

Machine Learning application for ∧ hyperon reconstruction in CBM at FAIR

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Strangeness in Quark Matter 17-22 May 2021





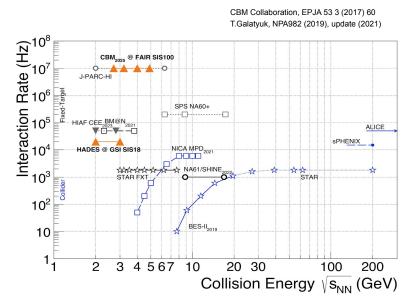


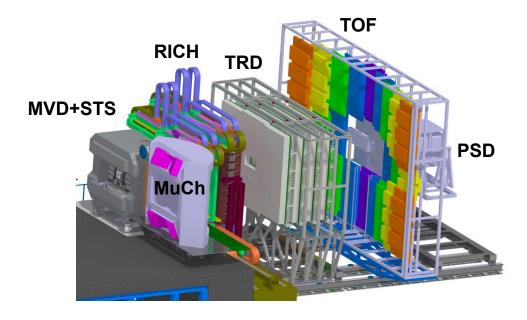






CBM physics goals and experimental challenges





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>
 - (Multi)-Strange particles
- Extend nuclei chart with hypernuclei 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

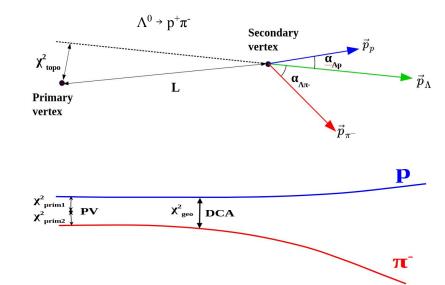
- A hyperons are the most abundant strange baryons produced at FAIR energies
- Collisions generated by URQMD and DCM-QGSM-SMM with Au+Au collisions at p_{heam} = 12A GeV/c (\langle s_{NN} = 4.93), mbias, 100k
- Using GEANT4 simulation, CA tracking within CbmRoot framework

A candidates reconstruction:

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

Variables:

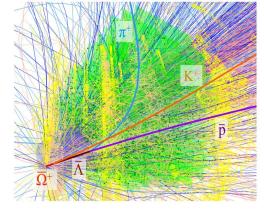
- x²_{prim} squared distance between the daughter track and the primary vertex divided by its Covariance Matrix (CV)
- DCA distance of closest approach between proton & pion tracks
- χ²_{geo} squared distance between daughter tracks divided by CV
- L/ΔL distance between primary and secondary vertex divided by CV
- $\cos \alpha_{p\Lambda}$, $\cos \alpha_{\Lambda\pi}$, χ^2_{topo} (future investigation)



Selection criteria are optimized multi-dimensionally, non-linearly and in an automatized way with Machine Learning algorithms

Machine Learning (ML)

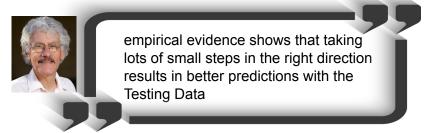
- ML algorithms can perform a specific task by analyzing examples and can learn from data
- Variables associated with decay tracks are analyzed by the algorithm to classify ∧ candidates
- Various ML algorithms tested: (SVM, Regression, MLP, Decision Trees, Gradient Boosting (GB), Extreme GB (XGB))
 -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



CBM Au+Au collisions @ 12A GeV/c

Gradient Boosting (GB)

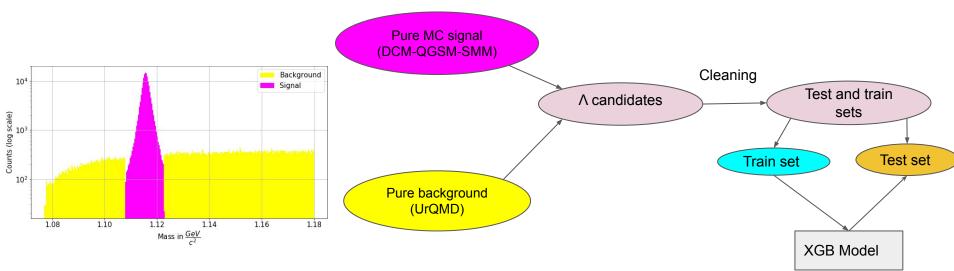
Jerome Friedman:



- 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
 - additional features

XGB implementation for Λ

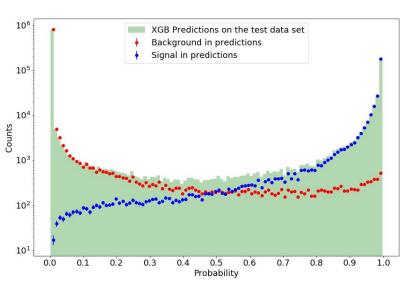
- UrQMD sample is taken as experimental data (pure background)
- DCM-QGSM-SMM sample as simulated data (pure signal)
- A candidates are cleaned by removing <u>nonphysical values</u>
- A candidates are divided into train and test samples



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

XGB Model evaluation

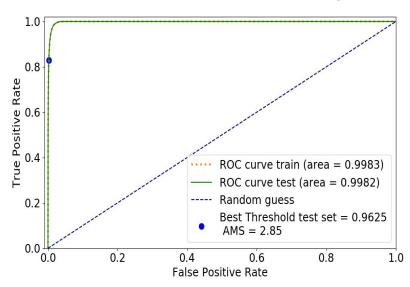
Model trained on the train sample is applied to the test sample



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

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

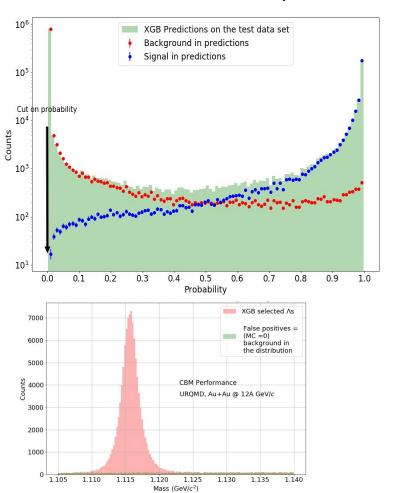
Optimize Λ candidates selection for significance

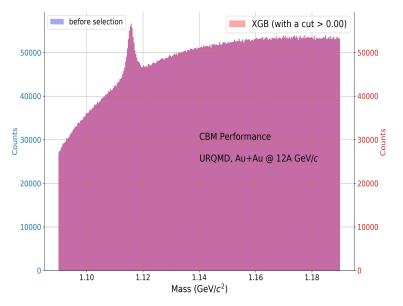


Threshold on the ROC (Receiver Operating Characteristic) curve which maximizes Approximate Median Significance (AMS) on the test sample is our Best Threshold

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

XGB performance for Λ candidates selection





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

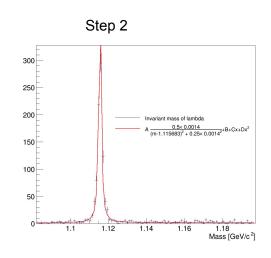
Yield Extraction: fitting procedure

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

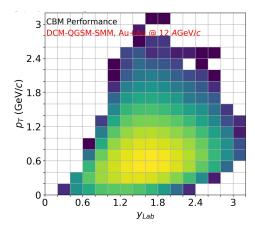
$$Fit(m) \ = \ A rac{(1/2)\Gamma}{(m-m_0)^2 + (\Gamma/2)^2} + pol2(m)$$

- 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 $m_o = 1.1156 GeV/c^2$ and width Γ =0.0014 GeV
- 3. Use fit parameters as initial values for unconstrained fit to the whole inv. mass range

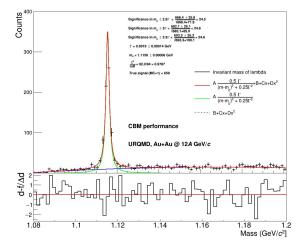
Step 1 25 Invariant mass without \(\Lambda \) peak B+Cx+Dx² Mass in [GeV/c²]



Divide (p_T,y) phase space into 15x15 bins



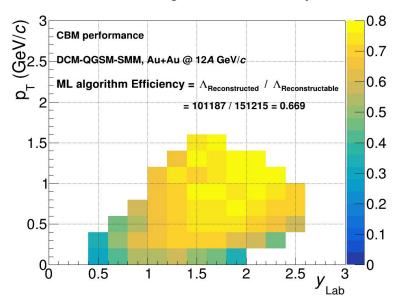




Fits for all (p_T,y) bins available here

Results: acceptance and efficiency of Λ reconstruction

XGB algorithm efficiency

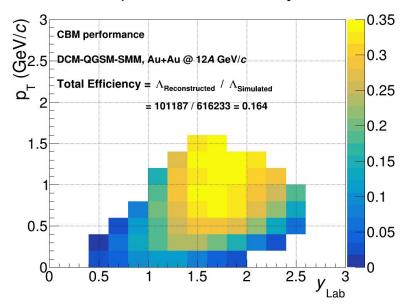


XGB algorithm shows high efficiency ~ 80%

Reconstructed = reconstructed + selected Λ

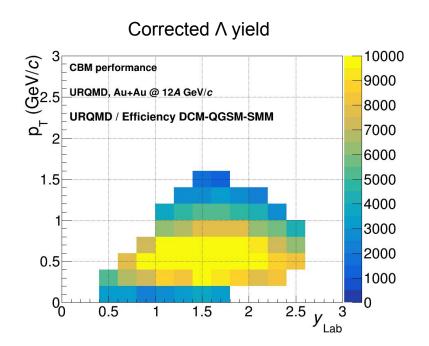
Reconstructable = both daughters are reconstructed

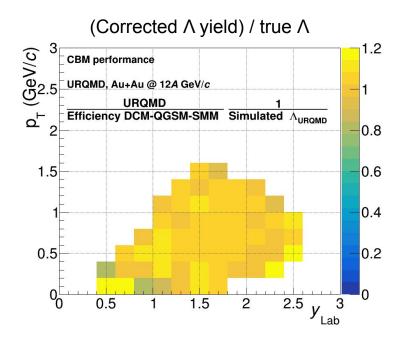
Acceptance and efficiency



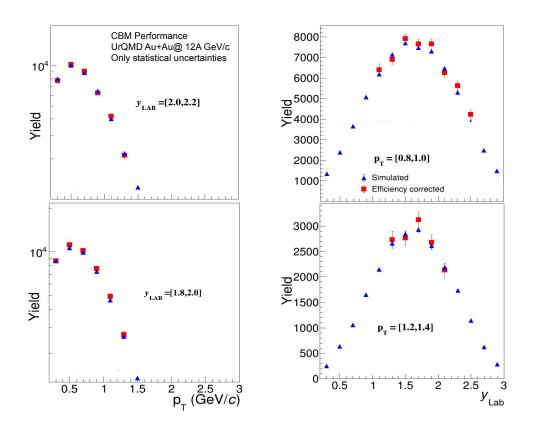
Total reconstruction (acc x efficiency) $\sim 35\%$

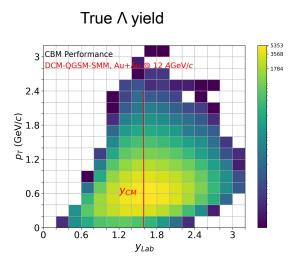
Results: efficiency and acceptance corrected Λ yield





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





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, preserved smooth background shape
- 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 Λ samples to minimize overfitting and investigate stability
 - 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



Backup slides



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

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.
Tübingen Univ.

Wuppertal Univ.

ZIB Berlin

India

Aligarh Muslim Univ.
Bose Inst. Kolkata
Panjab Univ.
Univ. of Jammu
Univ. of Kashmir
Univ. of Calcutta
B.H. Univ. Varanasi
VECC Kolkata
IOP Bhubaneswar
IIT Kharagpur
IIT Indore
Gauhati Univ.

Korea

Pusan Nat. Univ.

Romania

NIPNE Bucharest Univ. Bucharest

Poland

AGH Krakow Jag. Univ. Krakow Warsaw Univ. Warsaw TU

Russia

IHEP Protvino
INR Troitzk
ITEP Moscow
Kurchatov Inst., Moscow
MEPHI Moscow
PNPI Gatchina
SINP MSU, Moscow

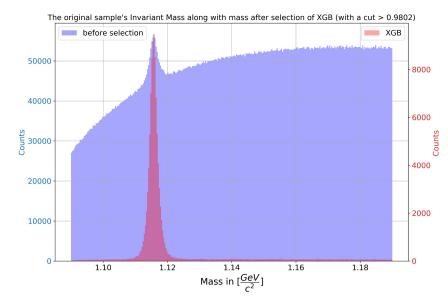
Ukraine

T. Shevchenko Univ. Kiev Kiev Inst. Nucl. Research

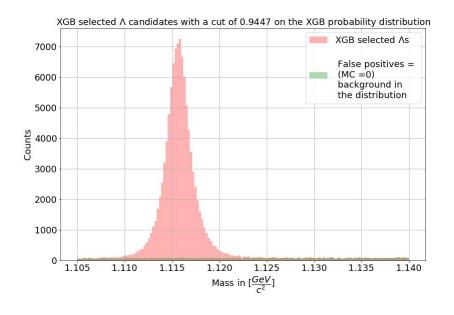
JINR

VBLHEP, Dubna LIT, Dubna

Applying the model on the URQMD data set



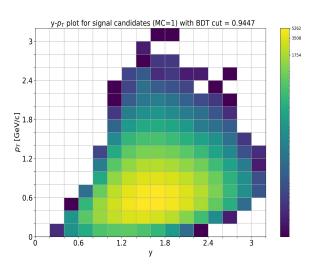
The threshold on the ROC curve which maximizes AMS on the test data set is applied on the URQMD 100k events data set

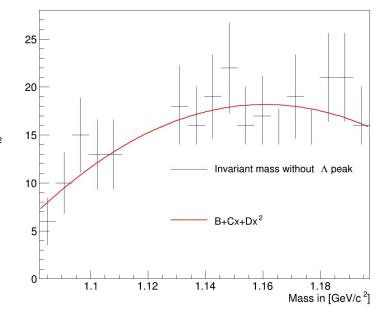


ML does not cut the background in an unexpected way, therefore, not introducing any bias

Yield Extraction: The Fitting Procedure

- Divide the data into p_{τ} -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
- ullet The final fit function $Arac{0.5\Gamma}{(m-m_0)^2+0.25\Gamma^2}+B+Cx+Dx^2$

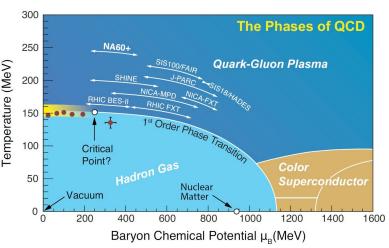




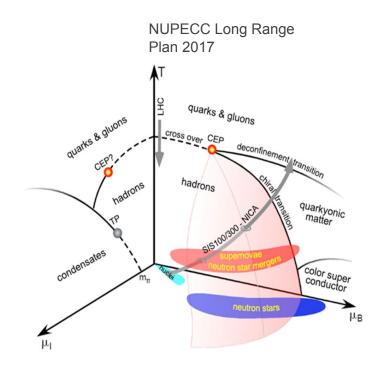
pdf of pT rapidity bins divided data, with fitting

Phase diagram

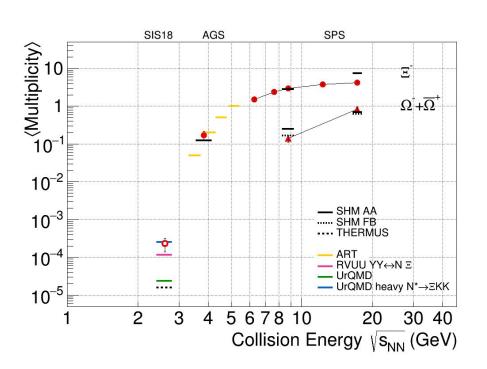
Alessandro D. Falco, CPOD-2021



https://indico.cern.ch/event/985460/contributions/4264615/attachments/2211234/3742919/adf cpod.pdf



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	8x10 ⁴	1.5x10 ⁴
СВМ	3.8 GeV	1 wk	1000	4x10 ⁹	5x10 ⁶	3.3x10 ⁵

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

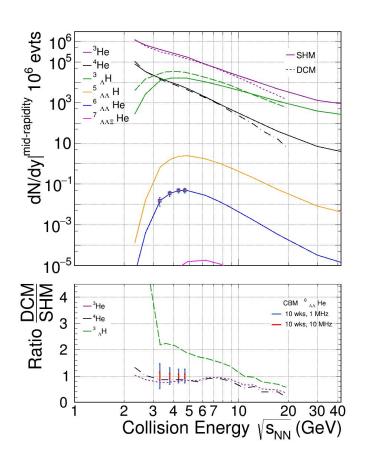
ART: C.M. Ko et al., PLB595 (2004) 158-164

A. Andronic et al., NPA 772 (2006)

F. Becattini et al., PRC69 (2004) 024905

E. Seifert et al., PRC97 (2018)

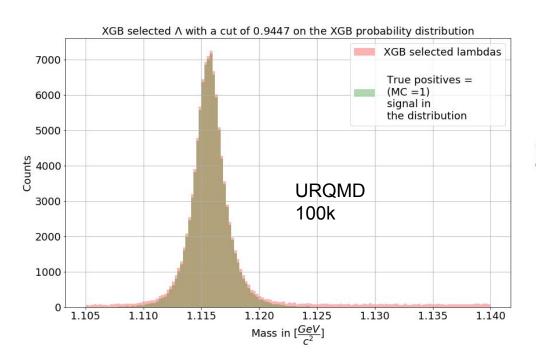
Hypernuclei yield: CBM projections

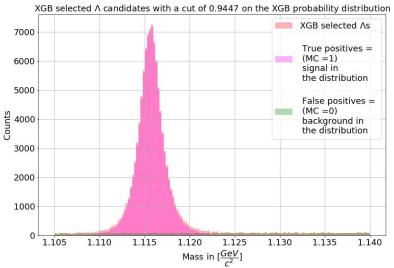


СВМ	√s _{nn}	Run time	e %	R _{int}	Duty F %	Yield
³ _L H	4.7 GeV	1 wks	19	10 MHz	50	5.5x10 ⁹
⁴ _L He	4.7 GeV	1 wks	15	10 MHz	50	2.7x10 ⁸
⁶ LLHe	4.7 GeV	10 wks	1	10 MHz	50	146

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>	cosinepos > 0.5	0 <i di<8000<="" td=""></i>
pT<3	-1< z <80	0 <x<sup>2geo < 10³</x<sup>	cosineneg>0.1	abs (x) < 50
1 <eta<6.5< td=""><td>0<distance< 100<="" td=""><td>0<x<sup>2primneg<3x10⁷</x<sup></td><td>Remove nan</td><td>abs (y) < 50</td></distance<></td></eta<6.5<>	0 <distance< 100<="" td=""><td>0<x<sup>2primneg<3x10⁷</x<sup></td><td>Remove nan</td><td>abs (y) < 50</td></distance<>	0 <x<sup>2primneg<3x10⁷</x<sup>	Remove nan	abs (y) < 50
1.07 <mass<2.5< td=""><td>I<80</td><td>0<x<sup>2topo< 10⁵</x<sup></td><td>infinite values</td><td>0<i di<8000<="" td=""></i></td></mass<2.5<>	I<80	0 <x<sup>2topo< 10⁵</x<sup>	infinite values	0 <i di<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

Gradient boost: regressor, in simple words

 GB: Trees predicting residuals and a learning rate to prevent overfitting

variance

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	X ₁	X
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			•	ı	_
Jsing these		→ variables	Variable ₁	1	2
			Variable ₂	3	4
	nese		Variable ₃	5	6
-	ıng tı	target	у	1	0
Ns		irst prediction	Average of y = y'	0.5	0.5
	Ps	eudo residuals	Residual = y - y'	0.5	-0.5
Predicts this		s this (_Tree=h ₁	0.5	-0.5
		2nd prediction	Predicted=y" =y'+tree	1	0
91	r fits:	controls	_Learning rate=0.1	0.1	0.1
۷ h	bias	3rd prediction	New prediction=y""=y'+ 0.1* (tree)	0.55	0.45

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

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 $_{ extsf{o}}$ (x)

2. Fit m = 1 upto m=M number of trees

Detailed Explanation GB

- Fit m = 1 upto m=M trees
 - a. Compute $r_{im}=-\left\lceil \frac{\partial L(y_i,F(x_i))}{\partial F(x_i)} \right\rceil$ 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 $j=1,...,J_{im}$ (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 F $_0$ so

$$F_m(x) = F_{m-1}(x) +
u iggl[\sum_{j=1}^{J_m} \gamma_{jm} I(x \epsilon R_{jm}) iggr]$$

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

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