



## Machine learning: simulation and data



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## Before we start



- Go to my binder: <a href="https://binderhub.ssl-hep.org/v2/gh/rafaellopesdesa/hsfindia-generative/HEADgpu\_false">https://binderhub.ssl-hep.org/v2/gh/rafaellopesdesa/hsfindia-generative/HEADgpu\_false</a>
- Go to the reweighting exercise and run the pip command.
- Let it run while I talk about what we will do...

Run this cell and let's talk about physics while it installs

#### Reweighting MC simulation to data using a NN

Reweighting MC simulations to data is a common task used to improve the modelling. The most common practice is to reweight a single used to perform the reweighting task by considering multiple variables together, which improves the modelling across multiple variables

Many thanks to Michele Faucci Giannelli, Marilena Bandieramonte and Martina Javurkova

: !pip install -U tensorflow !pip install -U tensorflow-probability !pip install -U matplotlib !pip install -U scipy



The goal of LHC analysis is to compare the data to a probability model for different hypothesis, usually Standard Model (SM) vs New Physics (NP)

$$p(x^{\text{data}}|H_1)$$
 and  $p(x^{\text{data}}|H_0)$ 

Building the p(x|H) models is **really complicated**.

In most cases p(x|H) are approximated by histograms where the number of events (I will call it v) in each bin comes from simulation

 $\rightarrow$  Counting MC (simulation) events.

# How do we count MC events?



In practice, MC events have weights

$$p(x|H_0) = \frac{v_{\rm I}^{\rm bkg}}{v^{\rm bkg}} = \frac{\sum_{\rm bkg}^{\rm bin\,I} w_i}{\sum_{\rm bkg} w_i}$$

$$p(x|H_1) = \frac{v_{\mathrm{I}}^{\mathrm{bkg}} + v_{\mathrm{I}}^{\mathrm{sig}}}{v^{\mathrm{bkg}} + v^{\mathrm{sig}}} = \frac{\sum_{\mathrm{bkg}}^{\mathrm{bin}\,I} w_i + \sum_{\mathrm{sig}}^{\mathrm{bin}\,I} w_i}{\sum_{\mathrm{bkg}} w_i + \sum_{\mathrm{sig}} w_i}$$

But what is  $w_i$  again? In simulation, the probability of a given event is given by the differential cross section

 $\frac{d\sigma}{dz}\simeq w_i$ 

Sometimes, we can "unweight" events, but not always...



Cross sections are all positive. So why do we have negative weights?

#### Perturbation theory, parton shower, and interference



introduced by parton showers. Negative weights are a common way to remove double counting.







# Simulation





#### **Full simulation**

- Common software framework (usually Geant4, but others exist)
- Experiments provide additional code (digitization, reconstruction, ...)
- Explicit modeling of detector geometry, materials, interactions w/ particles

Add thousands of additional variables. Sequential sampling

$$z_i \sim p(z_i | z_{j < i})$$
 and  $x \sim p(x | z_i)$ 

#### **Fast simulation**

- Usually experiment-specific framework
- Explicit modeling of detector geometry
- Add approximations: analytical shower shapes, truth-associated track reconstruction, ...





#### **Parametrized simulation**

- Does not describe the detector
- Replaces entire chain ("end-to-end")
- Can be done with analytical function or machine learning methods

 $x \sim p(x|z)$ 

# **Simulation landscape**

How can we use ML with simulation?

- Augment the full simulation
  - · Improve the MC weights with data
  - Calibrate your simulation
- Replace (part of) full simulation
- Create an "end-to-end" parametrized simulation

#### Goals:

- 1. Increase speed while preserving accuracy
- 2. Preserve speed while increasing accuracy



Accuracy



# Augment full simulation



• Usually deterministic.

#### **Classification based**

Uses a classification loss, like the one you tried in Gordon's lectures

$$L = -\frac{1}{N} \sum_{i} w_i [y_i \log s_i + (1 - y_i) \log(1 - s_i)]$$

The minimum of this loss function is achieved at:

$$s(x) = \frac{p_1(x)v_1}{p_0(x)v_0 + p_1(x)v_1}$$
  
If  $v_0 = v_1$  (balanced)  $\frac{p_1(x)}{p_0(x)} = \frac{s}{1-s}$ 

#### **Regression based**

Uses a regression loss, for instance MSE (there are others):

$$L = -\frac{1}{N} \sum_{i} w_i (y_i - s_i)^2$$

Creates a calibration function  $s_i(x_i)$ 

But it only calibrates the average (conditional on  $x_i$ ), not full distributions.

# Classification-based reweighting





- Reweight between CR and SR need to be validated carefully.
- Try the activity in binder!
  - <u>https://binderhub.ssl-hep.org/v2/gh/rafaellopesdesa/hsfindia-generative/HEADgpu\_false</u>

underfitting



High variance

overfitting

rror

Low bias, low variance

Good balance

some observable

## Intermezzo



- Now let's do the second part...
- Go to the generative directory in binder and run the pip cell (it assumes you are using the same session as the reweight one)



Introduction

This exercise is based on a normalizing flow exercise designed by T.Quadfasel, M.Sommerhalder and S.Diefenbacher, https://github.com/uhh-pd-ml/flow-exercise

Broadly speaking the exercise is organized into two parts.

- The first part takes a look at examples of some of the introduced generative models using the Two Moons data set
- The second part focuses on normalizing flows and how to build them using the nFlows package

# **Optimal transport**

- Moving points instead of reweighting histograms
- "Optimal" : Transport minimize some cost (L<sup>2</sup>)
  - Order preserving transformation between P and Q
- Easily scalable to higher dimensions
- Correct higher order correlation









# 1D optimal transport



$$p = 2$$
, i.e.  $c(x, y) = |x - y|^2$ 

#### For 1-dimensional distributions:

The optimal transport solution performs quantile-matching (works for all convex cost functions!)

 $\hat{T}(x) = Q^{-1}(P(x))$ Cumulative distributions of p(x), q(y): Generically:  $F(x) = \int_{0}^{x} dx' f(x')$ 





# ML optimal transport



Idea: Why not move the simulation instead of reweighting it?

- Optimal Transport
- Continuous calibration without histograms
- Easily scales to higher dimensions and cheap
- Integral of sample unchanged



$$\hat{T} = \arg \min_{T} \int dx \ p(x) \ c(x, T(x))$$
$$\pi(x, y) = p(x) \ \delta[y - T(x)] \qquad q(y) = p(x) \left(\frac{dT}{dx}\right)^{-1}$$

$$\hat{\pi} = \arg \min_{\pi} \int dx \, dy \, \pi(x, y) \, c(x, y)$$
$$\int dy \, \pi(x, y) = p(x) \qquad \int dx \, \pi(x, y) = q(y)$$

$$\hat{f}, \hat{g} = \arg \max_{f,g} \int dy \, q(y) \, f(y) + \int dx \, p(x) g(x) + \int dx \, p(x) g(x)$$

# **Optimal transport**



- Very recently, a solution on how to train multi-dimensional OT with ML has been found.
- Brand new area of ML that is just now finding applications



against data side bands)

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## Generative methods

- Generative models ("replace"):
  - Usually stochastic
  - Generative Adversarial Networks (GANs)
  - Variational Autoencoders(VAEs)
  - Normalizing Flows (NFs)







## **Generative Adversarial Network**

- Generator Network G(z) = x
- Maps noise *z* to *x*



## Generative Adversarial Network

- Generator Network G(z) = x
- Maps noise *z* to *x*
- Discriminator D(G(z)) and D(x)
- Learns difference between real and fake
- *D*(*G*(*z*)) is differentiable function measuring performance
- Use D(G(z)) as loss to update G



Machine learning: simulation and data



## **Generative Adversarial Network**

```
BCEloss = nn.BCELoss()
for ep in range(epochs):
   for i batch in range(max batches):
       # select the current batch from the dataset
       x_real = X_moons[i_batch * batch_size : (i_batch + 1) * batch_size]
       x real = torch.tensor(x real, device=device).float()
       DiscriminatorOpt.zero grad()
       with torch.no_grad():
           noise = torch.randn((batch_size, 8), device=device).float()
           x fake = GeneratorNet(noise)
       y real = torch.ones((batch size, 1), device=device)
       y fake = torch.zeros((batch size, 1), device=device)
       y = torch.cat((y_real, y_fake), 0)
       x = torch.cat((x_real, x_fake), 0)
       Discriminator_loss = BCEloss(DiscriminatorNet(x), y)
       Discriminator loss = Discriminator loss.mean()
       Discriminator loss.backward()
       DiscriminatorOpt.step()
       GeneratorOpt.zero grad()
       noise = torch.randn((batch size, 8), device=device).float()
       x_fake = GeneratorNet(noise)
       Generator loss = BCEloss(DiscriminatorNet(x fake), y real)
       Generator loss = Generator loss.mean()
       Generator loss.backward()
       GeneratorOpt.step()
```



## Upsides

- Intuitive approach
- Easy to introduce additional constraints
- Well explored with several improvements (WGANs, normalizations)

## Downsides

- Difficult to train
- Gen. and disc. needs to be balanced
- Can fail to converge
- Prone to mode collapse

# Simulation of showers in ATLAS calorimeter





# Simulation of showers in ATLAS calorimeter

#### FastCaloGAN V2

Different GAN for different type of particles and for different eta slices.

Prediction of deposit of energy in "voxels" which allow HITS reconstruction.





## How do we use this in a fast MC?



	Inner Detector	Calorimeters	Muon Spectrometer
Electrons Photons		FastCaloSimv2	
Hadrons	Geant4	Geant4 pions: $E_{kin} < 200 \text{ MeV}$ Other hadrons: $E_{kin} < 400 \text{ MeV}$ FastCalo Sim V2FastCalo GANFastCalo Sim V2 $K_{kin} < 200 \text{ MeV}$ $E_{kin} < (8-16) \text{ GeV}$ $K_{kin} < (8-16) \text{ GeV} < E_{kin}$ $K_{kin} > (256 - 512) \text{ GeV}$	Muon Punchthrough +Geant4
Muons		Geant4	Geant4

# Variational AutoEncoders (VAE)



- Encoding function *E*(*x*) = *z* map high dimensional data *X* to low dimensional latent space *Z*
  - Decoding function D(z) = x map latent space Z back to data X
  - Compare Input and Output with mean squared error
- Sample for Z and pass it to  $D(Z) \rightarrow$ Generate new samples
- Latent space: Series of Gaussians, regularised match  $N(\mu = 0, \sigma = 1)$ 
  - Using Gaussians lets us use Kullback– Leibler divergence
  - $\sum_{i=1}^{n} \sigma_i^2 + \mu_i^2 \log(\sigma_i) 1$



# **Training Variation AutoEncoders**

```
MSEloss = nn.MSELoss()
for ep in range(epochs):
    for i_batch in range(max_batches):
        EncoderOpt.zero grad()
        DecoderOpt.zero grad()
        # select the current batch from the dataset
        x_real = X_moons[i_batch * batch_size : (i_batch + 1) * batch_size]
        x real = torch.tensor(x real, device=device).float()
        latent = EncoderNet(x_real)
        mu = latent[:, ::2]
        log_var = latent[:, 1::2]
        KLD = torch.mean(-0.5 * torch.sum(1 + log_var - mu**2 - log_var.exp(), dim=1), dim=0)
        std = torch.exp(0.5 * log_var)
        eps = torch.randn_like(std, device=device)
        reparameterized = eps * std + mu
        x recon = DecoderNet(reparameterized)
        MSE = MSEloss(x_real, x_recon)
        loss = KLD * beta + MSE
        loss.backward()
        EncoderOpt.step()
        DecoderOpt.step()
```



## Upsides

- Directly evaluates log likelihood
- Stable in training

## Downsides

- MSE loss insufficient for certain data sets
- Needs to balance KLD and MSE loss terms

# Normalizing flows



- Variational AutoEncoder: map data to normal distribution and back using two networks
- Can we do this with a single network instead?



# Normalizing flows



- Train invertible model  $T^{*-1}$  to map data to Normal distribution
- Well understood loss function:









## Upsides

- Directly evaluates log likelihood
- Stable in training
- High generative quality
- Easy to train and use

## Downsides

- Fixed dimensionality through entire flow
- Slow generation times for large models/data

## How do we use this?



- Normalizing flow to predict high-level analysis quantities from generator-level information
- Reproduces correlations even in ML btagging algorithm scores
- Very promising solution for end-stage analyses
  - Effectively infinite MC  $\rightarrow$  minimize statistical fluctuations





# Simulation-based inference

• Remember that in my first slide I said that the purpose of an analysis was to calculate

 $p(x^{\text{data}}|H_1) \text{ and } p(x^{\text{data}}|H_0)$ 

• The methods presented here allows us to approximate these probabilities densities with much more precision than simple histograms



