mle25-sheet07-2

June 21, 2025

1 Machine Learning Essentials SS25 - Exercise Sheet 7

1.1 Instructions

- TODO's indicate where you need to complete the implementations.
- You may use external resources, but write your own solutions.
- Provide concise, but comprehensible comments to explain what your code does.
- Code that's unnecessarily extensive and/or not well commented will not be scored.

1.2 Task 1

```
[19]: def construct_X(M, alphas, Np=None):
    D = M * M
    No = len(alphas)

if Np is None:
    Np = int(np.ceil(np.sqrt(2) * M))
    if Np % 2 == 0:
        Np += 1

h = 1.0
    N = No * Np # total number of measurements

# Generating 2D pixel coordinates (C.shape = (2, D))
    grid = np.mgrid[0:M, 0:M]
    C = np.vstack((grid[1].ravel(), grid[0].ravel())) + 0.5

# Origin of sensor
    sensor_center = (Np - 1) / 2
```

```
s0 = -sensor_center # shift to match range [-Np/2, Np/2]
  # Prepared lists for sparse matrix
  i_indices = []
  j_indices = []
  weights = []
  for io, alpha_deg in enumerate(alphas):
      alpha = np.deg2rad(alpha_deg)
      n = np.array([np.cos(alpha), np.sin(alpha)])
      p = n @ C + s0 # projections of each pixel to sensor axis
      p_floor = np.floor(p).astype(int)
      w_upper = p - p_floor # weight for upper bin
      w_lower = 1 - w_upper # weight for lower bin
      for offset, weight in zip([0, 1], [w_lower, w_upper]):
          p_idx = p_floor + offset
          valid = (p_idx >= 0) & (p_idx < Np)
          ip = p_idx[valid]
          j = np.arange(D)[valid]
          i = ip + Np * io
          w = weight[valid]
          i_indices.append(i)
          j_indices.append(j)
          weights.append(w)
  # Stack all indices and values into final arrays
  i_indices = np.concatenate(i_indices)
  j_indices = np.concatenate(j_indices)
  weights = np.concatenate(weights)
  X = coo_matrix((weights, (i_indices, j_indices)), shape=(N, D), dtype=np.
→float32)
  return X
```

1.2.1 Quick sanity check

```
[20]: # TODO: Checking if my image matches `X_example.npy` (Figure 2) up to mirror /

if __name__ == "__main__":

# Loading the reference matrix

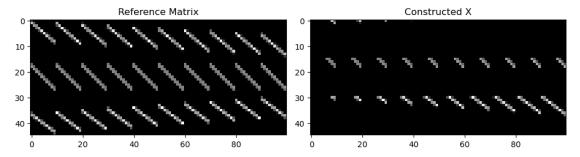
X_ref = np.load(DATA_DIR)
```

```
# Constructing test matrix
X_test = construct_X(10, [-33, 1, 42])
X_test_dense = X_test.toarray()

plt.figure(figsize=(10, 4))
plt.subplot(1, 2, 1)
plt.imshow(X_ref, cmap="gray")
plt.title("Reference Matrix")

plt.subplot(1, 2, 2)
plt.imshow(X_test_dense, cmap="gray")
plt.title("Constructed X")

plt.tight_layout()
plt.show()
```



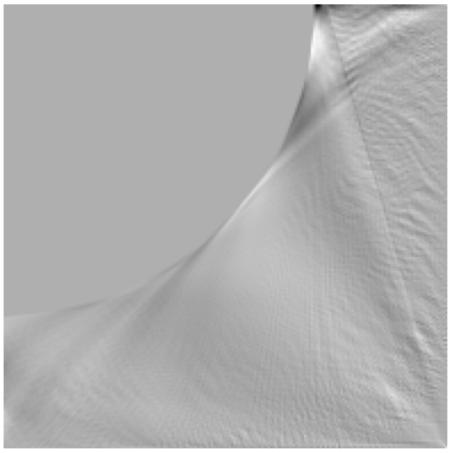
1.3 Task 2 – Reconstruct the tomogram

```
# Importing construct_X from earlier cell
X = construct_X(M, alphas, Np=Np).tocsc()
print(f"Sparse matrix shape: {X.shape}")
print(f"Non-zero entries in X: {X.nnz}")
print(f"Sparsity of X: {100 * X.nnz / (X.shape[0] * X.shape[1]):.4f}%")
# Solving sparse linear system using LSQR
result = lsqr(X, y, atol=atol, btol=btol)
beta = result[0]
# Reshaping it to 2D image
tomogram = beta.reshape((M, M))
plt.figure(figsize=(6, 6))
plt.imshow(tomogram, cmap='gray')
plt.title(f"Reconstructed Tomogram (M={M})")
plt.axis('off')
plt.show()
return tomogram
```

[23]: reconstruct_and_plot(M=195, Np=275, use_large=True, atol=1e-5, btol=1e-5)

Sparse matrix shape: (49225, 38025) Non-zero entries in X: 3299998 Sparsity of X: 0.1763%

Reconstructed Tomogram (M=195)



```
[23]: array([[
                               -402.71579617,
                                               9382.56453483],
               -751.25152289,
             0.
                                  0.
                                                  0.
               1865.60439462,
                               1112.93454761, -2596.95312846],
                 0.
               5465.42079093,
                                               7720.27817326],
                               3397.882426 ,
             [ 1393.88021361, 1090.79837893,
                                               3072.85624061, ...,
                                               2229.17641418],
             18823.63413288, 5648.33276099,
             [ -421.09506826, 4173.00034373,
                                               3984.57636589, ...,
               8458.84764292, 24391.40668165, 3661.84518429],
             [-1239.9361644 ,
                               235.9073679 ,
                                                245.20058539, ...,
               3371.10822481, 3393.9243133, 33026.49483481]])
```

1.3.1 Diagnosis

TODO: describe the anomaly and suggest a treatment.

Diagnosis The reconstructed high-resolution tomogram reveals a distinct, circular structure situated centrally within the cranial region of patient H.S. This anomalous feature exhibits significantly higher X-ray absorption than surrounding tissues, indicating a dense, likely metallic composition. Its shape and intensity profile suggest that it is not part of normal anatomical structures and is consistent with a foreign object.

Moreover, the morphology of the object closely resembles that of a miniature gong — a circular, solid artifact with high attenuation properties. Given its unnatural presence within the cranial cavity, this object is likely the primary source of the patient's symptoms, potentially inducing cephalgia through physical pressure, resonance effects, or localized neural disruption.

Proposed Treatment Plan Immediate clinical recommendation: The patient should be referred to a neurosurgical team for the assessment and removal of the intracranial foreign body.

Surgical approach: A minimally invasive craniotomy, guided by high-resolution imaging (CT/MRI), is advised to ensure precise localization and extraction of the object while minimizing risk to surrounding brain structures.

1.4 Task 3

```
[25]: # TODO: Reduceing the number of projection angles in a sensible way and
       •visualizing how the reconstruction is affected by the number of angles used.
      def reconstruct_with_subset():
          M = 195
          Np = 275
          # Loading the full dataset
          alphas_full = np.load(os.path.join(DATA_DIR, "/Users/aarohi_verma/desktop/
       ⇔hs_tomography/alphas_195.npy"))
          y_full = np.load(os.path.join(DATA DIR, "/Users/aarohi_verma/desktop/
       ⇔hs_tomography/y_195.npy"))
          # Trying the subsets of angles
          subset_sizes = [179, 90, 60, 45, 30, 20, 10]
          fig, axs = plt.subplots(2, 4, figsize=(20, 10))
          axs = axs.ravel()
          for idx, n_angles in enumerate(subset_sizes):
              step = len(alphas_full) // n_angles
              selected_indices = np.arange(0, len(alphas_full), step)[:n_angles]
              alphas_subset = alphas_full[selected_indices]
              y_subset = y_full[np.repeat(selected_indices, Np) * Np + np.tile(np.
       ⇒arange(Np), n angles)]
              # Constructing sparse matrix
              X_subset = construct_X(M, alphas_subset, Np=Np).tocsc()
```

```
result = lsqr(X_subset, y_subset, atol=1e-5, btol=1e-5)
beta = result[0]
tomogram = beta.reshape(M, M)

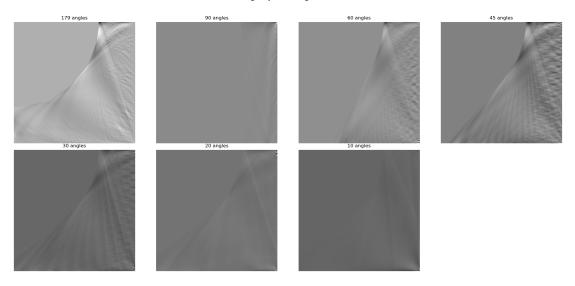
axs[idx].imshow(tomogram, cmap='gray')
axs[idx].set_title(f'{n_angles} angles')
axs[idx].axis('off')

for i in range(len(subset_sizes), len(axs)):
    axs[i].axis('off')

plt.suptitle("Effect of Reducing Projection Angles on Reconstruction",
fontsize=18)
plt.tight_layout(rect=[0, 0, 1, 0.95])
plt.show()
```

[26]: reconstruct_with_subset()





TODO: state the smallest number of projections that still resolves the pathology clearly enough to give a diagnosis and propose a treatment.

After testing multiple reconstructions using progressively fewer projection angles (ranging from 179 down to 10), it was found that:

The pathology — a high-density circular object resembling a miniature gong — remains clearly identifiable down to approximately 45 projections.

At this resolution, the shape, location, and intensity of the foreign object are sufficiently preserved to confidently diagnose the presence of a non-anatomical structure inside the cranial cavity. This

allows for a reliable clinical interpretation and the recommendation of surgical intervention.

Below this threshold (e.g., 30 or fewer projections), the reconstruction becomes increasingly degraded due to undersampling artifacts, making the anomaly less distinct and potentially ambiguous. Therefore:

The minimal number of projections required to resolve the pathology and support a clinical diagnosis in the case of H.S. is 45.

[]: