

Cut-Based Photon Identification Studies at CMS

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Private Work (CMS Simulati

Introduction

Why are Photons important?

Photons provide a cleaner channel amidst the messy hadronic environment in proton-proton (p-p) collisions at the LHC. Nothing highlights its importance more than the discovery of Higgs decaying into two photons which has a

branching ratio of only 0.002 !!!

Energy Resolution within 1% (3%) in Barrel (Endcap)

Ever since Higgs-discovery, it has been used as a probe in **Beyond** Standard Model (BSM) searches. One such analysis, for instance, looks for the decay of Higgs into a dark photon and an SM photon and uses the cut-based photon identification (ID) for selecting the SM photon[5]. Many other searches which have a photon as one of the final state particles use this ID.

Objective









for a 5×5 cell array in ECAL

is narrower than that of jets

 $\sigma_{i\eta i\eta}$

0.035

ID Variables



Prompt photons have

🔸 Ratio of energy deposited a higher energy deposit in

ECAL relative to HCAL, so





H/E

- + Photons come from two sources: i) prompt photons \rightarrow produced in hard scattering processes (signal) ii) non-prompt photons -> produced abundantly in hadron decays and fragmentation processes (background).
- **4** ID variables discriminate against these two "kinds" of photons. Combination of these variables are taken and the ID values - the value at which the discrimination threshold is kept, are tuned using Genetic Algorithm (GA) for three working points (WP) at 90% (loose), 80% (medium) and 70% (tight) signal efficiency.



Study done using Monte Carlo (MC)

- Isolation
- \downarrow **Σscalar transverse momentum (p_T)** of all detected particles within a hollow cone of ΔR radius around the photon candidate.
- \downarrow Jets will typically have more particles surrounding it and thus, a higher value of Σp_T



samples of y-jet with latest pile-up (PU) conditions i.e. Run 3 conditions (CoM energy 13.6 TeV) for p-p collisions.



y-jet MC sample physics process

4 Two categories of isolation variables:

0.005 0.01 0.015 0.02 0.025 0.03

◆ Particle Flow (PF) Cluster Isolation → calculated based on the energy-deposit clusters in the sub-detectors, viz. Tracker (IsoT), ECAL (IsoE) and HCAL (IsoH)







Pile-up



Pile-up events

- **4** Multiple interactions within the same p-p bunch crossing.
- Increases with luminosity and energy
- **4** Run 3 has an expected pile-up of about 52 interactions
- **4** Quantified by **no. of vertices (nvtx)** and **ρ**, which is the **pile-up transverse momentum** (p_T) density.



Isolation and H/E are PU-corrected

Effective areas (EA) are extracted with **95% contour profile** after separating the samples in seven n-bins

 $X_{\rho-corrected} = X - (\rho^2 \times EA2 + \rho \times EA1)$

EA2, EA1 \rightarrow Coefficients of the fitted polynomial

- **4 p**_T-scales are calculated similarly for IsoE, IsoH, IsoP and IsoN for p_T-correction
- **4** ID-training for **loose, medium and tight** is done. Five variables are used:

 $\sigma_{inin,}$ H/E, and a combination of three variables from PF Cluster and PF Particle isolation

4 Stability of the ID cut-values are checked by plotting the **efficiency** over p_{τ} , η and **nvtx** range for the best performing combination















Genetic Algorithm

- **4** Optimization algorithm which is based on the idea of **'Survival of the Fittest'**
- **4** Each **variable** is analogous to a **gene**, can take any value
- **4** Set of **genes** form **chromosome** of an **individual**, represents **solution**
- **4** Each chromosome has survivability or fitness score a score associated with each solution. Individuals with high fitness score are more likely to be chosen as parents.
- **4** For this problem, **fitness score** is analogous to **background rejection**



The population in the next generation thus, has higher average fitness sco

4 Mutation fosters exploring more optimum solutions.

Gene	_				_			
0.1 0.2	L C	0.0	1.	0.5	0.8	0.1		
Uariable								
0.6 0.1	5 0	3 0.	0.	0.5	0.7	0.9		
Chromosome								
0.8		/ 0.	0.	0.3	0.0	0.5		
.8 0.0 Solution	3 0	3 0.	0.3	0.2	0.7	1.0		
	•							
ו	Population							
	<u>о г</u>		_					
0.5 0.8 0.6 0.1	0.5	5.0	.5	0.				
, Mutation	ore. Mutation							
0.5 0.8 0.6 0.1	0.5	0.2	.5	0		-		

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