# **Simple Analysis for LLPs** -Truth-level Analysis Framework

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**LLP13** 

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## **A timeline on Reinterpretation** – Development of Tools and the Progression of Collaboration Standard

- 2000: 1st PHYSTAT
- 2010: introduction of Workspaces
- 2010: First proposition of Recast
- 2012: ATLAS profile likelihoods scans released for Higgs coupling
- 2017: CMS Simplified Likelihoods
- 2018: Release of analysis preservation and systematic reinterpretation with Recast within ATLAS



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## **A timeline on Reinterpretation** – Development of Tools and the Progression of Collaboration Standard



\*ATLAS development highlighted in blue

- 2019: <u>First full likelihood release</u> in ATLAS for SUSY searches
  - Using json schema of Histfactory class of likelihood
- 2019: ATLAS publication of full statistical likelihoods PUB note released (JSON HF)
- 2020: First full likelihood release regularly in ATLAS for SUSY searches
  - 2022: "<u>Simple Analysis: Generator-level Analysis</u> <u>Framework</u>" pub note released — Framework going public!
- 2022: Snowmass: "Reinterpretation and Long-Term Preservation of Data and Code"



# **Different Reinterpretation Tools**

### Less Precision



### Level of Complexity

### **More information/Precision**

### Experiment specific

Simple Likelihood Pubnote

**HistFactory Json:** Storing data products from analyses for reinterpretation

About - Connect - Activities - Fellows Jobs



pyhf is a pure-python implementation of the widely-used HistFactory p.d.f. template described in [CERN-OPEN-2012-016]. It also includes interval estimation is based on the asymptotic formulas of "Asymptotic formulae for likelihood-based tests of new physics" [arxiv:1007.1727]. The aim is also to support modern computational graph libraries such as PyTorch and TensorFlow in order to make use of features such as autodifferentiation and GPU acceleration.



preservation of the original analysis pipeline for collaboration-approvable results





# **Simple Analysis** -Experiment-independent middle

- Preserve full analysis pipeline for manipulation/maximal information
- Open source
  - Experiment non-specific, not tie to detector reconstruction
  - Approximation can be added
- Allow for reinterpretation up to the truth level
  - Detector effect can be added
- Non-experiment input format
  - HEPMC, DAOD, ROOT-ntuple, Delphes etc.
- For Public use: can be used outside of ATLAS
- Additional information for reinterpretation

### 53 SimpleAnalysis, v1.1.0

### **Q** Search

ANA-SUSY-2018-06

ANA-SUSY-2016-09

ANA-SUSY-2016-24

ANA-SUSY-2018-04

ANA-SUSY-2016-11

ANA-SUSY-2016-06

ANA-SUSY-2016-20

ANA-SUSY-2017-03



### SimpleAnalysis: <u>Webpage</u>



### Source Code

SimpleAnalysis PubNote

Available list of Analyses

EwkThreeLeptonERJR2018

EwkTwoLeptonTwoJet2016

OneLeptonMultiJets2016

DisappearingTrack2016

EwkTwoLeptonRJ2016

PairedDijets2016

DirectStau2018

StopZ2016



# Simple Analysis Framework



- 1 h7;1 - 1 h8;1 - 1 h9;1 - 1 h10;1

h11;1

h14;1

Filter: All Files (\*.\*)

Command

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Command (local):

### Simple Analysis Routine

Complex kinematic variables Custom efficiency maps Additional utilities for serializing NN and multi-variate analysis /BDT

HEPData

Q Browse all 🛛 🗐

QSearch HEPData

### HEP data publishing

- Truth level acceptance and selection efficiencies for HEPData publishing
- Root flat-ntuple files
  containing
  histograms
- Smearing available to approximate detector effect

Kide Publication Information	🕹 Download All 🗸	Systematic tab	le for SR_Gtt_0L_B	10.17182/hepdata.95928.v1/t1	JSON
Search for supersymmetry in final states with missing transverse momentum and three or more <i>b</i> -jets in 139	🗠 View Analyses 🗸	Res	Resources ht	https://www.hepdata.net/rec	
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$fb^{-1}$ of proton—proton	process.	A summary of the uncertainties in the background estimates for SR-Gtt- 0L-B. The individual experimental and theoretical uncertainties are			
collisions at $\sqrt{s}=13$ TeV with the ATLAS detector	acceptance SR- > Gtt-0L-M2	assumed to be uncorrelated and are combined by adding in quadrature.			
	Data from auxiliary	cmenergies	observables	phrases	reactions
The ATLAS collaboration	material Figure 10a. 10.17182/hepdata.95928.v1/t8	♥ 13000	relative	SUSY	🗣 P P> GLU GLU X
	Acceptance for SR-Gtt-0L-			Supersymmetry	

🚯 About 🛛 🗘

Last updated on 2022-11-25 13

Search



# **Simple Analysis —Tutorial**

- Tutorial contains:
  - Building analysis selection

Analysis pipeline Preservation

 Reinterpreting results from previous SA pipelines.

**Reinterpretation Effort** 

Welcome feedback!



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Analysis implementation Debugging Running

SimpleAnalysis repository

### GitLab 4 Stars · 4 Forks SimpleAnalysis, v1.1.0 <≁ Q Search Analysis Skeleton Table of contents Creating an empty skeleton General structure Creating an empty skeleton Recompilation Let's go ahead and create an analysis from scratch. Touch a new file in × SimpleAnalysis/Root/MyAnalysis.cxx 1 cd \$TUTORIAL\_DIR/SimpleAnalysis/SimpleAnalysisCodes/src 2 touch MyAnalysis.cxx General structure In general, the structure of an analysis runner is very similar to any other event looping algorithm that you encounter in ATLAS. There is an AnalysisClass::Init() and an AnalysisClass::ProcessEvent(AnalysisEvent \*event) that you need to implement. The skeleton of an analysis always looks like this: #include "SimpleAnalysisFramework/AnalysisClass.h" DefineAnalysis(MyAnalysisName) void MyAnalysisName::Init() {} void MyAnalysisName::ProcessEvent(AnalysisEvent \*event) { return; }

The Init() method handles initialisation of any regions or histograms that you would like to define in your analysis. The ProcessEvent() method is, as the name would suggest, called for



## **Simple Analysis** -Current analysis list

- Most SUSY analyses on ATLAS publishe analysis selection at the level of generate
- Growing list can be found <u>here</u>.

### Analysis List





Source Code

SimpleAnalysis PubNote

### Available list of Analyses

ANA-SUSY-2016-11

ANA-SUSY-2016-06

ANA-SUSY-2016-20

ANA-SUSY-2017-03

ANA-SUSY-2017-03

OneLeptonMultiJets2016

DisappearingTrack2016

EwkTwoLeptonRJ2016

StopZ2016

EwkTwoLeptonRJ2016



# **Simple Analysis** -Pub note

Entries / 40 Ge

- Pub note described the tool and analysis pipelines for preservation
- Carlo samples at reconstruction level
- One lepton+ two b-jet analysis is used to demonstrate the validation



# Comparing the generator level analysis by SimpleAnalysis with the Monte

(b) Leading lepton  $p_{\rm T}$ 

### (f) Missing transverse momentum

# LLP analysis in Simple Analysis

- Currently few LLP analyses are available in SA
  - Disappearing track
  - ... More to come!
  - R22 is work in progress that could use more person input





Image from Toshiaki Kaji

## **Future of Simple Analysis** -A full reusing pipeline(Mapyde)

Fresh off the arrive today at 9am

Giordon Stark<sup>\*,1</sup>, Camila Aristimuno Ots<sup>1,2</sup>, and Mike Hance<sup>†,1</sup>

<sup>1</sup>University of California, Santa Cruz, Santa Cruz Institute for Particle Physics, 1156 High Street, Santa Cruz, CA 95064 <sup>2</sup>University of Southern California, Liquid Propulsion Laboratory, 854 Downey Way, Los Angeles, CA 90089

### REDUCE, REUSE, REINTERPRET: AN END-TO-END PIPELINE FOR **RECYCLING PARTICLE PHYSICS RESULTS**

A PREPRINT

# Mapyde

Mayede: Performing a full pipeline of truth level analysis from event generation to statistical analysis!

- MADGRAPH5\_AMC@NLO(MADGRAPH) [26, 27] (event generation)
- PYTHIA8 [28] (parton shower, hadronization, decays)
- DELPHES [29–31] (detector simulation)
- SimpleAnalysis [32, 33] (analysis description)
- pyhf [34, 35] (probability model fitting)

### Containing

# **Reinterpretation Results from Mapyde**



Figure 6: Expected (left) and observed (right) constraints on the slepton-wino-bino model. The model is parameterized by the slepton mass, the slepton- $\tilde{\chi}_1^0$  mass splitting, and the  $\tilde{\chi}_2^0$ - $\tilde{\chi}_1^0$  splitting, and compared against the slepton-bino results from Ref. [40].

# Simple Analysis pub note summary





# Summary

- Simple Analysis is fully **public**, available for theorists/experimentalists alike.
- Generator level pipeline preservation allow for analysis detail to be preserved.
- Reinterpretation made more straight forward.
- Reconstruction/detector effect can be added later on.
- Validation has been made on the generator/reco level kinematics plot.
- Mapyde, a new pipe that that would allow for experimental independent full reinterpretation is available today!





# Back up

## What is re-interpretation? —And its importance in the BSM context

- Reinterpretation: The reusing of experimental results for alternative physics hypothesis  $\bullet$
- **Advantages:** 1. Reuse of data/workflow/optimization 2. Less labor intensive 3. Time saving ullet



- Two types of Reinterpretation
- > 1. Re-interpretation (Single data-> Many theory)

 > 2. Combination (Many data -> Single theory) Abundance of theoretical ideas vs experimental results available "Combination of results (E.G. Higgs combination)



## What is re-interpretation? -And its importance in the BSM context

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# **Available reinterpretation tools**

### Within Experiment



Experiments preserve analyses in full fidelity internally. Re-interpret on-demand for models that are deemed sufficiently promising.

### **Outside of Experiment**

### Unfolded Reinterpretation(Without detector effect) Tools:



Other Homegrown **HepMC** based Analysis



### Folded Reinterpretation(With Detector Effect) Tools:



**Homegrown Toy Detector Approximation** 

**Delphes based Analysis** 



# **Tools used by ATLAS for reinterpretation** Introducing the players

Precision Level	<b>Preserved Item</b>	Tools/S
0	Data product	HEP
1	Likelihood	Simp
1	LINCIII IOOOG	JSON
2	Truth Level Pipeline	Simp
10	Full Reco Pipeline	Reca
10+	Preservation of ML Model/optimization	ONN
	Increase precision er	hable b

Service used by ATLAS

Data

ole likelihood

N HistFactory

leAnalysis/Rivet

++Increased level of Information/ Precision

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etter reinterpretation from original analysis!



## Level 0: HEPData Storing of experimental data product from analyses for reinterpretation

- Previously: Table-based webpage (40 yearold):
- New digital library : Paper
  - Database: Include Efficiency Map, yields, uncertainties, outflow tables
  - Reinterpretation tools: Rivet and SimpleAnalysis code, preservation of ML
  - Webservice
- Made an ATLAS requirement for publication
- Centralized tool across other HEP experiments



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	JSON

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# Level 1: Statistical Model (Simple Likelihood) — Saving the most of our experiment results in one variable

- Previously, often HEPData information is used to construct approximate likelihood
- Likelihood: Great bang for the buck: encapsulate information about Limits, data/MC plots, yield take, and systematics. ->Best object that can be provided!
- ATLAS started providing them on HEPData since 2019 (SUSY)



# Existing Exp. Likelihood (What We Provide)



### See <u>"Publishing statistical models</u> Level 1: HistFactory JSON discussion" on Tuesday" —pyhf as a pure-Python implementation of HistFactory with JSon

- <u>PyHF</u>: HistFactory in full python
- Increasingly used by ATLAS analyses (24) analyses published-up-to-date)
- Results in easy to read json format: Database
- Reinterpretation:
  - 1. Publishes statistical model (Likelihood) in json file format
  - 2. New version of <u>SModel</u> and MadAnalysis5 allow for automatic reinterpretation from pyHF json file





Source Code





## Level 2: SimpleAnalysis -Truth level pipeline preservation

- Preserves the analysis code pipeline
- Allow for reinterpretation up to the truth level  $\bullet$ 
  - Detector effect can be added: public  $\bullet$ simulation like Delphes or internally by ATLAS fast simulation
- Experimentalist friendly input format (HepMC, DAOD, ROOT n-tuple, DELPHES)
- Large list of <u>ATLAS SUSY analyses</u> available:
- For **public use**: can be used external to ATLAS

### See <u>tutorial by Giordon Stark</u> Wednesday

### 53 SimpleAnalysis, v1.1.0

### **Q** Search

### SimpleAnalysis, v1.1.0 Simplified ATLAS SUSY analysis framework Home Analysis List **ATLAS Internal** Holds a collections of SI Analysis List formats: N-Tuple Structure Analysis Name ATLAS Public Result page Tutorial N-Tuple Structure DAOD\_TRUTH (TRU) ZeroLeptonRJigsaw2016 ANA-SUSY-2016-07 xAOD (either truth-le SbottomMultiB2018 ANA-SUSY-2018-31 HepMC (uncompres DMbb2016 ANA-SUSY-2016-18 • slimmed ntuples (Re DELPHES (Converte) ANA-SUSY-2016-18 DMttZeroLepton2016 MultiJets2018 ANA-SUSY-2018-17 It provides the analysis a histograms or ntuples w EwkThreeLeptonResonance2018 ANA-SUSY-2018-36

### SimpleAnalysis: <u>Webpage</u>



Source Code

SimpleAnalysis PubNote

EwkThreeLeptonERJR2018 ANA-SUSY-2018-06 ANA-SUSY-2016-09 PairedDijets2016 EwkTwoLeptonTwoJet2016 ANA-SUSY-2016-24 DirectStau2018 ANA-SUSY-2018-04 OneLeptonMultiJets2016 ANA-SUSY-2016-11 DisappearingTrack2016 ANA-SUSY-2016-06 ANA-SUSY-2016-20 StopZ2016 ANA-SUSY-2017-03 EwkTwoLeptonRJ2016

### Available list of Analyses







# Level 10: Recast -Full Analysis pipeline preservation

- Preservation of original analysis pipeline for re-interpretaion
- Contain full detector simulation information + original code/ workflow
- Reinterpretation with full workflow + simulation effect
- Made a requirement for publication in Exotics, HDBS and SUSY
- Implementation Internal to ATLAS: some reinterpretation results published with the tool
- <u>Software</u> fully open source!  $\bullet$
- Future: Allow for an interface to submit reinterpretation request online



gen, sim & reco



### preservation of the original analysis pipeline for collaboration-approvable results







## Level 10+: ONNX/NN See talk by Dan Guest later today — Beyond usual pipeline preservation: Reuse of ML training results

- ML development significantly improve analyses sensitivity. Theorists would like to make use of the BDTs/NN.
- ONNX(Open Neural Network Exchange)
- BDT preservation -> convert to standalone c++ in SimpleAnalysis
- Non-BDT NN -> Preservation by Serializing through Onnx
- Can be preserved with SimpleAnalysis to be reused
- 2 analyses already have ML encapsulated in the workflow



3 b-jets+

Search for supersymmetry in final states with missing transverse momentum and three or more *b*-jets in 139 fb<sup>-1</sup> of proton–proton collisions at  $\sqrt{s} = 13$  TeV with the ATLAS detector

The ATLAS Collaboration

### **One Lepton MultiJets**

Search for R-parity-violating supersymmetry in a final state containing leptons and many jets with the ATLAS experiment using  $\sqrt{s} = 13$  TeV proton-proton collision data

The ATLAS Collaboration

ANA-SUSY-2018-30

SA with Onnx

ANA-SUSY-2019-04



Source Code

Two existing analyses already contain Onnx



