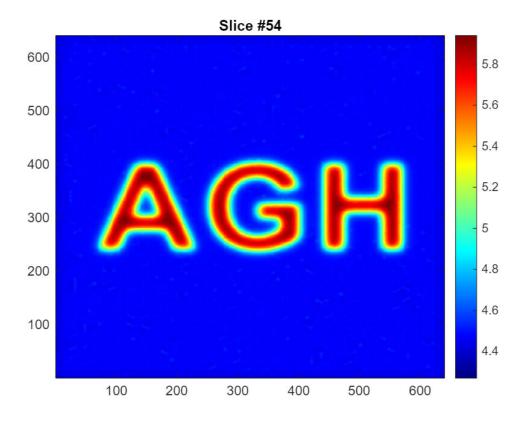
Lab by Mikołaj Suchoń

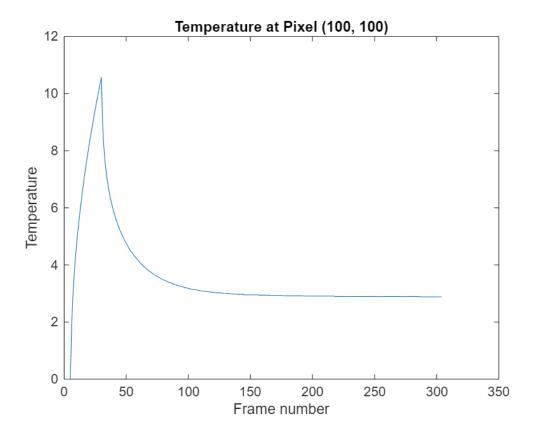
```
clear
load("termogram.mat")
size(data)

ans = 1×3
640 640 304
```

```
slice_index = 54; % Change this from 1 to 304 to view different slices

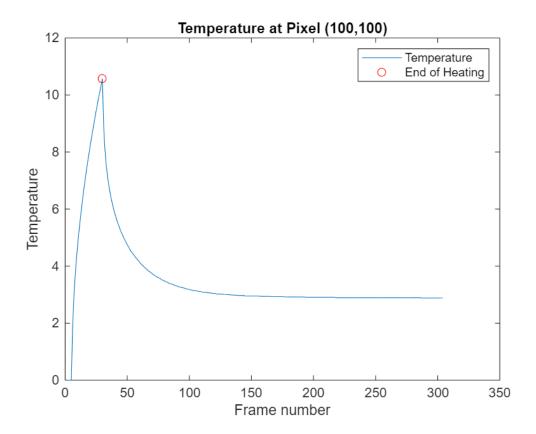
imagesc(data(:, :, slice_index)); % Display the selected 2D slice
colormap('jet'); % Choose your preferred colormap
colorbar; % Add colorbar to indicate data values
title(['Slice #' num2str(slice_index)]);
set(gca, 'YDir', 'normal'); % Set Y axis to normal (not reversed)
```





```
% Find the index of maximum temperature (end of heating)
[max_temp, max_idx] = max(temp_series);

% Optionally, show the result
plot(temp_series);
xlabel('Frame number');
ylabel('Temperature');
title('Temperature at Pixel (100,100)');
hold on;
plot(max_idx, max_temp, 'ro'); % Mark end of heating
legend('Temperature', 'End of Heating');
```



```
% Example: cutoff frame index (e.g., determined from previous max)
frame_cutoff = max_idx; % or assign manually

% Trim the data from frame_cutoff to the end
cooling_data = data(:, :, frame_cutoff:end);

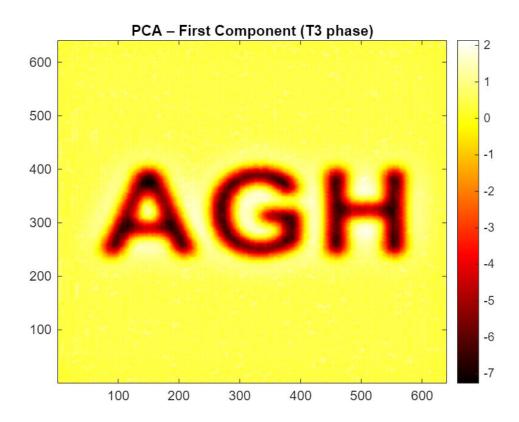
% Get new dimensions
[rows_cool, cols_cool, frames_cool] = size(cooling_data);

% Display results
disp(['Cooling data dimensions: ', num2str(rows_cool), ' x ', num2str(cols_cool), '
x ', num2str(frames_cool)]);
```

Cooling data dimensions: 640 x 640 x 275

```
% 7. PCA on cooling phase (T3)
reshaped = reshape(cooling_data, rows_cool * cols_cool, frames_cool);
% Centering data
mean_signal = mean(reshaped, 2); % mean for each pixel
centered_data = reshaped - mean_signal;
% PCA: columns = frames (variables), rows = pixels (observations)
[coeff, score, ~] = pca(centered_data);
% First PCA component
pca_image = reshape(score(:,1), rows_cool, cols_cool);
%%
```

```
% Displaying PCA
figure;
imagesc(pca_image);
colormap hot; colorbar; set(gca, 'YDir', 'normal'); title('PCA - First Component
(T3 phase)');
```



```
% Assume 'pca_image' is your PCA result (2D matrix)

% Step 1: Normalize to [0,1] for binarization
pca_norm = mat2gray(pca_image); % Normalize values to [0, 1]

% Step 2 (optional): Invert if the feature of interest (e.g., AGH string) is darker
pca_inverted = 1 - pca_norm; % Invert so that the feature becomes brighter

% Step 3: Compute threshold using Otsu's method
threshold = graythresh(pca_inverted); % Automatically finds optimal threshold

% Step 4: Binarize the image
binary_image = imbinarize(pca_inverted, threshold);

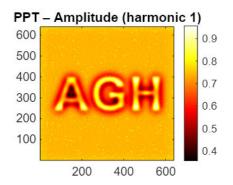
% Step 5: Display the result
imshow(binary_image);
title('Binarized PCA Image');
```

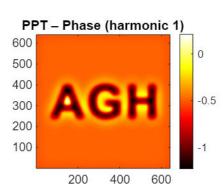


Defect Area = 945.2148 mm²

```
% 10. PPT (Pulse Phase Thermography)
% Define basic variables
L = size(cooling_data,3); % Length of the signal
% FFT
Y = fft(cooling_data,[],3);
```

```
% Single-sided spectrum
P2 = abs(Y/L);
P1 = P2(:,:,1:floor(L/2)+1);
P1(2:end-1) = 2*P1(2:end-1);
% Assign amplitude and phase
Amplitude = P1;
Phase = angle(Y(:,:,1:floor(L/2)+1));
% Choose harmonic for analysis (e.g., 2 = 1st harmonic)
harmonic idx = 2;
Amplitude img = Amplitude(:,:,harmonic idx);
Phase_img = Phase(:,:,harmonic_idx);
% Visualization
figure;
subplot(1,2,1);
imagesc(Amplitude_img); set(gca, 'YDir', 'normal');
colormap hot; colorbar;
axis image; title(['PPT - Amplitude (harmonic ', num2str(harmonic_idx-1), ')']);
subplot(1,2,2);
imagesc(Phase_img); set(gca, 'YDir', 'normal');
colormap hot; colorbar;
axis image; title(['PPT - Phase (harmonic ', num2str(harmonic_idx-1), ')']);
```





```
% Assume Phase_img is your phase image with values in [-pi, pi]
% Step 1: Normalize to [0, 1]
phase_norm = (Phase_img + pi) / (2 * pi); % Shift from [-pi, pi] to [0, 1]
```

Binarized Phase Image (PPT)



```
% Display result
disp(['Defect Area = ' num2str(defect_area_mm2) ' mm²']);
```

Defect Area = 904.5166 mm²