LIE GROUP-INDUCED DYNAMICS IN SCORE-BASED GENERATIVE MODELING

Anonymous authors

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ABSTRACT

We extend score-based generative modeling by incorporating Lie group actions on the data manifold into the denoising diffusion process. Our approach yields a Langevin dynamics whose infinitesimal transformations decompose as a direct sum of Lie algebra representations, enabling generative processes that align with the underlying symmetry properties of the data. Unlike equivariant models, which restrict the space of learnable functions by quotienting out group orbits, our method incorporates both global and local symmetries and can model any target distribution. Standard score-matching, which minimizes the Fisher divergence, emerges as a special case of our framework when the Lie group is the translation group in Euclidean space. We prove that our generalized generative processes arise as solutions to a new class of reverse-time stochastic differential equations (SDEs), introduced here for the first time. We validate our approach through experiments on diverse data types, demonstrating its effectiveness in real-world applications such as SO(3)-guided molecular conformer generation and modeling ligand-specific global SE(3) transformations for molecular docking. We show that an appropriate choice of Lie group enhances learning efficiency by reducing the effective dimensionality of the trajectory space and enables the modeling of transitions between complex data distributions, lifting the requirement of a Gaussian prior. Additionally, we demonstrate the universality of our approach by deriving how it extends to flow matching techniques.

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1 INTRODUCTION

033 Deep probabilistic generative modeling amounts to creating data from a known tractable prior distri-034 bution. Score-based models (Hyvärinen & Dayan, 2005; Sohl-Dickstein et al., 2015; Ho et al., 2020; Huang et al., 2021) achieve this by learning to reverse a corruption process of the data. The sampling process is realized by a Markov chain Monte Carlo (MCMC) using Langevin dynamics (Song et al., 2021; 2020b), producing a probabilistic trajectory $\{\mathbf{x}_t\}_{t=T,\ldots,0}$ in the data space X guided by the 037 score function $\mathbf{s}(\mathbf{x}) = \nabla_{\mathbf{x}} \log p(\mathbf{x})$: for each infinitesimal step $\mathbf{x}_{t+1} \mapsto \mathbf{x}_t - \mathbf{s}(\mathbf{x}_t) + \text{noise}$, the score induces a small translation in X (Figure 1a), and the noise prevents the dynamics from collapsing into local minima (Welling & Teh, 2011; Neal, 2010). The observation that translations are just one 040 of many transformations of Euclidean space leads to the central question of the present work: given 041 a group G acting on X, can we construct a score-based dynamics whose steps correspond to the 042 infinitesimal transformation of G? 043

We will prove that the answer is positive when G is a Lie group, as these are suitable to describe 044 local continuous transformations. Specifically, we construct a Langevin dynamics that decomposes as a direct sum of representations of the Lie algebra \mathfrak{g} of G, where each step is driven by tangent 046 vectors associated with elements of g (Figure 1)c. This construction employs the formalism of Gen-047 eralized Score Matching (GSM) (Lyu, 2009; Lin et al., 2016), whose aim is to estimate a probability 048 density $p(\mathbf{x}) = \exp(-U(\mathbf{x}))/Z$ through its generalized score function $\mathcal{L} \log p(\mathbf{x})$, where \mathcal{L} is a suitable linear operator. As Figure 1b suggest, a generalized score enables the dynamics to follow curved paths, a crucial feature for our scope since group orbits typically have nonzero curvature. 051 In this work we will systematically develop the connection between the transformation-generating Lie group G and the linear operator \mathcal{L} . We will also show that such generalized dynamics admits a 052 stochastic differential equations (SDE) interpretation: the G-induced generative process satisfies a continuous-(reverse-) time SDE involving the generalized score function.



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Figure 1: Standard score-matching (a) vs. generalized score-matching (b); (c) Trajectories for G = T(2) (dark blue, standard score-matching), and $G = SO(2) \times \mathbb{R}_+$ (dark orange). (d) Illustration of invariant, equivariant, standard and generalized score functions for a radial vector field in $X = \mathbb{R}^2$.

At this point, the attentive reader might wonder whether all we are achieving is to merely introduce multiple ways, according to the choice of G, to achieve the same goal of generating samples from a target distribution. So, what's the real benefit? This lies in the ability to tailor G to the specific transformation properties of the data and the learning task: the score components can be designed to align with the true degrees of freedom of the data, enhancing interpretability, sampling steerability, and enabling dimensionality reduction by leveraging the data's inherent symmetries.

Figure 1d illustrates this in a simple but instructive case. Consider $X = \mathbb{R}^2$ and let the true score be of the form $\mathbf{s}(\mathbf{x}) = f(r, \theta)\hat{\mathbf{r}}$. This is neither invariant nor equivariant since f depends on both r, θ , thus it cannot be learned by an invariant/equivariant network. A standard score function neglects the problem's symmetry and is required to inefficiently (and likely inaccurately) learn the correct pointdependent linear combination of both Cartesian components of the score. A SO(2) generalized score function, however, directly learns the radial component $f(r, \theta)$, effectively reducing the problem to a 1-dimensional formulation, as the angular component may be omitted from the leaning procedure.

In short, we propose an exact SDE-based diffusion framework that enables Lie group-guided curved dynamics while operating entirely in Euclidean space, thus combining the advantages of curved dynamics with the theoretical and practical effectiveness of Euclidean diffusion. Our framework in unconstrained, unlike equivariant models, but still leverages the information of the group action on the data space. Ours is the first methods we are aware of that realizes **simulation-free** training of Lie group-like diffusion models, and the **first result of denoising score-matching result for general non-Abelian groups** (unlike De Bortoli et al. (2022) and Huang et al. (2022)).

⁰⁹¹ More specifically, we summarize the main contributions of this manuscript as follows:

- **Generalized score matching via Lie algebras:** We develop the mathematical foundations for extending score-based generative modeling to incorporate Lie group-induced dynamics. We spell out the conditions for a suitable G (valid for any differentiable manifold X) and demonstrate that the dynamics decomposes as a linear combination of Lie algebra representations. We also show that standard score-matching is recovered as a specific case of our framework, corresponding to the group G = T(n) of translations on $X = \mathbb{R}^n$.
- Exact solution of a novel class of SDEs: We introduce a *new class of solvable* SDEs that govern symmetry-aware processes in Euclidean space, significantly expanding the range of processes that can be addressed using diffusion modeling techniques. Additionally, we show that our approach extends naturally to flow matching (see Appendix D).
- **Dimensionality reduction, bridging non-trivial distributions and trajectory disentanglement:** Through extensive experiments*, we demonstrate that: (1) our approach can estimate, regardless of the choice of G, any probability density (Sections 5(2,3,4d distributions) and 5)(QM9); (2) by appropriately selecting G to align with the data structure, the learning process is significantly simplified, effectively reducing its dimensionality (Section 5(MNIST)) (3) our framework enables solutions to processes that are challenging or unfeasible with standard score matching, such as bridging between



Figure 2: (a) Depiction of the fundamental vector field definition (1). Flow coordinates for a pair of commuting (b) and not-commuting ones vector fields (c).

2 DIFFUSION DYNAMICS THROUGH LIE ALGEBRAS

122 We start this section by setting up notation and review the connection between vector fields and Lie algebra actions on manifolds. A Lie group G is a group that is also a finite-dimensional differentiable 123 manifold, such that the group operations of multiplication $\cdot: G \times G \to G$ and inversion are C^{∞} -124 functions[†]. A Lie algebra g is a vector space equipped with an operation, the Lie bracket, [,]: 125 $\mathfrak{g} \times \mathfrak{g} \to \mathfrak{g}$, satisfying the Jacobi identity. Every Lie group gives rise to a Lie algebra as its tangent 126 space at the identity, $\mathfrak{g} = T_e G$, and the Lie bracket is the commutator of tangent vectors, [A, B] =127 AB - BA. In this work, we are interested in how Lie groups and Lie algebras act on spaces. Given 128 a manifold X, a (left) group action of G on X is an associative map $\rho_X : G \times X \to X$ such 129 that $\rho_X(e) = Id_X$. Fundamental concepts associated with a group action are the ones of orbits and 130 stabilizers. The **orbit** of $\mathbf{x} \in X$ is the set of elements in X which can be reached from \mathbf{x} through 131 the action of G, i.e., $G \cdot \mathbf{x} = \{\rho_X(g)(\mathbf{x}), g \in G\}$. The stabilizer subgroup of G with respect to \mathbf{x} 132 is the set of group elements that fix \mathbf{x} , $G_x = \{g \in G | \rho_X(g)(\mathbf{x}) = \mathbf{x}\}$. The action of a Lie algebra 133 on X, $\mathfrak{A} : \mathfrak{g} \to \operatorname{Vect}(X)$ is a Lie algebra homomorphism and maps elements of \mathfrak{g} to vector fields on X such that the map $\mathfrak{g} \times X \to TX$, $(A, \mathbf{x}) \mapsto \mathfrak{A}(A)(\mathbf{x})$ is smooth. Given $A \in \mathfrak{g}$ and a group action 134 ρ_X , the flow on X induced by ρ_X is given by $\xi_A : X \times \mathbb{R} \to X, (\mathbf{x}, \tau) \to \rho_X (\exp(\tau A))(\mathbf{x})$, 135 where the map exp : $\mathfrak{g} \to G$ is defined by $\exp(A) = \gamma_A(1)$, where $\gamma_A \colon \mathbb{R} \to G$ is the unique 136 one-parameter subgroup of G whose tangent vector at the identity is A. The **infinitesimal action** of 137 g on X is defined as the differential of the map ρ_X , that is 138

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$$d\rho_X : \mathfrak{g} \to \operatorname{Vect}(X) : A \mapsto \frac{d}{d\tau} \Big|_{\tau=0} \rho_X(\exp(\tau A))(\mathbf{x}) \equiv \Pi_A(\mathbf{x}) .$$
 (1)

 Π_A is called the **fundamental vector field** corresponding to $A \in \mathfrak{g}$. Given a fixed point $\mathbf{x}_0 \in X$, 142 we denote $\tau = \xi_A(\mathbf{x}_0)^{-1}(\mathbf{x})$ the **fundamental flow coordinate**, which is the parameter such that 143 applying the flow to x_0 gives x. Central to our discussion is the fact that any smooth vector field 144 $V: X \rightarrow TX$ on X can be interpreted as a differential operator acting on smooth functions 145 $f: X \to \mathbb{R}$. The operator V(f) represents the directional derivative of f at $\mathbf{x} \in X$ in the direction 146 of $V(\mathbf{x})$. We denote $\mathcal{L}_A = \prod_A \cdot \nabla$ the differential operator corresponding to \prod_A . In the following 147 we will use both Π_{τ} and Π_A interchangeably, when no potential confusion arises. When dim $\mathfrak{g} > 1$ 148 we indicate as $\mathbf{\Pi}(\mathbf{x}) = (\Pi_{A_1} \quad \Pi_{A_2} \quad \cdots)$ the matrix of the collection of fundamental vector fields. 149 Let us work out the example for $X = \mathbb{R}^2$ and G = SO(2), the group of rotations in the plane. 150 The Lie algebra $\mathfrak{so}(2)$ consists of all matrices of the form $A_{\alpha} = \begin{pmatrix} 0 & -\alpha \\ \alpha & 0 \end{pmatrix}$, where $\alpha \in \mathbb{R}$, and 151 152 the Lie bracket is identically zero. The flow on X induced by ρ_X is given by the exponential map 153 $\rho_{\mathbb{R}^2}(\exp(\tau A_\alpha))(\mathbf{x}) = \begin{pmatrix} \cos(\alpha\tau) & -\sin(\alpha\tau)\\ \sin(\alpha\tau) & \cos(\alpha\tau) \end{pmatrix} \mathbf{x}, \text{ and without loss of generality we can set } \alpha = 1.$ 154 155 The infinitesimal action is computed as 156

$$d\rho_{\mathbb{R}^2}(A) = \frac{d}{d\tau} \bigg|_{\tau=0} \begin{pmatrix} \cos\tau & -\sin\tau\\ \sin\tau & \cos\tau \end{pmatrix} \begin{pmatrix} x_1\\ x_2 \end{pmatrix} = \begin{pmatrix} -x_2\\ x_1 \end{pmatrix}.$$
 (2)

^{*}The code to reproduce our results will be open-sourced upon publication.

^{\dagger}We restrict ourselves here to real Lie groups. It would be interesting to extend the analysis presented here to the complex case as well (Le et al., 2021).

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and thus the fundamental vector field defines the derivation $\mathcal{L}_A(\mathbf{x}) = -x_2 \frac{\partial}{\partial x_1} + x_1 \frac{\partial}{\partial x_2}$. Let $\mathbf{x}_0 \in \mathbb{R}^2$ be a fixed point, then the flow equation $\mathbf{x}(\tau) \equiv \xi_A(\mathbf{x}_0, \tau) = \rho_{\mathbb{R}^2}(\exp(\tau A), \mathbf{x}_0)$ gives a system of two equations, which we can solve to find the expression of the fundamental flow coordinate

$$\begin{aligned} \mathbf{x} \cdot \mathbf{x}_0 &= |\mathbf{x}_0|^2 \cos \tau , \\ \mathbf{x} \times \mathbf{x}_0 &= |\mathbf{x}_0|^2 \sin \tau , \end{aligned} \Rightarrow \quad \tau = \arctan \frac{\mathbf{x} \times \mathbf{x}_0}{\mathbf{x} \cdot \mathbf{x}_0} . \end{aligned}$$
(3)

168 169 where $\mathbf{x} \times \mathbf{y} = y_2 x_1 - x_1 y_2$. Note that $\frac{\partial}{\partial \tau} = \frac{\partial x_1}{\partial \tau} \frac{\partial}{\partial x_1} + \frac{\partial x_2}{\partial \tau} \frac{\partial}{\partial x_2} = -x_2 \frac{\partial}{\partial x_1} + x_1 \frac{\partial}{\partial x_2} = \Pi_A(\mathbf{x})^\top \nabla =$ 170 \mathcal{L}_A .

2.1 INTUITION BEHIND LIE GROUP-INDUCED GENERALIZED SCORE MATCHING

Score matching aims at estimating a (log) probability density $p(\mathbf{x})$ by learning to match its score function, i.e., its gradient in data space. Generalized score matching replaces the gradient operator with a general linear operator \mathcal{L} . The learning objective is given by minimizing the generalized Fisher divergence

$$D_{\mathcal{L}}(p||q_{\theta}) = \int_{X} p(\mathbf{x}) \left| \mathcal{L} \log p(\mathbf{x}) - \mathbf{s}_{\theta}(\mathbf{x}) \right|^{2} d\mathbf{x} , \qquad (4)$$

where $\mathbf{s}_{\theta} = \mathcal{L} \log q_{\theta}$. The requirement on the choice of \mathcal{L} is that it preserves all the information about the original density. Formally, we require \mathcal{L} to be *complete*, that is, given two densities $p(\mathbf{x})$ and $q(\mathbf{x}), \mathcal{L}p(\mathbf{x}) = \mathcal{L}q(\mathbf{x})$ (almost everywhere *) implies that $p(\mathbf{x}) = q(\mathbf{x})$ (almost everywhere).

Given a Lie group G acting on X, the collection of fundamental fields Π corresponding to a choice of basis $\mathbf{A} = (A_1, A_2, ...)$ of \mathfrak{g} is a linear operator, thus potentially suitable for score-matching. It is then natural to set \mathcal{L} to the derivation associated with the fundamental fields Π , i.e., $\mathcal{L} = \Pi(\mathbf{x})^\top \nabla$. It then follows that $\mathcal{L} \log p(\mathbf{x})$ computes the directional derivatives of $\log p(\mathbf{x})$ with respect to the fundamental flow coordinates τ , and provided that Π meets some consistency conditions (which we will address in the next section), we can employ $\mathcal{L} \log p(\mathbf{x})$ to sample from $p(\mathbf{x})$ using Langevin dynamics:

$$\mathbf{x}_{t+1} = \mathbf{x}_t - \mathbf{s}_{\theta}(\mathbf{x}_t) d\rho_X(\exp(\boldsymbol{\tau} \mathbf{A}))(\mathbf{x}_t) = \mathbf{x}_t - \sum_i \underbrace{\mathcal{L}_i \log p_t(\mathbf{x}_t)}_{\text{generalized scores } A_i \text{ directions}} \underbrace{\Pi_{A_i}(\mathbf{x}_t)}_{A_i \text{ directions}} \Delta t , \qquad (5)$$

where Δt is the step size and we have temporarily set aside stochasticity and denoising aspects. This process mirrors the example depicted in Figure 1c: each infinitesimal step of the dynamics corresponds to infinitesimal transformations along the flow on X induces by the G-action, and each component of the generalized score is learned through maximum likelihood over the orbits ξ_{A_i} of the corresponding transformations.

2.2 Sufficient conditions for Lie group-induced generalized score matching

We now address the properties our setup $(X, G, \mathfrak{g} \Pi)$ must satisfy to meet the sufficient conditions for score-matching and Langevin dynamics. We note that these result hold for any differentiable manifold X. Proofs for these results can be found in Appendix B.

Condition 1: Completeness of II. We start by establishing an algebraic-geometric condition for Π 's completeness:

Proposition 2.1. The linear operator induced by Π is complete if Π is the local frame of a vector bundle E over X whose rank is $n \ge \dim X$ almost everywhere. If rank E = n everywhere, then E = TX, the tangent bundle of X.

The following result specifies which Lie groups yield operators Π satisfying the above proposition:

Proposition 2.2. The operator Π induced by \mathfrak{g} is complete if and only if the subspace $U \subseteq X$ such that $\dim \frac{G}{G_{\mathbf{x}}} < n$ for $\mathbf{x} \in U$, where $n = \dim X$, has measure zero in X.

^{*}Almost everywhere means everywhere except for a set of points of measure zero, where we assume the standard Lebesgue measure.

As an example, consider standard score-matching on mass-centered point clouds. Here $X = \mathbb{R}^{3N-3}$, since the points' coordinates satisfy $\sum_{i=1}^{N} \mathbf{x}_i = 0$. Without loss of generality, X can be parametrized by $\mathbf{x}_{1,\dots,N-1}$, with \mathbf{x}_N determined by the center of mass condition. The group G = T(3N) acts transitively on X, with a 3-dimensional stabilizer subgroup $G_X = \{(0,\dots,0,\mathbf{a})^\top \in \mathbb{R}^{3N}\}$ fixing the space. Thus, dim $G/G_X = n$ for all $\mathbf{x} \in X$, satisfying Proposition 2.2.

222 **Condition 2: Homogeneity of** X. While the completeness of the operators is necessary for esti-223 mating the target density, it is not sufficient to ensure that the Langevin dynamics (5) will behave ap-224 propriately, as the following example illustrates. Let $X = \mathbb{R}$, and $G = \mathbb{R}^{+}_{+}$, the multiplicative group 225 of non-zero positive real numbers. The orbits under the action $\rho_X(a, x) = ax$ are $\mathcal{O}_+ = (0, \infty)$, $\mathcal{O}_{-} = (-\infty, 0)$, and $\mathcal{O}_{0} = \{0\}$. If the dynamics begins within \mathcal{O}_{+} , it will be never be able to reach 226 values in \mathcal{O}_{-} , as G-transformations cannot move the system outside its initial orbit. We therefore 227 ask that each pair of points of X is connected through the G action. This amounts to require that 228 X is homogeneous for G, that is, $\forall \mathbf{x}, \mathbf{y} \in X$ there exists a $g \in G$ such that $\rho_X(g)\mathbf{x} = \mathbf{y}$. We 229 note that this condition solely ensures the generation outcome is independent of the initial sampling 230 condition. Beyond this, the formalism remains fully applicable in the non-homogeneous case. 231

232 **Condition 3: Commutativity of II.** The final requirement is that Π forms a (locally) commuting 233 frame of vector fields, $[\mathcal{L}_A, \mathcal{L}_B]f(\mathbf{x}) = 0 \ \forall A, B$ and $\forall f \in C^{\infty}(X)$. In this case, the coordinates 234 τ_i 's are orthogonal, and their flows commute, meaning the orbits parametrized by τ_i correspond 235 to $\{\tau_i = 0\}_{i \neq i}$. For non-commuting flows this is not the case, as Figure 2b-c illustrates: (b) 236 $V_1 = x_1\partial_{x_1} + x_2\partial_{x_2}, V_2 = x_1\partial_{x_2} - x_2\partial_{x_1}$ satisfy $[V_1, V_2] = 0$, and the orbits parametrized by $\tau_1 = r$ correspond to subspaces with constant $\tau_2 = \dot{\theta}$; (c) $\vec{W}_{1,2} = V_{1,2}/|\mathbf{x}|$ do not commute, and 237 238 the loci θ = const no longer coincide with the r-orbits, causing θ to vary along these, despite the 239 fact that r, θ are still orthogonal at each point. This last condition ensures that the updates governed by the different elements A_i of g in (5) remain independent of one another. Notably, this does not 240 exclude non-Abelian groups; even if $A_{1,2} \in \mathfrak{g}$ do not commute in the Lie algebra, their flows on X 241 can, as shown in the $g = \mathfrak{so}(3)$ example in Appendix A.3. 242

3 LIE ALGEBRA SCORE-BASED GENERATIVE MODELING VIA SDES

In this section we formalize the framework we developed above from the point of view of SDEs and
derive a Lie group curved dynamics in Euclidean space. Namely, we show that there exists a class of
SDEs, which, when reversed, can generate data according to dynamics similar to (5), guided by the
generalized score of the fundamental vector fields of the Lie algebra g. Our main result is provided
by the following

Theorem 3.1. Let G be a Lie group acting on X satisfying the conditions of Section 2.2, and let g be its Lie algebra. The pair of SDEs

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$$d\mathbf{x} = \left[\beta(t)\mathbf{\Pi}(\mathbf{x})\mathbf{f}(\mathbf{x}) + \frac{\gamma(t)^2}{2}\rho_X(\Omega)\right]dt + \gamma(t)\mathbf{\Pi}(\mathbf{x})d\mathbf{W}, \qquad (6)$$

$$d\mathbf{x} = \left[\beta(t)\mathbf{\Pi}(\mathbf{x})\mathbf{f}(\mathbf{x}) - \frac{\gamma^2(t)}{2}\rho_X(\Omega) - \gamma^2(t)\mathbf{\Pi}(\mathbf{x})\nabla^\top \cdot \mathbf{\Pi}(\mathbf{x}) - \gamma(t)^2\mathbf{\Pi}(\mathbf{x})\mathcal{L}\log p_t(\mathbf{x})\right]dt + \gamma(t)\mathbf{\Pi}(\mathbf{x})d\mathbf{W}, \quad (7)$$

where $\beta, \gamma : \mathbb{R} \to \mathbb{R}$ are time-dependent functions, $\Pi : \mathbb{R}^n \to \mathbb{R}^{n \times n}$ the fundamental vector fields, $\mathbf{f} : \mathbb{R}^n \to \mathbb{R}^n$ the drift, $\Omega = \sum_i A_i^2$ is known as the quadratic Casimir element of \mathfrak{g} , and $\mathcal{L} = \Pi(\mathbf{x})^\top \nabla$, is such that

1. The forward-time SDE (6) is exactly solvable, with solution

$$\mathbf{x}(t) = \left(\prod_{i} O_i(\tau_i(t))\right) \mathbf{x}(0) = \left(\prod_{i=1}^n e^{\tau_i(t)A_i}\right) \mathbf{x}(0) , \qquad (8)$$

where $O_i = e^{\tau_i(t)A_i}$ is the finite group action and $\boldsymbol{\tau}(t)$ is the solution to the SDE

$$d\boldsymbol{\tau}(\mathbf{x}) = \beta(t)\mathbf{f}(\mathbf{x})dt + \gamma(t)d\mathbf{W}.$$
(9)

2. The SDE (7) is the reverse-time process of (6).

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316 317 3. The Langevin dynamic of the above SDEs decomposes as a direct sum of \mathfrak{g} infinitesimal actions (1), each defining an infinitesimal transformation along the flows ξ_{τ} .

274 We refer to Appendix C for the proof of the above result. Here we limit 275 ourselves to a few comments regarding the extra terms that appear in 276 the SDEs. The appearance of the Casimir element (we assume the 277 identity as bilinear form on g (Kac & Kac, 1983)) compensates for the 278 deviation of the tangent vector from the orbit due to the curvature of 279 the flow coordinates. This can be seen in the example of SO(2) acting 280 on \mathbb{R}^2 (which will be discussed thoroughly below). An infinitesimal transformation along the θ direction, represented by Π_{θ} , moves any 281 point x along a vector tangent to its SO(2) orbit, a circle of radius 282 $r = \sqrt{x_1^2 + x_2^2}$. Due to the orbit's non-zero curvature, this movement 283 would shift the point to an orbit of radius r' > r. The term $\rho_X(\Omega)$ 284 compensates for this displacement, ensuring the final point remains 285 close to the original orbit. This is illustrated in Figure 3. 286



Figure 3: Quadratic Casimir for G = SO(2).

287 With this result at hand we can formulate our procedure for our Lie 288 group-induced score-based generative modeling with SDEs.

Perturbing data through the SDE. The forward-time SDE (6) defines a noising diffusion process 290 respecting the decomposition of the Lie algebra \mathfrak{g} infinitesimal actions on X. In fact, given a data 291 sample $\mathbf{x}(0) \sim p_0$, the solution (8) takes the form of a product of finite group element actions 292 \mathcal{O}_i on $\mathbf{x}(0)$, where the specific order is irrelevant since the Lie algebra generators commute. For 293 each factor, we first determine $\tau(0) = \tau(\mathbf{x}(0))$, and employ these as initial conditions for the forward SDE (9). By choosing appropriately the drift terms $f'_i s$, for instance, to be affine in the 295 flow coordinates τ_i , we can solve for $\tau(t)$ with standard techniques (Särkkä & Solin, 2019), as 296 this will follow a Gaussian distribution. Alternatively, we can sample from $\tau(t)$ by first simulating 297 (9), then performing sliced score matching Song et al. (2020a); Pang et al. (2020) to sample from 298 $p_t(\mathbf{x}(\boldsymbol{\tau}(t))|\mathbf{x}(0)).$ 299

300 Generating samples through the reverse SDE. The time-reverse SDE (7) guides the generation 301 of samples $\mathbf{x}(0) \sim p_0(\mathbf{x})$ starting from samples $\mathbf{x}(T) \sim p_T(\mathbf{x})$, provided we can estimate the generalized score $\mathcal{L} \log p_t(\mathbf{x})$ of each marginal distribution. To sample from p_T , we use the fact that 302 the distribution in the flow coordinates τ is tractable (with an appropriate choice of the drift terms 303 and time-dependent functions β , γ in (6)), and that (since $p_t(\mathbf{x})d\mathbf{x} = p_t(\boldsymbol{\tau})d\boldsymbol{\tau}$) 304

$$p_t(\mathbf{x}) = p_t(\boldsymbol{\tau}) \left| \frac{\partial \boldsymbol{\tau}}{\partial \mathbf{x}} \right| = p_t(\boldsymbol{\tau}) \left| \boldsymbol{\Pi}^{-1}(\mathbf{x}) \right| , \qquad (10)$$

307 where the extra term corresponds to the determinant of the Jacobian of the coordinate transformation 308 induced by the fundamental flow coordinates. In particular, when $f(\tau)$ is affine, it follows that 309 $p_T(\boldsymbol{\tau}) = \mathcal{N}(\boldsymbol{\tau} \mid \mathbf{0}, \boldsymbol{\Sigma})$, where $\boldsymbol{\Sigma} = \text{diag}(\sigma_1^2, \sigma_2^2, \dots, \sigma_n^2)$. Thus, we can sample $\boldsymbol{\tau}(T) \sim p_T(\boldsymbol{\tau})$ 310 simply as a collection of independent Gaussian random variables, and use the flow map to obtain 311 $\mathbf{x}(T) = \boldsymbol{\xi}_{\mathbf{A}}(\boldsymbol{\tau}(T), \mathbf{x}_0)$, which will follow the distribution (10) for t = T. 312

313 Estimating the generalized score. Analogously to standard score-matching, we train a timedependent neural network $\mathbf{s}_{\theta}(\mathbf{x}(t),t)$: $\mathbb{R}^n \times \mathbb{R} \to \mathbb{R}^n$ to estimate the generalized score 314 $\mathcal{L} \log p_t(\mathbf{x}(t)|\mathbf{x}(0))$ at any time point, that is, we minimize the objective 315

$$\mathbb{E}_{t}\left\{w(t)\mathbb{E}_{\mathbf{x}(0)\sim p_{0}(\mathbf{x})}\mathbb{E}_{\mathbf{x}(t)\sim p_{t}(\mathbf{x}|\mathbf{x}(0))}\left[\left|\mathbf{s}_{\boldsymbol{\theta}}(\mathbf{x}(t),t)-\mathcal{L}\log p_{t}(\mathbf{x}(t)|\mathbf{x}(0))\right|^{2}\right]\right\},\qquad(11)$$

318 where $w : [0,T] \rightarrow \mathbb{R}_+$ is a time-weighting function. Now, from Condition 3 above and the 319 property that \mathcal{L}_{A_i} computes the direction derivative along the flow of $\Pi_{A_i}(\mathbf{x})$, it follows that 320 $\mathcal{L}\log p_t(\mathbf{x}(t)|\mathbf{x}(0)) = \nabla_{\boldsymbol{\tau}(t)}\log p_t(\mathbf{x}(\boldsymbol{\tau})(t)|\mathbf{x}(\boldsymbol{\tau})(0))$. Under the above assumptions, $p_t(\boldsymbol{\tau}) = \mathbf{x}_t(t)$ $\mathcal{N}(\boldsymbol{\tau}|\boldsymbol{\mu}(\mathbf{x}(0),t),\boldsymbol{\Sigma}(t))$, where the form of the mean and the variance depends on the explicit form 321 of (9). Using the parametrization $\tau(t) = \mu(\mathbf{x}(0), t) + \sqrt{\Sigma(t)}\eta_t$, where $\eta_t \sim \mathcal{N}(0, I)$, we obtain 322 323

$$\mathcal{L}\log p_t(\mathbf{x}(t)|\mathbf{x}(0)) = -\boldsymbol{\Sigma}^{-1}(\boldsymbol{\tau}(t) - \boldsymbol{\mu}(\mathbf{x}(0), t)) = -\sqrt{\boldsymbol{\Sigma}(t)}^{-1}\boldsymbol{\eta}_t.$$
 (12)



Figure 4: Lie algebra $\mathfrak{so}(2) \subset \mathfrak{so}(3)$ dynamics for torsion (a,b) and bond angles (c,d) in molecular conformers.

3.1 EXAMPLES

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Standard Score Matching. Standard score matching can be recovered as a special case of our formalism by choosing $X = \mathbb{R}^n$ and G = T(n). As we show explicitly in Appendix A.1, we have $\mathcal{L} = \nabla$ and the Lie algebra action $\Pi(\mathbf{x}) = I$, the identity on X. Since Π is x-independent, its divergence vanishes, as well as the quadratic Casimir (T(N)) is Abelian), so that the SDEs (6) take the known form

$$d\mathbf{x} = \beta(t)\mathbf{f}(\mathbf{x})dt + \gamma(t)d\mathbf{W}, \quad d\mathbf{x} = \left[\beta(t)\mathbf{f}(\mathbf{x}) - \gamma(t)^2\nabla\log p_t(\mathbf{x})\right]dt + \gamma(t)d\mathbf{W}.$$
 (13)

346 $G = SO(2) \times \mathbb{R}_+$. A simple but non-trivial case in given by $G = SO(2) \times \mathbb{R}_+$ describing 347 rotations and dilations acting on $X = \mathbb{R}^2$. A basis for the 2-dimensional Lie algebra $\mathfrak{g} = \mathfrak{so}(2) \oplus \mathbb{R}$ 348 is given by $A_r = I$ and $A_\theta = \begin{pmatrix} 0 & -1 \\ 1 & 0 \end{pmatrix}$, yielding $\Pi(\mathbf{x}) = \begin{pmatrix} x & -y \\ y & x \end{pmatrix}$, which satisfies all the 350 conditions of section 2.2. Following our discussion above and in Appendix A.1 we have (since 351 $\rho(\Omega) = A_r^2 + A_\theta^2 = I - I = 0$)

$$d\mathbf{x} = \beta(t) \left(f_r(r) A_r \mathbf{x} + f_\theta(\theta) A_\theta \mathbf{x} \right) dt + \gamma(t) \left(dW_r A_r \mathbf{x} + dW_\theta A_\theta \mathbf{x} \right) , \qquad (14)$$

and we see that the SDE splits into contributions from the two Lie algebra summands. To find an explicit solution, let $\gamma(t) = \sqrt{\beta(t)}$ and $f_r = -\frac{1}{4}\log(x^2 + y^2)$, $f_{\theta} = -\frac{1}{2}\arctan\frac{y}{x}$. This choice corresponds, in the flow coordinates system, to a 2d Ornstein-Uhlenbeck system (Gardiner, 1985) which has a Gaussian solution with mean $\binom{r(0)}{\theta(0)}e^{-\int_0^t\beta(s)ds}$ and variance $(1 - e^{-\int_0^t\beta(s)ds})I$. Let us define $\sigma(t) = \sqrt{1 - e^{-\int_0^t\beta(s)ds}}$, such that $r(t) = r(0) + \lambda(t) = r(0) - r(0)\sigma(t)^2 + \sigma(t)\eta_r$

Let us define $\sigma(t) = \sqrt{1 - e^{-\beta_0 \beta(3) ds}}$, such that $r(t) = r(0) + \lambda(t) = r(0) - r(0)\sigma(t)^2 + \sigma(t)\eta_r$ and similarly $\theta(t) = \theta(0) + \varphi(t) = \theta(0) - \theta(0)\sigma(t)^2 + \sigma(t)\eta_{\theta}$ where $\eta_r, \eta_{\theta} \in \mathcal{N}(0, 1)$, then it is an easy calculation to show that

$$\begin{pmatrix} x(t) \\ y(t) \end{pmatrix} = \begin{pmatrix} e^{r(t)}\cos(\theta(t)) \\ e^{r(t)}\sin(\theta(t)) \end{pmatrix} = \begin{pmatrix} e^{\lambda(t)} & 0 \\ 0 & e^{\lambda(t)} \end{pmatrix} \begin{pmatrix} \cos\varphi(t) & -\sin\varphi(t) \\ \sin\varphi(t) & \cos\varphi(t) \end{pmatrix} \begin{pmatrix} x(0) \\ y(0) \end{pmatrix} .$$
(15)

We can look at the asymptotic behavior of the solution. Assuming that $\beta(t)$ is a monotonous increasing function, that is, $\beta(t + \epsilon) > \beta(t)$ for $\epsilon > 0$, then $\lim_{t\to\infty} \sigma(t) = 1$ and hence

$$\lim_{t \to \infty} \mathbf{x}(t) = e^{-r_0 + \eta_r} \begin{pmatrix} \cos \theta_0 & \sin \theta_0 \\ -\sin \theta_0 & \cos \theta_0 \end{pmatrix} \begin{pmatrix} \cos \eta_\theta & \sin \eta_\theta \\ -\sin \eta_\theta & \cos \eta_\theta \end{pmatrix} \begin{pmatrix} e^{r_0} \cos \theta_0 \\ e^{r_0} \sin \theta_0 \end{pmatrix} = \begin{pmatrix} e^{\eta_r} \cos \eta_\theta \\ e^{\eta_r} \sin \eta_\theta \end{pmatrix},$$

where $\theta_0 = \theta(0)$, $r_0 = r(0)$. Note that, even if (16) is not Gaussian, we can still easily draw samples from it by sampling the two Gaussian variables $\eta_{r,\theta}$.

Dihedral and bond angles. The above formalism can be applied to obtain transformations of physically meaningful quantities, as bond and torsion angles for molecules' conformations. Let γ_i be the dihedral angle between the planes identified by the points $\{\mathbf{x}_{i-1}, \mathbf{x}_i, \mathbf{x}_{i+1}\}$ and $\{\mathbf{x}_i, \mathbf{x}_{i+1}, \mathbf{x}_{i+2}\}$, respectively (Figure 4a). The Lie algebra element corresponding to an infinitesimal change in γ_i is given by a $3N \times 3N$ -dimensional 3×3 -block diagonal matrix, whose $j = 1, \ldots, N$ block is given by $H(j - (i + 1))\hat{\mathbf{x}}_{i+1,i} \cdot \mathbf{A}$, where $\mathbf{A} = (A_x, A_y, A_z)$ is the vector of the Lie algebra basis for $\mathfrak{so}(3), \hat{\mathbf{x}}_{i+1,i} = (\mathbf{x}_{i+1} - \mathbf{x}_i)/|\mathbf{x}_{i+1} - \mathbf{x}_i)|$ and H(i) = 1 if i > 0 and 0 otherwise is the Heaviside



Figure 5: (a) 2d mixture of Gaussians (top: ground truth, bottom: generated); (b) generating process using single scores for the subgroups SO(2), \mathbb{R}_+ with the corresponding vector field generating scores (c) (d,e) one-dimensional learning for a symmetric distributions; 3*d*-distributions: torus (f) and Möbius strip (g) (top: ground truth, bottom: generated);(f) 4d mixture of Gaussian for the group $G = SO(4) \times \mathbb{R}_+$.

step function. For bond angles β_i (Figure 4c) we construct the corresponding $\mathfrak{so}(2) \in \mathfrak{so}(3)$ algebra element blocks as $H(j-i)(\mathbf{x}_{i+1,i} \times \mathbf{x}_{i-1,i}) \cdot \mathbf{A}$. Examples of the dynamics generated by these operators are presented in Figure 4(b,d). This shows that the approach of Corso et al. (2023); Jing et al. (2022); Ketata et al. (2023) can be recovered as a particular case of our general formalism. Notably, our approach performs diffusion in a flat, rather than a curved Riemannian manifold.

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4 RELATED WORK

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Representation theory applied to neural networks has been studied both theoretically (Esteves, 2020; 400 Chughtai et al., 2023; Puny et al., 2021; Smidt, 2021) and applied to a variety of groups, architecture 401 and data type: CNNs (Cohen & Welling, 2016; Romero et al., 2020; Liao & Liu, 2023; Finzi et al., 402 2020; Weiler & Cesa, 2019; Weiler et al., 2018), Graph Neural Networks (Satorras et al., 2021), 403 Transformers, (Geiger & Smidt, 2022; Romero & Cordonnier, 2020; Hutchinson et al., 2021), point 404 clouds (Thomas et al., 2018), chemistry (Schütt et al., 2021; Le et al., 2022a). On the topic of dis-405 entanglement of group action and symmetry learning, Pfau et al. (2020) factorize a Lie group from 406 the orbits in data space, while Winter et al. (2022) learn through an autoencoder architecture in-407 variant and equivariant representations of any group acting on the data. Fumero et al. (2021) learns disentangled representations solely from data pairs. Dehmamy et al. (2021) propose an architec-408 ture based on Lie algebras that can automatically discover symmetries from data. Xu et al. (2022) 409 predict molecular conformations from molecular graphs in an roto-translation invariant fashion with 410 equivariant Markov kernels. 411

412 Partially related to our study is the field of diffusion on Riemannian manifolds. De Bortoli et al. (2022) propose diffusion in a product space – a condition which is not a necessary in our framework – 413 defined by the flow coordinates in the respective Riemannian sub-manifolds. When the Riemannian 414 manifold is a Lie group, their method yields dynamics similar to ours, as illustrated in an example 415 in Section 3.1. In fact, our formalism could be integrated with their approach to create a unified 416 framework for diffusion processes on the broader class of Riemannian manifolds admitting a Lie 417 group action. These techniques has been applied in a variety of use cases (Corso et al., 2023; Ketata 418 et al., 2023; Yim et al., 2023) for protein docking, ligand and protein generation. The works Zhu 419 et al. (2024); Kong & Tao (2024) leverage trivialized momentum to perform diffusion on the Lie 420 algebra (isomorphic to \mathbb{R}^n) instead of the Lie group, thereby eliminating curvature terms, although 421 their approach is to date only feasible for Abelian groups. An interesting connection with our work 422 is the work of Kim et al. (2022): the authors propose a bijection to map a non-linear problem to a linear one, to approximate a bridge between two non-trivial distributions. Our case can be seen as a 423 bijection between the (curved) Lie group manifold and the (flat) Euclidean data space. 424

In the context of interpreting the latent space of diffusion models, Park et al. (2023) explores the
local structure of the latent space (trajectory) of diffusion models using Riemannian geometry. Similarly, Haas et al. (2024) propose a method to uncover semantically meaningful directions in the
semantic latent space (*h*-space) (Wang et al., 2023) of denoising diffusion models (DDMs) by PCA.
Wang et al. (2023) propose a method to learn disentangled and interpretable latent representations
of diffusion models in an unsupervised way. We note that the aforementioned works aim to extract
meaningful latent factors in traditional DDMs, often restricting to human-interpretable semantic
features and focusing on image generation.

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5 EXPERIMENTS

2d, 3d and 4d distributions. In Figure 5 we illustrate the framework for a variety of d = 2, 3-446 dimensional distributions. In all cases we take $G = SO(d) \times \mathbb{R}_+$. Figure 5(a,b,c) displays a mixture 447 of Gaussians: in (a) (bottom) we see that our generalized score-matching can learn any distribution, 448 regardless of its inherent symmetry; (b) shows the output of the generation process using only one 449 score (top $\mathfrak{g} = \mathfrak{so}(2)$, bottom $\mathfrak{g} = \mathfrak{r}_+$), while (c) shows the vector fields corresponding to the 450 scores, where we color-coded the field directions. In Figure 5, we illustrate the framework for 451 various d = 2, 3-dimensional distributions, and we choose $G = SO(d) \times \mathbb{R}_+$. Figure 5(a,b,c) 452 shows a mixture of Gaussians: (a) demonstrates that our generalized score-matching can learn any 453 distribution, regardless of its symmetry; (b) shows the output of the generation process using a 454 single score (top $\mathfrak{g} = \mathfrak{so}(2)$, bottom $\mathfrak{g} = \mathfrak{r}_+$) and in (c) we display the corresponding vector fields. 455 Figures 5(d,e) depicts radial and angular distributions, where the score is learned using the respective 456 Lie algebra elements. This reflects the ability to leverage the symmetry properties of the data and perform diffusion in a lower-dimensional space. We also show in Figure 5f $(G = SO)(4) \times \mathbb{R}_+$ 457 that our method can be applied to higher dimensional Lie groups. 458

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t = T

 $\overline{t} = 0$

Figure 6: (a) Original and rotated MNIST samples with generated samples from our model and

BBDM. (b) Reverse diffusion trajectories of our model against BBDM. Intermediate samples from

BBDM resemble interpolation of mixed digits. For the last case, the 9-digit transitions into a 4-digit.

459 **Rotated MNIST.** In this experiment we show that our framework can be applied to effectively 460 learn a bridge between two non-trivial distributions, adopting however only techniques from score-461 matching and DDM. Let $p_T(\mathbf{x})$ be the rotated MNIST dataset and $p_0(\mathbf{x})$ the original (non-rotated) 462 MNIST dataset. We can learn to sample from p_0 starting from element of p_T by simply modeling 463 a SO(2) dynamic. Some examples of our results are shown in Figure 6. Notice that our formalism allows us to reduce the learning to a 1-dimensional score $\mathcal{L}_{\theta} = x_1 \partial_{x_1} - x_2 \partial_{x_2}$, which reflects 464 the true dimensionality of the problem. We trained the model with T = 100 time-steps, but for 465 sampling it suffices to set T = 10. As it can be seen in the example trajectories 6b, the model starts 466 converging already at $t/T \sim 0.5$. We employ a CNN which processes input images $\mathbf{x}(t)$, and the 467 resulting feature map is flattened and concatenated with a scalar input t, then passed through fully 468 connected layers to produce the final output. We compare our approach to the Brownian Bridge 469 Diffusion Model (BBDM) (Li et al., 2023). Unlike our method, BBDM operates unconstrained in 470 the full MNIST pixel space ($\mathbb{R}^{28 \times 28}$), where intermediate states represent latent digits. As shown 471 in Figure 6a, this can result in incorrect transitions, such as adding extraneous pixels or altering the 472 original digit, sometimes generating entirely different digits (Figure 6b). Further modeling details 473 can be found in Appendix E.

474 **QM9.** We use our framework to train a generative model $p_{\theta}(X|M)$ for conformer sampling of 475 small molecules M from the QM9 dataset (Ramakrishnan et al., 2014). We only keep the lowest en-476 ergy conformer as provided in the original dataset, that is, for each molecule only one 3D conformer 477 is maintained. Here $X = \mathbb{R}^{3N}$ and we choose $G = (SO(3) \times \mathbb{R}_+)^N$, where each factor acts on 478 the space \mathbb{R}^3 spanned by the Cartesian coordinates of the molecule's atoms, respectively. As Figure 479 7a shows, our generative process yields conformers that are energetically very similar to the ground 480 truth conformers, while showing some variability, as it can be seen in the last example where the tor-481 sion angle is differently optimized. We train another model $p_{\gamma}(X|M)$ via standard Fisher denoising score-matching, i.e., choosing $G = T(3)^N$ as in Sec. 3.1, and generate 5 conformers per molecule 482 for both models p_{θ}, p_{γ} . We then compute the UFF energy (Rappe et al., 1992) implemented in the 483 RDKit for all generated conformers and extract the lowest energy geometry as generated sample. 484 To compare against the reference geometry, we compute the energy difference $\Delta = U_{true} - U_{gen}$ 485 for both models. Figure 7b shows that both diffusion models tend to generate conformers that have



Figure 7: (a) Generated 3D conformer for several molecules from the QM9 validation set (top row) and ground truth conformer (bottom row). (b) Energy difference distribution between diffusion models (p_{θ}, p_{γ}) compared to the ground-truth energy. Both diffusion models generate conformers that have a similar Δ energy distribution.

lower energies than the ground true conformer according to the UFF parametrization, while the diffusion model that implements the dynamics according to $G = (SO(3) \times \mathbb{R}_+)^N$ (colored in blue) achieves slightly lower energy conformers, mean $\Delta_{\theta} = -0.2159$ against mean $\Delta_{\gamma} = -0.2144$ for the standard diffusion model (colored in orange).



Figure 8: (a) Generated SE(3) trajectories for molecular docking. (b) Comparison with RSGM.

CrossDocked2020: Global E(3) and Protein-Ligand Complexes. In this final experiment, we train a generative model for global SE(3) transformations acting on small molecules. Specifically, given a pair consisting of a compound and a protein pocket in space, our goal is to generate the trajectory by which the ligand best fits into the pocket. Importantly, the internal structure of the compound remains fixed, which presents a challenge when using standard diffusion processes. Thus, while the SE(3) transformations are global with respect to the ligand, they do not represent global symmetries of the overall system. We derive in appendix A.4 the relevant operators that guide the dynamics (6), (7). Figure 8a shows examples of docked molecules using SE(3)-guided scorematching diffusion. The true and generated molecules at different generation steps are visualized as point clouds, showing a good agreement. Figure 8b shows that our model achieves a lower RMSD $(2.9 \pm 1.0 \text{ Å vs } 5.6 \pm 1.2 \text{ Å})$ for the docked ligands than the method from RSGM (De Bortoli et al., 2022; Corso et al., 2023) (for implementation details we refer to Appendix E.2.1).

6 CONCLUSIONS AND OUTLOOK

We presented a framework for generative modeling that respects the action of any Lie group G and derived its mathematical foundation. Our framework generates a curved Lie group diffusion dy-namics in flat Euclidean space, thus without the need to transform the data and of performing group projections. Specifically, we introduced a new class of exactly-solvable SDEs that guide the corruption and generation processes. Traditional diffusion score-matching emerges as a special case when the group G is set to the translation group in Euclidean space. Thus, our framework does not merely complement existing methods, but *expands* the space of exactly solvable diffusion processes. Our framework is particularly relevant given recent findings (Abramson et al., 2024) showing that unconstrained models outperform equivariant ones: with our framework there is no need of a trade-off, as we retain the expressivity of unconstrained models with the benefits of group inductive bias. Moreover, our techniques descend quite straightforwardly to flow matching (Lipman et al., 2022) through the Diffusion Mixture Representation Theorem (Peluchetti, 2023; Brigo, 2008). We spell out the connection in appendix D and we plan to expand on this in future work.

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A EXAMPLES OF LIE GROUPS AND LIE ALGEBRA ACTIONS

In this appendix we list some important Lie groups and Lie algebra actions, their corresponding fundamental vector fields as well as the fundamental flow coordinates. These will be useful in the main text.

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1764 Let $X = \mathbb{R}^N$ and G = T(N), the group of translations in N-dimensional space. Element of T(N)1765 are represented by a vector $\mathbf{v} = (v_1, v_2, \dots, v_N)^\top \in \mathbb{R}^N$, where v_i are the translation components 1766 along the x_i axes for $i = 1, \dots, N$, thus $T(N) \simeq \mathbb{R}^N$. Explicitly, for a $\mathbf{x} \in X$ its action is given by 1767 $\rho_{\mathbb{R}^N}(\mathbf{v}, \mathbf{x}) = \mathbf{x} + \mathbf{v}$.

The corresponding Lie algebra $\mathfrak{t}(N)$ is also isomorphic to \mathbb{R}^N , and it consists of vectors $\mathbf{a} = (a_1, a_2, \ldots, a_N)^\top \in \mathbb{R}^N$. The Lie bracket of any two elements in $\mathfrak{t}(N)$ vanishes, as T(N) is Abelian.

To derive the infinitesimal action, we first note that the exponential map is trivial, $exp(\tau \mathbf{A}) = \tau \mathbf{A}$. Hence, we have

$$\Pi_A(\mathbf{x}) = \frac{d}{d\tau} \bigg|_{\tau=0} \rho_{\mathbb{R}^N}(\tau \mathbf{A}, \mathbf{x}) = \frac{d}{d\tau} \bigg|_{\tau=0} (\mathbf{x} + \tau \mathbf{A}) = \mathbf{A} .$$
(16)

Thus, the fundamental vector field Π_A corresponding to $\mathbf{A} \in \mathfrak{t}(N)$ is the constant vector field:

$$\Pi_A = a_1 \frac{\partial}{\partial x_1} + a_2 \frac{\partial}{\partial x_2} + \dots + a_N \frac{\partial}{\partial x_N} = \mathbf{A} \cdot \nabla \,.$$

A.2 $X = \mathbb{R}^N, G = \mathbb{R}^*_+$ (group of dilations)

Tet us consider $X = \mathbb{R}^N$ and $G = \mathbb{R}^*_+$, the group of dilations in *N*-dimensional space. The group \mathbb{R}^*_+ consists of all positive scaling factors. Each element of $G = \mathbb{R}^*_+$ can be represented by a scalar $\lambda > 0$ that scales all vectors in \mathbb{R}^N by this factor.

The action of $G = \mathbb{R}^*_+$ on \mathbb{R}^N is a dilation, meaning that every vector $\mathbf{x} = (x_1, x_2, \dots, x_N)^\top \in \mathbb{R}^N$ is scaled by the factor λ . Explicitly, the group action is given by

$$\rho_{\mathbb{R}^N}(\lambda, \mathbf{x}) = \lambda \mathbf{x} \,. \tag{17}$$

791 The Lie algebra $\mathfrak{g} = \mathbb{R}$ corresponding to the dilation group $G = \mathbb{R}^*_+$ consists of real numbers 792 representing the logarithm of the scaling factor. Specifically, an element $A \in \mathfrak{g}$ corresponds to a 793 generator of the dilation, and the exponential map $\exp : \mathfrak{g} \to G$ is given by: $\exp(\tau A) = e^{\tau A}$, where 794 τ is a real parameter.

The infinitesimal action corresponds to taking the derivative at $\tau = 0$. For a vector $\mathbf{x} \in \mathbb{R}^N$ and $A \in \mathfrak{g}$, the fundamental vector field Π_A is computed as:

$$d\rho_{\mathbb{R}^N}(A) = \frac{d}{d\tau} \bigg|_{\tau=0} \rho_{\mathbb{R}^N}(e^{\tau A}, \mathbf{x}) = \frac{d}{d\tau} \bigg|_{\tau=0} \left(e^{\tau A} \mathbf{x} \right) = A\mathbf{x} , \qquad (18)$$

and

$$\Pi_A(\mathbf{x}) = A\mathbf{x} \cdot \nabla$$

Now, solving the equation

$$\mathbf{x} = e^{\tau A} \mathbf{x}_0 \tag{19}$$

in terms of τ we obtain

$$\tau = \frac{1}{A}\log\frac{|\mathbf{x}|^2}{\mathbf{x}\cdot\mathbf{x}_0} = \frac{1}{A}\log\frac{|\mathbf{x}|^2}{|\mathbf{x}||\mathbf{x}_0|} = \frac{1}{A}\log\frac{|\mathbf{x}|}{|\mathbf{x}_0|} = \frac{1}{2A}\log\frac{|\mathbf{x}|^2}{|\mathbf{x}_0|^2} \,. \tag{20}$$

In the usual case of A = 1 (generator of the Lie algebra), $\mathbf{x}_0 = \frac{1}{\sqrt{N}}(1, 1, \dots, 1)^{\top}$ to be the unit vector we obtain the usual expression

$$\tau = \frac{1}{2}\log(x_1^2 + x_2^2 + \dots + x_N^2).$$
(21)

816 A.3
$$X = \mathbb{R}^3, G = SO(3) \times \mathbb{R}^*_+$$

The dilation part is solved in the previous section, so we actually just focus on the action of SO(3) on \mathbb{R}^3 . The orbits are given by spheres centered at the origin, and we can decompose the action of SO(3) by variying the azimuthal or the polar angle defined by a vector **x**. Namely, we have the two actions

$$\rho_{\mathbb{R}^{3}}(\varphi, \mathbf{x}) = \begin{pmatrix} \cos\varphi & -\sin\varphi & 0\\ \sin\varphi & \cos\varphi & 0\\ 0 & 0 & 1 \end{pmatrix} \begin{pmatrix} x\\ y\\ z \end{pmatrix},
\rho_{\mathbb{R}^{3}}(\theta, \mathbf{x}) = \begin{bmatrix} I + \sin\theta \begin{pmatrix} 0 & 0 & \cos\varphi\\ 0 & 0 & \sin\varphi\\ -\cos\varphi & -\sin\varphi & 0 \end{pmatrix} \\
+ (1 - \cos\theta) \begin{pmatrix} -\cos^{2}\varphi & -\cos\varphi\sin\varphi & 0\\ -\cos\varphi\sin\varphi & -\sin^{2}\varphi & 0\\ 0 & 0 & -1 \end{pmatrix} \end{bmatrix} \begin{pmatrix} x\\ y\\ z \end{pmatrix}. \quad (22)$$

If we take the differentials

$$d\rho_{\mathbb{R}^{3}}(\varphi, \mathbf{x})|_{\varphi=0} = \begin{pmatrix} -\sin\varphi & -\cos\varphi & 0\\ \cos\varphi & -\sin\varphi & 0\\ 0 & 0 & 0 \end{pmatrix} \begin{pmatrix} x\\ y\\ z \end{pmatrix}|_{\varphi=0} = \begin{pmatrix} 0 & -1 & 0\\ 1 & 0 & 0\\ 0 & 0 & 0 \end{pmatrix} \begin{pmatrix} x\\ y\\ z \end{pmatrix} = A_{z}\mathbf{x} ,$$

$$d\rho_{\mathbb{R}^{3}}(\theta, \mathbf{x}) = \begin{bmatrix} \cos\theta \begin{pmatrix} 0 & 0 & \cos\varphi\\ 0 & 0 & \sin\varphi\\ -\cos\varphi & -\sin\varphi & 0 \end{pmatrix} \\ -\sin\theta \begin{pmatrix} \cos^{2}\varphi & \cos\varphi\sin\varphi & 0\\ \cos\varphi\sin\varphi & \sin^{2}\varphi & 0\\ 0 & 0 & 1 \end{pmatrix} \Big]_{\theta=0} \begin{pmatrix} x\\ y\\ z \end{pmatrix}$$

$$= \begin{pmatrix} 0 & 0 & \cos\varphi\\ 0 & 0 & \sin\varphi\\ -\cos\varphi & -\sin\varphi & 0 \end{pmatrix} \begin{pmatrix} x\\ y\\ z \end{pmatrix} = (\cos\varphi A_{y} - \sin\varphi A_{x})\mathbf{x} , \qquad (23)$$

where

$$A_x = \begin{pmatrix} 0 & 0 & 0 \\ 0 & 0 & -1 \\ 0 & 1 & 0 \end{pmatrix} \qquad A_y = \begin{pmatrix} 0 & 0 & 1 \\ 0 & 0 & 0 \\ -1 & 0 & 0 \end{pmatrix} \qquad A_z = \begin{pmatrix} 0 & -1 & 0 \\ 1 & 0 & 0 \\ 0 & 0 & 0 \end{pmatrix}$$
(24)

form a basis for $\mathfrak{so}(3)$. The corresponding differential operators are

$$\mathcal{L}_{\varphi} = x\partial_y - y\partial_x , \qquad \qquad \mathcal{L}_{\theta} = \frac{1}{\sqrt{x^2 + y^2}} \left[zx\partial_x + zy\partial_y - (x^2 + y^2)\partial_z \right] , \qquad (25)$$

and it is an easy calculation to show that they commute $[\mathcal{L}_{\varphi}, \mathcal{L}_{\theta}] = 0$. The attentive reader might have noticed that the commutation does not hold at the matrices level. While this is expected, since there is no 2-dimensional commuting subalgebra in $\mathfrak{so}(3)$, it is nonetheless quite puzzling since ev-erything works out at the level of differential operators. This reflect the fact that the commutation properties are necessary at the level of the action of \mathfrak{g} on X, and not necessarily at the Lie algebra level. In this case, however, we can elegantly resolve the puzzle indem we found a matrix represen-tation for the action $d\rho_{\mathbb{R}^3}(\theta)\mathbf{x}$ which does commute with the φ action. To do this we note that we can rewrite

$$\mathcal{L}_{\theta} = \frac{\cos\theta}{\sin\theta} x \partial_x + \frac{\cos\theta}{\sin\theta} y \partial_y - \frac{\sin\theta}{\cos\theta} z \partial_z , \qquad (26)$$



Figure 9: (a) The coordinates $\hat{\mathbf{x}}_{\mu}$ are the coordinates in the coordinate system defined by \mathbf{x}_1 , the orthogonal projection of \mathbf{x}_2 with respect to x_1 . $\mathbf{x}_2 \perp \mathbf{x}_1 = \mathbf{x}_2 - \mathbf{x}_1 \cdot \mathbf{x}_2$, and $\mathbf{x}_2 \times \mathbf{x}_1$. (b) Graphical depiction of the global symmetry transformations parametrized by the three angles $\varphi_2, \theta_1, \varphi_1$.

which corresponds to simultaneous dilations, with different coefficient, in the z axis and x, y-plane. The finite action takes the form

$$\widetilde{\rho}_{\mathbb{R}^{3}}(\theta, \mathbf{x}) = \exp\left[\log\sin\theta \begin{pmatrix} 1 & 0 & 0\\ 0 & 1 & 0\\ 0 & 0 & 0 \end{pmatrix} + \log\cos\theta \begin{pmatrix} 0 & 0 & 0\\ 0 & 0 & 0\\ 0 & 0 & 1 \end{pmatrix}\right],$$
(27)

and computing the first order term we obtain

$$d\widetilde{\rho}_{\mathbb{R}^3}(\theta, \mathbf{x}) = \begin{pmatrix} \frac{\cos\theta}{\sin\theta} & 0 & 0\\ 0 & \frac{\cos\theta}{\sin\theta} & 0\\ 0 & 0 & -\frac{\sin\theta}{\cos\theta} \end{pmatrix} \mathbf{x} .$$
(28)

This matrix is diagonal and it trivially commutes with A_z . The price we had to pay to realize a system of commuting matrices is that in $\tilde{\rho}$ the flow parameter θ appear non-linearly, thus we traded-off commutativity at the level of the Lie algebra matrices for the linearity of the flow parameters at the group level. We remark that both give rise to the same differential operator on X, which is the relevant object for our purposes.

A.4
$$X = \mathbb{R}^{3N}$$
 and global SO(3)

Let $X = \mathbb{R}^{3N}$ be parametrized by $\mathbf{x}_{i=1,\dots,N}$. We can describe a global SO(3) action as follows

$$\mathbf{x}_1 = R_{\mathbf{e}_z}(\varphi_1) R_{\mathbf{e}_y}(\theta_1) \begin{pmatrix} 0\\0\\\hat{z}_1 \end{pmatrix}$$

$$\mathbf{x}_{2} = R_{\mathbf{e}_{z}}(\varphi_{1})R_{\mathbf{e}_{y}}(\theta_{1})R_{\mathbf{e}_{z}}(\varphi_{2})\begin{pmatrix}\hat{x}_{2}\\0\\\hat{z}_{2}\end{pmatrix},$$

> $\mathbf{x}_{\mu=3,\ldots,N} = R_{\mathbf{e}_z}(\varphi_1) R_{\mathbf{e}_y}(\theta_1) R_{\mathbf{e}_z}(\varphi_2) \widehat{\mathbf{x}}_{\mu} ,$ (29) where $R_{\mathbf{a}}(\omega)$ represents a rotation of an angle ω around the axis **a**. We can then derive the operator

 $\Pi \in \mathbb{R}^{3N \times 3N}$ as follows. Let $R'(\omega)$ be the matrix where we take the partial derivative with respect to ω of all elements of R. Then

$$\Pi_{\varphi_{1}} = \left(\begin{bmatrix} A_{z} \mathbf{x}_{1} \end{bmatrix}^{\top} & \begin{bmatrix} A_{z} \mathbf{x}_{2} \end{bmatrix}^{\top} & \cdots & \begin{bmatrix} A_{z} \mathbf{x}_{N} \end{bmatrix}^{\top} \right)^{\top} \\
\Pi_{\theta_{1}} = \begin{pmatrix} (\cos \varphi_{1} A_{y} - \sin \varphi_{1} A_{x}) \mathbf{x}_{1} \\ (\cos \varphi_{1} A_{y} - \sin \varphi_{1} A_{x}) \mathbf{x}_{2} \\ \vdots \\ (\cos \varphi_{1} A_{y} - \sin \varphi_{1} A_{x}) \mathbf{x}_{N} \end{pmatrix} \\
\Pi_{\varphi_{2}} = \begin{pmatrix} \mathbf{0} \\ (\sin \theta_{1} \cos \varphi_{1} A_{x} + \sin \theta_{1} \sin \varphi_{1} A_{y} + \cos \theta_{1} A_{z}) \mathbf{x}_{2} \\ \vdots \\ (\sin \theta_{1} \cos \varphi_{1} A_{x} + \sin \theta_{1} \sin \varphi_{1} A_{y} + \cos \theta_{1} A_{z}) \mathbf{x}_{N} \end{pmatrix}$$
(30)

Notice that these do represent global rotations since it is easy to see that $(\sin \theta_1 \cos \varphi_1 A_x +$ $\sin \theta_1 \sin \varphi_1 A_y + \cos \theta_1 A_z \mathbf{x}_1 = \mathbf{0}$. Formally, the true Lie algebra elements are 3×3 matrices of the form

> $A_{\varphi} = \begin{pmatrix} A_{z} & 0 & 0 & \cdots & 0\\ 0 & A_{z} & 0 & \cdots & 0\\ 0 & 0 & A_{z} & \cdots & 0\\ \vdots & \vdots & \vdots & \ddots & \vdots\\ 0 & 0 & 0 & \cdots & A \end{pmatrix}$ (31)

and similarly for the other operators. Now, for the inverse relations we have

- $\theta_1 = \arccos \frac{z_1}{(x_1^2 + y_1^2 + z_1^2)^{1/2}}$, $\varphi_1 = \operatorname{sgn}(y_1) \operatorname{arccos} \frac{x_1}{(x_1^2 + y_1^2)^{1/2}} ,$ $\varphi_2 = \arctan \frac{\widetilde{y}_2}{\widetilde{\alpha}_1}$, (32)
- where $\widetilde{\mathbf{x}}_{2} = R_{\mathbf{e}_{u}}(\theta_{1})^{-1}R_{\mathbf{e}_{z}}(\varphi_{1})^{-1}\mathbf{x}_{2} = R_{\mathbf{e}_{u}}(-\theta_{1})R_{\mathbf{e}_{z}}(-\varphi_{1})\mathbf{x}_{2}.$

The Lie algebra elements corresponding to the SO(4) flow coordinates are

A.5 $X = \mathbb{R}^4, G = SO(4) \times \mathbb{R}_+$

Now we look at the case of a higher dimensional Lie group, namely $G = SO(4) \times \mathbb{R}_+$. The parametrization is given by

 $x_1 = e^r \cos \varphi_1 ,$ $x_2 = e^r \sin \varphi_1 \cos \varphi_2 ,$ $x_3 = e^r \sin \varphi_1 \sin \varphi_2 \cos \varphi_3$, $x_4 = e^r \sin \varphi_1 \sin \varphi_2 \sin \varphi_3 .$ (33)

 $A_{\varphi_1} = \begin{pmatrix} 0 & -\cos\varphi_2 & -\sin\varphi_2\cos\varphi_3 & -\sin\varphi_2\sin\varphi_3\\ \cos\varphi_2 & 0 & 0 & 0\\ \sin\varphi_2\cos\varphi_3 & 0 & 0 & 0\\ \sin\varphi_2\sin\varphi_3 & 0 & 0 & 0 \end{pmatrix} ,$ $A_{\varphi_2} = \begin{pmatrix} 0 & 0 & 0 & 0 \\ 0 & 0 & -\cos\varphi_3 & -\sin\varphi_3 \\ 0 & \cos\varphi_3 & 0 & 0 \\ 0 & \sin\varphi_3 & 0 & 0 \end{pmatrix},$ (34)

A.6 G = SO(N)We present here the formalism for the G = SO(N) for any $N \ge 4$. The parametrization is given by (Blumenson, 1960) $x_1 = e^r \cos \varphi_1 ,$ $x_2 = e^r \sin \varphi_1 \cos \varphi_2 ,$ $x_3 = e^r \sin \varphi_1 \sin \varphi_2 \cos \varphi_3 ,$ $x_{i} = e^{r} \sin \varphi_{1} \sin \varphi_{2} \sin \varphi_{3} \cdots \sin \varphi_{i-1} \cos \varphi_{i},$ $x_{n-1} = e^r \sin \varphi_1 \sin \varphi_2 \sin \varphi_3 \cdots \sin \varphi_{n-2} \cos \varphi_{n-1} ,$ (35) $x_n = e^r \sin \varphi_1 \sin \varphi_2 \sin \varphi_3 \cdots \sin \varphi_{n-2} \sin \varphi_{n-1} .$ The corresponding Lie algebra elements are given by $A_{\varphi_{n-1}} = \begin{pmatrix} 0 & \cdots & 0 & 0 \\ 0 & \cdots & 1 & 0 \\ 0 & \cdots & 1 & 0 \end{pmatrix}, A_{\varphi_{n-2}} = \begin{pmatrix} 0 & \cdots & 1 & 0 & 0 \\ 0 & \cdots & 0 & -\cos\varphi_{n-1} & -\sin\varphi_{n-1} \\ 0 & \cdots & \cos\varphi_{n-1} & 0 & 0 \\ 0 & \cdots & \sin\varphi_{n-1} & 0 & 0 \end{pmatrix}$ $-\sin\varphi_{n-2}\sin\varphi_{n-1}$ $\sin\varphi_{n-2}\sin\varphi_{n-1}$ $A_{\varphi_j} = \frac{1}{x_j} \begin{pmatrix} 0 & \ddots & \vdots & \vdots & \vdots & \vdots & \vdots \\ 0 & \cdots & 0 & -x_{j+1} & -x_{j+2} & \cdots & -x_n \\ 0 & \cdots & x_{j+1} & 0 & 0 & \cdots & 0 \\ 0 & \cdots & x_{j+2} & 0 & 0 & \cdots & 0 \\ 0 & \cdots & \vdots & \vdots & \vdots & \ddots & \vdots \\ 0 & 0 & \cdots & 0 & 0 & \cdots & 0 \\ 0 & \cdots & \vdots & \vdots & \vdots & \ddots & \vdots \end{pmatrix},$ $A_{\varphi_1} = \frac{1}{x_1} \begin{pmatrix} 0 & u_2 & u_3 & u_4 & \dots & u_n \\ x_2 & \ddots & \vdots & \vdots & \vdots & \vdots \\ x_3 & \cdots & 0 & 0 & 0 & 0 \\ x_4 & \cdots & 0 & 0 & 0 & 0 \\ \vdots & \cdots & 0 & 0 & 0 & 0 \end{pmatrix} .$ (36)В

PROOFS OF CONDITION FOR SUITABLE LIE GROUP

Here we provide the statements with proofs of the results in Section 2.2.

Proposition B.1. The linear operator induced by Π is complete if Π is the local frame of a vector bundle E over X whose rank is $n \geq \dim X$ almost everywhere. If rank E = n everywhere, then E = TX, the tangent bundle of X.

1026 *Proof.* We start by noting that, given the expression of the fundamental fields as derivations, we 1027 can write $\mathcal{L}(\mathbf{x}) = \mathbf{\Pi}(\mathbf{x})^\top \nabla$. Let $\pi : E \to X$ be the projection map, then rank $\pi^{-1}(\mathbf{x}) = \mathbf{x}$ 1028 min(rank $\Pi(\mathbf{x}), n$), since rank $\nabla = n$. Now, consider $\mathcal{L} \log p(\mathbf{x}) = \mathcal{L} \log q(\mathbf{x})$, which implies 1029 $\mathcal{L}\log \frac{p(\mathbf{x})}{q(\mathbf{x})} = 0$. Let $U \subseteq X$ such that rank $\mathbf{\Pi} \ge n \ \forall \mathbf{x} \in U$, and by assumption $X \setminus U$ has mea-1030 sure zero. Then the above holds if and only if $\nabla \log \frac{p(\mathbf{x})}{q(\mathbf{x})} = 0$, which implies $\frac{p(\mathbf{x})}{q(\mathbf{x})} = c$, constant 1031 $\forall \mathbf{x} \in U$. Now, $p(\mathbf{x})$ and $q(\mathbf{x})$ are probability densities by assumption, thus c = 1, which proves the 1032 claim. 1033

Proposition B.2. The operator Π induced by \mathfrak{g} is complete if and only if the subspace $U \subseteq X$ such that $\dim \frac{G}{G_{\mathbf{x}}} < n$ for $\mathbf{x} \in U$, where $n = \dim X$, has measure zero in X.

Proof. First, we recall that the dimension of an orbit $\mathcal{O}_{\mathbf{x}}$ of $\mathbf{x} \in X$ equals the dimension of the image of the map $d\rho_{\mathbf{x}} : \mathfrak{g} \to T_{\mathbf{x}}X : \mathbf{A} \mapsto \mathbf{\Pi}(\mathbf{x})$. Suppose first that $\mathbf{\Pi}$ is complete. Then, from Proposition B.1 the rank of $\mathbf{\Pi}(\mathbf{x})$ is $\geq n$ almost everywhere, and therefore $\dim G/G_{\mathbf{x}} \geq n$ almost everywhere, which implies one direction of the claim. The reverse is quite straightforward. Assume that the rank of $\mathbf{\Pi}(\mathbf{x})$ is $\geq n$ almost everywhere. As $\mathbf{\Pi}$ represent the action of the infinitesianal transformations of G, it means that locally G cannot fix points in X, thus proving the claim. \Box

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1044 C PROOF OF MAIN THEOREM

Here we provide the full proof of Theorem 3.1:

Theorem C.1. Let G be a Lie group acting on X satisfying the conditions of Section 2.2, and let g be its Lie algebra. The pair of SDEs

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$$d\mathbf{x} = \left[\beta(t)\mathbf{\Pi}(\mathbf{x})\mathbf{f}(\mathbf{x}) + \frac{\gamma(t)^2}{2}\rho_X(\Omega)\right]dt + \gamma(t)\mathbf{\Pi}(\mathbf{x})d\mathbf{W}, \qquad (37)$$

$$d\mathbf{x} = \left[\beta(t)\mathbf{\Pi}(\mathbf{x})\mathbf{f}(\mathbf{x}) - \frac{\gamma^2(t)}{2}\rho_X(\Omega) - \gamma^2(t)\mathbf{\Pi}(\mathbf{x})\nabla^\top \cdot \mathbf{\Pi}(\mathbf{x}) - \gamma(t)^2\mathbf{\Pi}(\mathbf{x})\mathcal{L}\log p_t(\mathbf{x})\right]dt + \gamma(t)\mathbf{\Pi}(\mathbf{x})d\mathbf{W}, \quad (38)$$

1056 where $\beta, \gamma : \mathbb{R} \to \mathbb{R}$ are time-dependent functions, $\mathbf{\Pi} : \mathbb{R}^n \to \mathbb{R}^{n \times n}$ the fundamental vector fields, 1057 $\mathbf{f} : \mathbb{R}^n \to \mathbb{R}^n$ the drift, $\Omega = \sum_i A_i^2$ is the quadratic Casimir element of \mathfrak{g} , and $\mathcal{L} = \mathbf{\Pi}(\mathbf{x})^\top \nabla$ is 1058 such that

1. The forward-time SDE (37) is exactly solvable, with solution

$$\mathbf{x}(t) = \left(\prod_{i} O_i(\tau_i(t))\right) \mathbf{x}(0) = \left(\prod_{i=1}^n e^{\tau_i(t)A_i}\right) \mathbf{x}(0) , \qquad (39)$$

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where $O_i = e^{\tau_i(t)A_i}$ is the finite group action and $\boldsymbol{\tau}(t)$ is the solution to the SDE

$$d\boldsymbol{\tau}(\mathbf{x}) = \beta(t)\mathbf{f}(\mathbf{x})dt + \gamma(t)d\mathbf{W}.$$
(40)

2. The SDE (7) is the reverse-time process of (6).

3. The Langevin dynamic of the above SDEs decomposes as a direct sum of \mathfrak{g} infinitesimal actions (1), each defining an infinitesimal transformation along the flows ξ_{τ} .

1071 1072 Proof. We start by proving 3. We start by rewriting (37) in terms of the fundamental flow coordi-1073 nates $\tau_i = \xi_{A_i}^{-1}(\mathbf{x}_0)(\mathbf{x}) : X \to \mathbb{R}$. For this we employ Itô's Lemma for the multivariate case: given the SDE (37) and a transformation $\tau(\mathbf{x})$, it is given by

$$d\boldsymbol{\tau}(\mathbf{x}) = (\nabla_{\mathbf{x}}\boldsymbol{\tau})^{\top} \left[\beta(t)\mathbf{\Pi}(\mathbf{x})\mathbf{f}(\mathbf{x}) + \frac{\gamma^{2}(t)}{2}\rho_{X}(\Omega) \right] dt + \frac{\gamma^{2}(t)}{2} \operatorname{Tr} \left[\mathbf{\Pi}(\mathbf{x})^{\top} \left(H_{\mathbf{x}}\boldsymbol{\tau} \right) \mathbf{\Pi}(\mathbf{x}) \right] dt + \gamma(t)(\nabla_{\mathbf{x}}\boldsymbol{\tau})^{\top} \mathbf{\Pi}(\mathbf{x}) d\mathbf{W}$$

$$= \beta(t)\mathbf{f}(\mathbf{x}) + \frac{\gamma^{2}(t)}{2} \left[(\nabla_{\mathbf{x}}\boldsymbol{\tau})^{\top} \Delta_{\boldsymbol{\tau}}\mathbf{x} + \operatorname{Tr} \left[\mathbf{\Pi}(\mathbf{x})^{\top} \left(H_{\mathbf{x}}\boldsymbol{\tau} \right) \mathbf{\Pi}(\mathbf{x}) \right] dt + \gamma(t) d\mathbf{W}$$
(41)

since $\nabla_{\mathbf{x}} \boldsymbol{\tau} = \boldsymbol{\Pi}^{-1}(\mathbf{x})$ as matrices. Now, the second term can be rewritten in components as

 $\{(\nabla_{\mathbf{x}}\boldsymbol{\tau})^{\top}\Delta_{\boldsymbol{\tau}}\mathbf{x} + \operatorname{Tr}\left[\boldsymbol{\Pi}(\mathbf{x})^{\top}(H_{\mathbf{x}}\boldsymbol{\tau})\boldsymbol{\Pi}(\mathbf{x})\right]\}_{\boldsymbol{\mu}_{I}}$

- $=\sum_{i}\sum_{j}\frac{\partial x_{j}}{\partial \tau_{k}}\left(\frac{\partial}{\partial x_{j}}\frac{\partial x_{i}}{\partial \tau_{l}}\right)\frac{\partial \boldsymbol{\tau}}{\partial x_{i}}+\sum_{i}\sum_{j}\frac{\partial x_{j}}{\partial \tau_{k}}\frac{\partial x_{i}}{\partial \tau_{l}}\frac{\partial^{2}\boldsymbol{\tau}}{\partial x_{i}\partial x_{j}}$
- $=\sum_{i}\frac{\partial x_{j}}{\partial \tau_{k}}\frac{\partial}{\partial x_{j}}\left(\sum_{i}\frac{\partial x_{i}}{\partial \tau_{l}}\frac{\partial \tau}{\partial x_{i}}\right)$
- $=\frac{\partial}{\partial\tau_k}\left(\sum_i\frac{\partial x_i}{\partial\tau_l}\frac{\partial\boldsymbol{\tau}}{\partial x_i}\right)$

$$= \frac{\partial}{\partial \tau_k} \frac{\partial \tau}{\partial \tau_l}$$

$$\begin{array}{l} \partial \tau_k \ \partial \tau_k \$$

which vanishes. Thus we proved that

 $d\boldsymbol{\tau}(\mathbf{x}) = \beta(t)\mathbf{f}(\mathbf{x})dt + \gamma(t)d\mathbf{W},$ (43)

(42)

and provided that is chosen so that $f_i(\mathbf{x}(\tau)) = f_i(\tau_i)$, this corresponds to a system of independent SDEs, as claimed.

Now, to prove 1, let $\tau(t)$ be a solution to (43) and $\mathbf{x}(t)$ as in (8). Then a Taylor expansion yields

$$\mathbf{x}(t) = \left[I + \sum_{i} \tau_i(t)A_i + \frac{1}{2} \left(\sum_{i} \tau_i(t)A_i\right)^2 + \mathcal{O}(\tau_i^3)\right] \mathbf{x}(0)$$
(44)

since $[A_i, A_j] = 0$ and where $\mathcal{O}(\tau_i^3)$ represents terms of third order in τ_i 's. Then taking the differ-ential and dropping higher order terms

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$$= \left[\beta(t)\mathbf{\Pi}(\mathbf{x})\mathbf{f}(\mathbf{x})dt + \gamma(t)\mathbf{\Pi}(t)d\mathbf{W}\right] + \frac{1}{2}\left(\sum_{i}\gamma(t)d\mathbf{W}A_{i}\right)^{2}\mathbf{x}(0)$$

$$= \left[\beta(t)\mathbf{\Pi}(\mathbf{x})\mathbf{f}(\mathbf{x})dt + \gamma(t)\mathbf{\Pi}(t)d\mathbf{W}\right] + \frac{\gamma(t)^2}{2}\left(\sum_i A_i^2 dt\right)\mathbf{x}(0)$$

$$= \left[\beta(t)\mathbf{\Pi}(\mathbf{x})\mathbf{f}(\mathbf{x}) + \frac{\gamma(t)^2}{2}\rho_X(\Omega)\right]dt + \gamma(t)\mathbf{\Pi}(\mathbf{x})d\mathbf{W}, \qquad (45)$$

which in the forward SDE (6), proving our claim, where we used the relations $dW_i^2 = dt$ and $dW_i dW_j = 0$ for $j \neq i$.

Finally, we prove 2. To do this it suffices to apply Anderson's result (Anderson, 1982)

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$$d\mathbf{x} = \begin{bmatrix} \beta(t)\mathbf{\Pi}_{i}(\mathbf{x})\mathbf{f}(\mathbf{x}) + \frac{\gamma^{2}(t)}{2}\rho_{X}(\Omega) - \gamma^{2}(t)\nabla\cdot(\mathbf{\Pi}(\mathbf{x})\mathbf{\Pi}(\mathbf{x})^{\top}) \\ -\gamma(t)^{2}\mathbf{\Pi}(\mathbf{x})\mathbf{\Pi}(\mathbf{x})^{\top}\nabla_{\mathbf{x}}\log p_{t}(\mathbf{x}) \end{bmatrix} dt + \gamma(t)\mathbf{\Pi}(\mathbf{x})d\mathbf{W}_{i}, \quad (46)$$

and note that $\mathbf{\Pi}(\mathbf{x})^\top \nabla_{\mathbf{x}} = \mathcal{L}$, the generalized score, and

$$\left[\nabla_{\mathbf{x}} \cdot (\mathbf{\Pi}(\mathbf{x})\mathbf{\Pi}(\mathbf{x})^{\top})\right]_{i} = \frac{\partial}{\partial x_{k}} \left(\Pi_{ij}\Pi_{kj}\right)$$

$$= \frac{\partial}{\partial x_k} (\Pi_{ij}) \Pi_{kj} + \Pi_{ij} \frac{\partial}{\partial x_k} \Pi_{kj}$$

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$$= \frac{\partial x_k}{\partial \tau_i} \frac{\partial}{\partial x_k} \left(\frac{\partial x_i}{\partial \tau_i} \right) + \Pi_{ij} [\nabla^\top \cdot \mathbf{\Pi}(\mathbf{x})]_j$$

Plugging this back in into the previous expression we obtain our claim

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$$\frac{\partial}{\partial \tau_j} \frac{\partial x_k}{\partial \tau_j} (\partial \tau_j) \to \mathbb{T}^*$$

$$= \frac{\partial}{\partial \tau_j} \left(\frac{\partial x_i}{\partial \tau_j} \right) + \Pi_{ij} [\nabla^\top \cdot \mathbf{\Pi}(\mathbf{x})]_j$$

$$= [\operatorname{Tr} H_{\boldsymbol{\tau}}(\mathbf{x})]_i + \Pi_{ij} [\nabla^\top \cdot \mathbf{\Pi}(\mathbf{x})]_j$$

1143
$$= \frac{\partial}{\partial \tau_j} \left(\right)$$

where we recall that the divergence of a matrix is a vector whose components are the divergence of its rows. Recalling the relationship between the trace of the Hessian and the Laplacian we can write in operator form

$$\nabla_{\mathbf{x}} \cdot (\mathbf{\Pi}(\mathbf{x})\mathbf{\Pi}(\mathbf{x})^{\top}) = \mathbf{\Pi}(\mathbf{x})\nabla^{\top} \cdot \mathbf{\Pi}(\mathbf{x}) + \rho_X(\Omega) , \qquad (48)$$

$$d\mathbf{x} = \left[\beta(t)\mathbf{\Pi}(\mathbf{x})\mathbf{f}(\mathbf{x}) - \frac{\gamma^2(t)}{2}\rho_X(\Omega) - \gamma^2(t)\mathbf{\Pi}(\mathbf{x})\nabla^\top \cdot \mathbf{\Pi}(\mathbf{x}) - \gamma(t)^2\mathbf{\Pi}(\mathbf{x})\mathcal{L}\log p_t(\mathbf{x})\right]dt + \gamma(t)\mathbf{\Pi}(\mathbf{x})d\mathbf{W}.$$
 (49)

(47)

¹¹⁸⁸ D LIE GROUP-INDUCED FLOW MATCHING MODELING

1190 1191 1192 1192 1193 1194 We briefly summarize the formalism of flow matching. Given a target distribution $p_0(\mathbf{x})$ and a vector field u_t generating the distribution $p_t(\mathbf{x})$, i.e., if it satisfies $p_t(\mathbf{x}) = [u_t]_* p_0(\mathbf{x})$ where $[u_t]_*$ is the push-forward map, the flow matching objective is defined as

$$\mathcal{L}_{\mathrm{FM}}(\theta) = \mathbb{E}_{t,\mathbf{x}_t \sim p_t(\mathbf{x})} |v_{t;\theta}(\mathbf{x}_t) - u_t(\mathbf{x}_t)|^2.$$
(50)

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Marginalizing over samples $\mathbf{x}_0 \sim p_0(\mathbf{x})$ we obtain the conditional flow matching objective

$$\mathcal{L}_{\text{CFM}}(\theta) = \mathbb{E}_{t,\mathbf{x}_0 \sim p_0(\mathbf{x}), \mathbf{x}_t \sim p_t(\mathbf{x}|\mathbf{x}_0)} \left| v_{t;\theta}(\mathbf{x}_t) - u_t(\mathbf{x}_t|\mathbf{x}_0) \right|^2.$$
(51)

1202 Now, under the assumptions for learning the generalized 1203 score through the objective 11 we have that $p_t(\tau(\mathbf{x})) = \mathcal{N}(\tau | \boldsymbol{\mu}(\tau(0), t), \boldsymbol{\Sigma}(t))$, where $\tau(0) = \mathbf{x}(\tau)(0)$. Then the solution 1205 of the SDE from Theorem 3.1

$$\mathbf{x}(t) = \left(\prod_{i=1}^{n} e^{\tau_i(t)A_i}\right) \mathbf{x}(0) , \qquad (52)$$



Figure 10: $\mathfrak{so}(2)$ (green and blue) vs. $\mathfrak{t}(2)$ (orange) induced flows.

is a flow inducing the distribution $p_t(\tau(\mathbf{x}))$. Thus, the vector field that generates the conditional probability path is obtained by differentiating the path above with respect to t, yielding

$$u_t(\mathbf{x}(t)|\mathbf{x}(0)) = \frac{d}{dt}\mathbf{x}(t) = \sum_i \frac{\partial \mathbf{x}(t)}{\partial \tau_i} \frac{\partial \tau_i}{\partial t}$$
$$= \sum_i A_i \mathbf{x}(t) \left(\mu'_{t,i}(\tau_i(0)) + \sigma'_t(\tau_i(0))\eta_i\right)$$
$$= \mathbf{\Pi}(\mathbf{x}(t)) \left[\mu'_t(\tau_i(0)) + \frac{\sigma'_t(\tau_i(0))}{\sigma_t(\tau_i(0))} \left(\boldsymbol{\tau}(t) - \boldsymbol{\mu}_t(\boldsymbol{\tau}(0))\right)\right], \quad (53)$$

1220 where we used the fact that

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$$\boldsymbol{\tau}(t) = \mu_t(\boldsymbol{\tau}(0)) + \sigma_t(\boldsymbol{\tau}(0))\boldsymbol{\eta} , \qquad (54)$$

1223 where $\eta \sim \mathcal{N}(0, 1)$. Thus, we see that the unique vector field that defines the flow (8) is again 1224 proportional to the fundamental vector field $\Pi(\mathbf{x})$ of the Lie algebra g of G. In figure 10 we illustrate 1225 the flow generated by our formalism in the case of SO(2) in comparison with the traditional flow matching of T(2). The orange path depicts the linear (in Euclidean metric) displacement given 1226 by the traditional flow matching, assuming G = T(2). In green and blue we depicted the orbits 1227 trajectories resulting from generalized flow matching with $G = SO(2) \times \mathbb{R}_+$. Although the start 1228 and end points are the same, the path is decomposed into transformations along the orbits of the two 1229 group factors. This is particularly useful when these correspond to meaningful degrees of freedom in 1230 the system. For example, when flowing between conformers of the same molecule, the intermediate 1231 states produced by traditional flow matching are often unphysical, as they involve linear interpolation 1232 between the Cartesian coordinates of the atoms. However, generalized score matching, following 1233 the degrees of freedom given by bond and torsion angles as described in Section 3.1, would not only 1234 yield efficient learning but also produce chemically meaningful intermediate states, as the path is 1235 broken down into updates of chemically relevant degrees of freedom.

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E EXPERIMENTS

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1240 In this final section we present some further details regarding our experiment in Section 5. We 1241 provide the code to replicate our experiments in the Supplementary Information (SI). Following publication we will open-source our code. 1242 E.1 MNIST

1244 We parametrize the noising process through the SDE

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 $d\boldsymbol{\tau} = \sqrt{\beta(t)} d\mathbf{W} \,, \tag{55}$

1247 where we set the drift term to zero. Notice that this choice is consistent with a 2d-rotation of a 1248 function over the grid $\mathbf{x}_{i,j}$, given by $f(\mathbf{x}_{i,j}) = f_{i,j}$, denoting the value of the pixel of image f1249 at the location i, j. We train a convolutional neural network (CNN) with three convolutional layers 1250 followed by fully connected layers that outputs a single value, being the score for the flow coordinate 1251 τ . For the specific details of the implementation we refer to the code-base in the SI. In sampling, we 1252 apply a smoothing function to compensate interpolation artifacts due to rotations on a discretized 1253 grid. We choose T = 100 time-steps in training but only need T = 10 time-steps during sampling.

1254 1255 E.1.1 BBDM

1256 We implement the Brownian Bridge Diffusion Model (BBDM) (Li et al., 2023) and train it on the 1257 rotated MNIST dataset. The BBDM operates on the full pixel space \mathbb{R}^{784} of the 28 \times 28 MNIST 1258 digits and indicates a continuous time stochastic process conditioned on the starting $\mathbf{x}(0)$ and end 1259 point $\mathbf{x}(T)$ which are pinned together as paired data. In this case, we assume $\mathbf{x}(T) \sim p(\mathbf{x}_T)$ to be a randomly augmented MNIST digit obtained from an original MNIST digit $\mathbf{x}(0)$. During train-1260 ing, we sample an intermediate point $\mathbf{x}(t) \sim N(x_t | \mu_t(x(0), x(T)), \Sigma_t)$ where the mean function 1261 $\mu_t(t)(\mathbf{x}(0), \mathbf{x}(T))$ is a linear interpolation between the endpoints $(\mathbf{x}(0), \mathbf{x}(T))$ and use the score-1262 network to predict the original data point $\mathbf{x}(0) = s_{\theta}(x_t, t, x_T)$ as opposed to the noise or difference 1263 paramterization proposed in the original BBDM paper. We noticed that predicting the original data 1264 point led to better sampling quality including the inductive bias that MNIST digits are represented 1265 as binary tokens. Furthermore, we observe that the sampling quality is also better when the prior 1266 image x_T is input as context into the score network, enforcing a stronger signal throughout the tra-1267 jectory. As opposed to our model, we trained the BBDM on T = 1000 diffusion timesteps using the 1268 sin-scheduler from BBDM. 1269

1270 E.2 QM9 & CROSSDOCKED2020

1272 QM9 The conformer generation tasks is about learning a conditional probabilistic map \mathbf{x} \sim $p_{\theta}(X|M)$, where $\mathbf{x} \in \mathbb{R}^{3N}$ for a molecule with N atoms. We implement a variant of EQGAT 1273 (Le et al., 2022b) as neural network architecture where input features for the nodes consist of atom 1274 types and atomic coordinates, while edge features are encoded to indicate the existence of a single-, 1275 double, triple or aromatic bond based on the adjacency matrix. We use L = 5 message passing 1276 layers with $s_{\text{dim}} = 128$, $v_{\text{dim}} = 64$ scalar and vector features, respectively. To predict the scores for 1277 each atom, we concatenate the hidden scalar and vector embeddings $s \in \mathbb{R}^{128}$, $v \in \mathbb{R}^{3 \times 64}$ into one 1278 output embedding $o = \mathbb{R}^{128+3*64}$ which is further processed by a 2-layer MLP with three output 1279 units. Notice that the predicted scores per atom are neither invariant nor equivariant since the scalar 1280 and vector features are transformed with an MLP. 1281

We choose the drift f with its scaling β and the diffusion coefficients γ in such way that the forward SDE for the flow coordinates τ in (9) has the expression

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$$d\boldsymbol{\tau} = -\frac{1}{2}\beta(t)\boldsymbol{\tau}dt + \sqrt{\beta(t)}d\mathbf{W}, \qquad (56)$$

where for clarity we have omitted the dependency between the flow coordinates and the original data in Cartesian coordinates, i.e. $\tau(\mathbf{x})$, since the coordinate transformations with Lie algebra representation are described in A.3. The forward SDE in (56) is commonly known as *variance-preserving* SDE (Song et al., 2020b). We use the cosine scheduler proposed by Dhariwal & Nichol (2021) and T = 100 diffusion timesteps.

1292 **CrossDocked** For this experiment we adopt again an SDE of the form (56) for the three SO(3) 1293 flow coordinates $\theta_1, \varphi_1, \varphi_2$ and the three T(3) center of mass Cartesian flow coordinates. The SO(3) 1294 flow coordinates are always computed and applied in the ligand center of mass. In this way there 1295 is no ambiguity regarding the non-commutativity of SE(3), as rotation around the origin commute 1296 with translations of the system. We train a variant of EQGAT as in the QM9 case, but now including also node and edge features of the protein pocket. Specifically, the adjacency matrix for the GNN is computed dynamically at each time step, according to the relative distance between ligand and protein. For this, we choose a cut-off of 5 Å. We also use in this experiment a cosine scheduler and T = 100 diffusion timesteps. Since this learning problem is 6-dimensional, we aggregate the last layer's node embeddings from the ligand atoms into a global representation through summation. This embedding is fed as input into a 2-layer MLP to predict the six scores.

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E.2.1 RSGM ON CROSSDOCKED

1304 We utilized the framework of Riemannian Score-Based Generative Models (RSGM) by (De Bortoli 1305 et al., 2022) to model rigid-body motions on $G = (SO(3) \times T(3))$, in similar fashion to (Corso 1306 et al., 2023; Yim et al., 2023) by choosing a variance exploding SDE for the rotation dynamics and 1307 variance preserving SDE for the global translations. The terminal distribution for the rotation is 1308 designed to converge to an isotropic Gaussian distribution on SO(3) (Leach et al., 2022), while the 1309 terminal distribution for the translation component converges to an isotropic Gaussian in \mathbb{R}^3 . To ob-1310 tain the tractable scores for rotation and translation, we use the code by the authors from DiffDock 1311 and SE(3)-Diffusion for Protein Backbone Modeling in https://github.com/gcorso/ DiffDock/blob/main/utils/so3.py and https://github.com/jasonkyuyim/ 1312 se3_diffusion/blob/master/data/se3_diffuser.py and make sure that the score 1313 outputs for rotation and translation are SO(3) equivariant using the same EQGAT model archi-1314 tecture. The (variance-preserving) scheduler for the translation dynamics is chosen in similar 1315 fashion to our experiment using the cosine scheduler, while the (variance-exploding) scheduler 1316 for the rotation dynamics is implemented as an linear increasing sequence in \log_{10} space with 1317 $\sigma_{min} = 0.001$ and $\sigma_{max} = 2.0$ and T = 100 discretized diffusion steps as $\sigma(t) = 10^t$ for 1318 $t \in (\log_{10}(\sigma_{min}), \log_{10}(\sigma_{max})).$ 1319

To compare both modeling approaches with respect to the dynamics using the same network architecture, we perform 5 dockings per protein-ligand complex in the CrossDocked test dataset comprising 100 complexes and compute the mean RMSD between ground-truth coordinates and predicted coordinates.

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1325 F FURTHER OUTLOOK

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In the context of generative chemistry, particularly for modeling interactions within protein pockets, our methods could be employed to decouple the intrinsic generation of ligands from the global transformations required to fit the ligand into the pocket. This approach can also be extended beyond 3D coordinates, for instance, by working with higher-order representations, such as modeling electron density (Rackers et al., 2023).

Moreover, for more complex problems, it is feasible that an optimal generation process can be 1332 achieved by combining different choices of G along the trajectory. In the context of ligand gener-1333 ation, we propose a time-dependent group action $G_t = tT(3N) + (1-t)(SO(3) \times \mathbb{R}_+)^N$: at 1334 the beginning of the diffusion process, when the point cloud is still far from forming a recognizable 1335 conformer, we can leverage the properties of a true Gaussian prior. As the point cloud is gradu-1336 ally optimized to "resemble a molecule", we progressively transition to a generalized score-guided 1337 process. This shift allows us to fine-tune chemically relevant properties, such as bonds and torsion 1338 angles, ensuring that the intermediate and final conformers are chemically valid and accurate. This 1339 will be the focus of our forthcoming work.

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