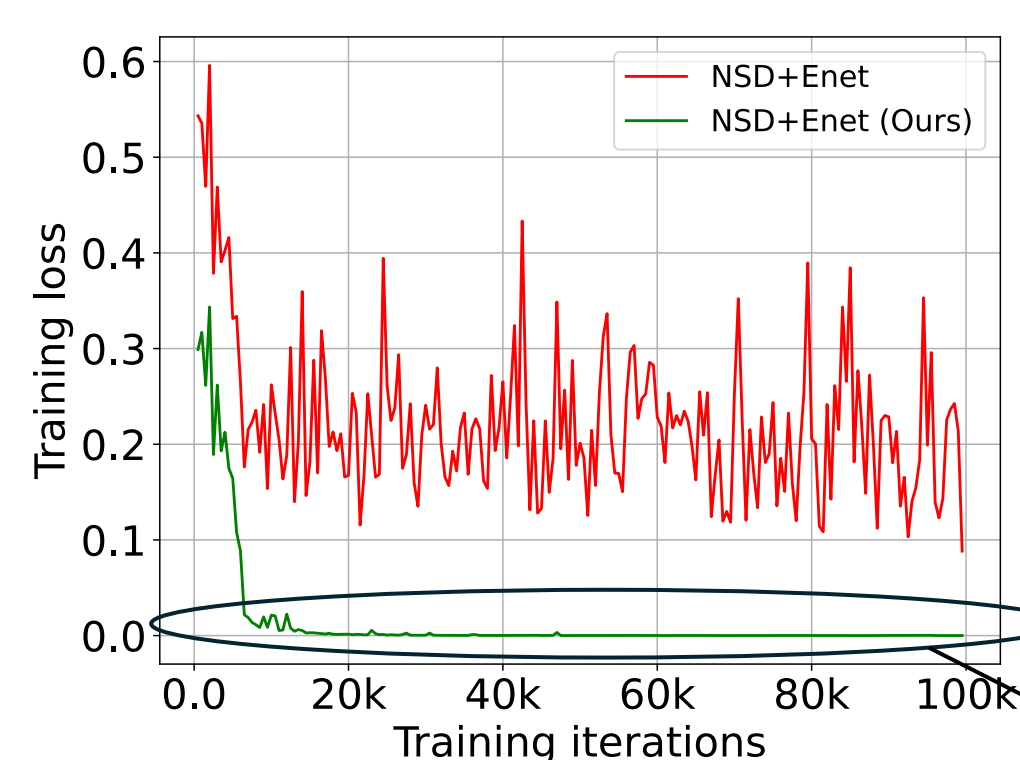
Algorithm 1 Approximation of $\phi_G^*(x_t, t)$

- 1: Draw M samples $x_0^m \sim \hat{q}_t(x_0)$, K group elements $g_k \sim \mu_G$
- 2: **for** each x_0^m, g_k **do**
- 3: $x_0^{mk} \leftarrow g_k \circ x_0^m, \quad R_{mk} \leftarrow \frac{-\|x_t - \alpha_t x_0^{mk}\|^2}{2\sigma_t^2}$
- 4: **end for**
- 5: $w_{mk} \leftarrow \frac{\exp(R_{mk})}{\sum_{j=1}^{MK} \exp(R_j)}, \quad \phi_G^*(x_t, t) \leftarrow \sum_{j=1}^{MK} w_j x_0^j$

Results

Reflection Group in 1D

Generate ± 1 from a single training sample using reflection symmetry.

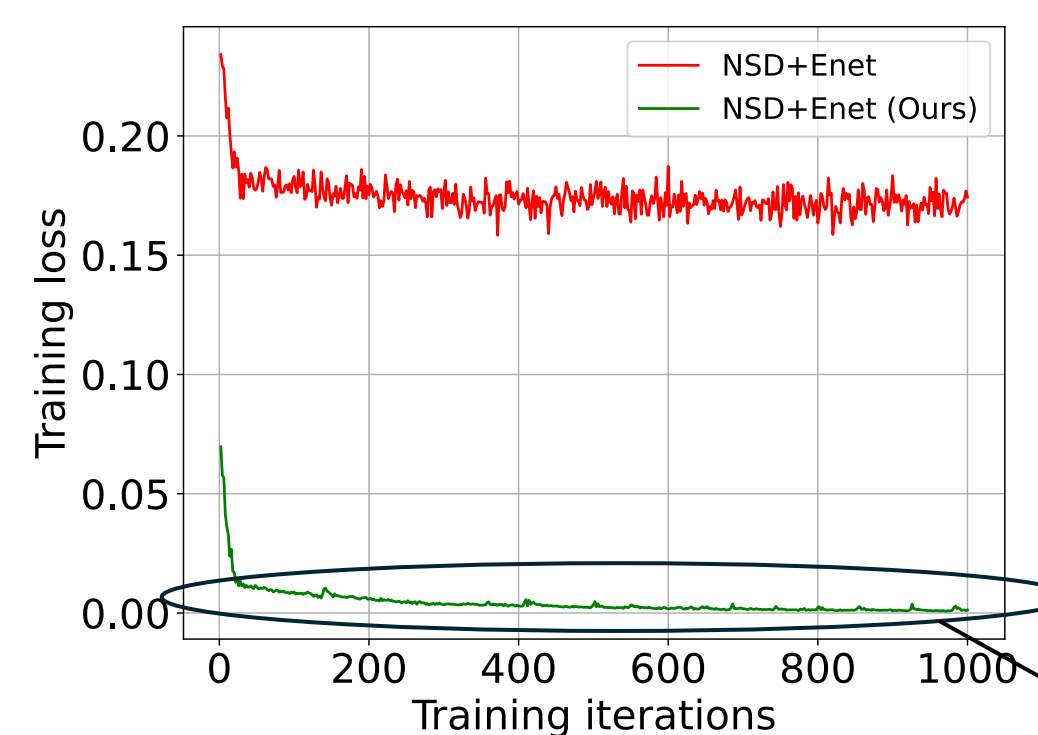


Our method converges faster with lower variance.

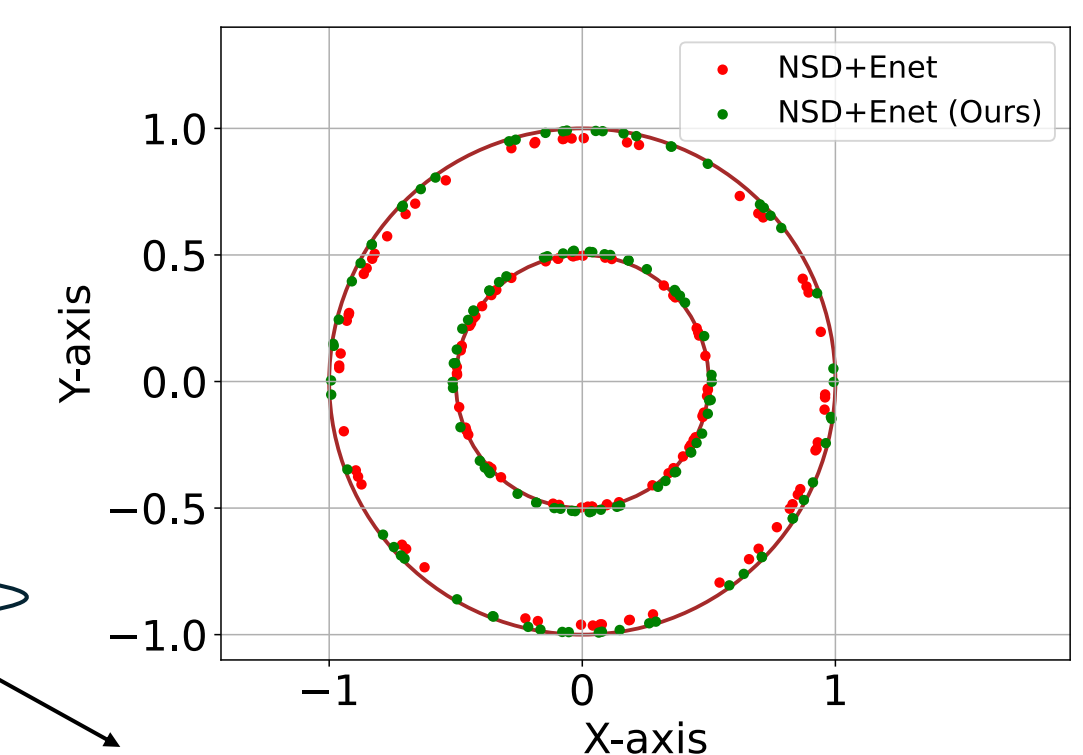
Models	RMSD ($\times 10^{-3}$)	W2 Distance ($\times 10^{-3}$)
NSD + Enet	8.392	0.014
NSD + Enet (Ours)	2.049	0.002

Rotation and Permutation Groups in 2D

Generate a 4-point 2D point cloud, ensuring equivariance to rotation and permutation.



Our method converges faster with lower variance.



Molecular Conformer Prediction

Models	Recall				Precision			
	Coverage $\uparrow$		AMR $\downarrow$		Coverage $\uparrow$		AMR $\downarrow$	
	M	Mdn	M	Mdn	M	Mdn	M	Mdn
ETDiff	80.7	100.0	0.092	0.038	71.4	75.0	0.162	0.093
ETDiff + Ours	81.0	100.0	0.089	0.037	72.2	80.0	0.151	0.077
ETFlow	83.8	100.0	0.082	0.032	77.7	90.0	0.130	0.049
ETFlow + Ours	84.7	100.0	0.079	0.028	79.1	92.5	0.122	0.046

Conclusion

Our method injects equivariance via a simple loss modification.

- No architectural changes.
- Minimal overhead.
- Strong theoretical backing.
- Better generalization on symmetric data.

