# $1\_PET\_CT\_WB\_FDG\_med\_IV\_Bolus.Adult\_PET4$

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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: 35
- kV: 120
- Topogram length: 1024 mm
- Tube position: Top

#### 1.2 Scan

- mA: 35
- kV: 120
- Delay: 4s
- Topogram length: 1024 mm
- Direction: Craniocaudal
- Tube position: Top
- API: API
- Kernel: 80
- Window: Topogram

# 2 PreMonitoring

## 2.1 Routine

- Eff. mAs: 20
- kV: 120
- CARE Dose4D: Off
- CareDoseType: CareDoseAEC
- CTDlvol: m 1=0.000000 1.18627mGy

• 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: 225

• Eff. mAs: 20

• kV: 120

 $\bullet$  Scan time: 0.500 s

• Rotation time: 0.500 s

• Delay: 2.000 s

• Slice: 10 mm

• Pitch: m 1=-1 0.00

• Direction: Craniocaudal

## 2.3 Recons

#### 2.3.1 Recon 1

 $\bullet$  Series description: m 1=12345MA m 2= PreMonitoring 10.0 eFoV

• Slice: 10

• Kernel: B19s

• Window: Abdomen

• Extended FoV: On

• 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

 $\bullet\,$  No. of images: 1

# 3 Monitoring

## 3.1 Routine

• Eff. mAs: 20

• kV: 120

• CARE Dose4D: Off

• CareDoseType: CareDoseAEC

• CTDlvol: m 1=0.000000 35.5881mGy

• 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: 225

• Eff. mAs: 20

• kV: 120

 $\bullet$  Scan time: 0.500 s

• Rotation time: 0.500 s

• Delay: 10.000 s

• Slice: 10 mm

• Pitch: m 1=-1 0.00

• Direction: Craniocaudal

#### 3.3 Recons

#### 3.3.1 Recon 1

• Series description: m 1=12345MA m 2= Monitoring 10.0 eFoV

• Slice: 10

• Kernel: B19s

• Window: Abdomen

• Extended FoV: On

• 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: 225

• kV: 120

• CARE Dose4D: On

• CareDoseType: CareDoseAEC

• CTDlvol: m 1=0.000000 16.4451mGy

• Scan time: 28.644 s

 $\bullet$  Delay: 25.000 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: 225

• Eff. mAs: 225

• kV: 120

• Scan time: 28.644 s

• Rotation time: 0.500 s

 $\bullet$  Delay: 25.000 s

• Slice: 2 mm

• Pitch: m 1=-1 0.80

• Direction: Craniocaudal

### 4.3 Recons

#### 4.3.1 Recon 1

- $\bullet$  Series description: m 1=12345MA m 2= CT WB RTD
- Slice: 4
- Kernel: B19f
- Window: Abdomen
- Extended FoV: Off
- 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: 213

#### 4.3.2 Recon 2

- Series description: m  $1=45 \mathrm{AC} \ \mathrm{CT} \ 2 \ \mathrm{mm} \ MA \ \mathrm{m} \ 2= \mathrm{AC} \ \mathrm{CT} \ 2 \ \mathrm{mm}$
- Slice: 2
- Kernel: B19f
- Window: Abdomen
- Extended FoV: On
- 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: 426

#### 4.3.3 Recon 3

 $\bullet$  Series description: m 1=CT WB2345MA m 2= CT WB 2.0 B30f

• Slice: 2

• Kernel: B30f

• Window: Abdomen

• Extended FoV: Off

• 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: 426

#### 4.3.4 Recon 4

- $\bullet$  Series description: m 1=CT WB245B30f MA m 2= CT WB 2.0 cor B30f
- Slice: 2
- Kernel: B30f
- Window: Abdomen
- Extended FoV: Off
- FoV: 1103
- 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: 185

#### 4.3.5 Recon 5

- $\bullet$  Series description: m 1=CT WB245B30f MA m 2= CT WB 2.0 sag B30f
- Slice: 2
- Kernel: B30f
- Window: Abdomen
- Extended FoV: Off
- FoV: 1103
- 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: 250

## 5 Pause

## 6 PET WB

## 6.1 Routine

• Isotope: F-18

• Pharm.: Fluorodeoxyglucose

• Inj. Dose: 0 Bequerels

• Scan mode: Sinogram

• Scan range: Do not match CT FOV

• No. of beds: Not given. Check recon range

• Scan duration/bed: 2.000000 Minutes

#### 6.2 Scan

• Autoload: On

• Rebinner LUT: Off

• Scan output: Sinogram

• Sinogram mode: false

• 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 HD
- Recon range (bed): 1 to 6
- Output image type: Corrected
- Recon method: TrueX
- Iterations: 2
- Subsets: 12
- Image size: 200
- Zoom: 1
- Filter: Gaussian
- FWHM (mm): 4.000000
- Offset X: 0 mm
- Offset Y: 0 mm
- Attenuation correction: Off ()
- Scatter correction: On
- Match CT slice location: On
- Save intermediate data: Off

#### 6.3.2 Recon 2

• Series description: PET WB Corr TrueX+TOF 2 mm

• Recon range (bed): 1 to 6

• 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 ()

• Scatter correction: On

• Match CT slice location: On

• Save intermediate data: Off