

JER/S

Hou Yingqi

2024/12/11



Now: TDR 24.12.0



Selection:

```
data_barrel = data[(abs(data["GEN_jet1_costheta"]) < 0.70) & (abs(data["GEN_jet2_costheta"]) < 0.70)]
data_barrel_match = data_barrel[ (data_barrel["jet1_GENMatch_id"] != data_barrel["jet2_GENMatch_id"]) ]

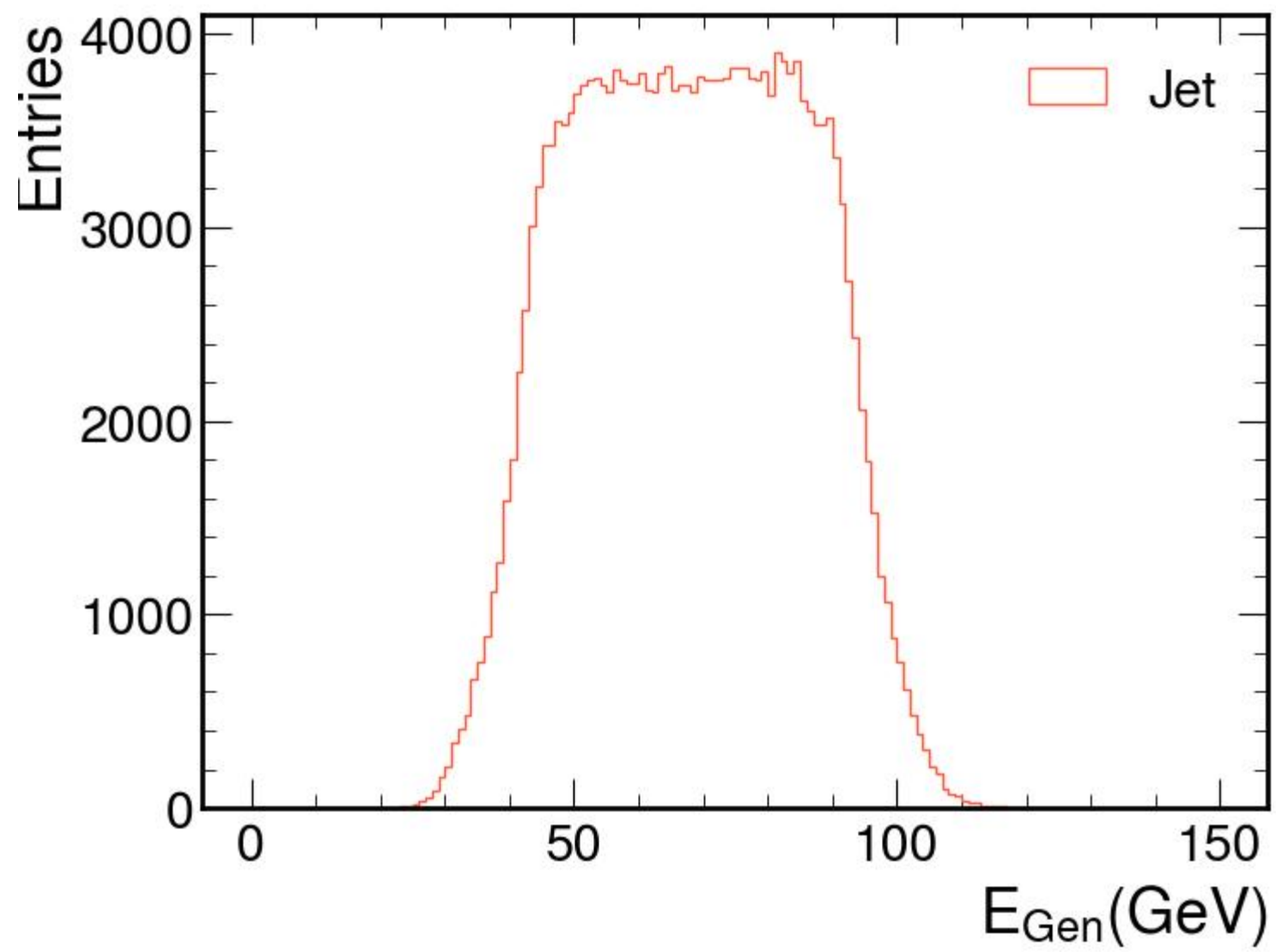
data_barrel_match["jet1_theta"] = np.arccos(data_barrel_match["jet1_costheta"])
data_barrel_match["jet2_theta"] = np.arccos(data_barrel_match["jet2_costheta"])
data_barrel_match["deltaR_j1j2"] = np.sqrt((data_barrel_match["jet1_theta"] - data_barrel_match["jet2_theta"])**2 +
                                           (data_barrel_match["jet1_phi"] - data_barrel_match["jet2_phi"])**2)
dR_cut = 0.6
data_barrel_match_rmOverlape_cutdR = data_barrel_match[(data_barrel_match["jet1_GENMatch_mindR"]<dR_cut)
                                                       & (data_barrel_match["jet2_GENMatch_mindR"]<dR_cut) & (data_barrel_match["deltaR_j1j2"] >= 2)]
```

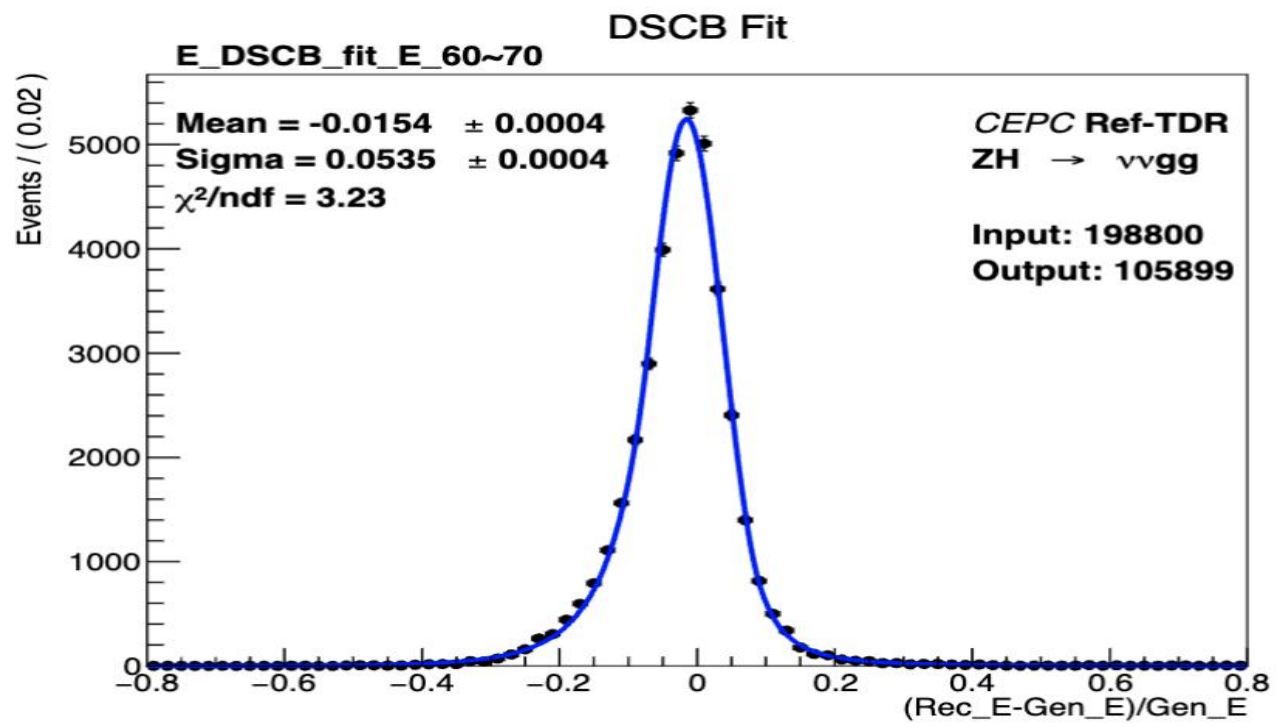
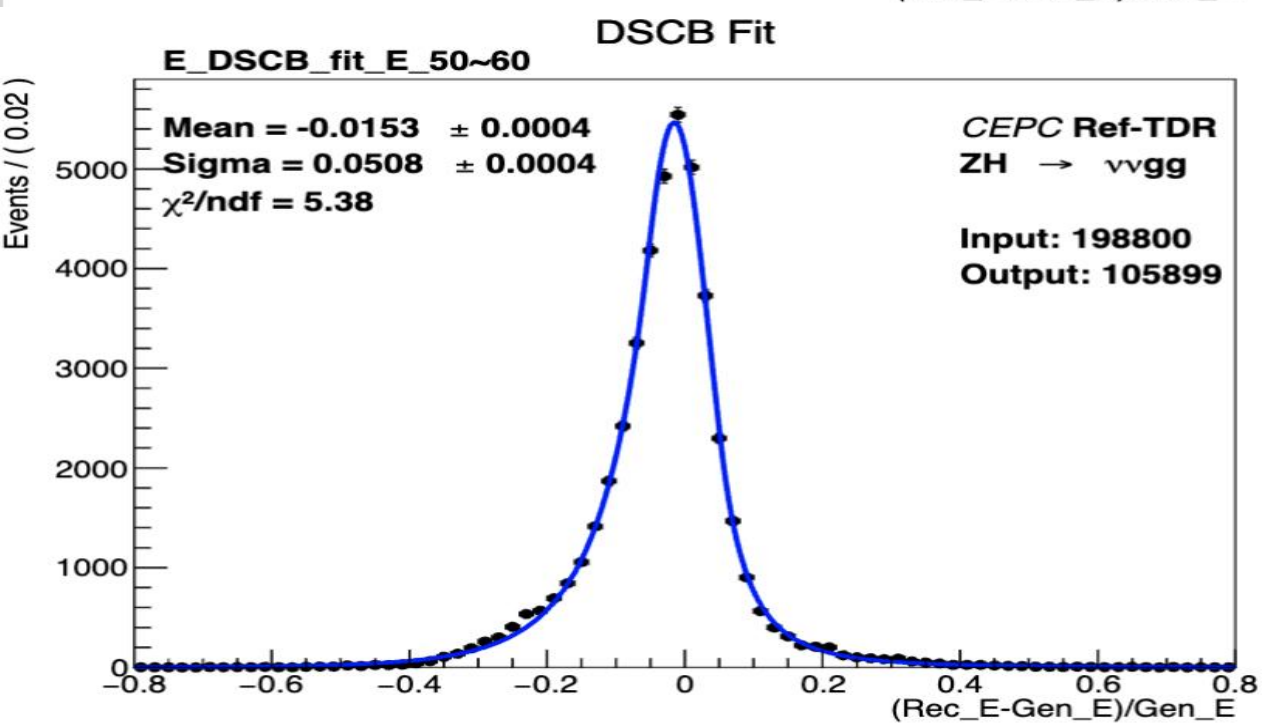
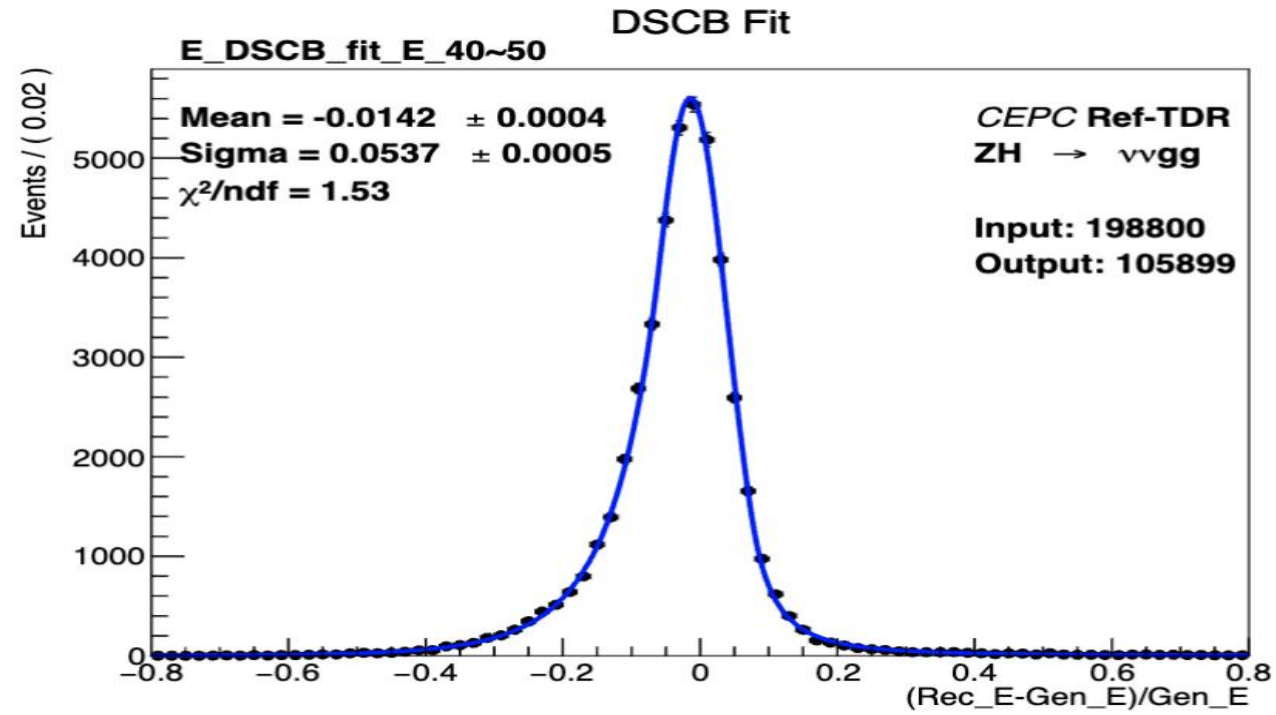
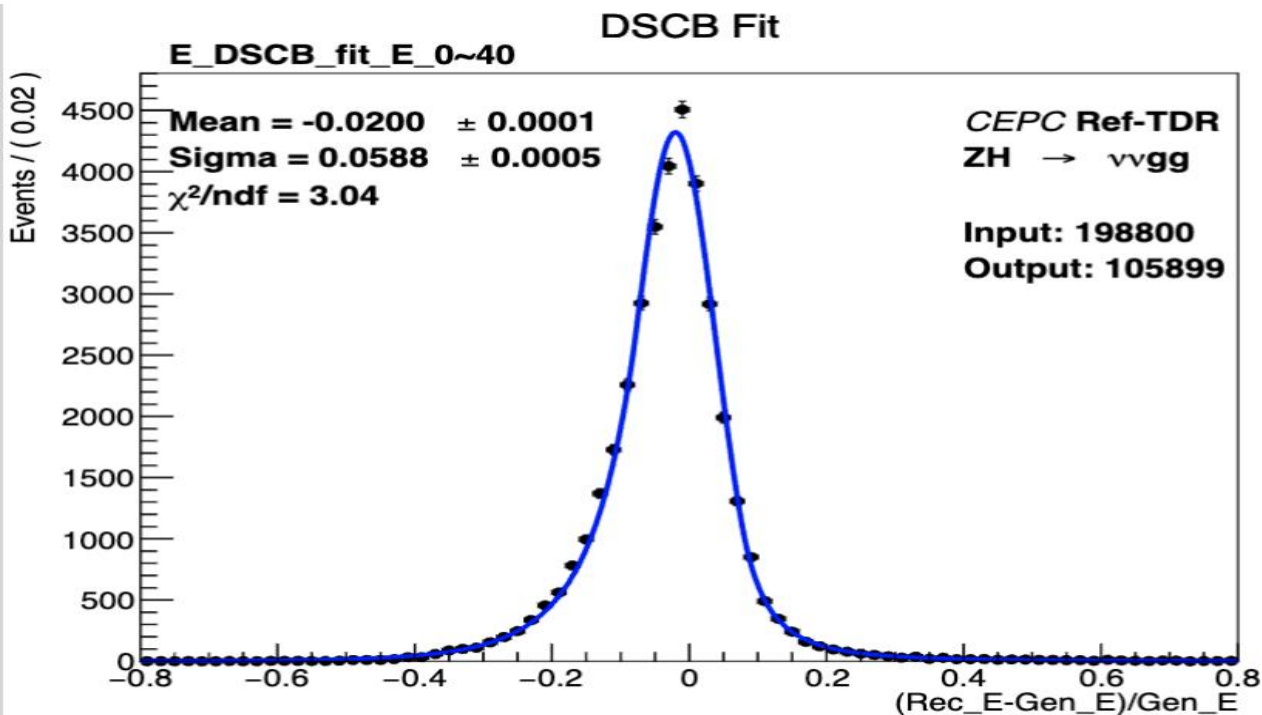
Process:

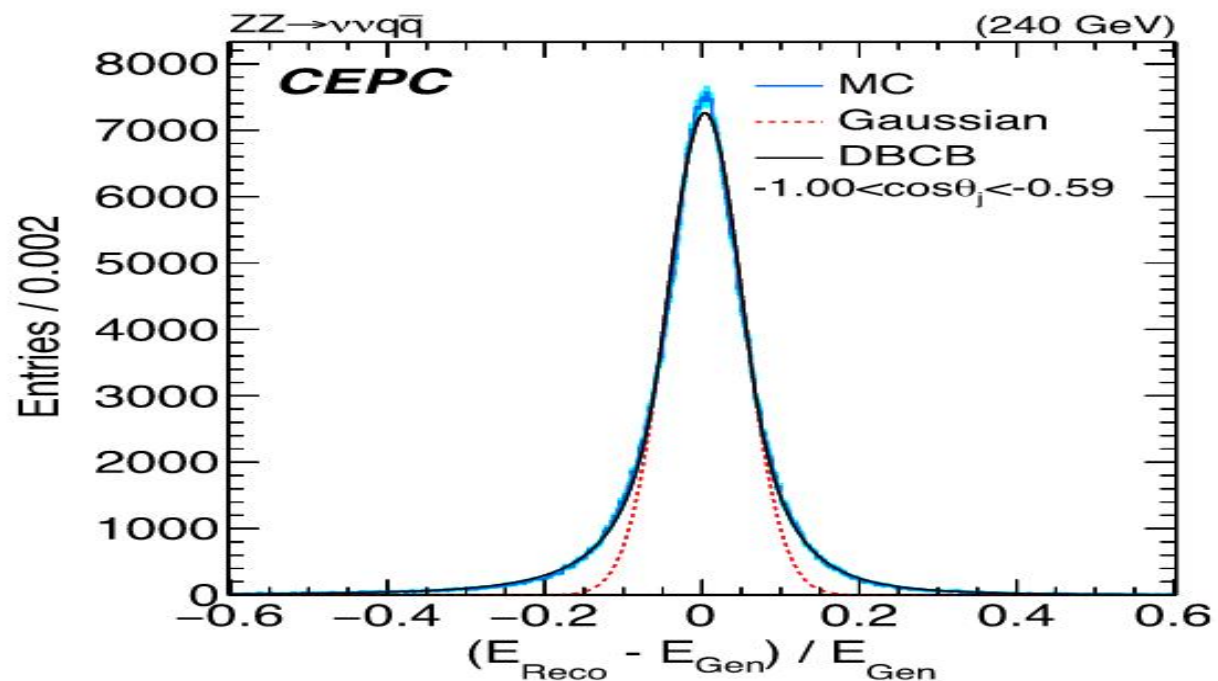
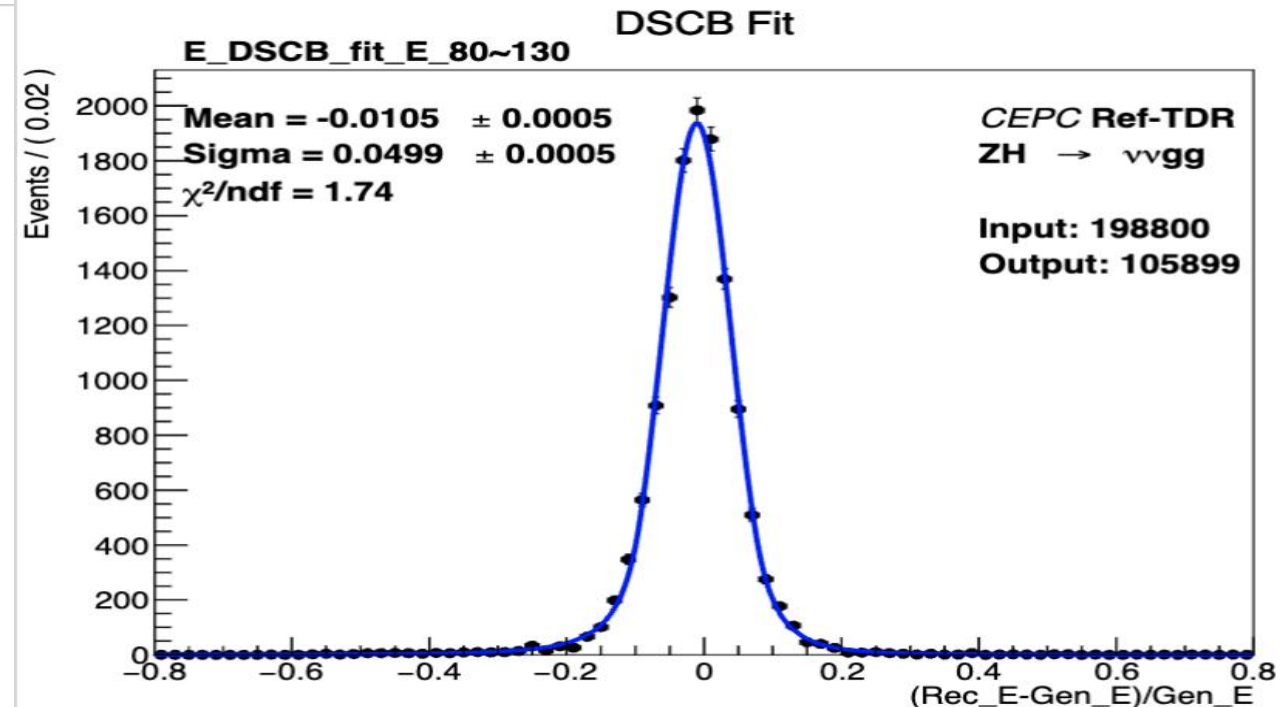
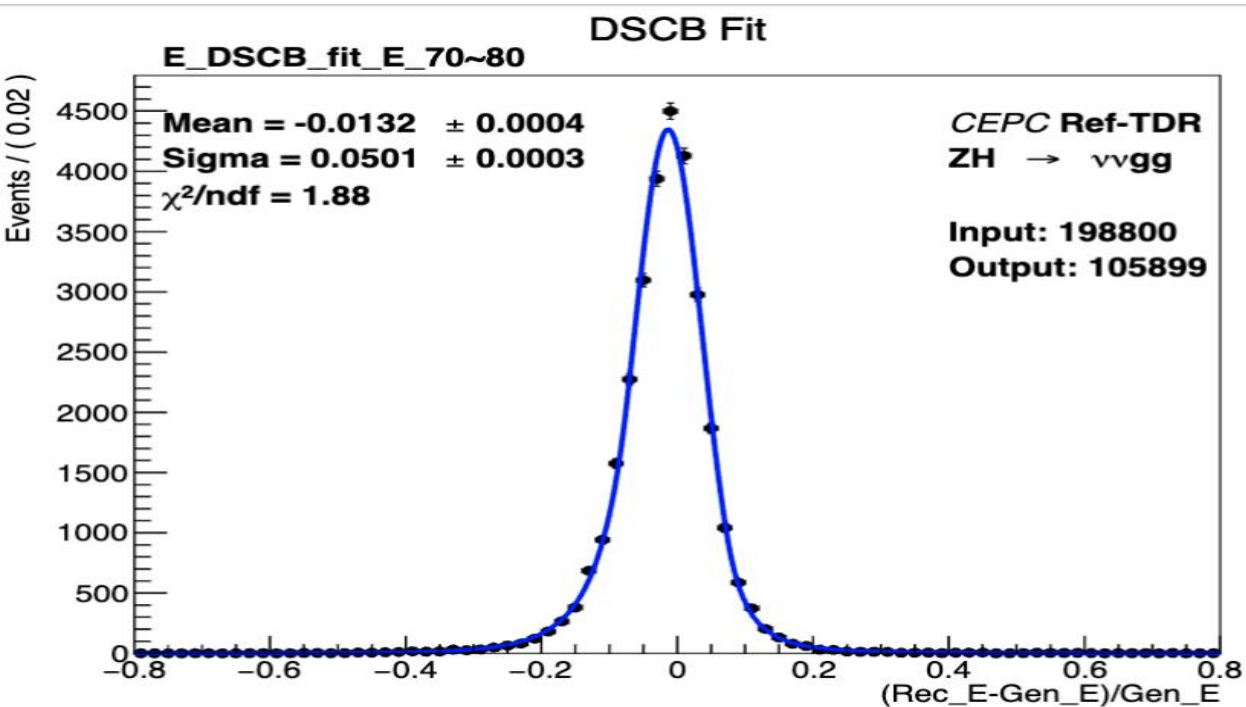
```
data = getEntries("/cefs/higgs/houyingqi/jet_E240_nnHgg/jet_E240_nnHgg.root", "jets", variables)
```



Check GenJet







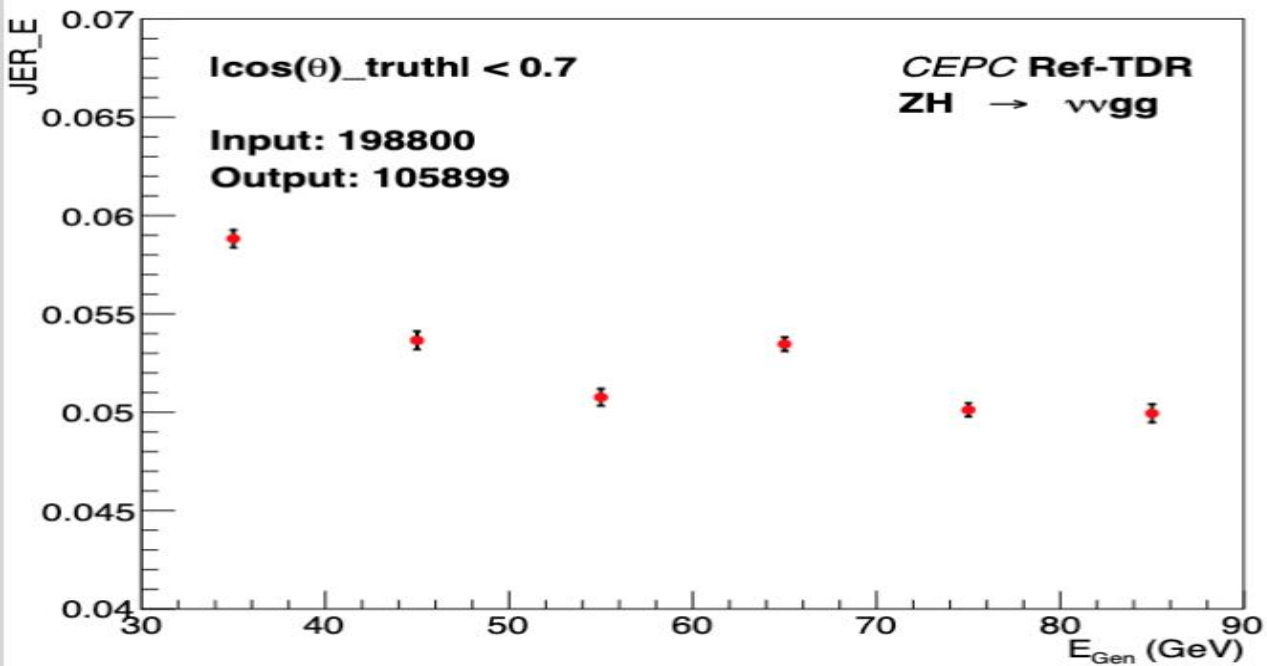
PAPER

Jet performance at the circular electron-positron collider

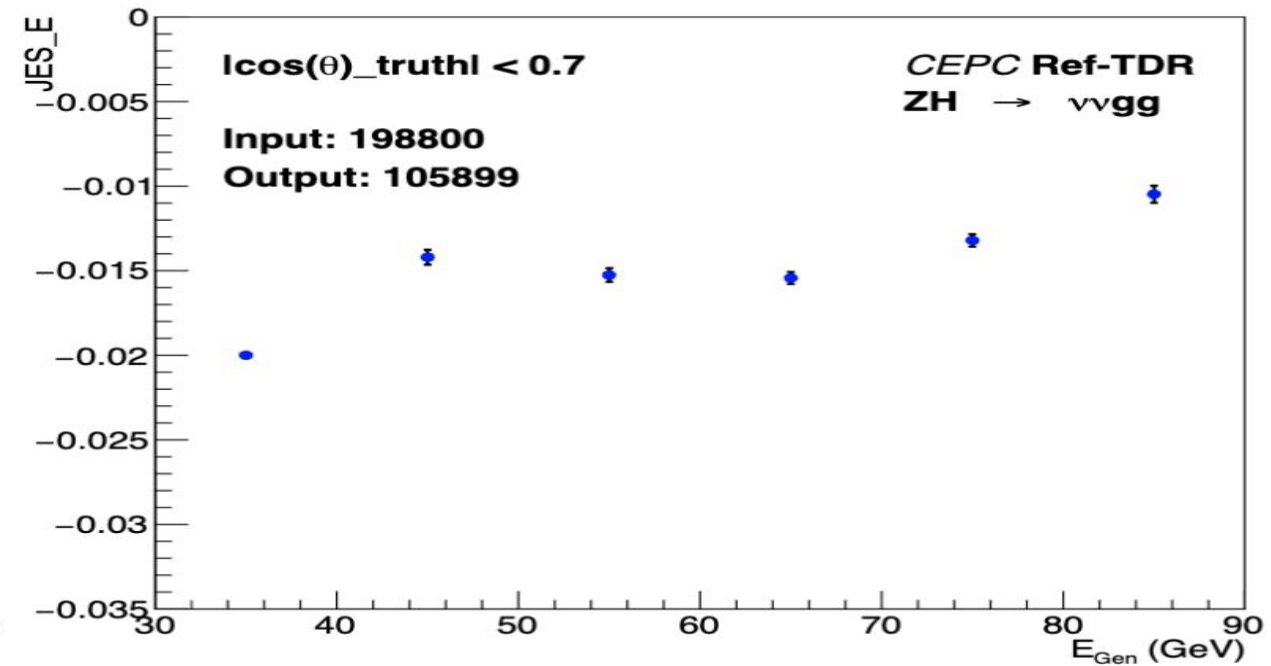
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✓ According to Peizhu's diagram, we can see that this is reasonable that some points of the peak do not fit well.

JER



JES



- ✓ JER generally decreases with the increase of energy.
- ✓ JER has a trip between energy of 60~70 GeV, doubt is a matter of Genmatch.

- JES has a tendency to increase with the increase of energy.
- Compared to the previous version, JES is negative now.

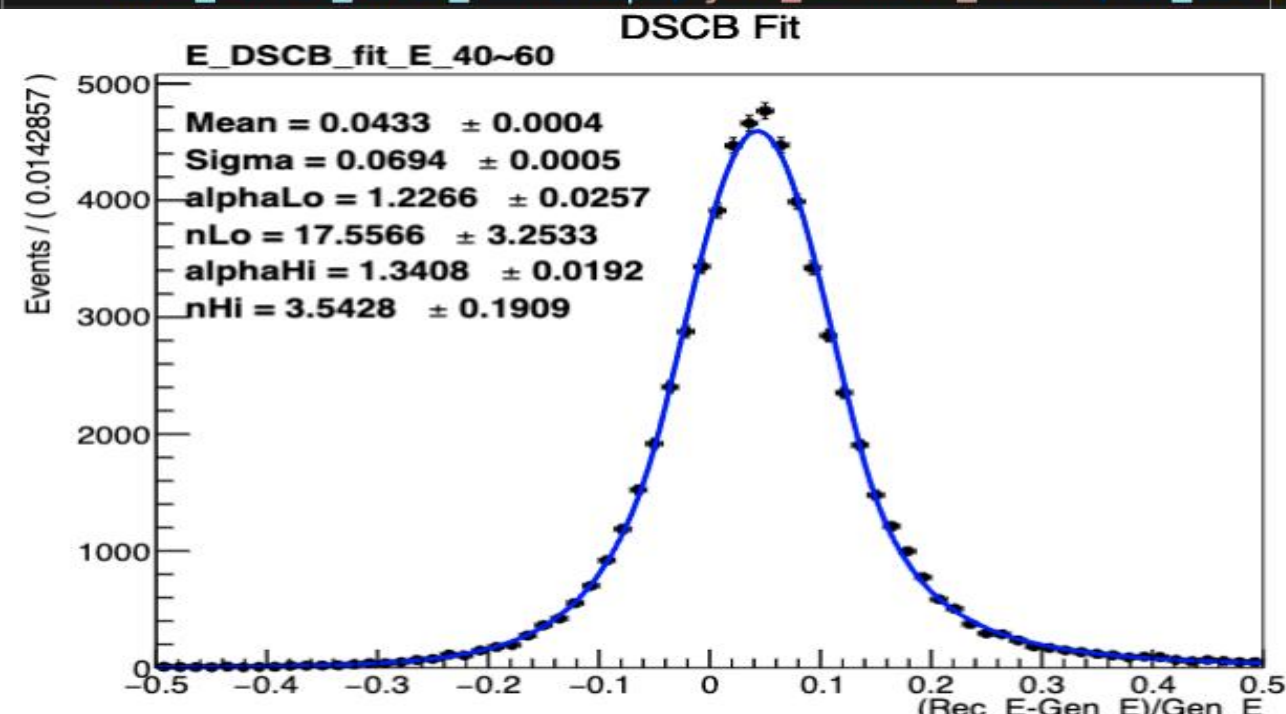
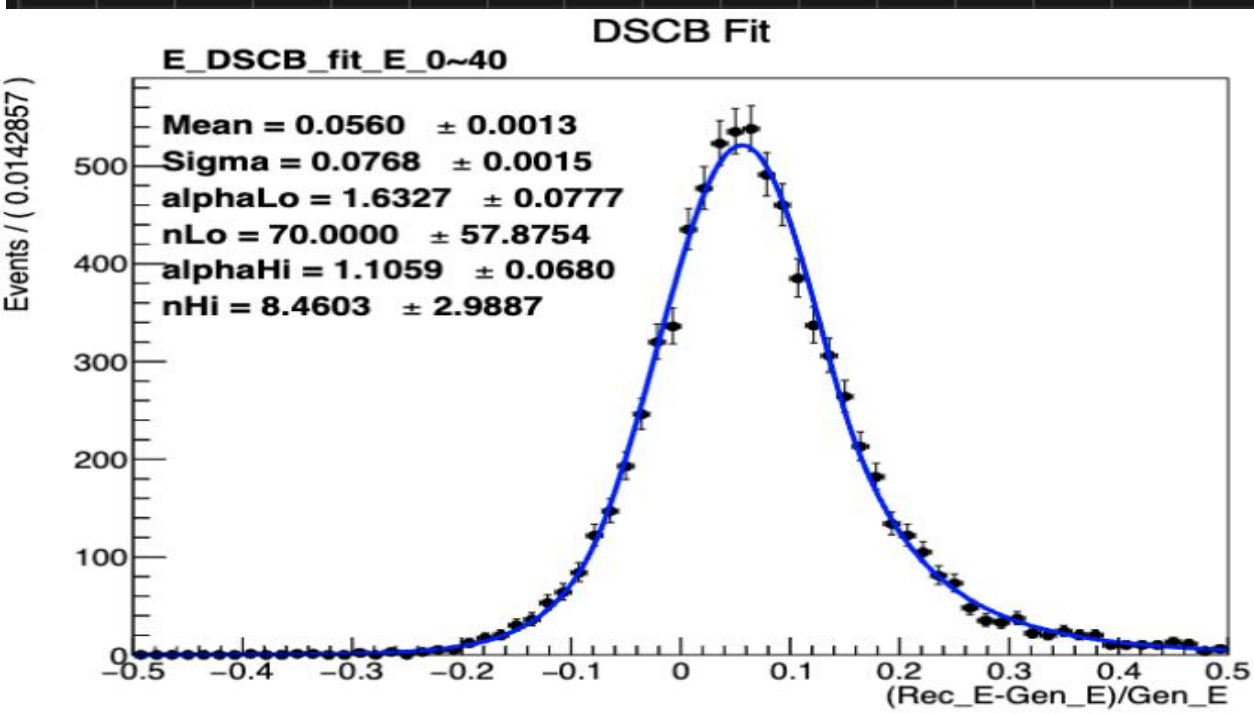


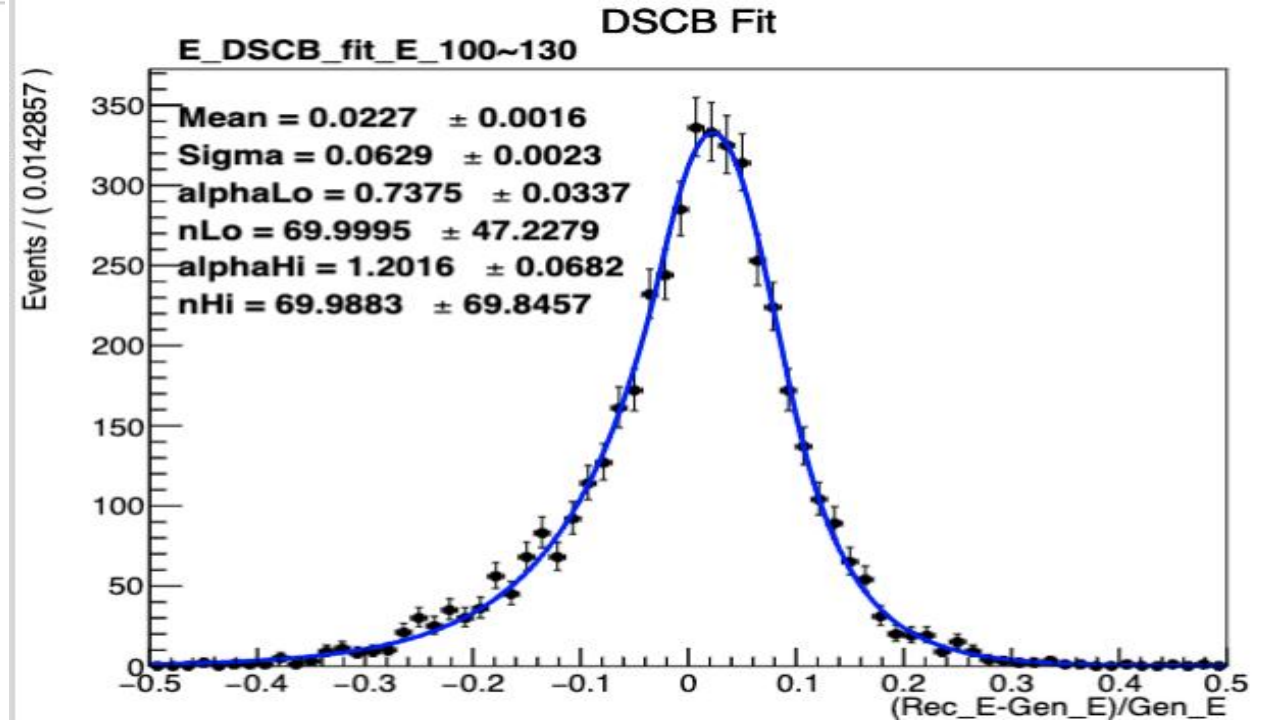
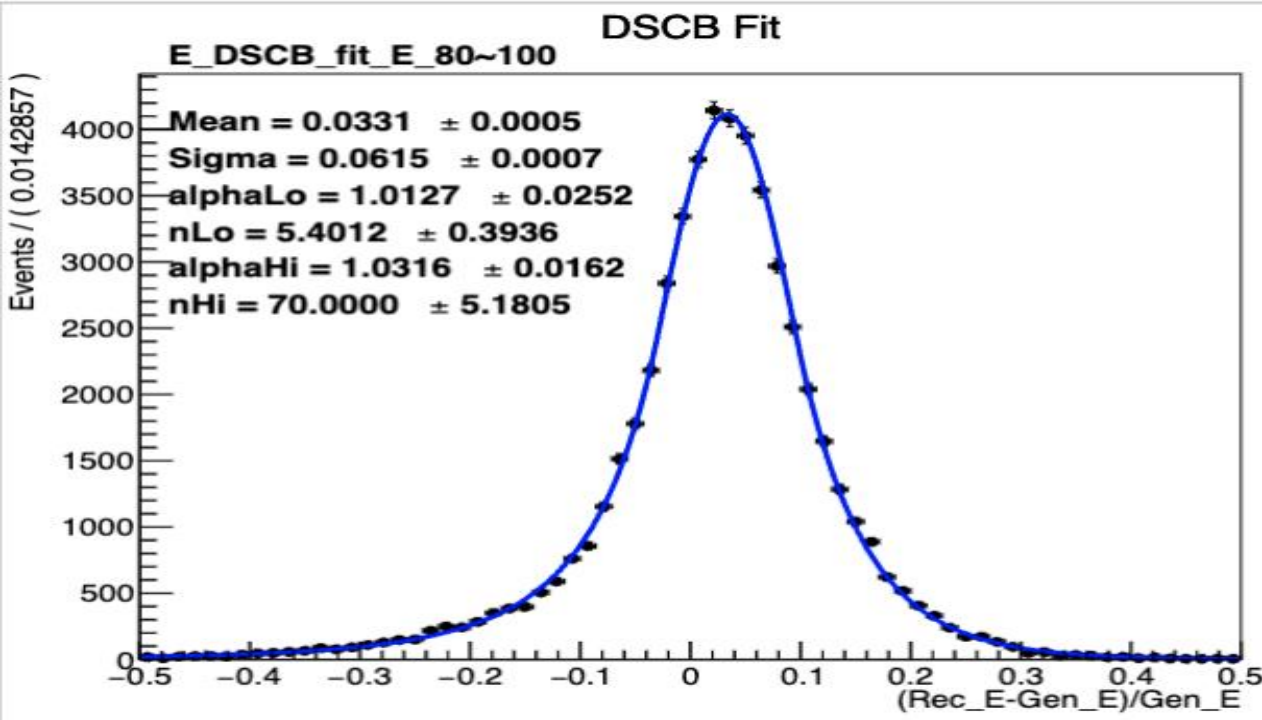
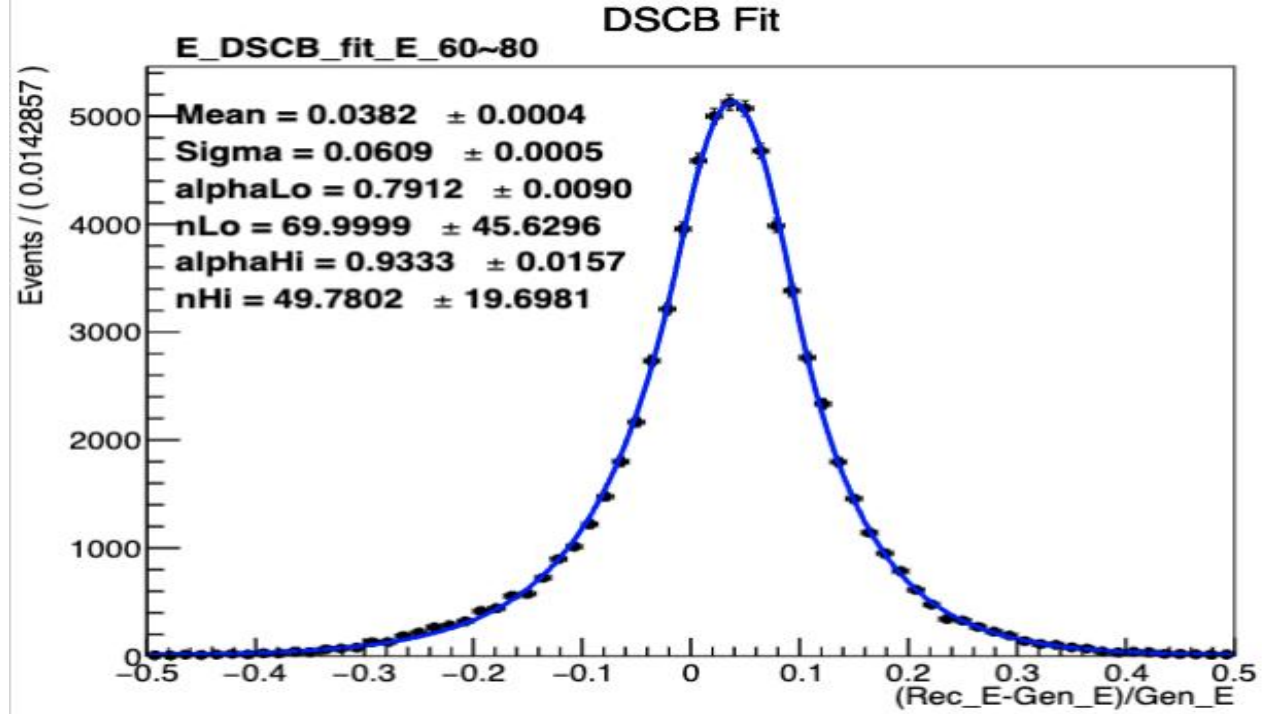
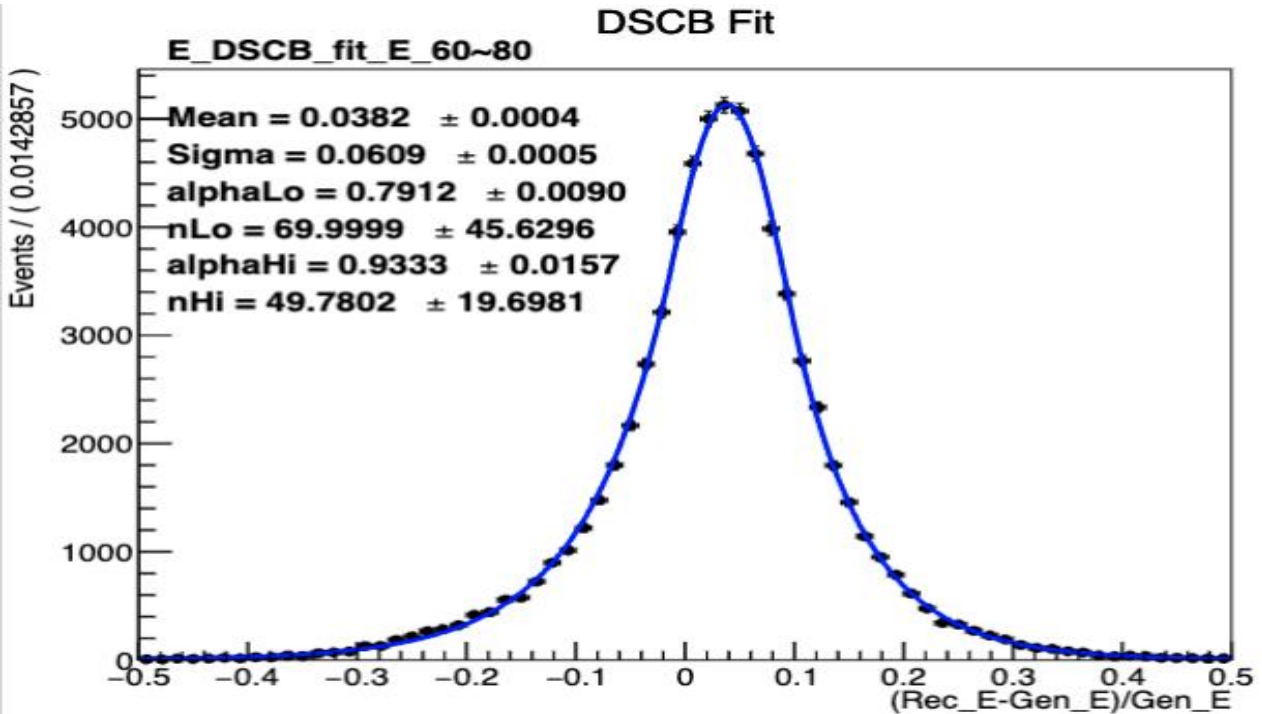
Before: TDR 24.10.0



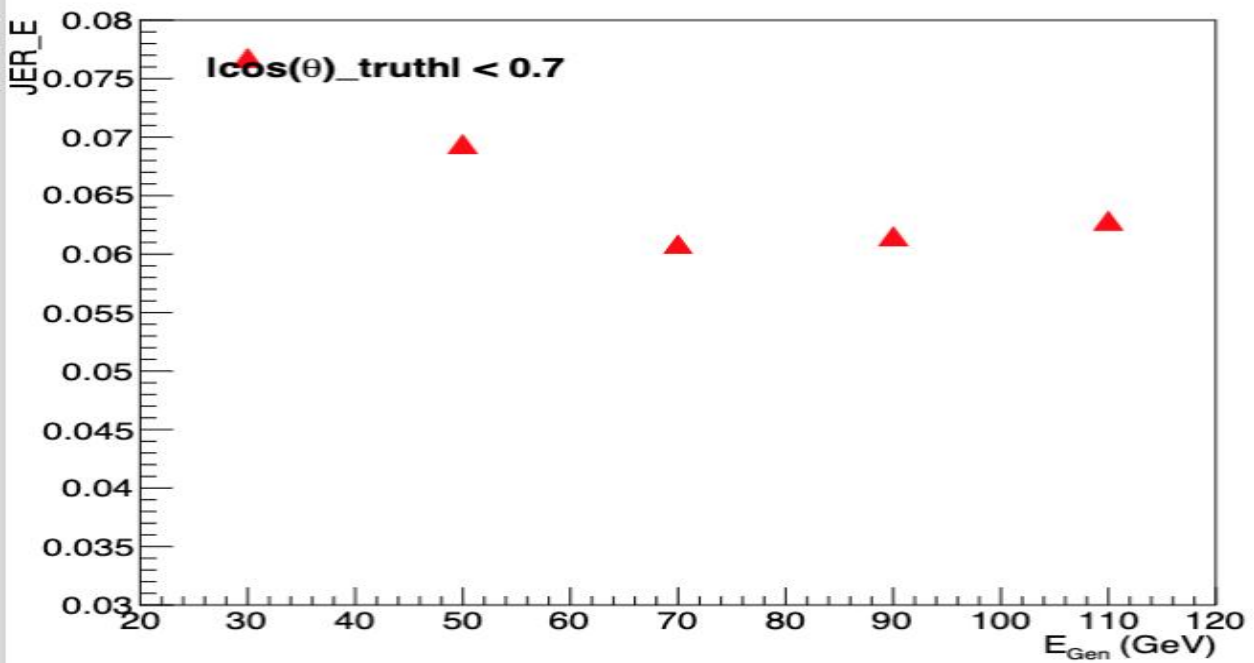
Selection:

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dR_cut = 0.6  
data_barrel_match_rmOverlape_cutdR = data_barrel_match_rmOverlape[(data_barrel_match_rmOverlape["jet1_GENMatch_mindR"] < dR_cut)  
                                                                    & (data_barrel_match_rmOverlape["jet2_GENMatch_mindR"] < dR_cut)]
```

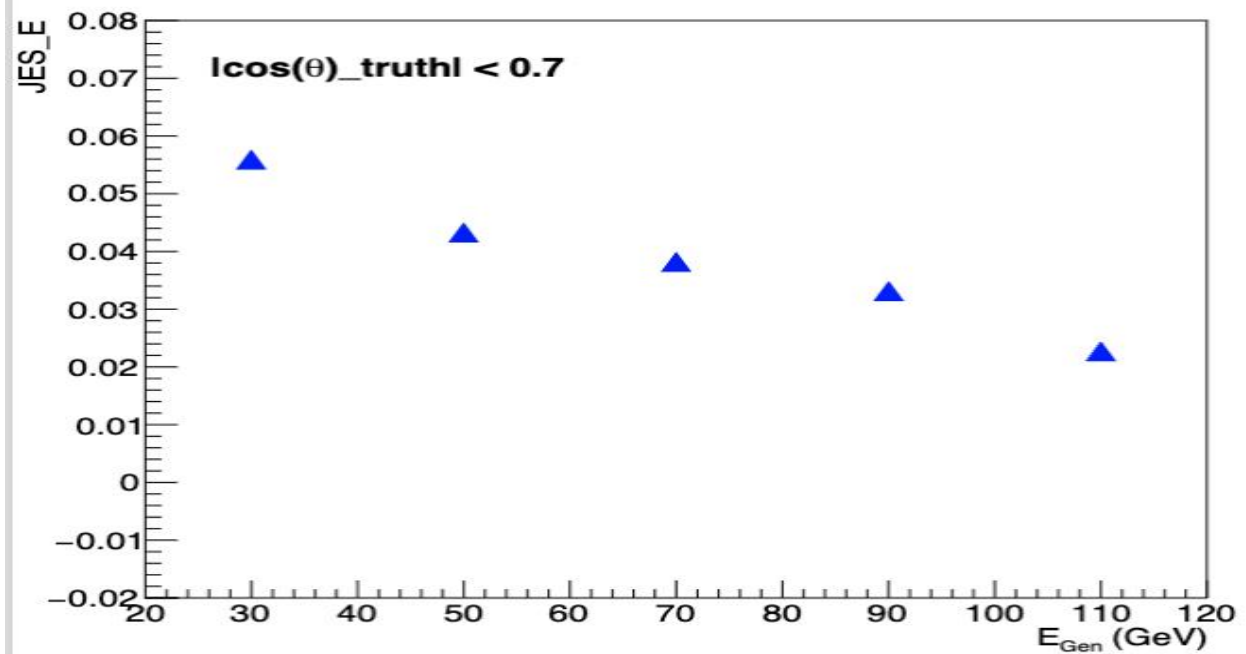


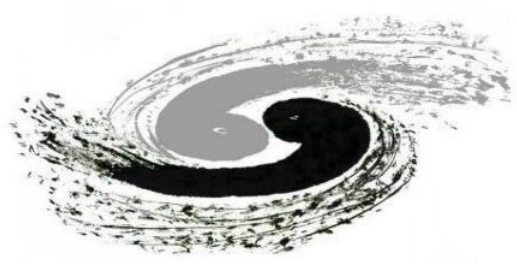


JER



JES





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