Sections Covered:
- CT Kidney Blob Analysis
- Dice Region Classifiers
- PCA on Breast Cancer Dataset
- Gradient Descent & Optimization
- Cell Label Matching and Distance
- Utility Function Design
Key Results:
1. **Blob Analysis (CT Scans)**:
- Labeled BLOBs after morphological opening and thresholding.
- Correct counting and border filtering reviewed.
2. **ROI Classifier Evaluation (Dice)**:
- Minimum distance classifier based on (B, C, D) found to match segmentation pattern.
- Threshold between D and E computed using:
- Manual Gaussian formula: ~142
- QDA Classifier: ~158.26 Correct Answer.

## 3. \*\*Breast Cancer PCA\*\*:

- Dataset shape: 569 observations, 30 features

- Accuracy of PCA classifier: 0.916

- Samples classified as positive: 349

- PC1 averages: -2.21 (cancer), 3.71 (no cancer)

4. **Gradient Descent**:
- Visualizations used to match iterative path.
- Cost function convergence tested with different initializations.
5. **Cell Landmark Distance**:
- Euclidean distance between average landmark positions computed: ~7.28
6. **Utility Function Suggestions**:
- Gaussian discriminant threshold (manual)
- QDA 1D classifier and threshold finder
- Blob analysis plotter with optional scaling
Tools and Libraries Used:
- scikit-learn (QDA, datasets)
- numpy (PCA via np.linalg.eig)
- skimage (morphology, labeling)
- matplotlib (visualizations)

- PCA scatter plot (color-coded) was used to visually validate clustering.