# $4\_PET\_CT\_WB\_FDG\_Terapi\_Barn\_IV\_Bolus.Child$

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## Indhold

T	Topogram			
	1.1	Routine		
	1.2	Scan		
<b>2</b>	Pre	Monitoring 1		
	2.1	Routine		
	2.2	Scan		
	2.3	Recons		
		2.3.1 Recon 1		
3	Mo	nitoring		
	3.1	Routine		
	3.2	Scan		
	3.3	Recons		
		3.3.1 Recon 1		
4	CT WB			
	4.1	Routine		
	4.2	Scan		
	4.3	Recons		
		4.3.1 Recon 1		
		4.3.2 Recon 2		
		4.3.3 Recon 3		
		4.3.4 Recon 4		
		4.3.5 Recon 5		
5	Pau	${f se}$		
6	PE	Γ WB		
	6.1	Routine		
	6.2	Scan		
	6.3	Recons		
		6.3.1 Recon 1		
		632 Recon 2		

## 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 PreMonitoring

### 2.1 Routine

• Eff. mAs: 20

• kV: 120

• CARE Dose4D: Off

• CareDoseType: CareDoseAEC

• CTDlvol: 1.186mGy

• Scan time: 0.500 s

• Delay: 2.000 s

• Slice: 10 mm

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

• Tilt: 0.0 grader

## 2.2 Scan

• Quality ref. mAs: 65

• Eff. mAs: 20

• kV: 120

 $\bullet$  Scan time: 0.500 s

• Rotation time: 0.500 s

• Delay: 2.000 s

• Slice: 10 mm

• Pitch: 0.00

• Direction: Craniocaudal

#### 2.3 Recons

#### 2.3.1 Recon 1

• Series description: PreMonitoring 10.0 eFoV

• Slice: 10

• Kernel: B19s

• 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: 10.000

• No. of images: 1

## 3 Monitoring

#### 3.1 Routine

• Eff. mAs: 20

• kV: 120

• CARE Dose4D: Off

• CareDoseType: CareDoseAEC

• CTDlvol: 35.58mGy

 $\bullet$  Scan time: 0.500 s

• Delay: 10.000 s

• Slice: 10 mm

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

• Tilt: 0.0 grader

## 3.2 Scan

• Quality ref. mAs: 65

• Eff. mAs: 20

• kV: 120

 $\bullet$  Scan time: 0.500 s

 $\bullet$  Rotation time: 0.500 s

• Delay: 10.000 s

• Slice: 10 mm

• Pitch: 0.00

• Direction: Craniocaudal

#### 3.3 Recons

#### 3.3.1 Recon 1

• Series description: Monitoring 10.0 eFoV

• Slice: 10

• Kernel: B19s

• 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: 10.000

• No. of images: 30

## 4 CT WB

#### 4.1 Routine

• Eff. mAs: 30

• kV: 120

• CARE Dose4D: On

• CareDoseType: CareDoseAEC

• CTDlvol: 2.184mGy

 $\bullet$  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

#### 4.2 Scan

• Quality ref. mAs: 30

• Eff. mAs: 30

• kV: 120

• Scan time: 11.070 s

• Rotation time: 0.500 s

 $\bullet$  Delay: 6.000 s

• Slice: 2 mm

• Pitch: 1.20

• Direction: Craniocaudal

#### 4.3 Recons

#### 4.3.1 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

#### 4.3.2 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

#### 4.3.3 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

#### 4.3.4 Recon 4

 $\bullet\,$  Series description: CT WB 2.0 SPO cor

• 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

#### 4.3.5 Recon 5

• Series description: CT WB 2.0 SPO sag

• 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

• Image order: LeftToRight

• Recon increment: 2.000

• No. of images: 251

## 5 Pause

## 6 PET WB

#### 6.1 Routine

• Isotope: F-18

• Pharm.: FDG

• Inj. Dose: 100000000.000000 Bequerels

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

• Inj. time: 14:45:45

• Scan mode: Sinogram

• Scan range: Match CT FOV

• No. of beds: 0

• Scan duration/bed: 2.000000 Minutes

#### 6.2 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

#### 6.3 Recons

#### 6.3.1 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: Off (1)
- Scatter correction: true
- Match CT slice location: true
- Save intermediate data: false

#### 6.3.2 Recon 2

• Series description: PET WB Corr-TrueX

• 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): 2.000000

• Offset X: 0 mm

• Offset Y: 0 mm

• Attenuation correction: Off (1)

• Scatter correction: true

• Match CT slice location: true

• Save intermediate data: false