

Finite Radon Transform

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Abstract

This paper provides a method for computing the forward Radon transform and the inverse Radon transform in the discrete case. Because the forward transform is casted as a linear algebra operation, the inverse transform can be broken down into a series of faster operations which make use of the fourier and inverse fourier transforms. Using the Cooley-Tukey algorithm, these fourier transforms are improved for efficiency, dramatically reducing the number of operations for bodies with large values of N , as the order of operations is of order $N \log N$, instead of N^2 .

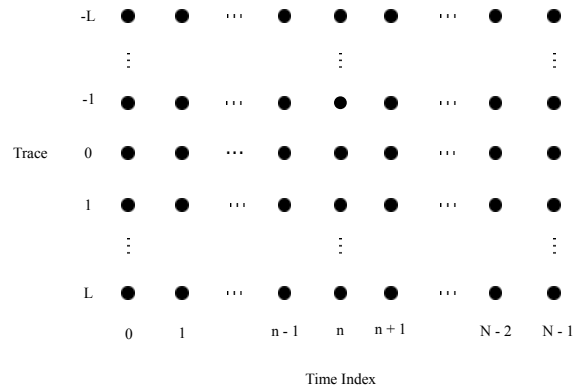
In addition, this paper also references a program written in Python that computes the forward and inverse Radon Transforms, as well as the transform R_m matrices associated with these transforms.

This paper exclusively addresses the discrete case, and outlines an algorithm written in Python for calculating the forward and inverse transforms of a given image.

1 Discrete Radon Transform

For simplicity, the Discrete Radon Transform will be abbreviated as DRT for the remainder of this paper.

Below is a lattice representation of the original body:



Let $x_l(n)$ be a 2-dimensional array, where $-L < l < L$ and $0 \leq n \leq (N - 1)$. The quantity $(2L + 1)$ gives the total number of traces in the body, which can be thought of as the thickness for simplicity. For a fixed time point n , $x(n)$ is denoted by the following vector:

$$x(n) = \begin{bmatrix} x_{-L}(n) \\ \dots \\ x_0(n) \\ \dots \\ x_L(n) \end{bmatrix}$$

We assume $x(n)$ is defined $\forall n \in \mathbb{Z}$, and that $x(n)$ is a periodic vector sequence with period N , so $x(n) = x(N + n)$. Stacking the column vectors in a row from $x(0)$ to $x(N - 1)$ produces a matrix with a one-to-one correspondence with the original lattice of the body, with the entry $x_{l,n}$ corresponding to the lattice point at (l, n) . The full matrix x is given as follows:

$$x = \begin{bmatrix} | & | & \dots & | \\ x(0) & x(1) & \dots & x(N - 1) \\ | & | & \dots & | \end{bmatrix}$$

Thus, to recover the original structure of the body, the inversion procedure attempts to recover these $x(n)$ vectors. This procedure is nontrivial because $x(n)$ cannot be directly determined from the body in question. Instead, scans of the body, whose values can be calculated, are used to recover the values in the $x(n)$ vectors, thus determining the physical makeup of the body in question. These scans are taken by (finish this section)

We define the forward Radon transform for x as follows:

$$y(n) = \sum_{m=-M}^{m=M} R_m \cdot x(n + m) \tag{1}$$

where $2M + 1 \leq N$ and R_m are $(2J + 1 + J') \times (2L + 1)$ transform matrices. J gives the maximum slope of scans used in the inversion procedure, J' gives the number of row sums used, and, for a fixed n , M gives the number of adjacent vectors to $x(n)$ used to compute the output vector, $y(n)$. For a fixed time point n , $y(n)$ is denoted by the following vector:

$$y(n) = \begin{bmatrix} y_{-J}(n) \\ \dots \\ y_0(n) \\ \dots \\ y_{J+J'}(n) \end{bmatrix}$$

where J is the magnitude of the maximum slope involved in the scan, and J' is the number of row sums taken in the scan. In the accompanying Python program, $J' = (2L + 1)$, which

accounts for scanning each row in the original body. If $J' = (2L + 1)$, then a $(2L + 1)$ identity matrix can be appended to the bottom of each R_m matrix.

Thus, y is a $(2J + 1 + J') \times N$ matrix, and in the special case involving all row scans, y is a $2(J + L + 1) \times N$ matrix.

2 Inverse Radon Transform

The DRT is inverted by treating the inversion process as a linear algebra problem. Consider the adjoint matrices R^* , defined as $R_i^* = \text{adj}(R_i)$ for $i = -J, \dots, J + J'$. The transform given by R^* is as follows:

$$z(n) = \sum_{m=-M}^{m=M} R_{-m}^* \cdot y(n + m) \quad (2)$$

We now introduce the matrix $\mathbf{H} = \mathbf{R}^* \cdot \mathbf{R}$. The transform associated with H mimics the operations of the DRT and adjoint transform. The transform is:

$$z(n) = \sum_{m=-2M}^{m=2M} H_m \cdot x(n + m) \quad (3)$$

where the H_m matrices are defined as follows:

$$H_m = \sum_{m'=-M}^{m=M-m} R_{m'}^* \cdot R_{m'+m} \quad (4)$$

and

$$H_{-m} = H_m^* \quad (5)$$

for $m = 0, \dots, 2M$.

Now, we introduce a key lemma that includes the Discrete Fourier Transforms (DFTs) of H and Z .

Lemma 1 If $z(n)$ is as defined in equation (3), and x is a periodic vector sequence with period N , then,

$$\hat{z}(k) = \hat{H}(k) \cdot \hat{x}(k) \quad (6)$$

for $k = 0, 1, \dots, N - 1$. The DFTs of z , x , and H are defined as follows:

$$\hat{z}(k) = \sum_{n=0}^{n=N-1} z(n) e^{2\pi i(nk/N)} \quad (7)$$

$$\hat{x}(k) = \sum_{n=0}^{n=N-1} x(n) e^{2\pi i(nk/N)} \quad (8)$$

$$\hat{H}(k) = \sum_{m=-2M}^{m=2M} H_m e^{-2\pi i(mk/N)} \quad (9)$$

Note: the difference in sign on the H is attributed to a physical property in seismology.

Proof (Beylkin) Given equation (3), we apply the DFT to both sides to get

$$\hat{z}(k) = \sum_{m=-2M}^{m=2M} H_m \sum_{n=0}^{n=N-1} z(n) e^{2\pi i(nk/N)}$$

or

$$\hat{z}(k) = \sum_{m=-2M}^{m=2M} H_m e^{-2\pi i(mk/N)} \sum_{\tilde{n}=m}^{\tilde{n}=N+m-1} z(\tilde{n}) e^{2\pi i(\tilde{n}k/N)}$$

The assumption that $x(n)$ is periodic with period N implies that

$$\sum_{\tilde{n}=0}^{\tilde{n}=m-1} z(\tilde{n}) e^{2\pi i(\tilde{n}k/N)} = \sum_{\tilde{n}=N}^{\tilde{n}=N+m-1} z(\tilde{n}) e^{2\pi i(\tilde{n}k/N)}$$

for $m \geq 1$. A similar identity holds when $m \leq -1$ by changing N on the right summation to $-N$, and thus

$$\hat{z}(k) = \sum_{m=-2M}^{m=2M} H_m e^{-2\pi i(mk/N)} \sum_{n=0}^{n=N-1} x(n) e^{2\pi i(nk/N)}$$

which is exactly Lemma 1 in equation (6).

Because $x(n)$ is a real vector sequence, DFT of x contains complex conjugation. Namely, $\hat{x}(N-k) = \bar{\hat{x}}(k)$, for $k = 0, 1, \dots, N-1$. This means only values of k in the range $[0, N/2]$ if N is even, $[0, (N-1)/2]$ if N is odd, must be considered. This dramatically cuts down the number of computations, especially for large values of N . This begins the discussion of invertible frequencies, k .

Definition The DRT given in equation (1) is uniquely invertible within the normalized frequency band $[k_{min}/N, k_{max}/N]$ for $0 \leq k_{min}/N$ and $k_{max}/N \leq .5$ if $\forall k = k_{min}, \dots, k_{max}$

$$\det(\hat{H}(k)) \neq 0 \quad (10)$$

In summary, the algorithm for inverting the DRT, y , of a given body, x , given transform matrices R is

1. Compute the R^* adjoint transform matrices of $R \forall m$

2. Find the adjoint transform of y using the formula

$$z(n) = \sum_{m=-M}^{m=M} R_{-m}^* \cdot y(n+m)$$

3. Compute the $(2L+1) \times (2L+1)$ H_m matrices from the R and R^* matrices for $m = -2M, \dots, 2M$

4. Compute the DFTs of z and H , \hat{z} and \hat{H}

5. Compute $\hat{H}^{-1}(k) \cdot \hat{z}(k) = \hat{x}(k)$ for invertible values of k

6. Compute the IDFT of $\hat{x}(k)$ for $k = 0, 1, \dots, (N-1)$ to recover the original image, x

3 Python Code

3.1 R_m Transform Matrix Maker

This first function, `Rm`, in this program takes two parameters, J and L , and $J, L \in \mathbb{Z}$. L is related to the number of distinct traces of the body, as $2L+1$ gives the total number of traces. J gives the bound for the maximum and minimum magnitude of the slopes, for a total of $2J+1$ slopes ranging from $-J$ to J . With these definitions, $M = J \times L$, where M is the number of adjacent vectors used on each side of the given vector for the set of scans. Thus, $2M+1$ gives the total number of vectors used for each set of scans, and it follows that there are $2M+1$ transform matrices, indexed as follows: $R_m, m = -M, \dots, 0, \dots, M$. These matrices are stored as a list of arrays, seen in the code as "R", with `R[0] = R-M`, ..., `R[M] = R0`, ..., `R[2M] = RM`. This "R" list is the returned product of the `Rm` function.

The `Rm` function implements the following formula

$$(R_m)_{j,l} = \delta_{m,jl} \tag{11}$$

where the second subscript of δ is the product of j and l . The R_m matrices have special indexing for rows and columns: rows are indexed top to bottom from $-J$ to J , columns are indexed left to right from $-L$ to L . The equation determining δ is

$$\delta_{m,jl} = \begin{cases} 1 & \text{if } m = j \times l \\ 0 & \text{otherwise} \end{cases} \tag{12}$$

Equation (12) is employed for each R_m matrix, $m = -M, \dots, M$. Once all entries are filled, the `Rm` function then returns the transform matrices, properly indexed as a list of arrays.

The program also contains a second function for outputting transform R_m matrices, called `Rmrows`. It takes the same slope and index parameters as the first `Rm` function, and its first step is identical to `Rm`'s. The key difference in `Rmrows` comes next, in the addition of row scans. This function assumes all row scans are taken, so each R_m matrix will have $(2J + 1) + (2L + 1)$ rows, but the same $(2L + 1)$ number of columns as before. These row scans are accounted for by simply appending a $(2L + 1) \times (2L + 1)$ identity matrix to the bottom of each R_m matrix. `Rmrows` returns a properly indexed list of $2(L+J+1) \times (2L+1)$ arrays.

3.2 Discrete Radon Transform

This program contains one function, `firt`, which takes two parameters, a and R . The a is the body to be transformed, and R is a list of arrays containing the transform matrices. For the function to work, the number of columns in the input array a must equal or exceed the number of transform matrices, staying consistent with the condition that $(2M + 1) \leq N$. After this check, the program uses equation (1) to calculate the transformed body, $y(n)$ for $n = 0, 1, \dots, N - 1$. Once complete, the function returns y .

3.3 Inverse Radon Transform

This program contains one function, `ifrt`, which takes two parameters: the transformed body y , and a list of arrays R of the transform matrices used in the operation.

First, the function computes the adjoint R^* matrices, and stores them in an array called "Rstar". Once computed, the function then uses equation (2) to calculate the adjoint transform of y , and stores it as z . Next, equation (4) is used to calculate H_m for $m = 0, \dots, 2M$, then H_m for $m = -2M, \dots, -1$ are calculated by equation (5).

The next step computes the DFTs of z and H . \hat{z} is calculated by the fast fourier transform in the numpy library, which uses the Tukey-Cooley algorithm to reduce the total number of computations. In the current version of the program, \hat{H} is calculated by means of a brute force fourier transform, as the numpy algorithm produced the incorrect output for the 3 dimensional array, H . In addition, though only values of $k = 0, \dots, N/2$ should be needed to successfully complete the transform, incorrect results were obtained when using this limited band of frequencies. As a consequence, the less efficient, brute force, complete computation of \hat{H} is used.

Once \hat{z} and \hat{H} are computed, the program uses the key lemma (6) to calculate \hat{x} for invertible frequencies of k . If $\hat{H}(k)$ is noninvertible, an error message with the problematic k frequency is printed to the console, and the \hat{x} in question is simply filled with a column of zeroes. This process is carried out for all values of k .

Finally, x is computed by taking the inverse fourier transform of \hat{x} . Like the fourier transform of z , this inversion also makes use of the numpy fast fourier transform library. Because the inverse fourier transform is normalized, \hat{x} must be multiplied by a factor of N

to receive the desired output.

4 Discussion and Open problems

Though this paper outlines an algorithm for calculating the forward and inverse DRTs of a given body, this algorithm has not been optimized for maximum efficiency. Open questions for maximizing this efficiency include:

- Is the 0 frequency recoverable without row scans?
- What is the smallest number of scans needed for successful inversion?
- Can the \hat{H} matrices be computed directly from R and R^* matrices, without computation of the H matrices?
- Optimizing the computation of \hat{H} in the current program, by successfully implementing the FFT
- Can one create a user friendly interface to select scans with unusual shapes (v's, x's, odd shapes) and output the corresponding transform R matrices?

5 References

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