## Learning parton densities with neural networks: The NNPDF methodology

IML Machine Learning Working Group, CERN

Zahari Kassabov June 16th 2017

University of Turin, University of Milan













• We want to study the Standard Model and eventually find deviations.

- We want to study the Standard Model and eventually find deviations.
- We need to compare theoretical predictions to experimental data.

- We want to study the Standard Model and eventually find deviations.
- We need to compare theoretical predictions to experimental data.
- Theory predictions at a *pp* collider:

$$\sigma_X(s, M_X^2) = \sum_{a,b} \int_{x_{min}}^1 dx_1 dx_2 f_a(x_1, M_X^2) f_b(x_2, M_X^2) \hat{\sigma}_{a,b \to X}(x_1 x_2 s, M_X^2)$$

- We want to study the Standard Model and eventually find deviations.
- We need to compare theoretical predictions to experimental data.
- Theory predictions at a *pp* collider:

$$\sigma_X(s, M_X^2) = \sum_{a,b} \int_{x_{min}}^1 dx_1 dx_2 f_a(x_1, M_X^2) f_b(x_2, M_X^2) \hat{\sigma}_{a,b \to X}(x_1 x_2 s, M_X^2)$$

•  $\hat{\sigma}_{a,b\to X}(x_1x_2s, M_X^2)$  Cross sections for *partons a*, *b* interacting at  $\hat{s} = x_1x_2M_X^2$  to produce the final state *X* at characteristic scale  $M_X$ . Can be calculated in perturbation theory.

- We want to study the Standard Model and eventually find deviations.
- We need to compare theoretical predictions to experimental data.
- Theory predictions at a *pp* collider:

$$\sigma_X(s, M_X^2) = \sum_{a,b} \int_{x_{min}}^1 dx_1 dx_2 f_a(x_1, M_X^2) f_b(x_2, M_X^2) \hat{\sigma}_{a,b \to X}(x_1 x_2 s, M_X^2)$$

- $\hat{\sigma}_{a,b\to X}(x_1x_2s, M_X^2)$  Cross sections for *partons a*, *b* interacting at  $\hat{s} = x_1x_2M_X^2$  to produce the final state *X* at characteristic scale  $M_X$ . Can be calculated in perturbation theory.
- $f_i(x, M_X^2)$  PDF of parton *i* carrying a fraction of momentum *x* at scale  $M_x$ . Needs to be *learned* from data.

#### The solution



Figure 1: NNPDF Collaboration, arxiv:1706.00428

I will describe the NNPDF methodology for determining PDFs

- As an application of Machine Learning: How it compares to other problems.
- Possible ways to improve it.

- We not only want to determine the PDFs, but also provide a sensible estimate of the uncertainty.
- Sources of uncertainty:
  - Uncertainties in input experimental data (part of the input: We have the covariance matrix).
  - Degenerate minima (+inefficiencies in the minimization).
  - Theoretical uncertainties (value of  $\alpha_{s}$ , fixed order calculations, etc)

- We not only want to determine the PDFs, but also provide a sensible estimate of the uncertainty.
- Sources of uncertainty:
  - Uncertainties in input experimental data (part of the input: We have the covariance matrix).
  - Degenerate minima (+inefficiencies in the minimization).
  - Theoretical uncertainties (value of  $\alpha_{s}$ , fixed order calculations, etc)
- Not a well researched topic in ML.

#### Experimental inputs (an oversimplification)



#### **Experimental Inputs**

$$\underbrace{\sigma_X(s, M_X^2)}_{Y} = \sum_{a, b} \int_{x_{min}}^1 dx_1 dx_2 f_a(x_1, M_X^2) f_b(x_2, M_X^2) \underbrace{\hat{\sigma}_{a, b \to X}(x_1 x_2 s, M_X^2)}_{X}$$

• Constraints come in the form of *convolutions*:

$$f(X) \rightarrow Y$$

#### **Experimental Inputs**

$$\underbrace{\sigma_X(s, M_X^2)}_{Y} = \sum_{a, b} \int_{x_{min}}^1 dx_1 dx_2 f_a(x_1, M_X^2) f_b(x_2, M_X^2) \underbrace{\hat{\sigma}_{a, b \to X}(x_1 x_2 s, M_X^2)}_{X}$$

• Constraints come in the form of *convolutions*:

$$f \otimes X \longrightarrow Y$$

#### **Experimental Inputs**

$$\underbrace{\sigma_X(s, M_X^2)}_{\mathbf{Y}} = \sum_{a, b} \int_{x_{min}}^1 dx_1 dx_2 f_a(x_1, M_X^2) f_b(x_2, M_X^2) \underbrace{\hat{\sigma}_{a, b \to X}(x_1 x_2 s, M_X^2)}_{\mathbf{X}}$$

• Constraints come in the form of *convolutions*:

 $f \otimes X \longrightarrow Y$ 

• 4285 data points points. Not a *big data* problem.

$$\underbrace{\sigma_X(s, M_X^2)}_{\mathbf{Y}} = \sum_{a, b} \int_{x_{min}}^1 dx_1 dx_2 f_a(x_1, M_X^2) f_b(x_2, M_X^2) \underbrace{\hat{\sigma}_{a, b \to X}(x_1 x_2 s, M_X^2)}_{\mathbf{X}}$$

• Constraints come in the form of *convolutions*:

$$f \otimes X \longrightarrow Y$$

- 4285 data points points. Not a *big data* problem.
- 7 physical processes from 14 experiments over ~30 years: Need to deal with inconsistencies.

$$\underbrace{\sigma_X(s, M_X^2)}_{\mathbf{Y}} = \sum_{a, b} \int_{x_{min}}^1 dx_1 dx_2 f_a(x_1, M_X^2) f_b(x_2, M_X^2) \underbrace{\hat{\sigma}_{a, b \to X}(x_1 x_2 s, M_X^2)}_{\mathbf{X}}$$

• Constraints come in the form of *convolutions*:

$$f \otimes X \longrightarrow Y$$

- 4285 data points points. Not a *big data* problem.
- 7 physical processes from 14 experiments over ~30 years: Need to deal with inconsistencies.
- Few data constrains on high and low *x*: Need to deal with extrapolation.

 $f_i(x_\beta, Q^2) = \Gamma(Q, Q_0)_{ij\beta\alpha} f_j(x_\alpha, Q_0^2)$ 

 $f_i(x_\beta, Q^2) = \Gamma(Q, Q_0)_{ij\beta\alpha} f_j(x_\alpha, Q_0^2)$ 

 $f_a(x_1, M_X^2)f_b(x_2, M_X^2)\hat{\sigma}_{a,b\to X}(x_1x_2s, M_X^2)$ 

 $f_i(x_\beta, Q^2) = \Gamma(Q, Q_0)_{ij\beta\alpha} f_j(x_\alpha, Q_0^2)$ 

 $f_j(x_{\alpha}, Q_0^2)f_k(x_{\delta}, Q_0^2)\Gamma(M_X^2, Q_0)_{aj\beta\alpha}\Gamma(M_X^2, Q_0)_{kj\beta\delta}\hat{\sigma}_{a,b\to X}(x_{\alpha}x_{\beta}s, M_X^2)$ 

 $f_i(x_\beta, Q^2) = \Gamma(Q, Q_0)_{ij\beta\alpha} f_j(x_\alpha, Q_0^2)$ 

 $f_{j}(x_{\alpha}, Q_{0})f_{k}(x_{\delta}, Q_{0}^{2})(\Gamma(M_{X}^{2}, Q_{0})_{aj\beta\alpha}\Gamma(M_{X}^{2}, Q_{0})_{kj\beta\delta}\hat{\sigma}_{a,b\rightarrow X}(x_{\alpha}x_{\beta}s, M_{X}^{2}))$ 

 $f_i(x_\beta, Q^2) = \Gamma(Q, Q_0)_{ij\beta\alpha} f_j(x_\alpha, Q_0^2)$ 

 $f_j(x_{\alpha}, Q_0)f_k(x_{\delta}, Q_0^2)(\Gamma(M_X^2, Q_0)_{aj\beta\alpha}\Gamma(M_X^2, Q_0)_{kj\beta\delta}\hat{\sigma}_{a,b\to X}(x_{\alpha}x_{\beta}s, M_X^2))$ 

• Can compute the DGLAP operator and apply to the partonic cross section. APFEL, [Bertone et al, arxiv:1310.1394].

 $f_i(x_{\beta}, Q^2) = \Gamma(Q, Q_0)_{ij\beta\alpha} f_j(x_{\alpha}, Q_0^2)$ 

 $f_j(x_{\alpha}, Q_0)f_k(x_{\delta}, Q_0^2)(\Gamma(M_X^2, Q_0)_{aj\beta\alpha}\Gamma(M_X^2, Q_0)_{kj\beta\delta}\hat{\sigma}_{a,b\to X}(x_{\alpha}x_{\beta}s, M_X^2))$ 

- Can compute the DGLAP operator and apply to the partonic cross section. APFEL, [Bertone et al, arxiv:1310.1394].
- Can store the result and perform much faster convolutions. APFELgrid [Bertone er al, arxiv:1605.02070].
- $\cdot$  Only need to deal with the x dependence at some initial scale

$$f(x,Q^2) \longrightarrow f(x,Q_0^2) := f(x)$$

#### Constraints on PDFs

- Sum rules:
  - $\sum_{i}^{\text{partons}} \int_{0}^{1} x f_{i}(x) dx = 1$
  - $\cdot \int_0^1 (u(x) \overline{u}(x)) dx = 2$
  - $\cdot \int_0^1 (d(x) \bar{d}(x)) dx = 1$
  - $\int_0^1 (q(x) \bar{q}(x)) dx = 0, \ q = s, b, t$
- Continuity:
  - $\cdot f(x) \xrightarrow{x \to 1} 0$
- ...and that's it!

To recapitulate, compared to a typical ML problem:

- We require a statistically sound uncertainty estimate.
- The problem is *regression* but the available data has a complex dependence on the PDFs.
- There are some physical constraints.

$$f(x) = Cx^{\alpha}(1-x)^{\beta}$$

$$f(x) = Cx^{\alpha}(1-x)^{\beta}$$

• Parameters can be chosen to satisfy the constraints.

$$f(x) = Cx^{\alpha}(1-x)^{\beta}$$

- Parameters can be chosen to satisfy the constraints.
- Can a simple model provide a reliable uncertainty? What is the "modelization" uncertainty? Is it possible to make any claims if the data doesn't fit?

$$f(x) = Cx^{\alpha}(1-x)^{\beta}$$

- Parameters can be chosen to satisfy the constraints.
- Can a simple model provide a reliable uncertainty? What is the "modelization" uncertainty? Is it possible to make any claims if the data doesn't fit?
- The NNPDF approach:

$$f(x) = Cx^{\alpha}(1-x)^{\beta}$$

- Parameters can be chosen to satisfy the constraints.
- Can a simple model provide a reliable uncertainty? What is the "modelization" uncertainty? Is it possible to make any claims if the data doesn't fit?
- The NNPDF approach:
  - Since we don't have constraints, we should have a very general parametrization:

 $f(x) = CNN(x)x^{\alpha}(1-x)^{\beta}$ 

NN(x)



- Fully connected.
- Two sigmoid hidden layers.
- One linear layer.
- ×8 PDF flavour combinations = 296 network parameters.

• We minimize the error function

$$\chi^2 = \sum_{ij} (D_i - O_i) \boldsymbol{\Sigma}_{i,j}^{-1} (D_j - O_j)$$

- $D_i$  experimental measurement for point i
- $O_i$  prediction for point  $i(=f \otimes \hat{\sigma})$ .
- Σ<sub>ij</sub> covariance between points *i* and *j* (corrected for normalization uncertainties [Ball et al 2009]).
- There is an additional penalty term for positivity observables.

#### Propagating experimental unccertainties

Perform N<sub>rep</sub> O(1000), fits, sampling *pseudodata replicas*:

$$\begin{aligned} D_i^{(r)} &\longrightarrow D_i^{(r)} + \operatorname{chol}(\Sigma)_{i,j} \xi_j \\ \xi_j &\sim \mathcal{N}(0,1) \\ i,j &= 1..N_{dat} \\ r &= 1..N_{rep} \end{aligned}$$

Obtain  $N_{rep}$  *PDF replicas*. All statistics of the PDFs (and functions thereof) can be computed from the ensemble of PDF replicas.

No assumptions at all about the Gaussianty of the errors. We also provide:

- Compressed Monte Carlo sets [Carrazza, et al, arxiv:1504.06469].
- Compressed [Carrazza et al, arxiv:1505.06736] and supercompressed [Carrazza et al, arxiv:1602.00005] Hessian sets.

#### Simple example



#### Simple example



#### (Discussed in detail in [Ball et al, arxiv:1410.8849])

The current approach is a genetic algorithm. At each iteration, select a node with P = 5%

$$w \to w + rac{\eta r_{\delta}}{N_{ite}^{r_{ite}}}$$

$$\eta = 15$$
  
 $r_{\delta} \sim U(-1, 1)$   
 $r_{\rm ite} \sim U(1, 0)$ 

At each iteration, generate 80 mutants, and select best mutant.

- Advantages:
  - Simple to implement and understand.
  - Good dealing with complex analytic behaviour.
  - Doesn't require evaluating the gradient.
- Disadvantages:
  - May not be close to a global minimum.
  - Requires many function evaluations (i.e. convolutions).
  - Needs tuning (discussed in later slides).

- We split the data in a training and validation set.
- Roughly 50%, different for each replica.
- We run the GA on the training set for a fixed number of iterations *O*(30000).
- We select the minimum of the validation set as the parameters from the replica.

#### Training-validation distribution



#### Preprocessing exponents

- We had  $f(x) = CNN(x)x^{\alpha}(1-x)^{\beta}$ .
- +  $\alpha$  and  $\beta$  chosen at random with ranges set from the results of a previous fit.
- We iterate the distribution of *effective exponents* doesn't change:

$$\alpha_{\text{eff}} = \left. \frac{\log f(x)}{\log x} \right|_{x \to 0}$$
$$\beta_{\text{eff}} = \left. \frac{\log f(x)}{\log(1-x)} \right|_{x \to 1}$$

- $\cdot$  We want to assess the validity of our procedure.
- We want to tune the parameters of the GA, architecture of the neural net, etc.
- Closure tests
  - Assume that the underlying PDF is known.
  - Generate data, fluctuating around the prediction of the true PDF.
  - Perform a fit and compare with assumed PDF.
  - Check that the results are consistent.

### Level 0 Fit predictions of the underlying PDF without fluctuations.

- Expect  $\chi^2/N_{\rm dat}=0.$
- $\cdot\,$  Tests the adequacy of the architecture and the GA.

Level 1 Fit fluctuations using the experiment covariance matrix.

- Expect  $\chi^2/N_{\rm dat} = 1$ .
- Test stability of central values.

Level 2 Generate pseudodata replicas on top of the replicas.

- Expect  $\chi^2/N_{dat} = 2$  for data replicas.
- Expect  $\chi^2/N_{dat} = 1$  for data central values.
- Validate the full procedure.

#### Closure test results (using MSTW2008 as Truth)



- Level 0 fit perfectly consistent: We reproduce the truth value as much as possible with the available data.
- At Level 2 (equivalent to our methodology) we reproduce exactly the fluctuated  $\chi^2 {\rm s.}$

#### Closure tests vs Truth



# If MSTW2008 was the truth, we would reproduce it within uncertainties!

- Parametrize the NNPDFs with neural networks.
- Propagate uncertainties by fitting to pseudodata replicas.
- Fit using a Genetic algorithm
- Use cross validation to avoid overfitting.
- Closure tests to validate the methodology.

### Thank you!

- In house C++ with increasing supporting code in Python.
- Core loop (convolution) written in assembly.
- Memory layout optimized.
- APFEL used for evolution.