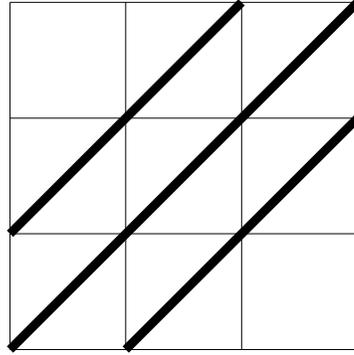
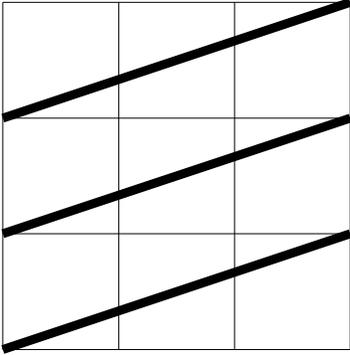


T3. Thin lines depict pixels and thick lines X-rays in this image:



Give a numbering to the nine pixels ($f \in \mathbb{R}^9$) and to the six X-rays ($m \in \mathbb{R}^6$), and construct the matrix A for the measurement model $m = Af$. The length of the side of a pixel is one.

You can work on these Matlab exercises (marked with M) in the exercise session.

- M1. Run the file `DC1_cont_data_comp.m`. Set the number of discretization points to 100 by writing `n = 100;` on line 12 in the file `DC2_discretedata_comp.m`. Then run the file `DC2_discretedata_comp.m` to produce discrete convolution data.

Compute the singular value decomposition of the 100×100 measurement matrix A using the code in the file `DC4_truncSVD_comp.m`. Determine the *condition number* of A , defined as the first (largest) singular value divided by the last (smallest) singular value.

Repeat the above with resolutions $n = 200, 300, 400$. What happens to the condition number when n grows?

- M2. Download from the course website all the Matlab routines whose name starts with `XRM`.

- Run in Matlab the routines `XRM1_matrix_comp.m`, `XRM2_naive_comp.m`, `XRM3_NoCrimeData_comp.m`, `XRM4_naive_comp.m`, `XRM5_SVD_comp.m` and `XRM6_truncSVD_comp.m`. Choose $N = 32$ in all of these files, leading to reconstruction size 32×32 .
- Compute the condition number $\text{cond}(A) := d_1/d_{\min\{k,n\}}$ of the measurement matrix A . You get the singular values from `XRM5_SVD_comp.m`.
- Record the minimum relative reconstruction error you see when running `XRM6_truncSVD_comp.m`.

- M3. Computational study of limited angle tomography with 90° view angle.

- In routine `XRM1_matrix_comp.m`, the projection directions are evenly distributed in the interval $[0, 180^\circ)$, excluding the value 180° . Modify line 23 so that the projection directions are evenly distributed in the interval $[0, 90^\circ)$. Run `XRM1_matrix_comp.m`.
- Run `XRM2_naive_comp.m`, `XRM3_NoCrimeData_comp.m`, `XRM4_naive_comp.m` and `XRM5_SVD_comp.m`. Compute $\text{cond}(A)$ and compare the result to that of M2(b).
- Run `XRM6_truncSVD_comp.m` and record the minimum relative reconstruction error. Compare the result to that of M2(c). How do the reconstructions and singular vectors differ from those you saw in M2(c)?