4_1_PET_CT_WB_FDG_Barn.Child

22. oktober 2012

Indhold

1	Top	Topogram														1							
	1.1	Routin	e														 						1
	1.2	Scan															 						1
2	\mathbf{CT}	WB																					1
	2.1	Routin	ie														 						1
	2.2	Scan															 						2
	2.3	Recons	3														 						2
		2.3.1	Recon 1																				2
		2.3.2	Recon 2																				3
		2.3.3	Recon 3																				4
		2.3.4																					4
		2.3.5	Recon 5																				5
3	Pau	se																					6
4	PET WB														6								
	4.1	Routin	ie														 						6
	4.2	Scan															 						6
	4.3		5																				7
		4.3.1	Recon 1																				7
		4.3.2	Recon 2																				7

1 Topogram

1.1 Routine

• mA: 30

• kV: 120

• Topogram length: 768 mm

• Tube position: Bottom

1.2 Scan

• mA: 30

• kV: 120

• Delay: 4s

• Topogram length: 768 mm

• Direction: Craniocaudal

• Tube position: Bottom

• API: API

• Kernel: 80

• Window: Topogram

2 CT WB

2.1 Routine

• Eff. mAs: 40

• kV: 120

• CARE Dose4D: On

• CareDoseType: CareDoseAEC

• CTDlvol: 2.912mGy

• Scan time: 11.070 s

• Delay: 6.000 s

• Slice: 2 mm

• No. of images: Samme som i foerste recon, slet?(y/n)

• Tilt: 0.0 grader

2.2 Scan

• Quality ref. mAs: 40

• Eff. mAs: 40

• kV: 120

 \bullet Scan time: 11.070 s

• Rotation time: 0.500 s

• Delay: 6.000 s

• Slice: 2 mm

• Pitch: 1.20

• Direction: Craniocaudal

2.3 Recons

2.3.1 Recon 1

• Series description: tmp CT WB RTD

• Slice: 4

• Kernel: B19f

• Window: Abdomen

• Extended FoV: false

• FoV: 500

• Center X: 0

• Center Y: 0

• Mirroring: None

• Extended CT scale: Standard

• Recon job: Axial

• Recon Axis: Axial

• Image order: Craniocaudal

• Recon increment: 4.000

• No. of images: 119

2.3.2 Recon 2

• Series description: AC CT WB

• Slice: 2

• Kernel: B19f

• Window: Abdomen

• Extended FoV: true

• FoV: 780

• Center X: 0

• Center Y: 0

• Mirroring: None

• Extended CT scale: Standard

• Recon job: Axial

• Recon Axis: Axial

• Image order: Craniocaudal

• Recon increment: 2.000

• No. of images: 237

2.3.3 Recon 3

• Series description: CT WB 2.0 B30f

• Slice: 2

• Kernel: B30f

• Window: Abdomen

• Extended FoV: false

• FoV: 500

• Center X: 0

• Center Y: 0

• Mirroring: None

• Extended CT scale: Standard

• Recon job: Axial

• Recon Axis: Axial

• Image order: Craniocaudal

• Recon increment: 2.000

• No. of images: 237

2.3.4 Recon 4

• Series description: CT WB 2.0 cor B30f

• Slice: 2

• Kernel: B30f

• Window: Abdomen

• Extended FoV: false

• FoV: 500

• Center X: 0

• Center Y: 0

- Mirroring: None
- Extended CT scale: Standard
- Recon job: 3D
- Recon Axis: Coronal
- Image order: AP
- Recon increment: 2.000
- No. of images: 251

2.3.5 Recon 5

- Series description: CT WB 2.0 sag B30f
- Slice: 2
- Kernel: B30f
- Window: Abdomen
- Extended FoV: false
- FoV: 500
- Center X: 0
- Center Y: 0
- Mirroring: None
- Extended CT scale: Standard
- Recon job: 3D
- Recon Axis: Sagittal
- ullet Image order: LeftToRight
- Recon increment: 2.000
- No. of images: 251

3 Pause

4 PET WB

4.1 Routine

• Isotope: F-18

• Pharm.: FDG

• Inj. Dose: 100000000.000000 Bequerels

• Inj. date (date/month - year): 01/04 - 2011

• Inj. time: 14:34:05

• Scan mode: Sinogram

• Scan range: Match CT FOV

• No. of beds: 0

• Scan duration/bed: 2.000000 Minutes

4.2 Scan

• Autoload: true

• Rebinner LUT: N/A

• Scan output: Sinogram

• Sinogram mode: PromptsRandoms

• Input trigger signal: None

• LLD (keV): N/A

• ULD (keV): N/A

4.3 Recons

4.3.1 Recon 1

- Series description: PET WB image check
- Recon range (bed): 1 to 3
- Output image type: Corrected
- Recon method: TrueX
- Iterations: 2
- Subsets: 12
- Image size: 200
- Zoom: 1
- Filter: Gaussian
- FWHM (mm): 3.000000
- Offset X: 0 mm
- Offset Y: 0 mm
- Attenuation correction: On (1)
- Scatter correction: true
- Match CT slice location: true
- Save intermediate data: false

4.3.2 Recon 2

- Series description: PET WB Corr-TrueX + TOF 2 mm
- Recon range (bed): 1 to 3
- Output image type: Corrected
- Recon method: TrueXTOF
- Iterations: 2
- Subsets: 21

• Image size: 400

• Zoom: 1

• Filter: Gaussian

• FWHM (mm): 3.000000

• Offset X: 0 mm

• Offset Y: 0 mm

• Attenuation correction: On (1)

• Scatter correction: true

• Match CT slice location: true

• Save intermediate data: false