Learning (from) High-dimensional Models with PhenoAI and iDarkSurvey

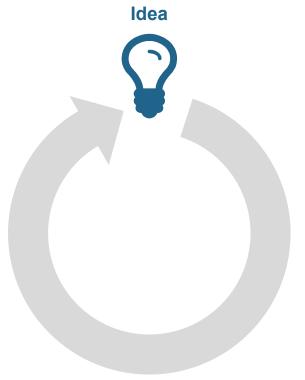
Jisk Attema Tom Heskes Roberto Ruiz de Austri Sascha Caron Sydney Otten Jong Soo Kim Faruk Diblen Krzysztof Rolbiecki <u>Bob Stienen</u>

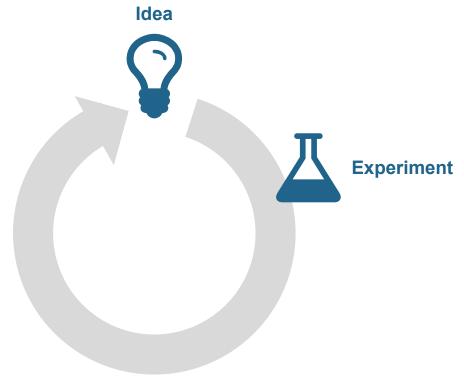
netherlands

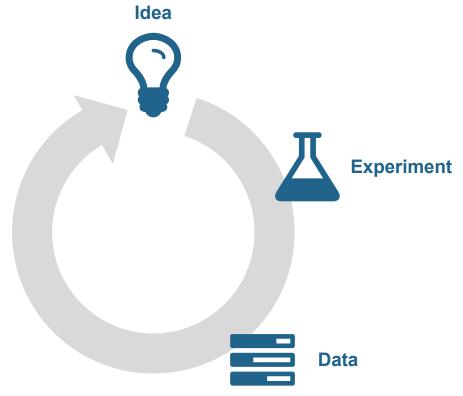


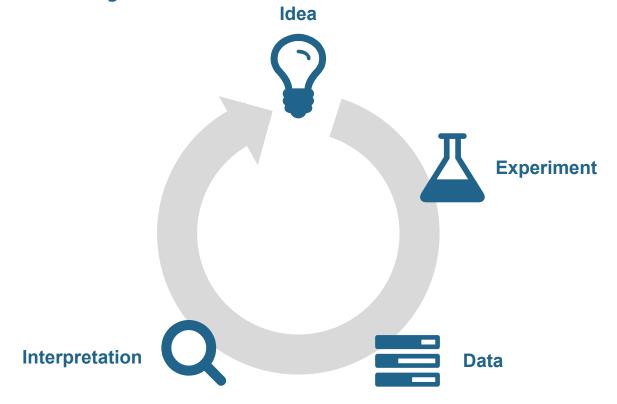


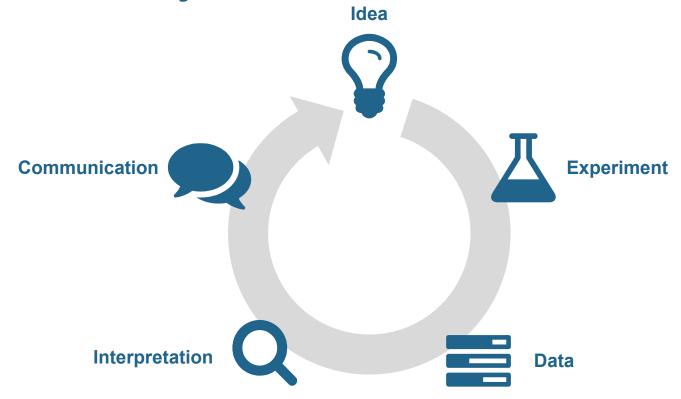
How do we do (particle) physics?

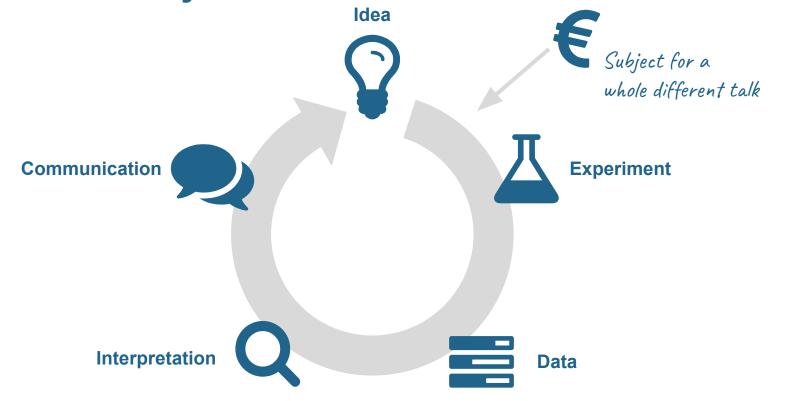


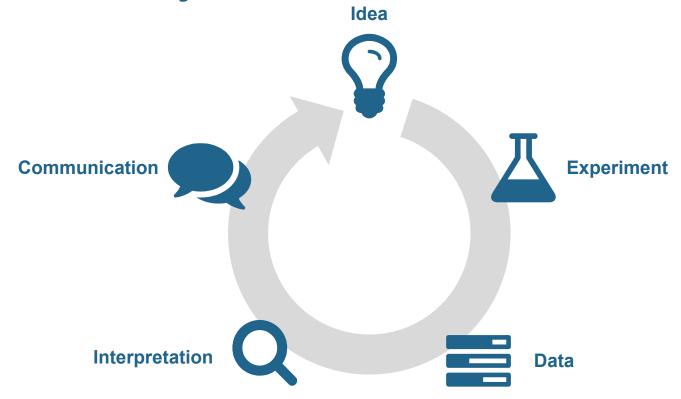




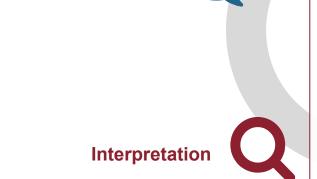








Communication



Idea

- Inherently model dependent
 - → different model = different interpretation

Experiment

- Interpretation of the results in the context of a single model point is computationally very expensive
 - → Simplified models are often used, but



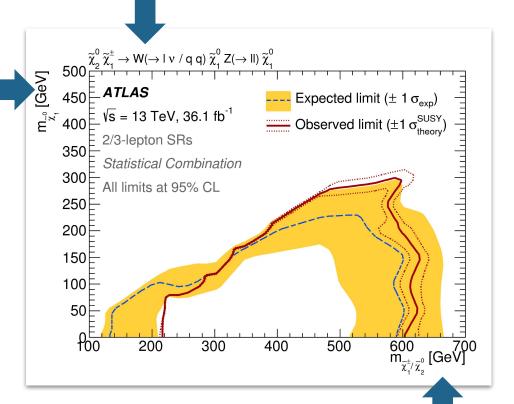


Interpretation

Idea

- Images in papers are inherently 2-dimensional
 - → displaying more than 4 dimensions in a plot is difficult

- Simplified models are often used, but at the cost of information loss
- Raw data can be published (e.g. model points + evaluations)
 - → Individual results are not extremely useful



What if...

- i don't have a 100% BR to the specified final state?
- i want to know the exclusion in another projection?
- i have the other free parameters set differently?

Core of the problem: Plotting N>2 dimensions is hard

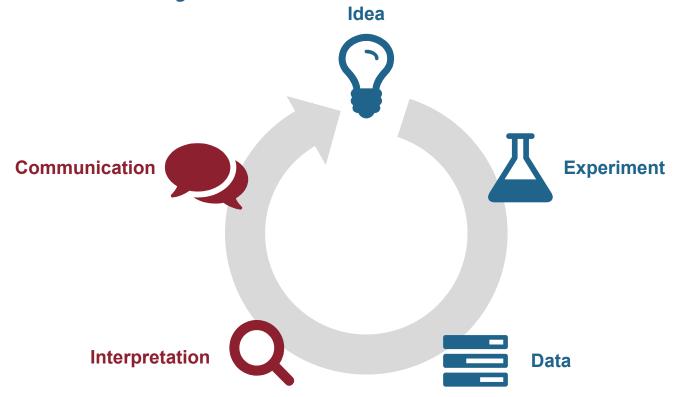


Interpretation

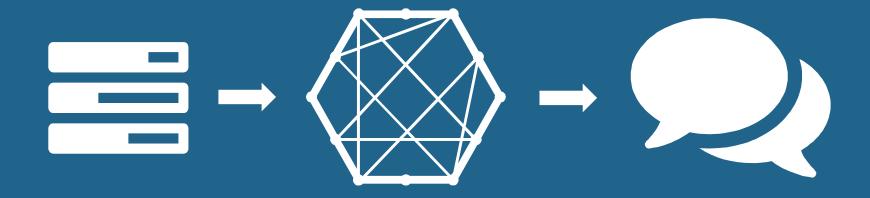
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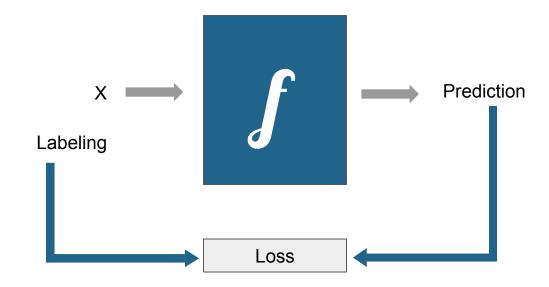
How to manage our information to retain most of it?



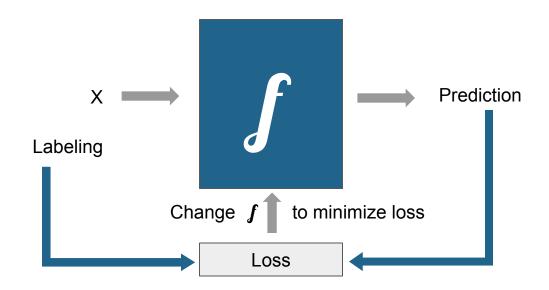




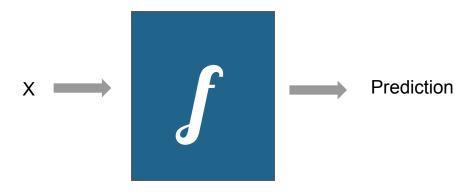








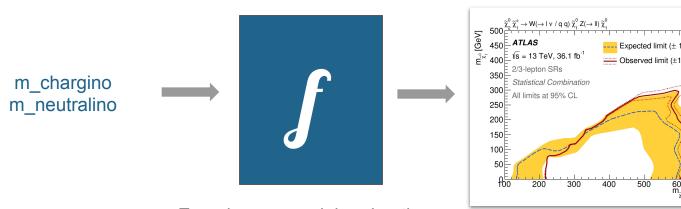




Encodes our model and entire analysis workflow

Example





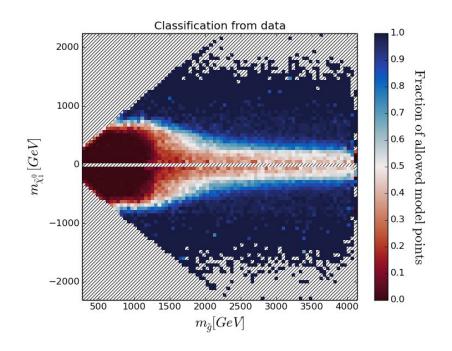
Encodes our model and entire analysis workflow

But... can be N>2...

SUSY-AI as proof-of-principle

DOI: 10.1140/epjc/s10052-017-4814-9





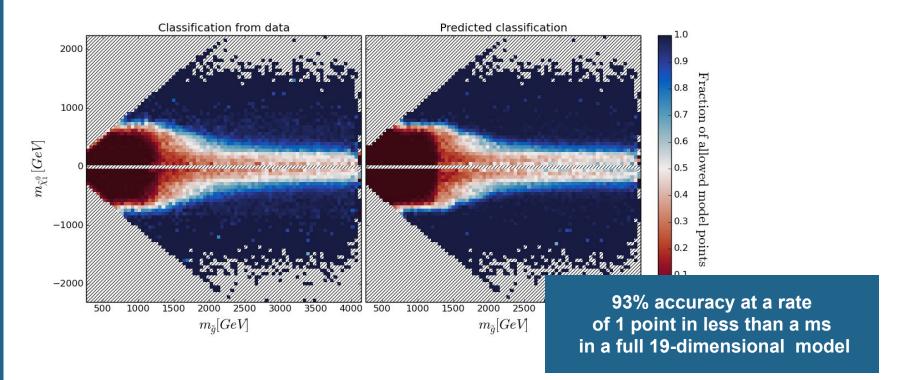
- pMSSM19
- 300,000 training points 10.1007/JHEP10(2015)134

- Exclusion determined by 22 different analyses
- RandomForest (for the connaisseurs)

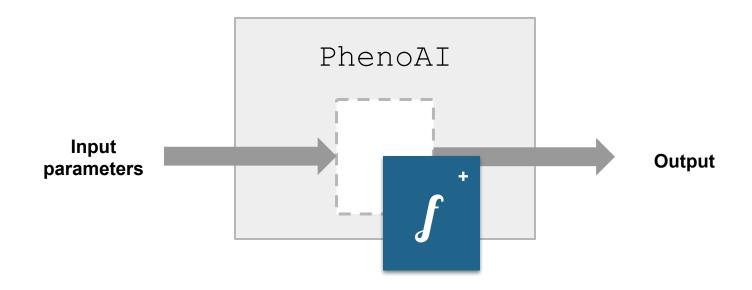
SUSY-AI as proof-of-principle

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PhenoAI as natural evolution

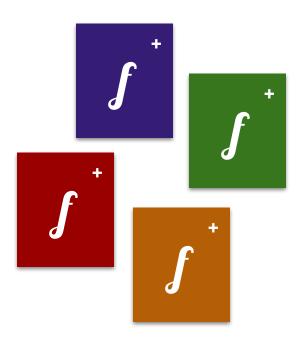


Machine Learning is abstracted away: anyone with Python knowledge can use the trained models

Communication of high-dimensional results becomes possible:

publish a trained algorithm

PhenoAlnalyses

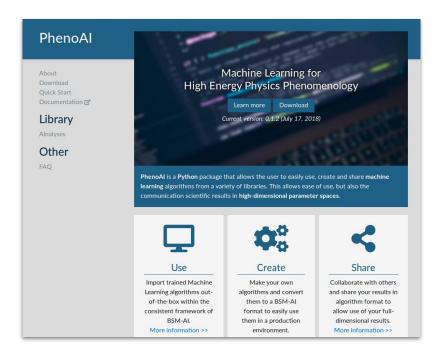


- Trained algorithms (**Alnalyses**) still need to be made. You can do this yourself, or ...
- ... download one from the Alnalysis library on the PhenoAl website
- Currently working on Alnalyses for:

Cross Sections << *See S. Otten's talk tomorrow*Electroweakino
Likelihoods from Gambit

Pheno for the masses"

- PhenoAl is available via pip3
 (phenoai) and via the website
 http://hef.ru.nl/~bstienen/phenoai
- Extensive documentation available
- Feedback is more than welcome!
- Paper will be out soon



But what about data?

Data publishing

- Individual data points (e.g. model points) are not really informative on their own
- Data can be published on HEPData, but...
 - ... lacks an easy interface to navigate and explore the data
 - ... data sets can not be easily compared

Result: Publishing information like model point evaluations is still not extremely common in our field.

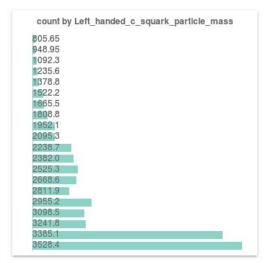
iDarkSurvey for Data Publishing

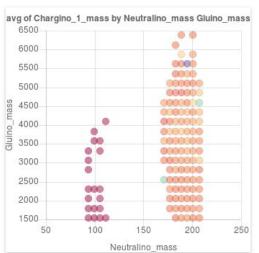


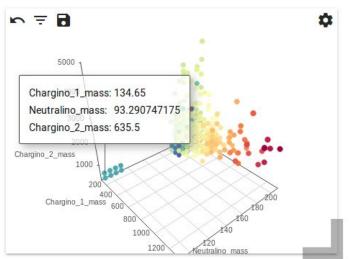
- Online data storage for high energy physics data
- Has online plotting interface to explore data
- Multiple data sets can easily be compared within the same plots
- Own data can be viewed alongside the data in the database
- Online demo at http://www.idarksurvey.org/

iDarkSurvey for Data Publishing



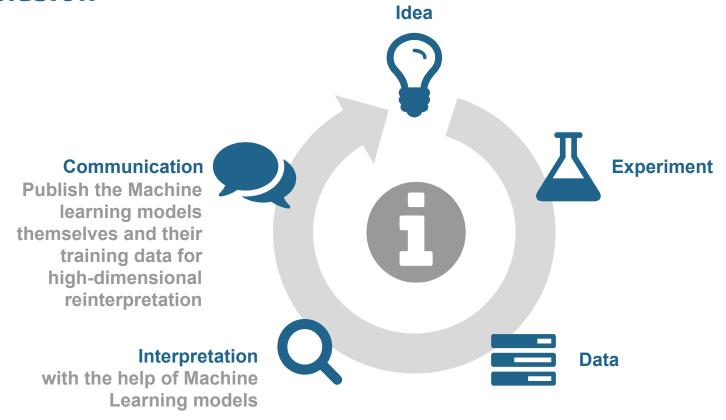






http://www.idarksurvey.org/

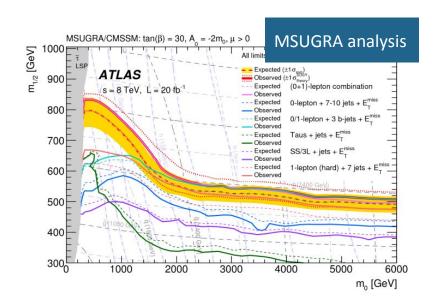
Conclusion

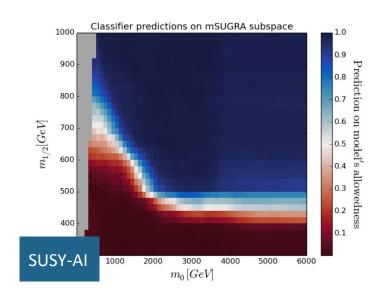


Extra slides

What about my simplified model?

Training on a full model still allows access to submodels. SUSY-AI was trained on the pMSSM19, of which MSUGRA/CMSSM is a submodel.

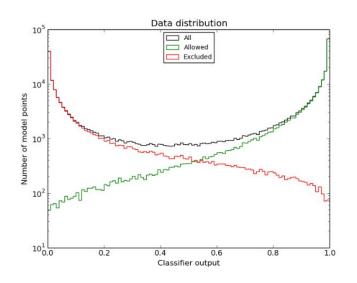


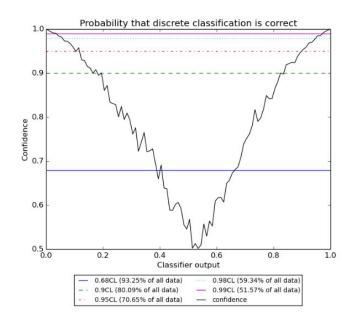


Comparison not entirely fair: the dedicated MSUGRA/CMSSM scan combined signal regions in a smart way, whereas the exclusion of the SUSY-AI dataset uses the simple: "if excluded by any analysis -> excluded"

Confidence construction from SUSY-AI

SUSY-AI is a classifier, but outputs a continuous value between 0 (excluded) and 1 (allowed). It can *not* be interpreted as a probability, but can be transformed into one.





Is PhenoAI really that simple?

```
1 from phenoai.phenoai import PhenoAI
2
3 master = PhenoAI()
4 master.add("./example_ainalysis", "example")
5 result = master.run(X)
```

Learning to use PhenoAl

PhenoAl aims to be as easy to use as possible. To this end we have created:

- online documentation
- in-code documentation
- example scripts
- a quick start manual

We are busy optimizing the learning experience of PhenoAI even further, making material as a tutorial and a cheat sheet.

Supported ML libraries

All estimators and models created with Keras/tensorflow and scikit-learn are supported within PhenoAl. We are in the process of adding support for ROOT TMVA models as well.



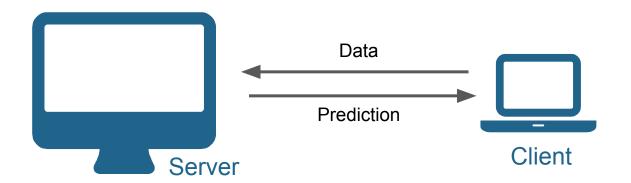




Server-client structure

PhenoAI has a built-in ability to create a server-client structure. The server has the Alnalyses loaded, the client can be added to any script and will query the server for prediction on a specific data set. In this way, the loading and configuration overhead are needed only once.

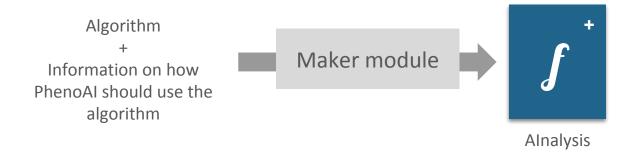
Server and client can of course just be the same machine



Maker module

In order to use a trained algorithm within PhenoAI, it needs to be stored within a folder with a PhenoAI configuration file. This collective as files is called an AInalysis and can, in principle, be made by hand. It is however more convenient to use the phenoai.maker module. Which will indicate if errors are made.

Example scripts on how to use the maker module are availble.



DarkMachines

PhenoAI is connected to the DarkMachines initiative as well, a research collective aiming to unravel the mystery that is dark matter with the help of machine learning. See <u>darkmachines.org</u> for more information.

