

Machine Learning application for Λ hyperon reconstruction in CBM at FAIR

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CBM physics goals and experimental challenges



Main CBM physics cases:

- QCD matter equation-of-state at large baryon densities
- The production of strange quarks is sensitive to the properties of created matter in high energy nuclear <u>collisions</u>
 - (<u>Multi)-Strange</u> particles
- Extend nuclei chart with <u>hypernuclei</u> measurements



- Tracking: Micro-Vertex Detector (MVD), Silicon Tracking System (STS)
- Particle identification: Muon Chamber (MuCh), Ring Imaging Cherenkov (RICH), Transition Radiation Detector (TRD), Time of Flight (TOF)
- Collision geometry: Projectile Spectator Detector (PSD)



(Multi)-Strange reconstruction via weak decays

- A hyperons are the most abundant strange baryons produced at FAIR energies
- Collisions generated by URQMD and DCM-QGSM-SMM with Au+Au collisions at p_{beam} = 12A GeV/c (4s_{NN} = 4.93), mbias, 100k
- Using GEANT4 simulation, CA tracking, KFParticle within CbmRoot framework

 $\Lambda^0 \rightarrow p^+ \pi^-$

 Λ candidates reconstruction:

- Combine all proton and pion tracks
- Signal from a lambda decay
- Combinatorial background

Variables :

- χ²_{prim} squared distance between the daughter track and the primary vertex divided by its Covariance Matrix (CV)
- DCA distance of closest approach between proton & pion tracks
- $\chi^2_{\alpha eo}$ squared distance between daughter tracks divided by CV
- L/ΔL distance between primary and secondary vertex divided by CV
- $\cos \alpha_{p\Lambda}^{}$, $\cos \alpha_{\Lambda\pi}^{}$, $\chi^{2}_{topo}^{}$ (future investigation)



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Two implementations of **KFParticle** based reconstruction

KFParticleFinder

Software design driven by requirement of very fast online particle selection to fit CBM operation at the 10⁷ interaction rate

Implementation:

- vectorized libraries (SIMD technic)
- Fixed set of decays selection criteria
- all-in-one approach:

>150 decay channels different decay topologies, etc.

PFSimple

Software design oriented on flexibility and modularity for systematic performance studies and physics analysis

Implementation:

- User controlled reconstruction algorithm
- Stream reconstruction information and parameters from/to other analysis tools
- Manual selection criteria optimization
- Offers the possibility of flat trees for python
- Optimize selection criteria using Machine Learning techniques for individual decays,

or/and phase space (e.g. p_T - rapidity - centrality)

PFSimple



More details on Git: Analysis tree, KFParticle, PFSimple

X,Y,Z (track position), $T_x=dx/dz$, $T_y=dy/dz$

Distribution of signal & background in the variables

Background

Signal

800

Background

Signal

2.5

3.0

1e7

1000



Distribution of MC signal (pure signal) and background plotted for the variables used in this study



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Machine Learning (ML)

- ML algorithms can perform a specific task by analyzing examples and can learn from data
- Variables associated with decay tracks are analyzed by the algorithm to classify ∧ candidates
- Various ML algorithms tested: (SVM, Regression, MLP, Decision Trees, Gradient Boosting (GB), Extreme GB (XGB))
 - XGB works better in terms of performance



CBM Au+Au collisions @ 12A GeV/c

Gradient Boosting (GB)

Jerome Friedman:



empirical evidence shows that taking lots of small steps in the right direction results in better predictions with the Testing Data

- Boosting combines weak learners (error rate <50%) to make a strong learner (error rate <25%)
- Decision trees (weak learners) are combined together to make a GB algorithm
- In each step a new tree is used to improve the previous prediction
- XGB is an extension of GB with:
 - better control over overfitting
 - parallel processing
 - o additional features



XGB implementation for Λ

- UrQMD sample is taken as experimental data (pure background)
- DCM-QGSM-SMM sample as simulated data (pure signal)
- A candidates are cleaned by removing <u>nonphysical values</u>
- A candidates are divided into train and test samples



Background is selected $\pm 5\sigma$ away from the Λ peak mean

XGB Model evaluation

Model trained on the train sample is applied to the test sample

Optimize Λ candidates selection for significance



True positive rate = tpr; Signal = S ; Background = B

$$tpr = rac{S \ classified \ as \ S}{S \ classified \ as \ S+S \ classified \ as \ B}$$
 $fpr = rac{B \ classified \ as \ B+B \ classified \ as \ S}{B \ classified \ as \ B+B \ classified \ as \ S}$



Threshold on the ROC (Receiver Operating Characteristic) curve which maximizes Approximate Median Significance (AMS) on the test sample is our Best Threshold

 $AMS = \sqrt{2} \left[(tpr + fpr) \log(1 + tpr/fpr) - tpr] \right]$

XGB performance for Λ candidates selection





- Preserve smooth background shape after XGB selection
- Optimal XGB probability (0.96) is applied

Distribution of True signal and background in XGB Selected Signal

1e7



The distribution of MC signal • (pure signal) and background in the XGB selected signal



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Yield Extraction: fitting procedure

Lorentzian function is used for signal and 2nd order polynomial for background:

$$Fit(m) \ = \ A rac{(1/2)\Gamma}{(m-m_0)^2+(\Gamma/2)^2} + pol2(m)$$

- 1. Exclude signal region (m < 1.108 & m > 1.13) and fit background with pol2(m)
- 2. Use background fit parameters as initial values for next iteration, where signal (Lorentzian) fit function has fixed $m_0 = 1.1156 \text{GeV/c}^2$ and width Γ =0.0014 GeV
- 3. Use fit parameters as initial values for unconstrained fit to the whole inv. mass range







Results: acceptance and efficiency of Λ reconstruction



XGB algorithm efficiency

XGB algorithm shows high efficiency $\sim 80\%$

- Reconstructed = reconstructed + selected Λ
- Reconstructable = both daughters are reconstructed



Acceptance and efficiency

Total reconstruction (acc x efficiency) ~ 35%

Results: efficiency and acceptance corrected Λ yield





XGB selection, yield extraction procedure, and efficiency correction allow to recover true Λ yield

Results: Efficiency and acceptance corrected yield (p_T /y projections)







Reconstructed input signal without introducing any bias due to XGB model

Summary and outlook

- A baryon reconstruction in CBM@FAIR with Machine Learning techniques
 - Optimization of selection criteria performed via XGB
 - High signal purity and efficiency achieved
- A yield extraction and efficiency
 - Yield, extracted after XGB selection and (acceptance x efficiency) corrected is compatible with initial model spectra

Outlook

- Include more variables to improve XGB selection and signal to background ratio
- Study different A samples to minimize overfitting and investigate stability
 - \circ multi-differential (p_T, y, centrality) XGB selection, test and training
- Evaluate systematic uncertainties
 - XGB selection variation
 - Yield extraction: variation of fit ranges, background and signal fit functions
- Apply developed procedure for multi-strange hadrons and hyper-nuclei

The CBM Collaboration



56 institutions, 12 countries, ~450 members

Germany Darmstadt TU FAIR Frankfurt Univ, IKF Frankfurt Univ. FIAS Frankfurt Univ. ICS **GSI** Darmstadt Giessen Univ. Heidelberg Univ. P.I Heidelberg Univ. ZITI HZ Dresden-Rossendorf **KIT** Karlsruhe Münster Univ. Tübinden Univ. Wuppertal Univ ZIB Berlin

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Duba

Backup slides





- Well go and check out the following
- Two easy to use jupyter notebooks are available on the following links
 - <u>https://colab.research.google.com/drive/10fD3XNnf_0qt12DiAzlQunbW7IVEqqIE?usp=sharing</u>
 - <u>https://colab.research.google.com/drive/1yV3xboB67trorfOKy1-VLT1kLxYdN6dn?usp=sharing</u>
- <u>our code on github</u>

Applying the model on the URQMD data set



The threshold on the ROC curve which maximizes AMS on the test data set is applied on the URQMD 100k events data set



ML does not cut the background in an unexpected way, therefore, not introducing any bias

Yield Extraction: The Fitting Procedure

- Divide the data into p_{T} -y bins
- Applied fitting to all the bins individually
- Apply a mass cut of 1.13<m<1.108 for a 2nd order pol background fit
- Get the fit parameters and use them as initial fit parameters for the whole mass range, the fitting function is $A \frac{0.5 \times 0.0014}{(m-1.115683)^2 + 0.25 \times 0.0014^2} + B + Cm + Dm^2$
- Get the fit parameters and use them as initial parameters
- The final fit function $A rac{0.5\Gamma}{\left(m-m_0
 ight)^2+0.25\Gamma^2}+B+Cx+Dx^2$





pdf of pT rapidity bins divided data, with fitting

Phase diagram





Multi-strange yields



	√s _{nn}	Run time	R _{int,} kHz	X-	X⁺	Ω⁺
HADES (Ag)	2.6 GeV	4 wks	10	2.5x10 ³		
MPD S1	11 GeV	10 wks	5	1.5x10 ⁶	8x104	1.5x10⁴
СВМ	3.8 GeV	1 wk	1000	4x10 ⁹	5x10 ⁶	3.3x10⁵

Compilation TG, QM2018

C. Blume, C. Markert, PPNP 66 (2011) HADES Coll., PLB 778 (2018) HADES Coll., PRL 103 (2009) 132301 RVUU: F. Li et al., PRC 85 (2012) 064902 UrQMD: J. Steinheimer et al., J.Phys. G43 (2016) 015104 ART: C.M. Ko et al., PLB595 (2004) 158-164 A. Andronic et al., NPA 772 (2006) F. Becattini et al., PRC69 (2004) 024905 E. Seifert et al., PRC97 (2018)

Hypernuclei yield: CBM projections



СВМ	√s _{nn}	Run time	e %	R _{int}	Duty F %	Yield
³ LH	4.7 GeV	1 wks	19	10 MHz	50	5.5x10 ⁹
⁴ _He	4.7 GeV	1 wks	15	10 MHz	50	2.7x10 ⁸
⁶ _{LL} Не	4.7 GeV	10 wks	1	10 MHz	50	146

Blue and red lines are the precision with which we can measure yields assuming various scenario

Compilation TG, QM2018



Data cleaning/skimming

The data contains some entries which does not make sense, so we clean it by pre cuts

p<20	pz > 0	0 <x<sup>2primpos < 1x10⁶</x<sup>	0 <l dl<8000<="" th=""></l>
abs (x, y) < 50	-1< z <80	0 <x<sup>2geo < 10³</x<sup>	cosineneg>0.1
1 <eta<6.5< td=""><td>0<distance< 100<="" td=""><td>0<x<sup>2primneg<3x10⁷</x<sup></td><td>Remove nan, infinite</td></distance<></td></eta<6.5<>	0 <distance< 100<="" td=""><td>0<x<sup>2primneg<3x10⁷</x<sup></td><td>Remove nan, infinite</td></distance<>	0 <x<sup>2primneg<3x10⁷</x<sup>	Remove nan, infinite
1.07 <mass<1.3< td=""><td>I<80</td><td>0<x<sup>2topo< 10⁵</x<sup></td><td>0<i dl<8000<="" td=""></i></td></mass<1.3<>	I<80	0 <x<sup>2topo< 10⁵</x<sup>	0 <i dl<8000<="" td=""></i>

Removes 3.2 % signal candidates from a set of 10k events (AU 12AGeV mbias URQMD)

But also removes 57 % background

mass can't be negative and we select mass greater or equal to the mass of proton and pion

Fixed target experiment, target position: (0,0,0)

X² can't be negative

Gradient boost: regressor, in simple words

 GB: Trees predicting residuals and a learning rate to prevent overfitting 			2 samples $X_1 X_2$		
			Variable ₁	1	2
	varia	► variables	Variable ₂	3	4
ing these	hese		Variable ₃	5	6
	b b c target	rget	У	1	0
	First prediction		Average of y = y'	0.5	0.5
	Pseudo resi	duals	Residual = y - y'	0.5	-0.5
Predicts this			_Tree=h ₁	0.5	-0.5
	2nd pred	iction	Predicted=y" =y'+tree	1	0
Ove	r fits: cont	rols	Learning rate=0.1	0.1	0.1
Low High	y bias 3rd prec	diction	New prediction=y'''=y'+ 0.1* (tree)	0.55	0.45
varia	ance			-	

New residuals=y-y'''	0.45	-0.45
Tree 2 = h_2	0.45	-0.45
Newest prediction= y'''' = y'''+0.1 * (tree 2)	0.595	0.405
Goes on		

Step towards the main target

Further reading https://xgboost.readthedocs.io/en/latest/tutorials/model.html

Detailed Explanation GB

1. Input: Data $\{(x_i, y_i)\}_{i=1}^n$ and a differentiable Loss Function $L(y_i, F(x))$ If we choose L = $\frac{1}{2} \{ y_i - F(x) \}^2$

Then

 $d/dF(x) \{\frac{1}{2} \{ y_i - F(x) \}^2 \} = (-(y_i - F(x))) = F(x) - y_i = -(residuals)$

We minimize this $F(x)-y_i$ for all values

$$\sum_{i}^{n}F(x)-y_{i}=0$$

A predicted value which can minimize this sum is the average

$$F(x) = rac{\sum_{i}^{n}y_{i}}{n}$$
 = average = F_o(x)

2. Fit m = 1 upto m=M number of trees
a. Compute
$$r_{im} = -\left[\frac{\partial L(y_i, F(x_i))}{\partial F(x_i)}\right]$$
 at $F(x) = F_{m-1}(x)$ for i = 1,...,n
since first iteration so $F(x)=F_o(x)$ $-d/dF(x) \{\frac{1}{2} \{y_i - F(x)\}^2\} = -(-(y_i - F(x))) = y_i - F(x)$ Residuals

Detailed Explanation GB

• Fit m = 1 upto m=M trees

a. Compute
$$r_{im} = -\left[rac{\partial L(y_i, F(x_i))}{\partial F(x_i)}
ight]$$
 at F(x) = F_{m-1}(x) for i =1,...,n

- b. Fit a regression tree to the r_{im} values and create terminal regions R_{im} , for j=1,..,J_m (leaves but not with output values)
- c. Determine the output value for each leaf:

for j=1,...,J_m compute

again will turn out to be average if $L = \frac{1}{2} \{ y_i - F(x) \}^2$

d. Update $\gamma_{jm} = argmin \sum_{x_i \in R_{ij}} L(y_i, F_{m-1}(x_i) + \gamma)$ v is learning rate and the equation in the box is the tree we just made We started with F₀ so

$$F_m(x) = F_{m-1}(x) +
u \left| \sum_{j=1}^{J_m} \gamma_{jm} I(x \epsilon R_{jm})
ight|$$

• Output $F_M(x)$ (The final classifier)

$$F_1(x) = F_0(x) +
u \sum_{j=1}^{J_m} \gamma_{jm} I(x \epsilon R_{jm})$$