

# Reverse search in Mandelbrot set

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- 2 CNN approach
- 3 Feature Matching approach
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# Mandelbrot set

The Mandelbrot set is a two-dimensional set described by the iteration of a simple function on the complex plane.

Within this set, there are convergent and divergent values, with the former laying inside of the set while the latter outside of it.

$$\text{Mandelbrot set: } f_c(z) = z^2 + c$$

$$\text{Iteration: } f_c(0), f_c(f_c(0)), f_c(f_c(f_c(0)))...$$

# Mandelbrot set

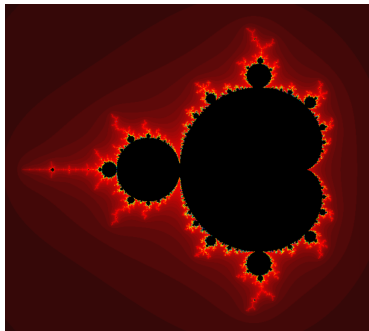


Figure: Mandelbrot set.

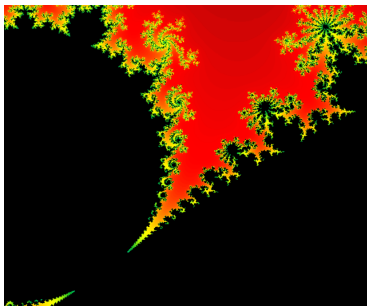


Figure: Mandelbrot fragment and fractal curves.

# Mandelbrot set - Assignment.

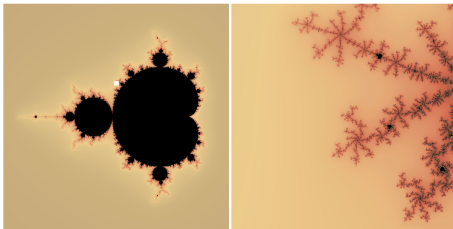


Figure: Assignment example.

- Convolutional Neural Network.
- Featuring matching approach.

# Convolution Neural Network.

In CNN, all neurons share the same weights and biases → All neurons can identify the same features.

CNN can have multiple layers → each layer can focus on a different feature.

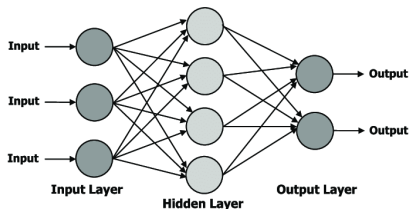


Figure: Typical Neural Network.

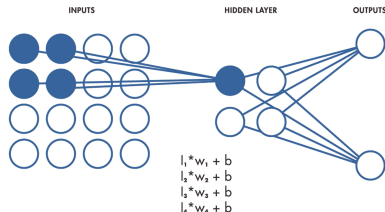
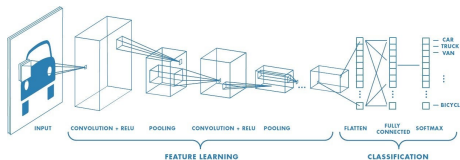


Figure: Convolution neural network.

# Convolution Neural Network.

Activation and pooling are further measures to reduce the learning workload in terms of dimensions and feature identification.



**Figure:** Reduction of dimensionality at CNN with activation and pooling.

## Let's create and train the model

- 1 Create and train the dataset.
- 2 Load and train the model.
- 3 Search and zoom functions.
- 4 Output (GIF)



# CNN - Create Dataset.

- Define Mandelbrot.  $f_c(z) = z^2 + c$
- Generate random sections of it:
  - `x_min = np . random . uniform ( -2 , -0.8 )`
  - `x_max = np . random . uniform ( x_min + 0.1 , x_min + 0.6 )`
  - `y_min = np . random . uniform ( -1.4 , 1.4 - ( x_max - x_min ) )`
  - `y_max = y_min + ( x_max - x_min )`
- Generate 1000 images. (Image size = 480\*480 pixels. Approx. 30min.)

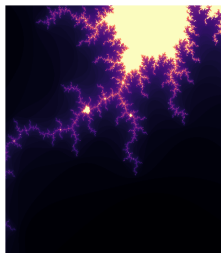


Figure: Example of image generated.

- 1 Load the data.
- 2 Train the model.
- 3 Store the model.

## Characteristics.

- 1 Image size.
  - 64\*64 pixels
  - 480\*480 pixels
- 2 3 layers.
- 3 Activation and pooling.  
(Number of filters and reduction of spatial dimensions by 2).
- 4 Dropouts (10 or 20 %)

## Zoom

- Overlap of 30 images
- Each image is created by Mandelbrot function, closing in the coordinates defined by the model.
- Generate GIF as an output.

## Output

- Zoom functions runs the model.
- Identifies the coordinates and approximates the location.

2 models trained on the same dataset:

- Updated model: load\_data(480pixels) and dropouts.
- Model: load\_data(64pixels) and no dropouts.

	Updated model	Model
Training Mae	0.7521	0.1867
Validation Mae	0.7393	0.3359
Training Loss	0.7624	0.0800
Validation Loss	0.7286	0.1974

[Table](#): Final training and validation metrics.

# CNN - Results (Training Metrics)

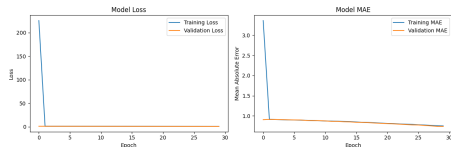


Figure: Training metrics updated model.

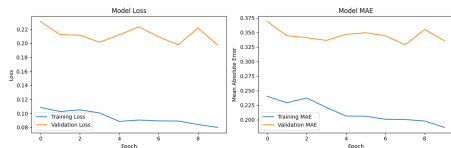


Figure: Training metric original model. (Overfitting)

# CNN - Results (Training Metrics)

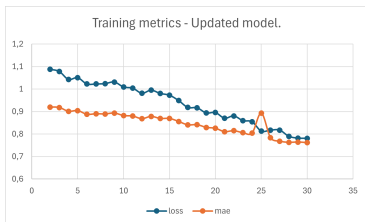


Figure: Training metrics excluding the first odd, epoch.

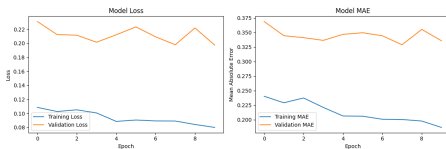


Figure: Training metric original model. (Overfitting).

# CNN - Results

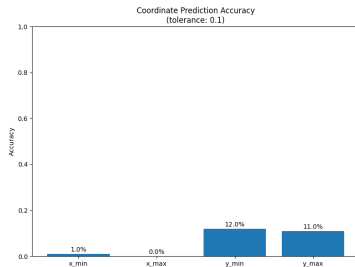


Figure: Updated model.

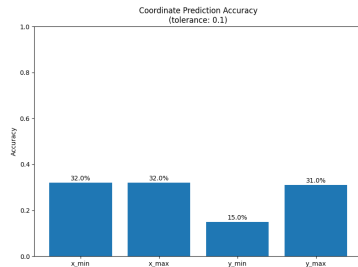


Figure: Model.

# CNN - Example in the three models.

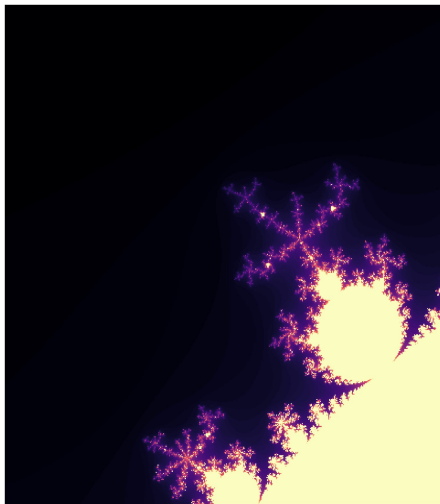


Figure: Fragment used to compared the results in different models.



# Feature Matching Overview

The core idea of this method is to identify locations within the Mandelbrot set by comparing visual features between an input image and a database of reference images.

- Uses SIFT (Scale-Invariant Feature Transform)
- Matches distinctive points between images
- Database of pre-processed reference images

## ① Feature Detection

- Extract SIFT keypoints and descriptors
- Each point has location, scale, orientation

## ② Feature Matching

- FLANN-based matching algorithm
- Fast search through reference database

## ③ Location Prediction

- Best match determines position and zoom
- Coordinates from reference image

SIFT operates in two main steps:

## ① **Keypoint Detection:**

- Identifies distinctive points in the image
- Each keypoint has:
  - Location ( $x, y$ )
  - Scale (size of the region)
  - Orientation (dominant gradient direction)

## ② **Descriptor Computation:**

- For each keypoint, computes a 128-dimensional descriptor
- Descriptor captures gradient information around the keypoint
- These descriptors are what get matched between images

**FLANN** (Fast Library for Approximate Nearest Neighbors).

**Algorithm:** FLANN with kd-tree index

- 1 For each input feature, find  $k=2$  nearest neighbors
- 2 Apply Lowe's ratio test:

$$ratio = \frac{distance_{best\_match}}{distance_{second\_best}} < 0.7$$

- 3 If  $ratio < threshold$  - it's a good match

It is essentially an efficient search algorithm for nearest neighbor search problems, it:

- ① Creates multiple kd-trees to organize points for our SIFT descriptors
- ② For each query point:
  - Searches these trees in parallel
  - Checks a specified number of leaf nodes
  - Returns the closest matches found

- **Database**

- 1000 reference images
- 80-20 train-test split

- **Matching Process**

- SIFT for feature detection
- FLANN with kd-trees for efficient search
- Ratio test for match filtering

# Results

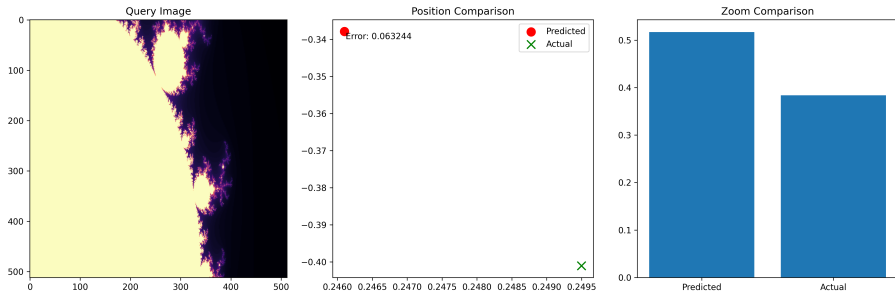


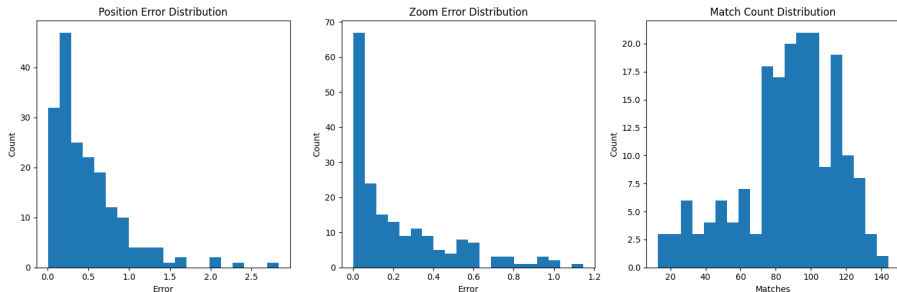
Figure: Example matching result showing position and zoom prediction.

## Test Set Results:

- Total test images: 200
- Successful matches: 186
- Failed matches: 14
- Success rate: 93.0%
- Average position error: 0.494662
- Average zoom error: 0.226827
- Average matches found: 87.7



# Results



**Figure:** Distribution analysis of feature matching performance. On the left we have position error distribution showing histogram of different errors in coordinate prediction, in the center - zoom error distribution showing the accuracy of zoom level predictions, and on the right match count distribution displaying the number of successful SIFT feature matches found per image.

Figure: Feature matching-based zoom animation

# The End