Real-time AI-based dead hot map in the ePIC detector: a selfadaptive alternative to traditional big data calibration pipelines

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#### University of Debrecen – 13 faculties

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- Engineering
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- Music
- Pharmacy
- Science and Technology





# Solid background in high-energy physics and Al

- Host of the
  - CMS (CERN) group, 4 FTE
  - EPIC group, 5 FTE
- High computational capacity for AI, NVIDIA DGX (A100, H100, H200) servers
- Official industrial Al-curricula (NVIDIA DLI, MS Learn Partner, CISCO, ...)
- Industrial exam centers













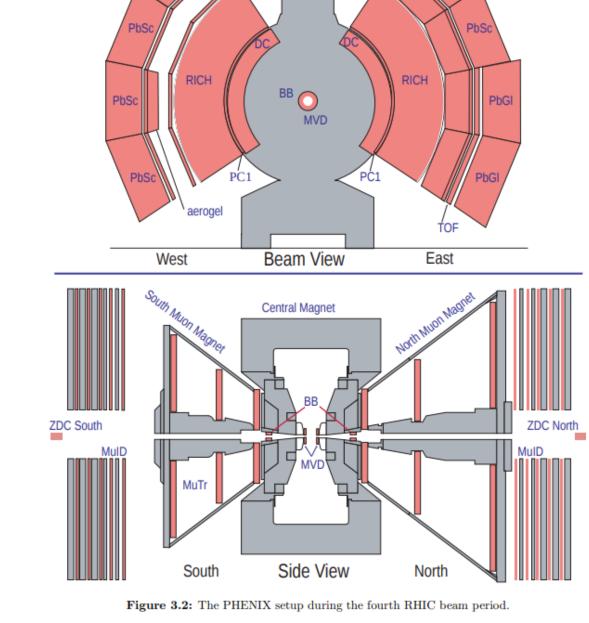






## Background

- PHENIX detector
- direct photon analysis
- DHM@EMCal (25k towels)
- Strong energy dependence
- Patterns from HV issues

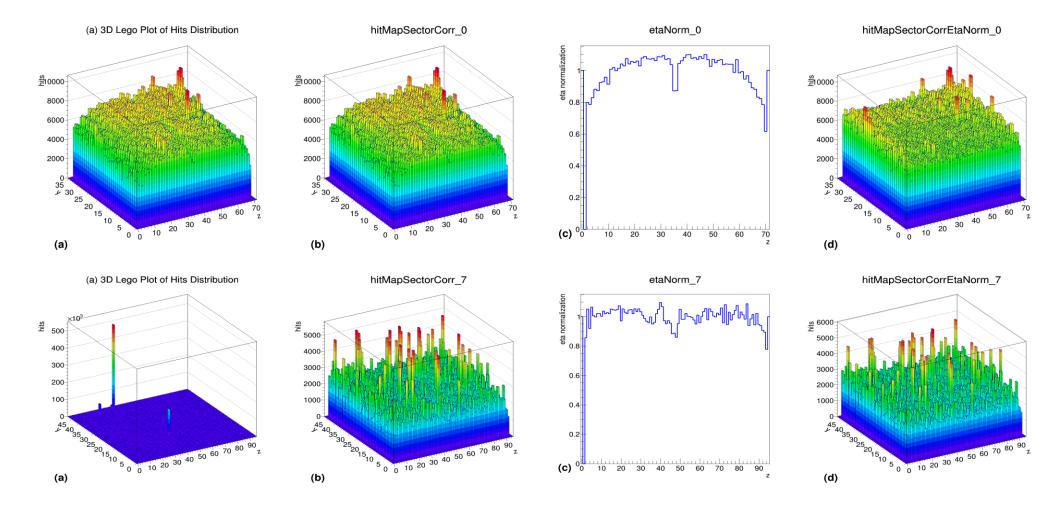


**PHENIX Detector** 

Central Magnet



# Raw maps, eta norm, 8 sectors, 40 energy bins, 1000 runs

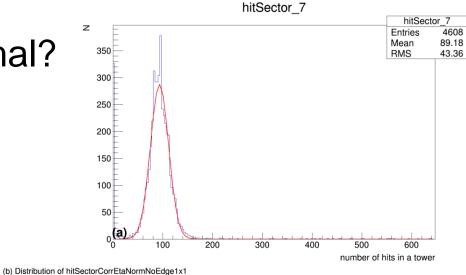


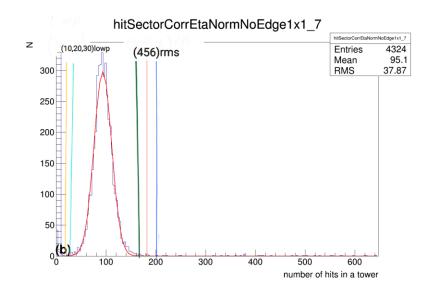
#### Hit distributions

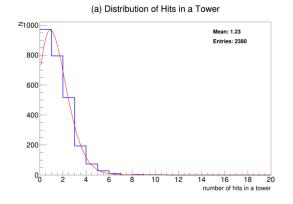
• What is the normal?

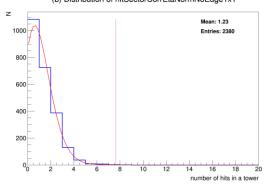
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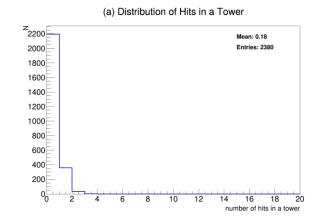
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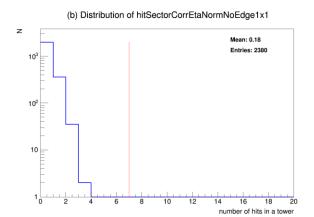






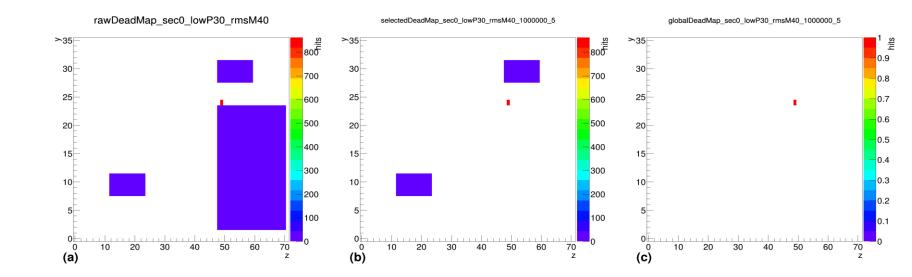






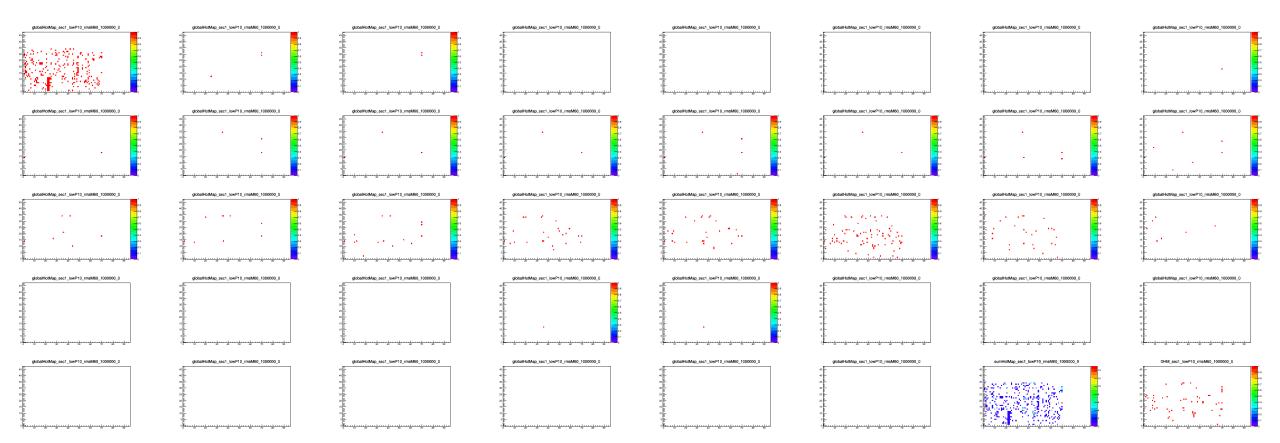
### Run16 level DHM in one energy bin

- 800 good runs (low quality runs was removed from the 1000 runs)
- plot the dead towers from each run
- Keep the stable ones

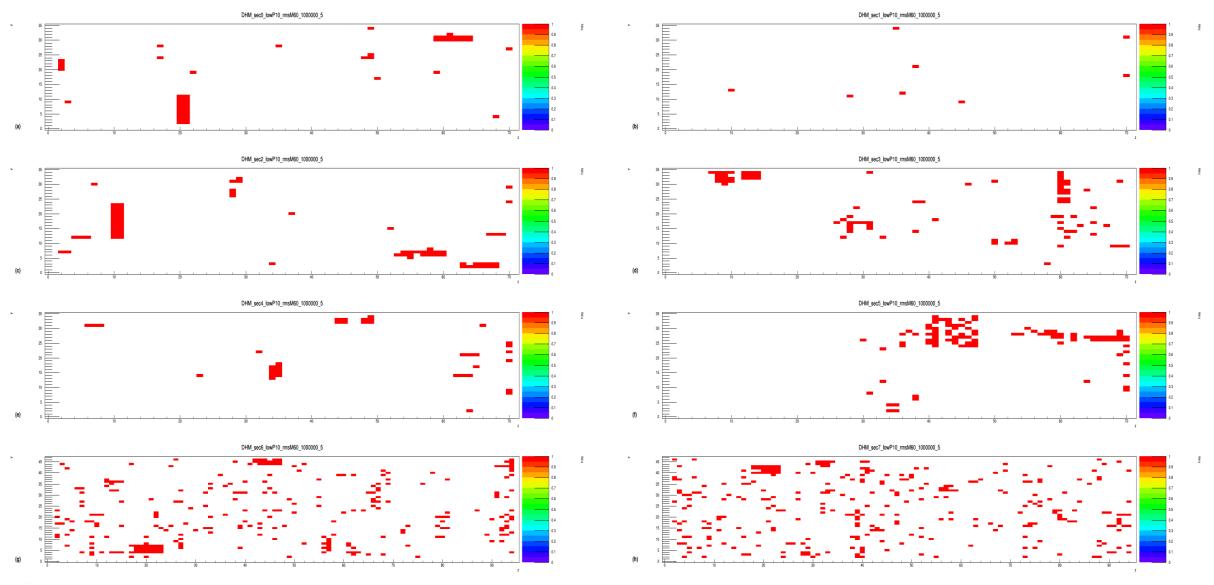


### In every energy bin

Phyisics is back, direct photon at high pT

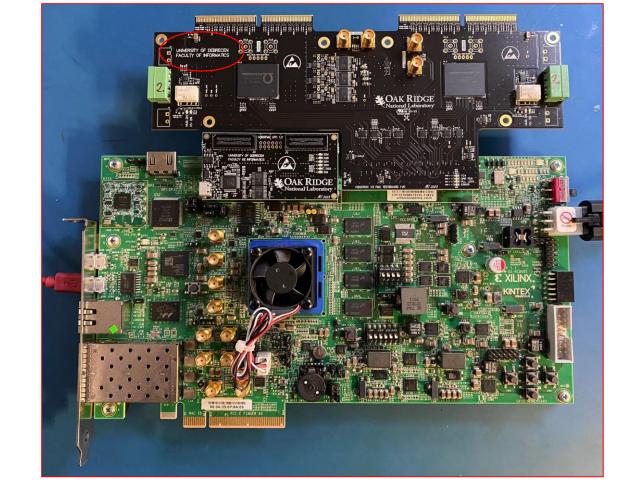


#### Final DHM



#### ePIC IfHCal

- IfHCal front-end development with ORNL
- IfHCal SiPM test
  - Prototype at Yale (100/day), final system
  - (~200k SiPM/years) at Yale/or and ORNL
  - We have to check at least 5%

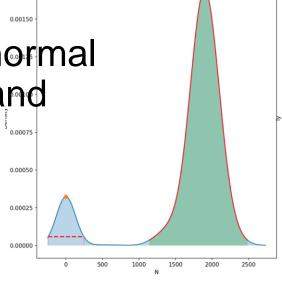




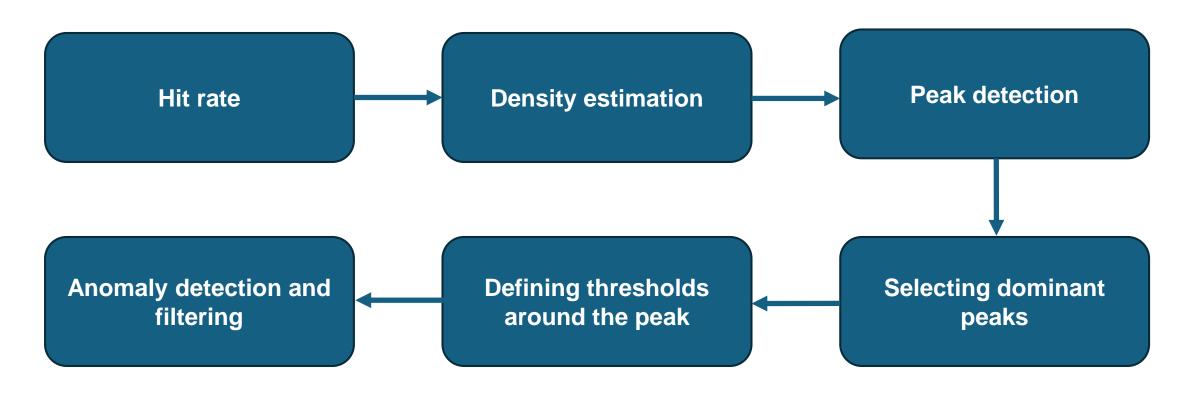
#### Density function

- A probability density function (PDF) is a function that shows how frequently a continuous random variable takes values in different ranges.
- The density function indicates how densely the values are distributed within an interval.
- The peak of the function corresponds to the most likely value, where observations tend to cluster.
- The method analyzes the distribution of hit rate, defines normal ranges based on the dominant peaks of the distribution, and treats any values that deviate from these as anomalies.





#### Description



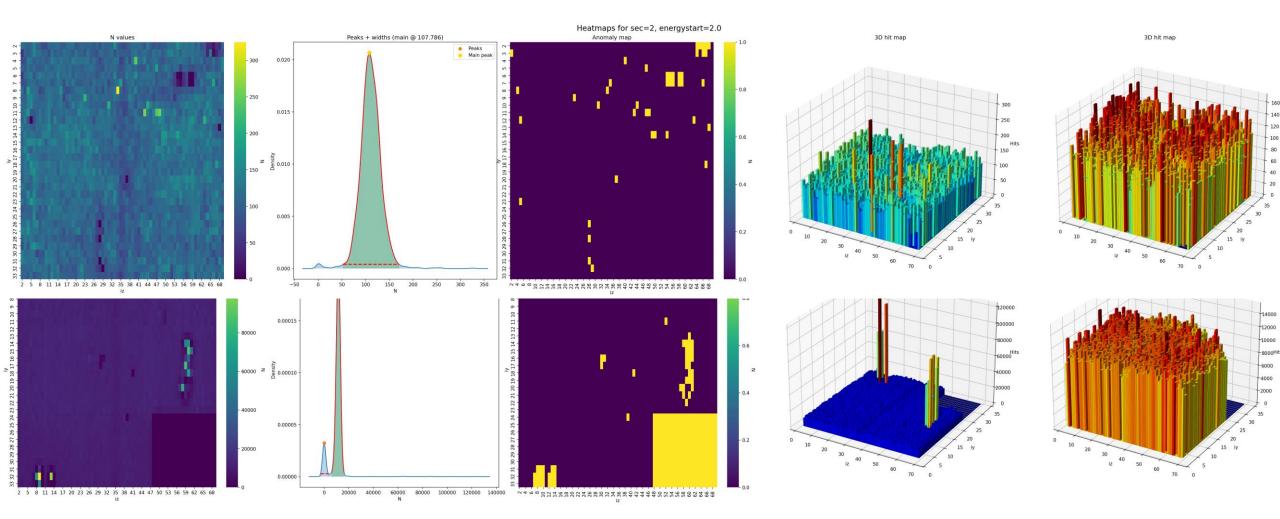


#### Description

- 1. Extracting hit rate
- 2. Density estimation: Examining the distribution of keystroke counts using kernel density estimation (KDE), which reveals how densely values occur along the number line.
- 3. Peak detection in the density function: Identifying local maximum in the density curve, representing frequent and typical value ranges (concentration regions).
- 4. Selecting dominant peaks: Retaining only statistically significant modes. These define the system's normal operating ranges. Currently, only the most dominant peak is used.
- 5. Defining thresholds around the peak: Determining cutoff values by analyzing the drop-off of the density curve. The tails beyond the peak boundaries represent anomaly zones.
- 6. Anomaly detection and filtering



#### Result



Average

calculation

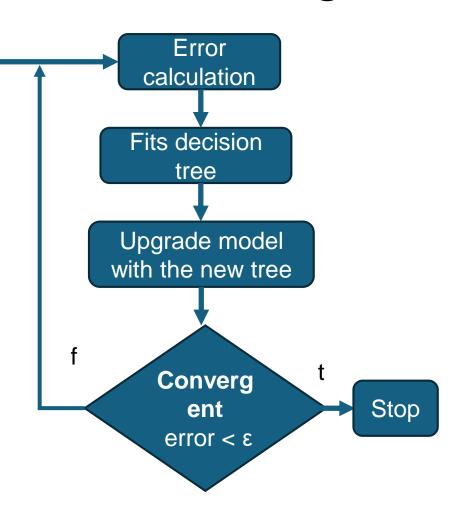
#### **XGBoost**

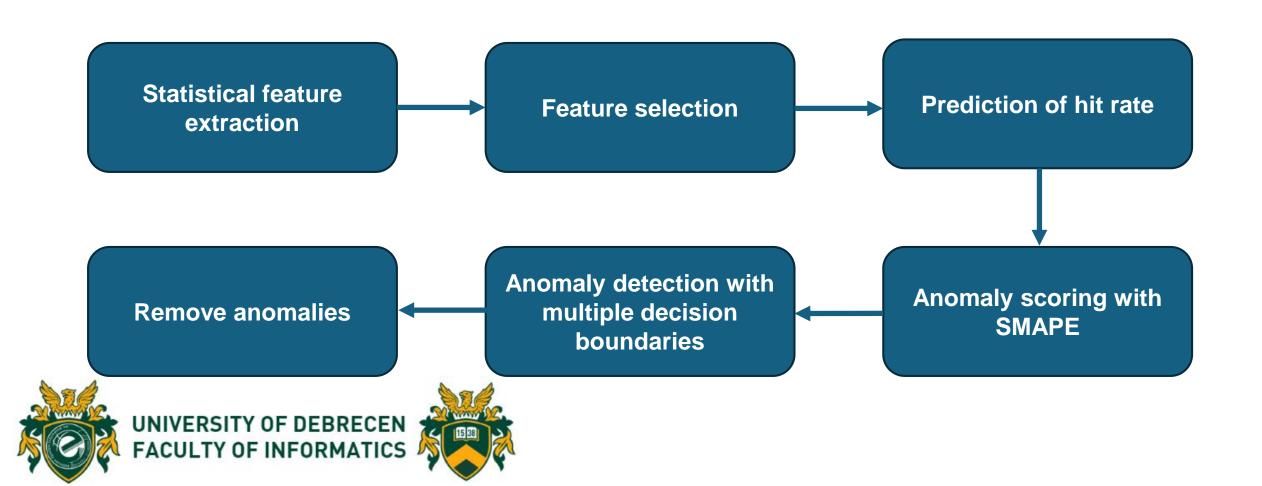
eXtreme Gradient Boosting

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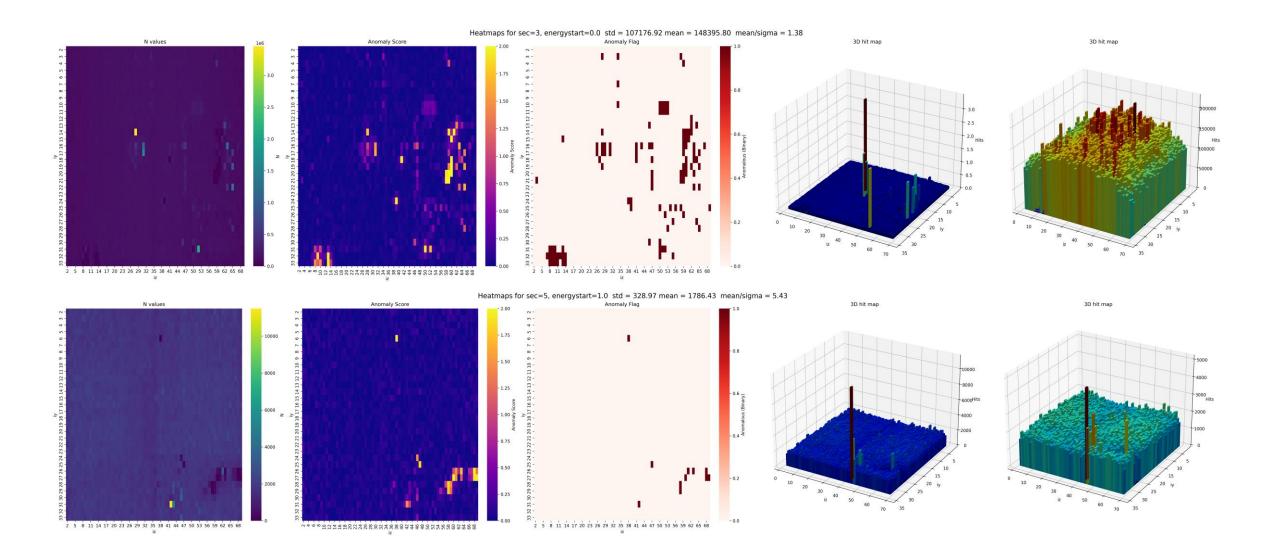
- Boosting is a technique in which many weak models (usually decision trees) are combined to create one strong model.
- Goal: continuously reduce errors through a sequence of learning steps, where each new model corrects the mistakes of the previous ones.





- 1. Statistical feature extraction: Computing descriptive metrics of the data: percentiles (1%, 95%, 99%), median, mean, standard deviation, skewness, kurtosis, entropy, ratio of non-zero values.
- 2. Feature selection: Sector, iy coordinate, iz coordinate, initial energy level, 99th percentile, median, 1st percentile, mean, standard deviation, skewness, kurtosis, entropy, ratio of non-zero values.
- 3. Anomaly scoring using SMAPE (Symmetric Mean Absolute Percentage Error) metric
  Determining the degree of anomaly based on the difference between predicted and actual hit rate using SMAPE.
- 4. Anomaly detection with multiple decision boundaries
  Applying a mean- and standard deviation-based threshold to identify actual anomalies.
- 5. Removing detected anomalies from the data





#### **Preprocessing:**

- Suspicious (anomalous) pixels are marked in the grids according to statistical rules:
  - dead: values lower than a given proportion of the grid average
  - hot: values that exceed the average by several standard deviations
  - extra hot: extremely large outliers
- We replace the marked pixels with the neighborhood average (imputation).

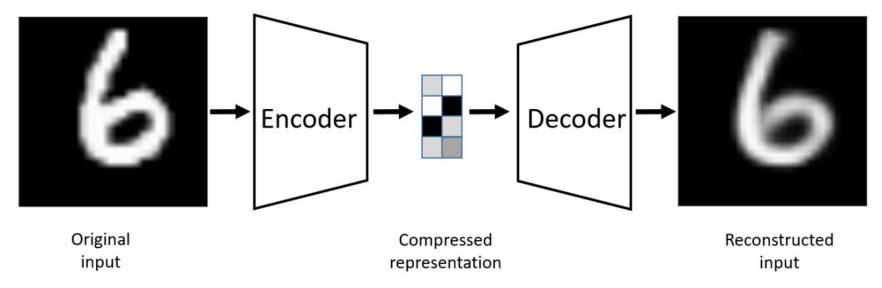
#### **Robust scaling and normalization**

- We normalize the grid reducing the impact of extreme values.
- We standardize the size of different grids (padding).



#### **Convolutional Auto Encoder**

A small convolutional autoencoder:



• Goal: to learn typical, "normal" spatial patterns.



#### The training – Masked Reconstruction + Huber

- We set part of the input (mask ratio) to zero, and the network only tries to reconstruct the omitted pixels.
- Loss function: Huber (Smooth L1) robust to outliers.
- Early stopping: if the validation loss does not improve, we stop training.

#### Loss function (Huber / Smooth L1):

$$L_{\delta}(r) = egin{cases} rac{1}{2}r^2, & ext{if } |r| \leq \delta \ \delta(|r| - rac{1}{2}\delta), & ext{otherwise} \end{cases}$$

$$r = X - \hat{X}$$



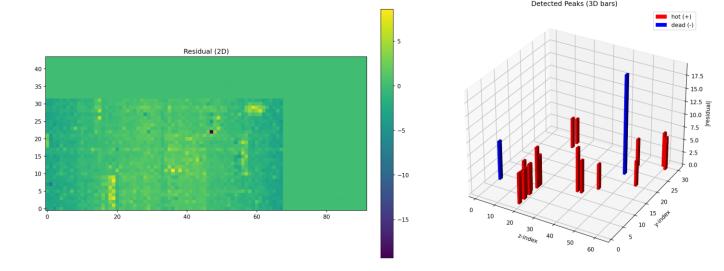
#### Inference - Residual and Thresholding

• The learned model performs reconstruction, the residual:

$$R = X - \hat{X}$$

- Based on the distribution of the residual, we mark anomalies with percentile thresholds:
  - hot: above the upper percentile of the positive range
  - dead: below the upper percentile of the negative range
  - extra hot: extreme cases
- The top 1% of the largest errors is the peak.





## Thank you for your attention.

Questions?