STATISTICALLY LEARNING THE NEXT STANDARD MODEL FROM LHC DATA

Wolfgang Waltenberger (ÖAW and Uni Wien)

Seminar for HEPHY AI group leader position Vienna, September 2022

Who Am I

Priv. Doz. Dr. Wolfgang Waltenberger

Born december 10th 1974

Father of Fabian (2000) and Yannick (2005)

2001 - now: CMS member 2004: Finished PhD at TU Wien 2008 - now: staff member at HEPHY Wien 2014/15: Guest Professor at USP São Paulo (Brazil) since 2016: lecturing at TU Wien, Uni Wien 2018: Habilitation at Uni Wien 2021/22: Guest Professor in Grenoble (France)

Worked on: tracking, vertexing, *b*-tagging, search for SUSY, interpretation, simplified models, SModelS, ml/stats

inverse problem of particle physics

MY CURRENT DATA SCIENCE Involvements

At the Austrian Academy of Sciences

Founded ÖAW AI Graduate School, directed first installment (2019), now preparing to be again director of third iteration (january 2023)

At HEPHY

(Co-)supervising three PhD students in three different groups [see next slide] Work on learning (as opposed to postulating) the Next Standard Model from data

At TU Wien and Uni Wien

Teaching regular courses in statistics (see [youtube.com/WolfgangWaltenberger\)](https://youtube.com/WolfgangWaltenberger)

Generally – as AI/ML is still trending: currently many invited lectures, lecture series, and seminars on AI/ML/stats related topics

Eg. WMLQ2022, September 2022, Poland, Vietnam School of Physics, July 2022, Vietnam In the past: MCNet Summer School, Sweden, VDSP Winter School, Austria, DKPI Graduate School, Austria, ….

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Co-supervising three PhD students:

common denominator is data science!

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PRELUDE:

Fun With Meta-Statistics

- Our SModelS v2.2.0 database summarizes the results of almost 1000 signal regions of about 100 CMS and ATLAS publications of searches for new physics
- For each signal region, we know the number of observed events ending up in this signal region, alongside with the number of expected Standard Model "background" events and its error. Assuming a simplified statistical model, we can compute *p*-values for the Standard Model hypothesis
- **If there is no new physics is in the data, the distribution of** *p***-values should look like this:**

 $(p = 0$ means huge excess of observed events)

<https://smodels.github.io/docs/ListOfAnalyses220>

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- **Given that the background errors are conservatively, systematically overestimated by ~ 30%, we expect the following distribution for the** *p***values:**

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- For each signal region, we know the number of observed events ending up in this signal region, alongside with the number of expected Standard Model "background" events and its error. Assuming a simplified statistical model, we can compute *p*-values for the Standard Model hypothesis
- If dispersed new physics were slowly seeping in, it would look like this:

(peak moving to smaller values)

 $(p = 0$ means huge excess of observed events)

<https://smodels.github.io/docs/ListOfAnalyses220>

Here's with the actual, real data:

SModelS database v2.2.0, all topologies, all analyses

Lookin' good! No obvious *p*-hacking in our search programme.

Random fluke? Selection bias? New physics slowly seeping in?

[*] searches that target chargino/neutralino productions in RPC SUSY scenarios. Decays via W,Z,h bosons + dark matter candidate $[*!]$ some data are used more than once in this plot. We cannot – and do not pretend to -- make too serious frequentist statements

If the trend were (conjunctive form!) due to new physics, how would we go from such dispersed signals ….

… to identifying the One True Point [*] (a.k.a. the Next Standard Model) in Hitoshi Murayama's landscape of theories?

→ **our Inverse Problem!**

[*] obviously in reality we would actually want a posterior density, not one point

STATISTICALLY LEARNING THE NEXT STANDARD MODEL FROM LHC DATA

image courtesy of Jon Butterworth, Chris Wormell

Wolfgang Waltenberger (ÖAW, Uni Wien)

[presenting work in collaboration with Andre Lessa and Sabine Kraml]

<https://arxiv.org/abs/2012.12246>

Seminar, HEPHY AI group leader Vienna, September 2022

- too difficult a task for us humans (we are neither smart nor creative enough)
- Let the machine solve it!

Our mission statement:

- Given the SModelS database of simplified models results, we **let a machine find the simplest possible model** that identifies **the largest possible violation of the Standard Model hypothesis** in the results, while evading all constraints from the negative search results in our database.
- Actually, we don't just want a single model, we want posteriori probabilities in these theory landscapes.
- The models are allowed to be "**incomplete**", we want precursor theories, "proto-models", whose construction is driven by data, not by abstract principles. UV-completing such models will be a separate step (of course also partly executed by machines)

PROTOMODELS

Instead of testing BSM scenarios one-by-one against the experimental data:

- identify potential dispersed signals in the slew of published LHC analyses
- build candidate "protomodels" from them.

PROTOMODELS

- Protomodels can be thought of as **consistent sets of simplified models**.
- **Caveat:** The (variable-, or trans-dimensional) protomodels space is restricted by the SModelS software and database: currently restricted to models exhibiting a **Z2 symmetry** (i.e. SUSY- and UED-like):

X ("Xeno"-) particles, X_{q} is squark-like, X_z is neutralinolike, etc

Ongoing work: Next iteration will likely not anymore require a Z_2 symmetry [Andre Lessa, UFABC Brazil]

Building the Next Standard Model

"MCMC-type walk" over model+parameter space

After many iterations/steps, the builder "learns" the best BSM model

Building the Next Standard Model

- In each step of this random walk, the following changes to the existing protomodel are allowed:
	- randomly add or remove a particle
	- randomly change a branching ratio, or the mass of a particle
	- randomly change a production cross section of a particle
- after each step a test statistic *K* is computed that quantifies how well the protomodel describes the data.
- *K* got much worse? \rightarrow Revert to old protomodel
- *K* stayed the same or got better? \rightarrow keep new protomodel

<https://smodels.github.io/protomodels/videos> and a series of the ser

THE TEST STATISTIC

The test statistic K^c is a likelihood-ratio test that quantifies how much better the proto-model describes the data than the Standard-Model (plus a penalty for model complexity).

We search for proto-models and combinations of results / likelihoods that maximize K^c *while remaining compatible with all negative results in our database*.

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INPUT DATA

The test statistic is based on likelihoods.

- likelihood computation based on simplified models results in SModelS database
- vast number efficiency and upper limit maps from \sim 50 CMS and \sim 50 ATLAS publications.
- Assume simplified statistical models "behind" the data \rightarrow simplified likelihoods

<https://smodels.github.io/docs/ListOfAnalyses124> ₂₂

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THE COMBINER

As we are chasing dispersed signals, we need to allow the machine to combine (i.e. multiply) likelihoods. Simplified, binaric "inter-analyses *exclusivity* matrix":

green: approximately uncorrelated \rightarrow combinable

red: correlated, not combinable

White: cannot construct a likelihood

Signal regions within each analysis: correlated

In this publication: "educated guesses" from description of signatures in signal regions.

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Recent work:

TACO collaboration (members from gambit, MA5, rivet, SModelS including WW): use recasting tools and SModelS to systematically study signal region overlaps, i.e. approximate combinability of signal regions

In this publication: "educated guesses" from description of signatures in signal regions.

The Penalty Term

For every legal combination, we define a test statistic *K^c*

$$
K^{c}:=-2\ln\frac{\mathrm{L}_{\mathrm{SM}}^{c}\cdot\pi(\mathrm{SM})}{\mathrm{L}_{\mathrm{BSM}}^{c}(\hat{\mu})\cdot\pi(\mathrm{BSM})}
$$

 π (BSM) is the prior of the BSM model. We use it to "regularize" the model, i.e. impose the *law of parsimony*:

$$
\pi(\text{BSM}) \approx \exp[-n_{\text{particles}}^{\text{BSM}}]
$$

Resulting in a test statistic that resembles an "Akaike Information Criterion" (AIC):

$$
K^c \approx \Delta \chi^2 - 2 n_{\rm particles}^{\rm BSM}
$$

An additional BSM particle will have to increase the (delta-)chi-square by approximately two units.

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Resulting in a test statistic that resemble and according to the **Ongoing work:**

An additional BSM particle approximately two units.

Extend penalty term of test statistic to not only correctly penalize for model complexity, but also for number of experimental results

With: Jamie Yellen (PhD student, Glasgow)

THE CRITIC

For every legal combination *c*, we define a test statistic *K^c*

$$
K^{c} := -2 \ln \frac{L_{\text{SM}}^{c} \cdot \pi(\text{SM})}{L_{\text{BSM}}^{c}(\hat{\mu}) \cdot \pi(\text{BSM})}
$$
\n
$$
\Rightarrow K = \max \{ K^{c} \mid \forall \text{ combinations} \}
$$
\n
$$
\hat{\mu} \text{ is the signal strength of the model that maximizes the likelihood.}
$$

• By limiting its support we guarantee compatibility with all negative

results in the SModelS database.

• In allusion to adversarial setups, we also call this feature the critic

And Then We Ran the Algorithm ...

We defined a "run" as 50 parallel walkers, making 1,000 steps each. We performed 10 such runs on the SModelS database. Total computing resources spent: \sim 1,000,000 CPU hours

… and Obtained Results

We performed 10 such runs on the SModelS database:

All 10 runs introduced a top partner as well as a light quark partner. The cross sections are compatible with values expected from the MSSM. The best test statistic was *K*=6.9.

Global p -Value

- Because we have statistical models of the search results, we can synthesize statistically correct databases of results that are "typical", if no new physics is in the data.
- From this we can compute a p-value for the Standard Model hypothesis: that is the chances that – under the SM hypothesis – we would obtain a result as extreme as ours or more extreme.

Since we did not correct for the conservativeness of the experimental results, we assume our result to also be conservative.

By construction, no Look-Elsewhere Effect applies. (within the database, the machine does look everywhere)

TOWARDS UV COMPLETION

Work from protomodels to UV complete theories has recently begun (John Gargalionis, PostDoc in Valencia) – lot's of combinatorics!

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ITERATIONS

We of course envisage the procedure to be iterative:

The pheno community can point out models and combinations of interest, interpret results, make predictions about potential channels, provide feedback to the experiments

Only the experimental communities can claim potential (dispersed) discoveries.

FUTURE: FULLY DIFFERENTIABLE CHAIN OF Inductive Reasoning

If we had gradients we could perform gradient descent to find the best model, and we could use e.g. the Fisher information to infer the error on its parameters (or, alternatively we can then MCMC-sample).

Not yet required (theory space as well as space of measurements are still low-dimensional enough).

SUMMARY, OUTLOOK

- In light of no clear evidence for new physics in the individual channels/results, a more global attempt at finding new physics seems appropriate
- First prototype run of a machine that builds protomodels with results from ~ 100 analyses resulted in p-value of SM hypothesis of \sim 0.2: a very small tension with the Standard Model hypothesis (but also some tension between some results)
- Working on next iteration with more results, better likelihoods, surrogate models as accelerators, covering more signatures, larger protomodels space, UV completion
- Extend to measurements, not only searches
- Can we make the entire chain differentiable?

Run-3 will continue the trends or REFUTE THEM

Backup

A situation unlike in the Past

Not a handful of experimental signatures. Hundreds of publications with a wide range of signatures!

The Inverse Problem

So how do we get from here to here?

Top-down versus bottom-up

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Top-down versus bottom-up

THE HISCORE PROTO-MODEL

 $r_{\rm obs}$

1.30

1.28

1.24

0.92

0.90

Data driving the protomodel

CMS 0l stop (CMS-SUS-16-050)

seem different!

Likelihoods

● **Only exclusion lines** If only exclusion lines are given, without upper limits, we can do nothing ● **Observed 95% CL upper limits only:** cannot construct likelihood, binary decision "excluded" / "not-excluded" only ("critic") **Expected and observed 95% CL upper limits** can construct an approximate likelihood with truncated Gaussian, cannot combine topologies, very crude approximation **Efficiency maps** can construct a likelihood as Gaussian (for the nuisances) * Poissonian (for yields), can work per SR, and combine topologies in each SR [*] **Efficiency maps + correlation matrices** can combine signal regions via multivariate Gaussian * Poissonians **Efficiency maps + full likelihoods** full realism, correct statistical model Likelih oo $\overline{\mathbf{C}}$ <u>(</u> \bigcup o Ξ $\mathbf{\Omega}$ o $\boldsymbol{\mathsf{C}}$

[*] if efficiency maps are not supplied, we can try to produce them with recasting frameworks

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The Combiner

we allow the machine to combine likelihooods.

- Approximately uncorrelated are analyses that are:
- from different runs, and/or
- from different experiments, and/or
- looking for (clearly) different signatures

Fig. 2

A combination "c" of analyses is "legal" if the following conditions are met:

- all results are mutually uncorrelated (= "combinable")
- if a result can be added, it has to be added (any subset of a legal combination is not itself legal)
- combined likelihood: $L_c = \prod L_i$ $i \in \mathbb{C}$

THE TEST STATISTIC

For every legal combination, we define a test statistic *K*

$$
K^{c} := -2 \ln \frac{\mathcal{L}_{\mathbf{SM}}^{c} \cdot \pi(\mathbf{SM})}{\mathcal{L}_{\mathbf{BSM}}^{c}(\hat{\mu}) \cdot \pi(\mathbf{BSM})}
$$
 Eq. 6

(Remember, we have a database of results from \sim 100 CMS+ATLAS searches. We want to find the most interesting combinations of these results, i.e. the ones that maximally violate the SM hypothesis)

Of all "legal" combinations of experimental results, the builder chooses the one combination "c" that maximizes *K*:

$$
K := \max_{\forall c \in C} K^c \qquad \qquad \text{Eq. 7}
$$

μ denotes an global signal strength multiplier – the production cross sections are free parameters

$$
\forall i, j : \sigma (pp \to X_i X_j) = \mu \bar{\sigma} (pp \to X_i X_j)
$$

 It is maximized in the denominator, but its support is confined such that no limits in the SModelS database are violated (the "critic"),

$$
\hat{\iota} \in [0, \mu_{\max}]
$$

The Walker

The Walker takes care of moving in the protomodel space with varying dimensionality by performing the following types of modifications to the protomodel:

- add or remove particles from the protomodel
- change the masses of particles
- change the signal strengths of production modes
- change decay channels and branching ratios

At each step the test statistic *K* is computed. An MCMC-like procedure[*] is then applied in the sense that the step is reverted with a probability of

$$
\exp\left[\frac{1}{2}(K_i - K_{i-1})\right]
$$

if and only if *Ki* is smaller than *Ki-1*

* (note however, instead of ratios of unnormalized posteriors we have ratios of ratios of unnormalized posteriors)

WALKING OVER FAKE STANDARD MODEL **DATABASES**

- Produced 50 "fake" SModelS databases by sampling background models
- Corresponds to typical LHC results if no new physics is in data
- Determine 50 "fake" *K* values by running 50 walkers on each of the 50 databases (50 x 50 walkers in total) \rightarrow density of *K* under null SM-only hypothesis

The Walks

We define a "run" as 50 parallel walks, each taking 1000 steps.

We performed

- 10 runs on the SModelS database (Sec. 5.2)
- 50 runs on fake "Standard Model-like" databases (Sec 5.1) to be able to determine a global *p*-value under the SM hypothesis
- 2x10 runs on fake "Signal-like" databases (Sec 5.3) to show closure of the method

WALKING OVER DATABASES WITH FAKE **SIGNALS**

To show closure of our method, we inject the winning protomodel as a signal in fake databases, and see if the algorithm can reconstruct the injected signal.

Sec 5.3

Technical closure test **Physics closure test**

Sampling turned on

No sampling of the models for the SRs, i.e. observed events := expected SM + expected signal events

FUTURE IMPROVEMENTS

Improvements of the SModelS database:

- add latest full run-2 CMS and ATLAS publications (Moriond!)
- produce efficiency maps for existing results
- enlarge mass range of older efficiency maps

Improvements in speed:

- learn the SModelS database
- make everything differentiable

Improvements in procedure:

- improve the "analyses correlation matrix", automate the determination
- ponder relationship between proto-models and effective field theories
- connect proto-models with complete theories

Reinforcement Learning, Cryogenic **DETECTORS**

The situation with cryogenic detectors

by Felix Wagner

Reinforcement Learning, Cryogenic **DETECTORS**

A framework for policy optimization: *reinforcement learning*

by Felix Wagner

Reinforcement Learning, Cryogenic **DETECTORS**

Actor critic

Policy Actor **TD** Critic error Value state action Function reward Environment

Soft Actor-Critic: Off-Policy Maximum Entropy Deep Reinforcement Learning with a Stochastic **Actor**

Tuomas Haarnoja, Aurick Zhou, Pieter Abbeel, Sergey Levine

Model-free deep reinforcement learning (RL) algorithms have been demonstrated on a range of challenging decision making and control tasks. However, these methods typically suffer from two major challenges: very high sample complexity and brittle convergence properties, which necessitate meticulous hyperparameter tuning. Both of these challenges severely limit the applicability of such methods to complex, real-world domains. In this paper, we propose soft actor-critic, an offpolicy actor-critic deep RL algorithm based on the maximum entropy reinforcement learning framework. In this framework, the actor aims to maximize expected reward while also maximizing entropy. That is, to succeed at the task while acting as randomly as possible. Prior deep RL methods based on this framework have been formulated as Q-learning methods. By combining off-policy updates with a stable stochastic actor-critic formulation, our method achieves state-of-the-art performance on a range of continuous control benchmark tasks, outperforming prior on-policy and off-policy methods. Furthermore, we demonstrate that, in contrast to other off-policy algorithms, our approach is very stable, achieving very similar performance across different random seeds.

$$
\pi^* = \arg \max_{\pi} \mathbb{E}_{\pi} \left[\sum_{t=1}^T \frac{R(s_t, a_t) + \alpha \operatorname{H}(\pi(\cdot \mid s_t))}{\text{reward}} \right]
$$

 $\delta_t = r_{t+1} + \gamma V(s_{t+1}) - V(s_t),$

55

by Felix Wagner

GRAPH NEURAL NETWORKS

- ❖ using pytorch geometric
- ❖ consists of ConvBlocks, a linear model and a pooling layer
- ❖ GCNConv
	- \triangleright updates nodes according to neighbors via adjacency matrix
	- \blacktriangleright symmetric normalization

Input: (5, n), Output: (5,n)

Input: (5, n), Output: (1,n) Input: (1, n), Output: (1,1)

 $X' = \hat{D}^{-1/2} \hat{A} \hat{D}^{-1/2} X \Theta$

[https://arxiv.org/abs/1609.0](https://arxiv.org/abs/1609.02907) [2907](https://arxiv.org/abs/1609.02907)

by Mark Matthewman

GRAPH NEURAL NETWORKS

Hackathon - Convolution Operations

by Mark Matthewman

GRAPH NEURAL NETWORKS

Hackathon - Results

by Mark Matthewman