

Recent progress in M for LHC phenom

	Convolution	Max-Pool
Jet Image		

Benjamin Nachman

Lawrence Berkeley National Laboratory

bpnachman.com bpnachman@lbl.gov







vs for an image-





Virtual Houches June 17, 2021

Data analysis in particle physics 2 Theory of everything Nature **Physics simulators** Experiment Detector-level observables Detector-level observables Pattern recognition Pattern recognition



Data analysis in particle physics 4 Theory of everything Nature **Physics simulators** Experiment Detector-level observables Detector-level observables Pattern recognition Pattern recognition

This is where most machine learning is being applied.

Representing our data



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Larkoski, Moult, and Nachman, 1709.04464, with images from Komiske, Metodiev, Thaler, 1810.05165; ATLAS, PUB-2017-003; T. Cheng, 1711.02633; Henrion et al. MLPS @ NeurIPS 2017

Data analysis in particle physics 6 Theory of everything Nature **Physics simulators** Experiment Detector-level observables Detector-level observables Pattern recognition Pattern recognition →

The growing toolkit of generative models are being developed to accelerate or augment simulations.

Deep Generative Models



Speeding up slow simulation

Generating Phase space

Estimating SM backgrounds

Measurements and Inference

BSM searches

N.B. being comprehensive with citations would fill up the slide - please see my link to the Living Review at the end for a comprehensive list

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Measurements and Inference

CaloFlow: Krause and Shih, 2106.05285

CaloGAN: Paganini, Oliveira, Nachman, 1705.02355 not quite a fair comparison, but the state-of-the-art accuracy is highly non-trivial and very impressive!

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<u>See also the LHC</u> <u>Olympics 2020</u>

BSM searches



Bortolato et al., 2103.06595

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Data analysis in particle physics 16 Theory of everything Nature **Physics simulators** Experiment Detector-level observables Detector-level observables Pattern recognition Pattern recognition -----

Simulators are a unique and powerful aspect of particle physics, but, they do not allow us to go "backwards" !!

The Inference Challenge

Want this

Measure this





(or sometimes the parameters of the model that generate this) If you know p(meas. I true), could do maximum likelihood, i.e.

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(or sometimes the parameters of the model that generate this) If you know p(meas. I true), could do maximum likelihood, i.e.

unfolded = argmax p(measured | true)

Challenge: **measured** is hyperspectral and **true** is hypervariate ... p(**meas.** | **true**) is **intractable** !

If you know p(meas. I true), could do maximum likelihood, i.e.

unfolded = argmax p(measured | true)

Challenge: **measured** is hyperspectral and **true** is hypervariate ... *p(meas.* | *true)* is *intractable* !

However: we have **simulators** that we can use to sample from *p(meas.* | *true)*

→ Simulation-based (likelihood-free) inference !

...an area of machine learning were particle physics is making key contributions!

See Cranmer, Brehmer, Louppe, 1911.01429 for a recent overview

What if we could unfold all particles simultaneously? We could then compute observables (and their bins) AFTER doing the measurement (!)

...stick around for the second part of this session for more discussions on this point

Andreassen et al., 1911.09107

Anomaly detection

Current Search Paradigm

SUSY = Supersymmetry

23

(well-motivated) theory-biased & low-dimensional observables

Current Search Paradigm

SUSY = Supersymmet

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Can we relax model assumptions and explore highdimensional feature spaces?

(well-motivated) theorybiased & low-dimensional observables

Current Search Paradigm

What if we are not looking in the right place for the new phenomena?!

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Can we relax model assumptions and explore highdimensional feature spaces?

Supervision refers to the type of label information provided to the ML during training.

Unsupervised = no labels Weakly-supervised = noisy labels Semi-supervised = partial labels Supervised = full label information

These categories are not exact and the boundaries are not rigid!

Unsupervised = no labels

Typically, the goal of these methods is to look for events with low *p(background)*

One strategy (autoencoders) is to try to compress events and then uncompress them. When x = uncompres(compress(x)), then x probably has low p(x).

Farina, Nakai, Shih, 1808.08992; Heimel, Kasieczka, Plehn, Thompson, 1808.08979; + many more including the recent LHC Olympics (Kasieczka et al., 2101.08320) and Dark Machines (Aarrestad et al., 2105.14027) reports

Weakly-supervised = noisy labels

Typically, the goal of these methods is to look for events with high *p(possibly signal-enriched)/p(possibly signal-depleted)*

e.g. Classification Without Labels (CWoLa), events in a signal region are labeled "signal" and events in a sideband are labeled "background". These labels are "noisy" but a classifier trained with them can detect the presence of a signal.

Metodiev, Nachman, Thaler, 1708.02949; Collins, Howe, Nachman, 1805.02664 + many more including the recent LHC Olympics (Kasieczka et al., 2101.08320)

Solutions: Weakly-supervised

Metodiev, Nachman, Thaler, 1708.02949; Collins, Howe, Nachman, 1805.02664 + many more including the recent LHC Olympics (Kasieczka et al., 2101.08320)

Solutions: Semi-supervised

Semi-supervised = partial labels

Typically, these methods use some signal simulations to build signal sensitivity

S. Park, D. Rankin, S.-M. Udrescu, M. Yunus, P. Harris, 2011.03550 + many more including the recent LHC Olympics (Kasieczka et al., 2101.08320)

Overview: Particle physics and ML Theory of everything Nature Parameter Fast estimation / simulation / unfolding phase space Online **Physics simulators** Experiment processing & quality control **Detector-level observables Detector-level observables** Data curation Pattern recogn Pattern recognition calibration **Classification to** clustering enhance tracking sensitivity noise mitigation particle identification

"signal" versus "background"

....

Conclusions and or

Deep learning has a great potential to **enhance**, **accelerate**, and **empower** discoveries in particle physics.

We have some unique challenges that require dedicated solutions.

With these new tools, we will be able to fully exploit the data in their natural high dimensionality enhancing the potential for **discovery**!

Conclusions and ou

My apologies to everyone who's exploit the awesome plot(s) I could not show... ality enhancing the potential for **discovery**.

Upcoming ML Workshops

ML4Jets hybrid July 6-8 2021

INSTITUTE FOR THEORETICAL PHYSICS

UNIVERSITÄT HEIDELBERG

ZUKUNFT SEIT 1386

Where can I learn more?

HEPML-LivingReview

A Living Review of Machine Learning for Particle Physics

Modern machine learning techniques, including deep learning, is rapidly being applied, adapted, and developed for high energy physics. The goal of this document is to provide a nearly comprehensive list of citations for those developing and applying these approaches to experimental, phenomenological, or theoretical analyses. As a living document, it will be updated as often as possible to incorporate the latest developments. A list of proper (unchanging) reviews can be found within. Papers are grouped into a small set of topics to be as useful as possible. Suggestions are most welcome.

download review

The purpose of this note is to collect references for modern machine learning as applied to particle physics. A minimal number of categories is chosen in order to be as useful as possible. Note that papers may be referenced in more than one category. The fact that a paper is listed in this document does not endorse or validate its content - that is for the community (and for peer-review) to decide. Furthermore, the classification here is a best attempt and may have flaws - please let us know if (a) we have missed a paper you think should be included, (b) a paper has been misclassified, or (c) a citation for a paper is not correct or if the journal information is now available. In order to be as useful as possible, this document will continue to evolve so please check back before you write your next paper. If you find this review helpful, please consider citing it using \cite{hepmllivingreview} in HEPML.bib.

https://iml-wg.github.io/HEPML-LivingReview/ https://github.com/iml-wg/HEPML-LivingReview https://arxiv.org/abs/2102.02770

Unbinned diffe cross section meas Jet Image towards a commo

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Differential Cross Section Measurements

Differential Cross Section Measurements 39 This is the bread and butter of the measurement program at the LHC! The key component of this is the correction of detector effects. This allows for the data to be compared with other experiments and with predictions. correction of detector effects = unfolding

Want this

Measure this

There were some early proposals for unbinned unfolding* but as far as I am aware, they were not used for any measurements.

However, recent innovations in machine learning and resulted in new methods for unbinned unfolding, which are being used for data analysis+ (!)

The goal of this discussion is to propose a common way for publishing unbinned results to maximize their science potential

We need input from both experimentalist and theorists (!)

*see L. Lindemann and G. Zech, NIM A 354 (1995) 516 & related +see https://www-h1.desy.de/h1/www/publications/htmlsplit/H1prelim-21-031.long.html

How do we publish binned results?

	Repository for publication-re	EPData elated High-Energy Physics data	
	Search on 9427 publicat Search for a paper, author, experiment e.g. reaction P P> LQ LQ X, title has "pl	tions and 96116 data tables. t, reaction Search Advanced hoton collisions", collaboration is LHCf or D0.	
ATLAS View Data	Data fre LICE View Data	om the LHC CMS View Data	LHCb View Data

YAML with resource files YAML YODA ROOT CSV

YAML with resource files
YAML
YODA
ROOT
CSV

ndependent_variables: header: {name: '\$\Delta R(b,b)\$'} values: - {high: 0.25, low: 0.2} - {high: 0.3, low: 0.25} - {high: 0.4, low: 0.3} - {high: 0.5, low: 0.4}

- {high: 0.6, low: 0.5}
- {high: 0.7, low: 0.6}

YAML files with metadata, bin contents, and uncertainties If the data can be fit with a function, you could publish the function (e.g. if it is a NN, you could publish the architecture and weights). 49

Another natural representation that doesn't require a function fit is to publish data sampled from the unfolded result.

My proposal is based on this idea.

As in HEPData, I propose there is a "submission" YAML file with the same measurement metadata.

Each submeasurement* also has some metadata & points to a data file. In HEPData, the data file is itself a YAML file.

The files will have data with the "shape" $[(M+1) \times N(k+1)]$

...where N is the number of sampled events and M is the number of systematic uncertainties and k is the number of dimensions per event

*this could be a single observable, or many observables

The files will have data with the "shape" $[(M+1) \times N(k+1)]$

Each event has k floats* and 1 event weight

There are N events

This is repeated for each of the M systematic uncertainties

For representations that don't have weights, the weights will be set to 1. For representations that only use weights, there will be M copies of the original array.

I have not thought deeply about file formats (npy, root, hdf5) and would be happy to hear opinions.

*For variable-length measurements, perhaps should use variable-length arrays like awkward for storage

The submission YAML should give metadata about which uncertainties are included.

For statistical uncertainties, there should be Q replicas and the uncertainty in a given bin is computed by taking the standard deviation over replicas.

For systematic uncertainties, the difference between the nominal and varied bin content is the uncertainty.

There should be warnings in metadata and/or inflated uncertainties in regions of phase space that should not be studied with the data.

Zenodo is a very natural location. Maybe the submission YAML can also be hosted on HEPData and linked to Zenodo for each searching?

Proposal - example

In [1]: import numpy as np import pandas as pd import matplotlib.pyplot as plt

In [2]: x = np.random.normal(0,1,10000)

```
In [3]: w = np.abs(x)**0.2
w_syst_up = np.abs(x)**0.3
w_syst_dn = np.abs(x)**0.1
```

```
In [4]: plt.hist(x,bins=np.linspace(-3,3,10),alpha=0.5)
plt.hist(x,bins=np.linspace(-3,3,10),weights=w,histtype="step",color="black")
n_syst_up,b=np.histogram(x,bins=np.linspace(-3,3,10),weights=w_syst_up)
n_syst_dn,_=np.histogram(x,bins=np.linspace(-3,3,10),weights=w_syst_dn)
for i in range(len(b)-1):
    plt.fill_between([b[i],b[i+1]],n_syst_dn[i],n_syst_up[i],color="black",alpha=0.3)
plt.xlabel("X")
```

```
In [5]: d = {"nominal":x,
    "nominalw":w,
    "syst_up":x,
    "syst_upw":w_syst_up,
    "syst_dn":x,
    "syst_dnw":w_syst_dn}
```

In [6]: df = pd.DataFrame(data=d)

```
In [7]: df
```

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υ	u	L	ι/	1

	nominal	nominalw	syst_up	syst_upw	syst_dn	syst_dnw
0	0.731914	0.939490	0.731914	0.910622	0.731914	0.969273
1	0.146232	0.680783	0.146232	0.561711	0.146232	0.825096
2	-0.629654	0.911634	-0.629654	0.870423	-0.629654	0.954795
3	0.581001	0.897089	0.581001	0.849675	0.581001	0.947148
4	-0.321038	0.796730	-0.321038	0.711159	-0.321038	0.892597

