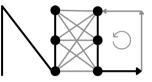
## Hands on NeuLat: A Toolbox for **Neu**ral Samplers in **Lat**tice Field Theory



#### NEULAT

#### Kim A. Nicoli, Christopher J. Anders et al.

December 06, 2024 - Taipei Lattice Field Theory and Machine Learning Workshop code: https://github.com/neulat/neulat preprint: LATTICE2023 PoS 286

## Sampling from Boltzmann Distributions

# $p(\phi) = \frac{\exp\{-S[\phi]\}}{Z}$

## Sampling from Boltzmann Distributions.... with Generative Models

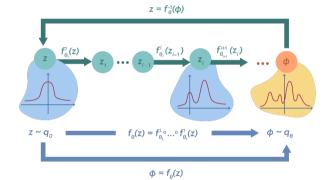
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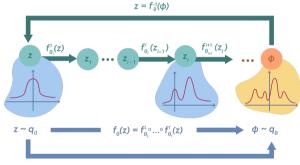
$$p(\phi) = \frac{\exp\{-S(\phi)\}}{Z} \xrightarrow{q_{\theta}} p(\phi) \approx q_{\theta} \sim \phi_i$$

Bare with me for now, and wait for Tej's Mathis' and Fernando's talks tomorrow!

## Sampling with Normalizing Flows



## Sampling with Normalizing Flows



 $\phi = f_{\theta}(z)$ 

$$q_{ heta}(\phi) = q_0(f_{ heta}^{-1}(\phi)) \left| \det \left( rac{\partial f_{ heta}}{\partial z} 
ight) \right|^{-1}$$



## Christopher J. Anders



**Christopher J. Anders** PostDoc at RIKEN AIP, Tokyo

software framework for machine-learning-based lattice field theory



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■ software framework for machine-learning-based lattice field theory
 ■ e.g., φ<sup>4</sup>-theory, U(1) gauge theory, up to 3 + 1D



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unifies existing tools into one framework



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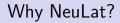
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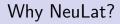
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Existing examples:

- SchNetPack Deep Neural Networks for Atomistic Systems
- **BGFlow** Boltzmann Generators (BG) and other sampling methods



Introduction to Normalizing Flows for LFT [Albergo et al., 2101.08176 (2021)]



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- Flows/HMC for LFT
  - GomalizingFlow.jl [github.com/AtelierArith/GomalizingFlow.jl] (A. Tomiya)
  - NeuMC [github.com/nmcmc/nmcmc-code] (P. Bialas)

## Why NeuLat?

There are already great tools available!

- Introduction to Normalizing Flows for LFT [Albergo et al., 2101.08176 (2021)]
- Flows enhanced HMC
  - **fthmc** (Sam Foreman et al.) [github.com/nftqcd/fthmc] (S. Foreman)
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But: We want to create a highly customizable reference implementation.

**Density Estimator**: Learn approximations of targeted Boltzmann distributions

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- Sampling:
  - various MCMC implementations (HMC, Cluster algorithms, NE-HMC, etc.)
  - Normalizing Flow framework
    - Neural Importance Sampling (NIS)
    - Neural HMC

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#### Estimation:

- Asymptotically unbiased estimators for physical observables [Nicoli et al. (2020)]
- Direct estimation of thermodynamic observables [Nicoli et al., (2021)]
- Sampling in the presence of mode-collapse [Nicoli et al., (2023)]

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#### Tutorials and Documentation:

- Step-by-step tutorials
- Extensive reference

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#### Tutorials and Documentation:

- Step-by-step tutorials
- Extensive reference
- Modularity and Customizability: Swiftly incorporate new actions/theories/models/techniques

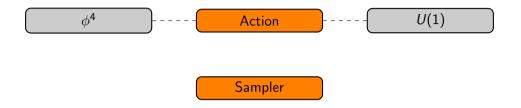
## NeuLat Overview



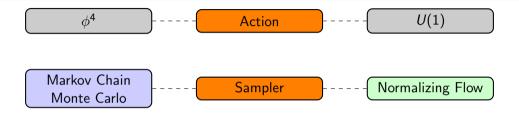
Actions S[U] define target Boltzmann distributions  $p(U) \propto e^{-S[U]}$ 



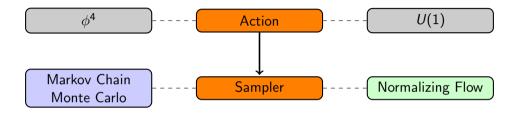
Actions are, e.g.,  $\phi^4$  and U(1)



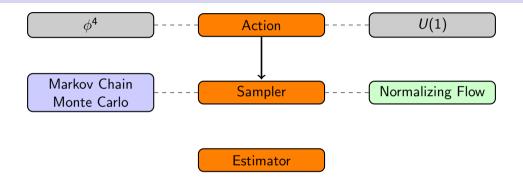
Samplers are anything that can be sampled from



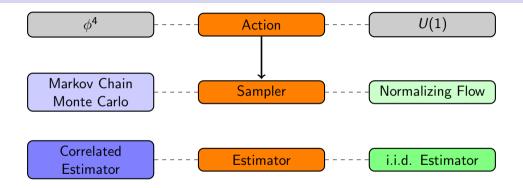
Samplers are, e.g., MCMCs, Flows,  $\mathcal{N}(0,1)$ 



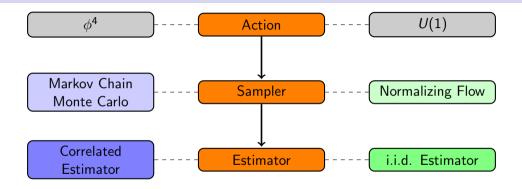
Samplers require Actions



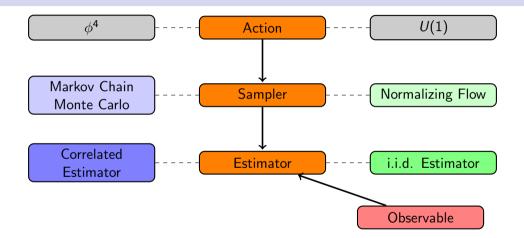
Estimators are used to estimate observables



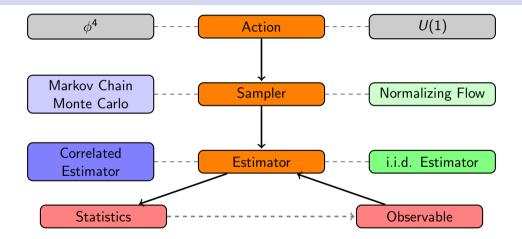
Estimators are, e.g., i.i.d or correlated, based on the samples



Estimators require samples from Samplers



Observables, e.g., Magnetization, Topological Charge, etc., are used by the Estimator



The resulting Statistics are estimations for the Observables

### Actions

Actions are classes (objects) and need to be instantiated.

```
1 import torch
2 from neulat.action.phi4 import Phi4Action
3 
4 # ndim_features is the number of dimensions in the lattice
5 action = Phi4Action(kappa=0.3, lamb=0.022, ndim_features=2)
```

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```

Action objects can be called to compute action values for configurations.

```
1 \quad config = torch.randn(8, 8)
```

```
2 unnormalized_prob = torch.exp(-action(config))
```

## **Defining Actions**

Actions are **very simple** to implement, for instance the  $\phi^4$  action reads:

```
from neulat.action.base import Action
1
2
    class Phi4Action(Action):
3
        name = 'phi4_action'
4
        def __init__(self, kappa, lambd, ndim_feature=2):
5
6
        def forward(self, config):
7
             dims = tuple(range(-1, -self.ndim_feature, -1))
8
             kinetic = (-2 * self.kappa) * config * sum(
q
                 torch.roll(config, 1, dim) for dim in dims)
10
            mass = (1 - 2 * \text{self.lambd}) * \text{config } ** 2
11
             inter = self.lambd * config ** 4
12
            return (kinetic + mass + inter).sum(dim=dims)
13
```



At the core of NeuLat are Samplers, which are anything from which we can **sample**.

 $\frac{1}{2}$ 

3

At the core of NeuLat are Samplers, which are anything from which we can sample.

For instance, the normal distribution is also a Sampler in Neulat:

```
from neulat.sampler.distribution import Normal
```

```
normal = Normal(loc=0., scale=1., feature_shape=(8, 8))
```

## Sampling

Samplers can sample, and may or may not support probability values.

```
1 samples = normal.sample(sample_shape=8)
2 logprobs = normal.logprob(samples)
3 
4 samples2, logprobs2 = normal.sample_with_logprob((2, 2))
```

## Sampling

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3 4 samples2, logprobs2 = normal.sample_with_logprob((2, 2))
```

In NeuLat, we assume configurations of shape (\*sample\_shape, \*feature\_shape).

- sample\_shape is the number of samples, supporting arbitrary shapes
- **feature\_shape** is the shape of the lattice (i.e., # of dimensions)

## Hamiltonian Monte Carlo

A more involved Sampler is the HMC:

```
from neulat.sampler.mc.hmc import HMCMarkovChain
1
2
    hmc = HMCMarkovChain(
3
        action. # action
4
        feature_shape=(8, 8), # lattice shape
5
        burn_in=5000, # equilibration steps
6
        skip_interval=1, # skipped samples in chain
7
        overrelax_interval=50, # steps between sign flips
8
        eps=0.05, # step size along a trajectory
q
        traj_steps=20, # number of steps in the trajectory
10
        bias=0.0. # bias in initialization
11
12
```

## HMC Sampling 1/2

As for any Sampler, we can sample from HMC

1 configs = hmc.sample(sample\_shape=13)

# HMC Sampling 1/2

1

1

2

3

As for any Sampler, we can sample from HMC

```
configs = hmc.sample(sample_shape=13)
```

However, HMC does not implement logprob and, by extension, sample\_with\_logprog, as no normalized probabilities are available. (unnormalized logprobs possible)

```
# both cause exceptions:
# logprobs = hmc.logprob(sample_shape=13)
# configs2, logprobs2 = hmc.sample_with_logprob(13)
```

One can also iterate over HMC chains to sample

```
1 configs = []
2 for n, config in zip(range(25), hmc):
3 configs.append(config)
4 print(f'Sampled config number {n}.')
5
6 # this gives a list of configs, combine them:
7 configs = torch.cat(configs)
```

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#### !! But be careful, HMC chains are infinite iterators. !!

# Normalizing Flows

Normalizing flows [Papamakarios et al., JMLR (2021)] require a base distribution, and a transform.

```
1 from neulat.sampler.flow import Flow, SequentialTransform
2
3 flow = Flow(
4 base_distribution=Normal(feature_shape=(8, 8)),
5 transform=SequentialTransform([]) # identity for demo
6 )
```

# Normalizing Flows

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Normalizing flows [Papamakarios et al., JMLR (2021)] require a base distribution, and a transform.

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```

Normalizing flows are (i.i.d.) Samplers supporting logprobs.

```
configs, logprobs = flow.sample_with_logprob(8)
```

## Normalizing Flows: Base Distributions

The base distribution can be any sampler that supports logprobs.

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1

 $\mathbf{2}$ 

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Commonly, simple distributions such as  $\mathcal{N}(0,1)$  are used.

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## Normalizing Flows: Base Distributions

1

2

3 4

1

2

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```
flow = Flow(
   base_distribution=Normal(feature_shape=(8, 8)),
   transform=SequentialTransform([]) # identity for demo
)
```

Flows themselves support logprobs, and can thus be base distributions.

```
flow2 = Flow(
   base_distribution=flow,
   transform=SequentialTransform([]) # identity for demo
)
```

# Normalizing Flows: Transforms 1/2

Transforms are (optionally) invertible PyTorch modules, and require (either or both) a forward and a inverse.

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E.g., implementation for transform  $f(\mathbf{x}) = -\mathbf{x}$ ,  $f^{-1}(\mathbf{x}) = -\mathbf{x}$ 

```
from sampler.flow.base import Transform, withlogdet
1
2
    class FlipSign(Transform):
3
        @withlogdet
\mathbf{4}
        def forward(self, input):
\mathbf{5}
             return -input. 0.
6
        @withlogdet
7
        def inverse(self, input):
8
             return -input, 0.
9
```

# Normalizing Flows: Transforms 1/2

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E.g., implementation for transform  $f(\mathbf{x}) = -\mathbf{x}$ ,  $f^{-1}(\mathbf{x}) = -\mathbf{x}$ 

from sampler.flow.base import Transform, withlogdet 1  $\mathbf{2}$ class FlipSign(Transform): 3 **@withlogdet** 4 def forward(self, input): 5 return -input. 0. 6 **@withlogdet** 7 def inverse(self, input): 8 return -input. 0. 9

Decorator @withlogdet ensures the logdet is accumulated between transforms.

# Normalizing Flows: Transforms 2/2

A useful transform is the SequentialTransform, which is used to apply transforms sequentially:

```
1 from sampler.flow.base import SequentialTransform
2
3 flip_a_bunch = SequentialTransform([
4 FlipSign(),
5 FlipSign(),
6 FlipSign(),
7 ])
```

# Normalizing Flows: Transforms 2/2

A useful transform is the SequentialTransform, which is used to apply transforms sequentially:

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1 from sampler.flow.base import SequentialTransform
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3 flip_a_bunch = SequentialTransform([
4 FlipSign(),
5 FlipSign(),
6 FlipSign(),
7 ])
```

For common *Coupling* Flows, e.g., NICE, ReaINVP, there is, however, a more convenient way.

## Coupling Flows

Coupling flows like NICE consist of two parts, a partitioner, and a net\_factory

```
1 from neulat.sampler.flow.coupling import NICE
2
3 coupling = NICE(
4 partitioner=partitioner,
5 net_factory=net_factory
6 )
```

The partitioner partitions (or masks) the input into active and passive components.

# **Coupling Flows**

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4 partitioner=partitioner,
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6 )
```

- The partitioner partitions (or masks) the input into active and passive components.
- The net\_factory is a function that constructs the conditioner, e.g., a neural network that acts on the partitioned input.

# Coupling Flows: Partitioners

A very simple partitioner is the AltFlatPartitioner, which stands for **alt**ernating **flat**tened partitioner

```
1 partitioner = AltFlatPartitioner(feature_shape=(2, 2)),
2 input = torch.tensor([[1., 2.],[3., 4.]])
3 active, passive = partitioner(input)
4 active += 10
5 output = partitioner(active, passive)
```

This will generate an output of 
$$\begin{pmatrix} 11 & 2\\ 13 & 4 \end{pmatrix}$$

# Coupling Flows: Partitioners

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```

This will generate an output of 
$$\begin{pmatrix} 11 & 2\\ 13 & 4 \end{pmatrix}$$

**N.B.** Inputs can be partitioned into more than two parts, e.g., active, passive, frozen.

Partitioners usually flip the active and passive elements. Such a partitioner can be created by calling .flip():

```
1 flipped = partitioner.flip()
2 input = torch.tensor([[1., 2.],[3., 4.]])
3 active, passive = flipped(input)
4 active += 1
5 output = flipped(active, passive)
```

Partitioners usually flip the active and passive elements. Such a partitioner can be created by calling .flip():

```
1 flipped = partitioner.flip()
2 input = torch.tensor([[1., 2.],[3., 4.]])
3 active, passive = flipped(input)
4 active += 1
5 output = flipped(active, passive)
```

This will generate an output of 
$$\begin{pmatrix} 1 & 12 \\ 3 & 14 \end{pmatrix}$$

# Coupling Flows: Net Factory (Conditioner)

The net\_factory defines the conditioner  $\Theta$  that transforms the passive input:

$$\mathbf{x}_{\text{active}}^{\prime+1} = h(\mathbf{x}_{\text{active}}^{\prime}, \Theta(\mathbf{x}_{\text{passive}}^{\prime})) \tag{1}$$

```
from functools import partial
1
    from neulat.sampler.flow.coupling.affine import NICE, MLP
2
3
    net_factory = partial(
4
        MLP,
5
        n_blocks=3.
6
        latent_size=1024.
7
        activation=torch.nn.Tanh,
8
        bias=False.
9
10
```

The coupling Transform itself is mostly only concerned with implementing the coupling function h. E.g. in NICE:  $h(\mathbf{x}_{active}, \mathbf{x}_{passive}) = \mathbf{x}_{active} + m_{\theta}(\mathbf{x}_{passive})$ 

```
class NICE(Coupling):
1
        @withlogdet
2
        Opartitioned
3
        def forward(self, active, passive):
4
            return active + self.net(passive), 0.
5
6
        @withlogdet
7
        Opartitioned
8
        def inverse(self, active, passive):
9
            return active - self.net(passive), 0.
10
```

# Coupling Flows: Defining Couplings

The coupling Transform itself is mostly only concerned with implementing the coupling function *h*. E.g. in NICE:  $h(\mathbf{x}_{active}, \mathbf{x}_{passive}) = \mathbf{x}_{active} + m_{\theta}(\mathbf{x}_{passive})$ 

```
class NICE(Coupling):
1
        @withlogdet
2
        Opartitioned
3
        def forward(self, active, passive):
4
             return active + self.net(passive), 0.
\mathbf{5}
6
        @withlogdet
7
        Opartitioned
8
        def inverse(self, active, passive):
q
             return active - self.net(passive), 0.
10
```

Recall: @withlogdet makes sure the log abs jacobian det is propagated.

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```

New: @partitioned automates the partitioning in subsequent couplings!

### Coupling Flows: Assembling the Flow

Putting all the previous parts together, we can create a Flow in the following way:

```
1 flow = Flow(
2 base_distribution=Normal(0.0, 1.0, feature_shape=(8, 8)),
3 transform=6 * NICE(
4 partitioner=AltFlatPartitioner(feature_shape=(8, 8)),
5 net_factory=partial(MLP, n_blocks=3, latent_size=1024,
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**Note:** the transform=6 \* NICE. This creates a sequential transform of 6 Couplings, with alternating masking/partitioning!

# Coupling Flows: Global Scaling

Global scaling is a Transform that scales all elements of the input tensor by the **same** scalar value.

This is necessary to enhance the convergence of the model:

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7
8
   )
```

**Note:** Transform has an add method that allows simple concatenatenation of different types of transforms.

# Coupling Flows: Traning

Training of the flow with, e.g., ReverseKL, is straightforward:

```
from neulat.loss import ReverseKLLoss
1
\mathbf{2}
    optim = torch.optim.Adam(flow.transform.parameters(), lr=5e-4)
3
    loss fn = ReverseKLLoss()
4
    for _ in range(1000):
5
        configs, log_probs = flow.sample_with_logprob(10)
6
        loss = loss_fn(action(configs), log_probs) # loss contains `mean` and `std`
7
        optim.zero_grad()
8
        loss.mean.backward() # we train only using the loss `mean`
q
        torch.nn.utils.clip grad norm (model.transform.parameters(), 1.0) # clipping
10
        optim.step()
11
```

#### Estimating Observables: IidEstimator

Observables themselves are classes in NeuLat. In order to estimate them, we additionally need an Estimator, and configurations. For instance:

```
1 from neulat.observable.base import AbsMagnetization, Magnetization
2 from neulat.estimator.base import IidEstimator
3
4 observables = [AbsMagnetization(), Magnetization(), action]
5 iid_estimator = IidEstimator(observables)
6 configs = flow.sample(1000)
7 flow_statistics = iid_estimator.named_evaluate(configs)
```

The dict flow\_statistics will contain one entry per observable:

{ 'absmag': Statistics(mean=0.6408, std=0.0473), 'mag': ...}

Estimation of Observables from correlated samples (e.g., from HMC) requires the use of the appropriate estimator:

```
1 from neulat.estimator.base import CorrelatedEstimator
2 
3 correlated_estimator = CorrelatedEstimator(observables)
4 configs = hmc.sample(1000)
5 hmc_statistics = correlated_estimator.named_evaluate(configs)
```

The dict hmc\_statistics will instead contain correlated statistics objects:

```
{'absmag': CorrelatedStatistics(mean=32.82628, std=1.4674,
tau_int=0.5909, tau_int_err=0.3162), ...}
```

# Estimating Observables: ImportanceSamplingEstimator

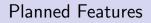
To obtain an unbiased estimator, Nicoli et al., (2020) proposed to use Neural Importance Sampling (NIS).

This additionally requires the logprobs from the flow, as well as the specific action:

```
1 from neulat.estimator.base import ImportanceSamplingEstimator
2
3 flow_configs, flow_logprobs = flow.sample_with_logprob(1000)
4 iw_estimator = ImportanceSamplingEstimator(observables, action)
5 flow_iw_stats = iw_estimator.named_evaluate(flow_configs, flow_logprobs)
```

The dict flow\_iw\_statistics will contain the same Statistics object returned by the IidEstimator:

{'absmag': Statistics(mean=2.6021, std=0.4674), ...}



Non-Equilibirum MCMC [Caselle et al., Phys. Rev. D (2016)]

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With the help of the community, we plan to extend NeuLat into many directions, including following features

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- YOUR FEATURE HERE!

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#### Conclusion

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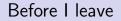
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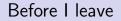
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And now it's time for a real demo...



NeuLat will be available soon at

# https://github.com/neulat/neulat



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# Thank you for your attention!