

Theoretical exercises:

T1. Study the following parts of the textbook:

- Section 3.1: the three conditions of Hadamard for a well-posed problem.
- Figure 3.3: the subspaces related to a matrix understood as a linear map.
- Section 4.1: the Moore-Penrose pseudoinverse.

Consider the measurement model $\mathbf{m} = \mathbf{A}\mathbf{f} + \varepsilon$ with A a $k \times n$ matrix. The inverse problem “given \mathbf{m} , find \mathbf{f} ” can be ill-posed as any of the three conditions of Hadamard may fail.

- (a) Explain how the Moore-Penrose pseudoinverse takes care of ill-posedness related to Hadamard’s conditions on existence and uniqueness.
- (b) Why does the Moore-Penrose fail with the stability condition of Hadamard?

Matlab exercises:

M1. **Stacked-form Tikhonov regularization** for the 1D deconvolution problem. Follow the procedure of Problem M2 of Exercise 2. Take a suitable $k = 128$ and simulate discrete convolution data $\tilde{\mathbf{m}}$ (with a little noise added) using the simulated continuum model. Furthermore, take $n = k$ and let A be the square-shaped measurement matrix from the computational model.

- (a) Choose $\alpha = 1$ and consider the matrix equation $A'\mathbf{f} = \tilde{\mathbf{m}}'$, where

$$A' = \begin{bmatrix} A \\ \sqrt{\alpha}I \end{bmatrix}, \quad \tilde{\mathbf{m}}' = \begin{bmatrix} \tilde{\mathbf{m}} \\ \mathbf{0} \end{bmatrix}, \quad (1)$$

and I denotes the $n \times n$ identity matrix and $\mathbf{0}$ is a vertical n -vector with all components equal to zero. Compute reconstruction \mathbf{f}_α in Matlab using the backslash operator:

$$\mathbf{f}_\alpha = A' \setminus \tilde{\mathbf{m}}'. \quad (2)$$

Plot original signal and \mathbf{f}_α in the same picture and compare.

- (b) Let the regularization parameter $\alpha > 0$ range over many values (for example $\dots, 10^{-6}, 10^{-5}, 10^{-4}, 10^{-3}, 10^{-2}, 10^{-1}, 10^0, 10^1, 10^2, \dots$). For each α , calculate the relative square norm error between the reconstruction and the original signal. Which value of α gives the smallest error? What seems to be the limit of the reconstruction as $\alpha \rightarrow \infty$?
- (c) Repeat (b) using a higher noise level. Is the optimal regularization parameter α smaller or greater than in (b)?

M2. Download *tomo01_RadonMatrix_comp.m* and *tomo02_firstTSVD_comp.m* from the course website. Let the image resolution be 32×32 and choose the number of angles in tomographic measurement to be 63. Construct the 32×32 “true object” \mathbf{f} using the routine *SqPhantom.m*.

- (a) Compute the sinogram using the `radon` command of Matlab. Then compute the “inverse crime” data $\mathbf{m} = A\mathbf{f}$. Test computationally that the sinogram and data are approximately equal. (Note: there are several reshaping operations involved.)
- (b) Plot some of the singular vectors, similarly to Figure 4.4 of the textbook.
- (c) Find numerically the optimal truncation index for TSVD for the 2D by comparing the reconstruction to ground truth and calculating relative square norm error. (Note: you do not need to compute the TSVD reconstruction for *all* possible truncation indices. Start from a small number and increase until the reconstructions become really bad.)
- (d) Repeat (b) for a higher noise level. Is the optimal truncation index smaller or larger than in (b)?

M3. Repeat M2 using only 13 projection angles. What is different in (b)-(d)?

M4. Simulate 2D tomographic data using higher resolution and a matrix-free formulation, thereby avoiding inverse crime. Here “matrix-free” means that you can just use the `radon` command for computing the higher-resolution sinogram; the matrix A is needed only for the low resolution.

You can use the routines on the page

<http://wiki.helsinki.fi/display/mathstatHenkilokunta/X-ray+tomography+with+matrices>

See the Matlab files there under title *Creating data without inverse crime*.

- (a) Construct the 32×32 “true object” \mathbf{f} using the routine *SqPhantom.m*. Let the number of angles in tomographic measurement to be 63. Compute two sinograms: first simply with the command `radon`, and second with the “inverse crime-free” method using the resolution 64×64 , applying `radon`, and downsampling. (This is what’s done in the routine *XRMC_NoCrimeData_comp.m* available on the aforementioned web-page.) What is the relative square norm error between the two sinograms? Plot the two sinograms and their difference.
- (b) Repeat (a) with the resolutions 128×128 and 256×256 . What is the relative square norm error between the two sinograms? Plot the two sinograms and their difference.