

# Riemannian Flow Matching for Brain Connectivity Matrices via Pullback Geometry



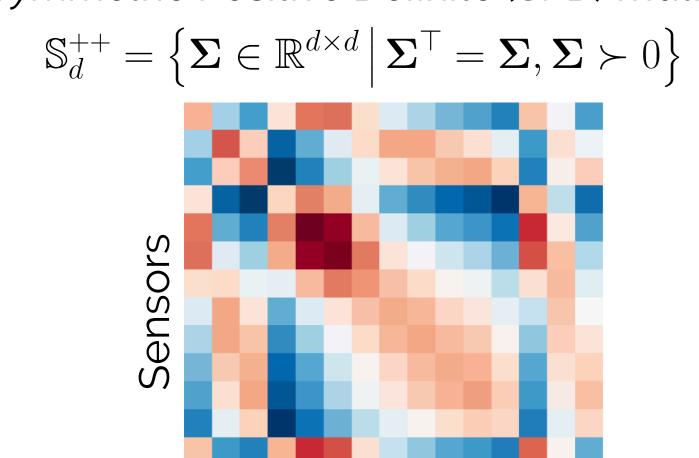
Inria, CEA, University Paris-Saclay



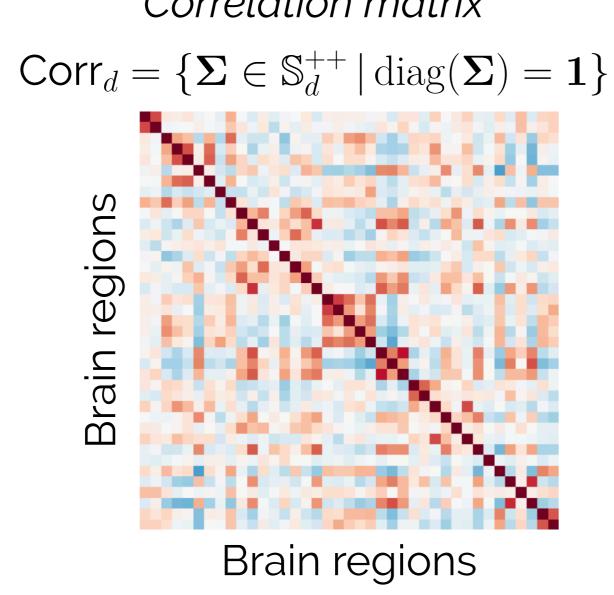
## **Brain Connectivity & Manifolds**

Modern neuroimaging analyzes signals via connectivity matrices:

# EEG Spatial Covariance Symmetric Positive Definite (SPD) matrix



# fMRI Functional Connectivity Correlation matrix



Challenges of brain connectivity generation:

Sensors

- 1. Euclidean methods (e.g., standard Flow Matching) generate non-SPD matrices.
- 2. Riemannian methods require expensive geodesics and Riemannian norms, making training slow ( $10 \times$  slower).

#### Contribution:

We propose DiffeoCFM that enables **Euclidean** Flow Matching on matrix manifolds with **pullback metrics** induced by global diffeomorphisms.

- Fast: Train and sample in Euclidean space.
- Exact: Equivalent to Riemannian Flow Matching on the pullback manifold.
- Valid: Guarantees manifold constraints by construction.

## Riemannian Flow Matching

Learn a vector field  $u_{\theta}^{\mathcal{M}}$  on the manifold  $\mathcal{M}$  to match geodesic velocities [1].

**Training:** Requires computing geodesics  $\gamma(t)$  and Riemannian norms  $\|\cdot\|_{\gamma(t)}$ :

$$\mathcal{L}(\theta) \triangleq \mathbb{E}_{t,x_0,x_1} \left\| u_{\theta}^{\mathcal{M}}(t,\gamma(t)) - \dot{\gamma}(t) \right\|_{\gamma(t)}^2$$

**Sampling:** Requires solving Riemannian ODEs on  ${\cal M}$ 

$$\dot{x}(t) = u_{\theta}^{\mathcal{M}}(t, x(t)), \quad x(0) = x_0$$

#### Pullback Manifold

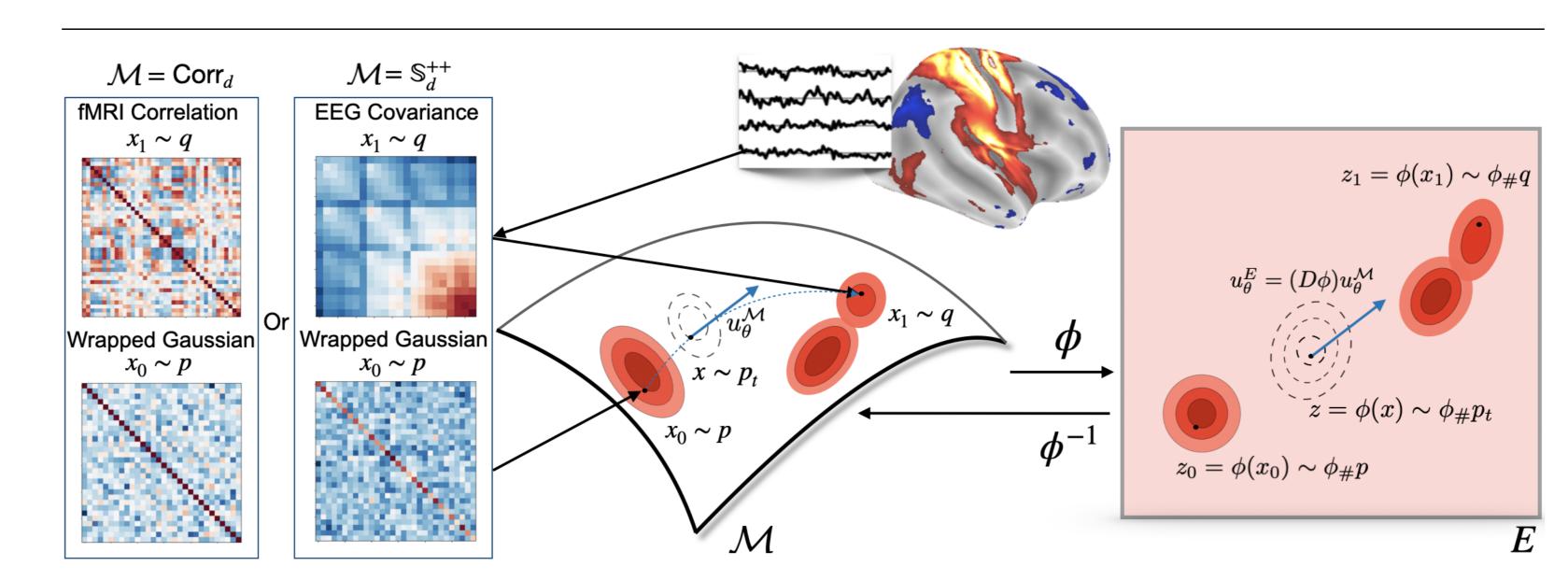
 $\phi: \mathcal{M} \to E$  a **global diffeomorphism** to a Euclidean space E,

$$g_x(\xi, \eta) \triangleq \langle \mathcal{D} \phi(x)[\xi], \mathcal{D} \phi(x)[\eta] \rangle, \quad \xi, \eta \in T_x \mathcal{M}$$

**Key Insight:** we can compute everything in E and map back to  $\mathcal{M}$  via  $\phi$ :

(Geodesic)  $\gamma(t) = \phi^{-1} ((1-t)\phi(x_0) + t\phi(x_1))$ (Distance)  $d_{\mathcal{M}}(x_0, x_1) = \|\phi(x_0) - \phi(x_1)\|_E$ 

#### DiffeoCFM: Overview



Assumption: There exists a global diffeomorphism  $\phi:\mathcal{M}\to E$  to a Euclidean space E.

- **Training**: Map data to E via  $\phi$  and learn a Euclidean vector field  $u_{\theta}^{E}$  to match straight-line velocities.
- Sampling: Solve Euclidean ODE  $\dot{z}(t) = u_{\theta}^{E}(t,z)$  in E and map back to  $\mathcal{M}$  via  $\phi^{-1}$ .

#### **Theoretical Guarantees**

We prove that training in Euclidean space is **exactly equivalent** to Riemannian Flow Matching on  $(\mathcal{M},g)$  by defining the pulled-back vector field:

$$u_{\theta}^{E}(t,z) \triangleq \mathcal{D} \phi(\phi^{-1}(z)) \left[ u_{\theta}^{\mathcal{M}}(t,\phi^{-1}(z)) \right]$$

Proposition 1 (Training Equivalence): The Riemannian loss simplifies to the standard Euclidean Flow Matching loss on transformed data  $z = \phi(x)$ :

$$\mathcal{L}(\theta) = \mathbb{E}_{t, z_0, z_1} \underbrace{\left\| u_{\theta}^E\left(t, (1-t)z_0 + tz_1\right) - \left(z_1 - z_0\right) \right\|_E^2}_{\text{Standard Euclidean Flow Matching Loss}}$$

Proposition 2 (Sampling Equivalence): Let z(t) be the solution to  $\dot{z}(t) = u_{\theta}^{E}(t,z)$  in E. Then

$$x(t) = \phi^{-1}(z(t))$$

is the exact solution to the Riemannian ODE on  $\mathcal{M}$ .

 $\rightarrow$  We avoid all manifold-specific operations (Exp, Log, Parallel Transport) during training and sampling while being exactly equivalent to Riemannian Flow Matching on  $(\mathcal{M}, g)$ .

# Diffeomorphisms for covariance and correlation matrices

 $\phi_{\mathbb{S}_d^{++}}(\mathbf{\Sigma}) = \operatorname{vec}(\log(\mathbf{\Sigma}))$ 

We use specific global diffeomorphisms for neuroimaging data:

- Covariance ( $\mathbb{S}_d^{++}$ ): Log-Euclidean map  $\phi_{\mathbb{S}_d^{++}}:\mathbb{S}_d^{++}\to\mathbb{R}^{d(d+1)/2}$ .
- Correlation (Corr $_d$ ): Normalized Cholesky map  $\phi_{\mathsf{Corr}_d}$ :  $\mathsf{Corr}_d \to \mathbb{R}^{d(d-1)/2}$ .  $\phi_{\mathsf{Corr}_d}(\mathbf{\Sigma}) = \mathrm{vec}(\mathrm{NormChol}(\mathbf{\Sigma}))$

### **Experiments**

#### **Datasets:**

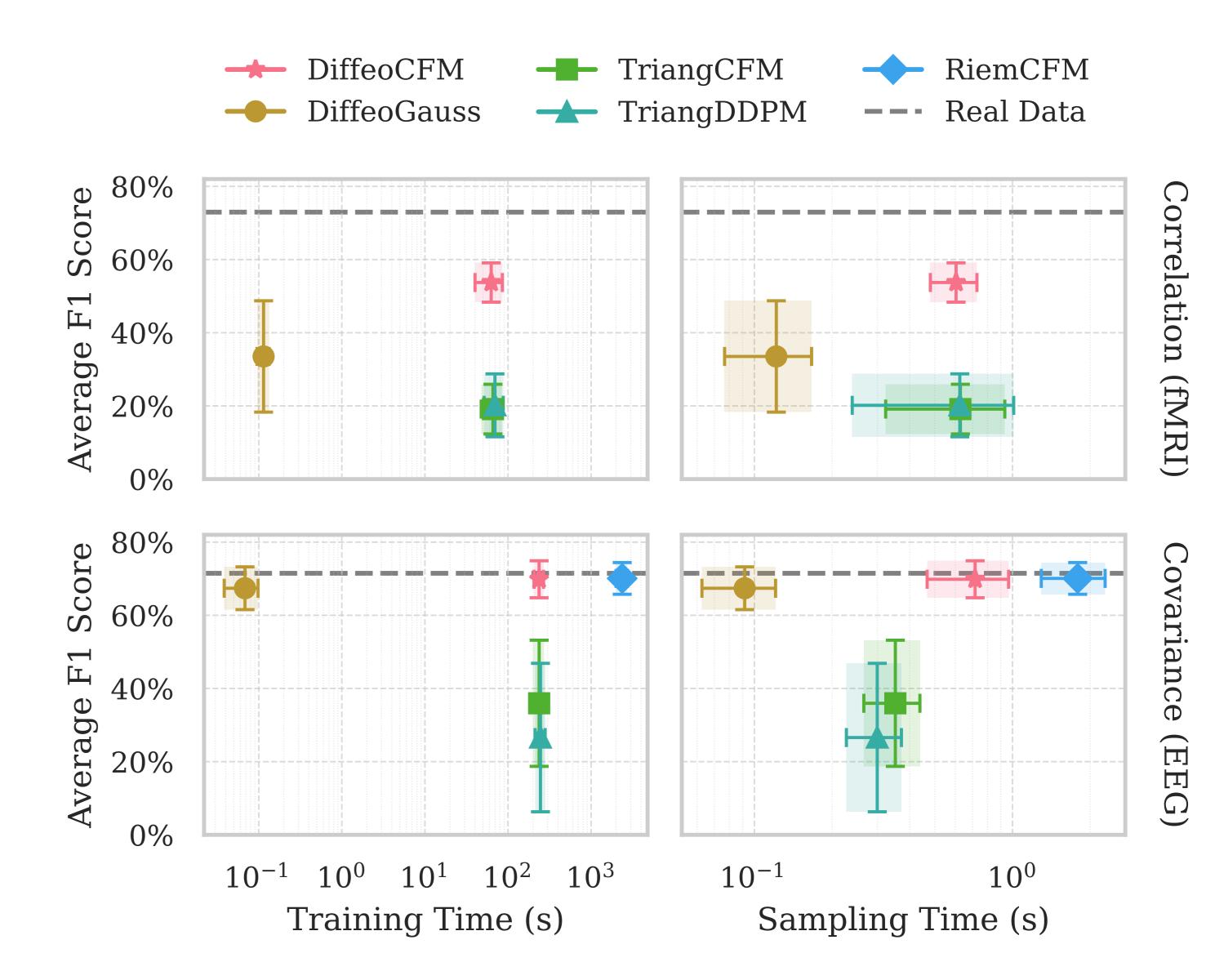
- fMRI (correlation): ABIDE, ADNI, OASIS-3, 4600 scans from 2800 subjects with disease labels (Autism, Alzheimer's, Control) and 10 subject-level splits.
- EEG (covariance): BNCI2014, BNCI2015, >30k trials from 26 subjects with right hand vs. feet motor imagery labels and cross-session splits.

#### **Baselines:**

- DiffeoGauss: one Gaussian per class in E with  $\phi$  [2].
- TriangDDPM / TriangCFM: generates triangular part with DDPM or FM + projection  $\Sigma \leftarrow (1 \alpha)\Sigma + \alpha I$ .
- RiemCFM: Riemannian Flow Matching (affine-invariant metric) [1].

#### **Evaluation Metrics:**

- Distribution Quality:  $\alpha$ -Precision (fidelity) &  $\beta$ -Recall (diversity) [3].
- Utility (CAS): train a classifier on *generated* data  $\rightarrow$  test on *real* data [4].
- Efficiency: training and sampling time.



#### References

- [1] Ricky TQ Chen and Yaron Lipman. "Flow matching on general geometries". In: *The Twelfth International Conference on Learning Representations*. 2024.
- [2] Thibault de Surrel et al. "Wrapped Gaussian on the manifold of Symmetric Positive Definite Matrices". In: Forty-second International Conference on Machine Learning. 2025.
- [3] Ahmed Alaa et al. "How faithful is your synthetic data? sample-level metrics for evaluating and auditing generative models". In: *International Conference on Machine Learning*. PMLR. 2022, pp. 290–306.
- [4] Suman Ravuri and Oriol Vinyals. "Classification accuracy score for conditional generative models". In: *Advances in neural information processing systems* 32 (2019).

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