# UPDATE ON THE ANALYSIS OF GSI2 <sup>16</sup>O DATA TAKING

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### Outline

• ECC reconstruction and improvements

- Improvements in tracking algorithms
- ECC reconstructions: GSI1, GSI2, GSI3 (NEW)
- Comparison between: true and reconstructed Monte Carlo - reconstructed Monte Carlo and DATA
- Cross section measurement for GSI1 and GSI2
  - Integrated
  - Differential for fragment's angles
  - Differential for charge

#### Summary



#### GSI1 DATA vs MC RECO Comparison @ XI General Meeting



In GSI1 (1mm C target) there were several discrepancies between data and reconstructed MC, that we didn't see in GSI2 (2mm  $C_2H_4$  target).

Hypothesis was that these discrepancies were due to alignment problems



## GSI1 - S1<sup>0</sup> / <sup>5</sup>B<sup>1</sup>F<sup>15</sup>O<sup>20</sup>R<sup>25+30</sup> <sup>5</sup>[0].eScaniD.ePlate



\_0.**5**000

• Problem not with coordinates alignment, but with angle distributions! The distribution of  $\theta_{X/Y}$  of "signal" tracks seemed to be centered on a different mean angle in each plate, probably due to scanning done on different microscopes  $10^{5}$ • Differences between consecutive base-tracks up to ~ 20 mrad, out of tolerances  $\xrightarrow{12000}$  match between consecutive base-tracks not found  $^{3000}$ 10<sup>4</sup> 10000 0.5 2500

### $GSI1 - S1 \frac{5}{4F^{5}} \frac{100}{15} \frac{15}{20} \frac{25}{25} \frac{25}{25} \frac{30}{25} \frac{30}{25} \frac{100}{15} \frac{15}{20} \frac{20}{25} \frac{25}{25} \frac{25}{20} \frac{30}{25} \frac{30}{25} \frac{30}{25} \frac{100}{25} \frac{100}{25$



Angles correction implemented in the last stepson glabal alignment
Nove all distributions are centered to the mean of the first plate 105







### Merging stacks

- Different reference systems in consecutive stacks both for coordinates and angles
- Efforts to align the two datasets
- New procedure (still to improve) **Example from GSI2 DATA:**



Impact Parameter

s.eTX:s.eScanID.ePlate {abs(s.eTX)<0.05&&abs(s.eTY)<0.05}

-0.02

-0.04

2500

2000

1500

1000

500

s.eScanID.ePlate





# Analysis of Oxygen at 200 MeV/n on C and C<sub>2</sub>H<sub>4</sub> GSI 1 & GSI2

#### Number of reconstructed vertices

		GSI1 C target	GSI2 C2H4 target		
MC	Beam particles	18990	19988		
	True vertices	4798	5567		
	<b>Reco vertices</b>	4031	4753		
DATA	Beam particles	19375	20625		
	Data vertices	4086	5136		

Cuts for vertices selection:

•  $n \ge 3$ , n = number of tracks (parent + daughters)

• At least 2 daughters with at least 3 segments

#### Daughters' impact parameters distribution

Distributions normalized to beam particles

#### Carbon



#### Vertex Plate distribution: MC RECO vs MC



Since at least 3 tracks are required, efficiencies increases when the probability of fully reconstruct the incoming oxygen is higher. Smearing at Bragg peak.

#### Vertex Plate distribution: DATA vs MC RECO



Trend is confirmed in DATA

#### Fragments' multiplicity distribution: MC RECO vs MC

Distributions normalized to beam particles, requirement of at least 3 tracks in the vertex.

Carbon



#### Fragments' multiplicity distribution: DATA vs MC reco

Distributions normalized to beam particles, requirement of at least 3 tracks in the vertex.

#### Carbon



#### Fragments' angular distribution: MC RECO vs MC

Distributions normalized to beam particles, requirement of at least 3 tracks in the vertex.

Carbon



#### Fragment's angular distribution: DATA vs MC RECO

Distributions normalized to beam particles, requirement of at least 3 tracks in the vertex.

Carbon



#### Mean Oxygen Kinetic Energy per layer distribution (MC)



#### **Cross Section Measurement**

#### Integrated cross section:

$$\sigma(E_{kin \ O})\Big|_{C \ or \ C_2H_4} = \frac{Y_i(E_{kin})}{N_B N_{TG} \epsilon^i_{reco}(E_{kin})}$$
$$\sigma(E_{kin \ O})\Big|_{H} = \frac{1}{4} \left( \sigma(E_{kin \ O})\Big|_{C_2H_4} - 2\sigma(E_{kin \ O})\Big|_{C} \right)$$

•  $Y_i = \#$  of fragments in the interval  $\Delta E_{kin \ Oxy}$ • $N_B = \#$  of ions colliding on the target • $N_{TG} = \#$  of particles in the target:  $\frac{\rho dN_A}{\Lambda}$ , with: •  $\rho$  = target density:  $\rho_{C} = 2.26 g/cm^{3}$  $\rho_{C_2H_4} = 0.94g/cm^3$  $\rho_{H} = 0.0708 g/cm^{3}$ • d =target thickness:  $d_C = 0.1 cm$  per layer  $d_{C_2H_4} = 0.2cm$  per layer •  $N_A = 6.022 \cdot 10^{23} / mol$  $\bullet A = molar mass:$  $A_C = 12g/mol$  $A_{C2H4} = 28g/mol$  $A_H = 1g/mol$ • $\epsilon_{reco}^{i}$  = reconstruction efficiency 22

#### Integrate cross section



Projectile atomic number	Projectile mass number	Target atomic number	Target mass number	Target chemical formula	Projectile kinetic energy (MeV/u)	Cross-section + type	Cross-section (mb)	Cross-section	Cross-section upper error (mb)	First author of the publication	Year of publicatio
8	16	6	12	С	288	сс	852	17	17	Yamaguchi	2011
8	16	6	12	С	290	СС	863	20	20	Zeitlin	2011
8	16	6	12	С	400	сс	842	22	22	Zeitlin	2011



#### Integrate cross section



https://crosssection-db.herokuapp.com/

#### Integrate cross section



#### **Cross Section Measurement**

Starting from kinematic distributions it's possible to evaluate the differential cross section:

$$\frac{d\sigma(\theta)}{d\theta}\Big|_{C \text{ or } C_2H_4} = \frac{Y_i(\theta)}{N_B N_{TG} \Delta \theta \epsilon_{reco}^i(\theta)}$$
$$\frac{d\sigma(\theta)}{d\theta}\Big|_{H} = \frac{1}{4} \left( \frac{d\sigma(\theta)}{d\theta} \Big|_{C_2H_4} - 2\frac{d\sigma(\theta)}{d\theta} \Big|_{C} \right)$$

• 
$$Y_i = \#$$
 of fragments in the interval  $\Delta \theta$   
•  $N_B = \#$  of ions colliding on the target  
•  $N_{TG} = \#$  of particles in the target:  $\frac{\rho dN_A}{A}$ , with:  
•  $\rho = \text{target density:}$   
 $\rho_C = 2.26g/cm^3$   
 $\rho_{C_2H_4} = 0.94g/cm^3$   
 $\rho_H = 0.0708g/cm^3$   
•  $d = \text{target thickness:}$   
 $d_C = 0.1cm \text{ per layer}$   
 $d_{C_2H_4} = 0.2cm \text{ per layer}$   
•  $N_A = 6.022 \cdot 10^{23}/mol$   
•  $A = \text{molar mass:}$   
 $A_C = 12g/mol$   
 $A_{C2H_4} = 28g/mol$   
 $A_H = 1g/mol$   
•  $\Delta \theta = \theta$  bin  
•  $\epsilon_{reco}^i = \text{reconstruction efficiency}$ 

#### Differential cross section for angle (Mean Energy)



#### Differential cross section for angle (Mean Energy)



### Charge distribution: MC RECO vs MC



- Ionization and refreshing not simulated in MC: true charge used
- Efficiency of merging S1 and S2 to improve

### Charge distribution: DATA vs MC RECO

#### Distributions normalized to beam particles



For charge evaluation see:

- https://doi.org/10.1515/phys-2021-0032
- V. Boccia's talk at Physics Meeting:

https://agenda.infn.it/event/29377/contributions/149216/attachments/90378/121729/ Update ChargeID GSI3 04 05 2022.pdf



# Analysis of Oxygen at 400 MeV/n on C GSI 3

### **GSI3 - Emulsions Quality Check S1**

s.eY:s.eX

s.eY:s.eX



- Bad emulsions quality
- S1 were made by Slavich company, S2 and following by Nagoya emulsions
- GSI3 and GSI4 batch different from GSI1' and GSI2' one

#### GSI3 - Emulsions Quality Check S2



• No problems in S2 (emulsions produced in Nagoya)

#### Local alignment and shrinkage corrections

• Trials on-going to recover tracks and vertices reconstructions

• Trying to local align (corrections for each cm<sup>2</sup>)



#### **GSI3** Vertex Plate distribution

Distributions normalized to beam particles



Still many fake or multiplied vertices

### **GSI3** Fragments' angular distribution

Distributions normalized to beam particles, requirement of at least 3 tracks in the vertex.



#### Conclusions

Oxygen @ 200 MeV/n on C and C2H4

- Integrated cross section evaluated
- Differential cross section evaluation on-going

#### Oxygen @ 400 MeV/n on C

• Bad emulsions quality: efforts on-going to recover data reconstruction



# **BACK UP SLIDES**

#### **Detector Structure**



#### **GSI1 - S1**



#### **GSI1 - S2**















#### npl {abs(t.eTX)<0.05&&abs(t.eTY)<0.05}





#### **GSI2 - S1**



#### **GSI2 - S2**





s.eTX:s.eScanID.ePlate {abs(s.eTX)<0.05&&abs(s.eTY)<0.05}



s.eTY:s.eScanID.ePlate {abs(s.eTX)<0.05&&abs(s.eTY)<0.05}



#### **GSI3 - S1**



s.eTX:s.eScanID.ePlate {abs(s.eTX)<0.05&&abs(s.eTY)<0.05}



s.eTY:s.eScanID.ePlate {abs(s.eTX)<0.05&&abs(s.eTY)<0.05}





s.eTX:s.eScanID.ePlate



s.eTY:s.eScanID.ePlate





#### npl {abs(t.eTX)<0.05&&abs(t.eTY)<0.05}





#### **GSI3 - S2**













s.eTX:s.eScanID.ePlate {abs(s.eTX)<0.05&&abs(s.eTY)<0.05}



s.eTY:s.eScanID.ePlate {abs(s.eTX)<0.05&&abs(s.eTY)<0.05}







