

### Medical Image Reconstruction Term II – 2010

## **Topic 1: Mathematical Basis Lecture 3**

Professor Yasser Mostafa Kadah

# **Topics Today**

- Projection-slice theorem
- Interlaced Fourier transform

- Also known as Central-Slice Theorem
- A property of the Fourier transform
- Relates the projection data in the spatial domain to the frequency domain
- States that the ID Fourier transform of the projection of an image at an angle θ is equal to the slice of the 2D Fourier transform at the same angle





D Fourier transformation:

$$F(k_x, k_y) = \iint f(x, y) \cdot e^{-j2\pi(k_x \cdot x + k_y \cdot y)} dxdy$$

• The slice of the 2D Fourier transform at  $k_x=0$  is given by:

$$F(0,k_y) = \int \left( \int f(x,y) dx \right) e^{-j2\pi k_y \cdot y} dy$$

and at  $k_y = 0$  is given by

$$F(k_x,0) = \int \left( \int f(x,y) dy \right) e^{-j2\pi k_x \cdot x} dx$$

For a general angle, the rotation property of the Fourier transformation can be used to generalize the mathematical result for a vertical projection to any angle



### **Projection-Slice Theorem: Application to CT**

- The projection data can be shown to correspond to radial sampling of the frequency domain
- It is not straightforward to numerically compute the image from this frequency domain representation
  - Limitation of the DFT to uniform sampled data
- Interpolation can be used in the frequency domain to regrid the radial sampling to uniform sampling



### **Projection-Slice Theorem: Application to MRI**

- Navigator echo motion estimation
  - Acquire a single k-space line in the middle to estimation linear translation in this direction
- Early MRI reconstruction based on backprojection algorithms

### **Interlaced Fourier Transform**

#### • A special case of nonuniform Fourier transform



### **Interlaced Fourier Transform**

Mathematical formulation

$$g(k_x, y) = \frac{-e^{\kappa 2\pi i\xi(k_x)}}{1 - e^{\kappa 2\pi i\xi(k_x)}} g_P(k_x, y) + \frac{e^{iy\eta(k_x)}}{1 - e^{\kappa 2\pi i\xi(k_x)}} g_N(k_x, y),$$
  
$$h(k_x, y) = \frac{1}{1 - e^{\kappa 2\pi i\xi(k_x)}} g_P(k_x, y) - \frac{e^{iy\eta(k_x)}}{1 - e^{\kappa 2\pi i\xi(k_x)}} g_N(k_x, y),$$

$$G(k_{x}, y) = h\left(k_{x}, y + \frac{L_{y}}{2}\right) \qquad -\frac{L_{y}}{2} \leq y < -\frac{L_{y}}{4},$$

$$G(k_{x}, y) = g(k_{x}, y) \qquad -\frac{L_{y}}{4} \leq y \leq \frac{L_{y}}{4},$$

$$G(k_{x}, y) = h\left(k_{x}, y - \frac{L_{y}}{2}\right) \qquad \frac{L_{y}}{4} < y \leq \frac{L_{y}}{2}.$$

### **Interlaced Fourier Transform**



FIG. 2. (a)  $g_P(k_x, y)$  at a specific  $k_x$ . (b)  $g(k_x, y)$  at same  $k_x$  as in (a). (c)  $h(k_x, y)$  at same  $k_x$  as in (a). (d)  $G(k_x, y)$  at same  $k_x$  as in (a), (b), and (c).

D

# **Shepp-Logan Phantom**

Numerical phantom used to simulate the human head to evaluate reconstruction algorithms in computed tomography

RECONSTRUCTING INTERIOR HEAD TISSUE FROM X-RAY TRANSMISSIONS L. A. Shepp and B. F. Logan Bell Laboratories Murray Hill, New Jersey 07974





### **Shepp-Logan Phantom**

D

TABLE 1											
Ellipses	Center	Major Axis	Minor Axis	Theta	Gray level						
a	(0,0)	.69	.92	0	2						
Ъ	(0,-0184)	.6624	.874	0	98						
с	(.22,0)	.11	.31	-18°	02						
đ	(22,0)	.16	.41	18°	02						
e	(0,.35)	.21	.25	0	.01						
f	(0,.1)	.046	.046	0	.01						
g	(0,1)	.046	.046	С	.02						
h	(08,605)	.046	.023	0	.01						
i	(0,605)	.023	.023	0	.01						
j	(.06,605)	.023	.046	0	.01						

### **Shepp-Logan Phantom: 3D**

Ellipsoid (i)	Center $(r_0)$			Half-Axis		Angle	Spin	Portion	Tissue	
	x	y	r.	a	b	c		Density	Subtracted	Type
1*	0	0	0	0.72	0.95	0.93	0	0.8	None	Scalp
2	0	0	0	0.69	0.92	0.9	0	0.12 [13]	2[Prop[1]]	Bone & Marrow
3*	0	-0.0184	0	0.6624	0.874	0.88	0	0.98 [13]	3[Prop[2]]	CSF
4**	0	-0.0184	0	0.6524	0.864	0.87	0	0.745 [14]	4[Prop[3]]	Gray Matter
5	-0.22	0	-0.25	0.41	0.16	0.21	-720	0.98	5[Prop[4]]	CSF
6	0.22	0	-0.25	0.31	0.11	0.22	720	0.98	6[Prop[4]]	CSF
7	0	0.35	-0.25	0.21	0.25	0.35	0	0.617 [14]	7[Prop[4]]	White Matter
8	0	0.1	-0.25	0.046	0.046	0.046	0	0.95 [6]	8[Prop[4]]	Tumor
9	-0.08	-0.605	-0.25	0.046	0.023	0.02	0	0.95	9[Prop[4]]	Tumor
10	0.06	-0.605	-0.25	0.046	0.023	0.02	-900	0.95	10[Prop[4]]	Tumor
11	0	-0.1	-0.25	0.046	0.046	0.046	0	0.95	11[Prop[4]]	Tumor
12	0	-0.605	-0.25	0.023	0.023	0.023	0	0.95	12[Prop[4]]	Tumor
13†*	0.06	-0.105	0.0625	0.056	0.04	0.1	-900	0.93 [6]	13[Prop[4]]	Tumor
14+*	0	0.1	0.625	0.056	0.056	0.1	0	0.98	14[Prop[4]]	CSF
15++	0.56	-0.4	-0.25	0.2	0.03	0.1	700	0.85 [15]	Not Used	Blood Clot

phantom only, <sup>††</sup> Optional region for original S-L phantom, not used herein. Portion subtracted: e.g., **2**[Prop[**1**]] means we subtract an ellipsoid with Ellipsoid 2's geometry (center and dimensions) but Ellipsoid 1's MR properties (relaxation and spin density). Scalp spin density is based on muscle/fat water content since skin water content is highly variable. Tumor spin density is based on its x-ray attenuation coefficient [6].

### **Shepp-Logan Phantom: k-Space**

- Using the known Fourier transformation of the Shepp-Logan phantom components (circles and ellipses), one can generate the analytical form of its Fourier transformation
  - Can be sampled arbitrarily to generate uniformly or nonuniformly sampled data for close to real data generation
  - Applications include radial sampling (e.g., CT and MRI), spiral and random sampling (MRI).
- This will be the standard for all evaluation procedures of image reconstruction methods.

# **Simulation of Medical Image Artifacts**

- Motion artifacts in MRI and CT
  - Different parts of k-space correspond to different subject positions
  - Can be simulated using Shepp-Logan phantom





### Exercise

- Write a program to verify the projection-slice theorem using a simple 2D phantom (e.g., a basic shape like a square).
- Perform interlaced sampling on a function of your choice with known analytical Fourier transform and verify the interlaced Fourier transform theorem.
- Write a Matlab program to implement the analytical Shepp-Logan phantom and test it using sampling on a uniform grid.