# 7\_PET\_AC\_FDG\_FINE.MlAdult\_PET6

20. februar 2013

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## 1 Topogram

## 1.1 Routine

- mA: 35
- kV: 120
- Topogram length: 0 mm
- Tube position: Top

#### 1.2 Scan

- mA: 35
- kV: 120
- Delay: 4s
- Topogram length: 0 mm
- Direction: Craniocaudal
- Tube position: Top
- API: None
- Kernel: 80s shar
- Window: Topogram Body

## 2 AC CT Femur

- Eff. mAs: 11
- kV: 120
- CARE Dose4D: Off
- CareDoseType: CareDoseAEC
- $\bullet$  CTDlvol: 0.7414mGy

• Scan time: 3.360 s

• Delay: 4.000 s

• Slice: 3.00 mm

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

• Tilt: 0.0 grader

## 2.2 Scan

• Quality ref. mAs: 12

• Eff. mAs: 11

• kV: 120

 $\bullet$  Scan time: 3.360 s

• Rotation time: 0.500 s

• Delay: 4.000 s

• Slice: 3.00 mm

• Pitch: 1.5000000

• Direction: Craniocaudal

## 2.3 Recons

#### 2.3.1 Recon 1

• Series description: AC CT Femur

• Slice: 3.00

• Kernel: B19f PET very smooth

• Window: Abdomen

• Extended FoV: On

• FoV: 700

• 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: 111

## 3 Pause

## 4 PET Femur LM

## 4.1 Routine

• Isotope: F-18

• Pharm.: FDG

• Inj. Dose: 200 MegaBequerels

• Scan mode: List mode

• Scan range: Match CT FOV

• No. of beds: 1

• Scan duration/bed: 60 Minutes

#### 4.2 Scan

• Autoload: On

• Rebinner LUT: Off

• Scan output: List mode

• Sinogram mode: Trues

- Input trigger signal: None
- LLD (keV): 435
- ULD (keV): 650

#### 4.3 Recons

#### 4.3.1 Recon 1

- Series description: PET WB
- $\bullet$  Recon range (bed): -1 to -1
- Output image type: NoRecon
- Recon method: Iterative
- Iterations: 4
- Subsets: 8
- Image size: 168
- Zoom: 1
- Filter: Gaussian
- FWHM (mm): 5
- Offset X: 0 mm
- Offset Y: 0 mm
- Attenuation correction: On (1)
- Scatter correction: On
- Match CT slice location: On
- Save intermediate data: Off

## 5 Pause

## 6 Topogram

## 6.1 Routine

- mA: 35
- kV: 120
- Topogram length: 0 mm
- Tube position: Lateral

## 6.2 Scan

- mA: 35
- kV: 120
- Delay: 4s
- Topogram length: 0 mm
- Direction: Craniocaudal
- $\bullet\,$  Tube position: Lateral
- API: None
- Kernel: 20s standar
- Window: Topogram Head

## 7 AC CT Hjerne

- Eff. mAs: 22
- kV: 120
- CARE Dose4D: Off

• CareDoseType: CareDoseAEC

• CTDlvol: 3.4496mGy

• Scan time: 9.280 s

• Delay: 4.000 s

• Slice: 3.00 mm

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

• Tilt: 0.0 grader

#### 7.2 Scan

• Quality ref. mAs: 14

• Eff. mAs: 22

• kV: 120

• Scan time: 9.280 s

• Rotation time: 1.000 s

• Delay: 4.000 s

• Slice: 3.00 mm

• Pitch: 1.5000000

• Direction: Caudocranial

#### 7.3 Recons

#### 7.3.1 Recon 1

• Series description: AC CT Hjerne

• Slice: 3.00

• Kernel: H19s PET very smooth

• Window: Cerebrum

• Extended FoV: Off

• FoV: 300

• Center X: 0

• Center Y: 0

• Mirroring: None

• Extended CT scale: Standard

• Recon job: Axial

• Recon Axis: Axial

• Image order: Caudocranial

• Recon increment: 3.000

• No. of images: 74

## 8 Pause

## 9 PET Brain LM

## 9.1 Routine

• Isotope: F-18

• Pharm.: FDG

• Inj. Dose: 200 MegaBequerels

• Scan mode: List mode

• Scan range: Match CT FOV

• No. of beds: 1

• Scan duration/bed: 15 Minutes

#### 9.2 Scan

• Autoload: On

• Rebinner LUT: Off

• Scan output: List mode

• Sinogram mode: Trues

• Input trigger signal: None

• LLD (keV): 435

• ULD (keV): 650

#### 9.3 Recons

#### 9.3.1 Recon 1

• Series description: PET FDG Brain corrected

• Recon range (bed): -1 to -1

• Output image type: NoRecon

• Recon method: Iterative3D

• Iterations: 6

• Subsets: 16

• Image size: 336

• Zoom: 3

• Filter: Gaussian

• FWHM (mm): 3

• Offset X: 0 mm

• Offset Y: 0 mm

• Attenuation correction: Off (1)

• Scatter correction: On

• Match CT slice location: On

• Save intermediate data: Off

## 10 Pause

## 11 Topogram

## 11.1 Routine

- mA: 35
- kV: 120
- Topogram length: 0 mm
- Tube position: Top

### 11.2 Scan

- mA: 35
- kV: 120
- Delay: 4s
- Topogram length: 0 mm
- Direction: Craniocaudal
- Tube position: Top
- API: None
- Kernel: 80s shar
- Window: Topogram Body

## 12 AC CT WB

- Eff. mAs: 11
- kV: 120
- CARE Dose4D: Off

• CareDoseType: CareDoseAEC

 $\bullet$  CTDlvol: 0.7414mGy

• Scan time: 10.310 s

• Delay: 4.000 s

• Slice: 3.00 mm

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

• Tilt: 0.0 grader

#### 12.2 Scan

• Quality ref. mAs: 14

• Eff. mAs: 11

• kV: 120

• Scan time: 10.310 s

• Rotation time: 0.500 s

• Delay: 4.000 s

• Slice: 3.00 mm

• Pitch: 1.5000000

• Direction: Craniocaudal

#### 12.3 Recons

#### 12.3.1 Recon 1

• Series description: AC CT WB

• Slice: 3.00

• Kernel: B19f PET very smooth

• Window: Abdomen

• Extended FoV: On

• FoV: 700

• Center X: 0

• Center Y: 0

• Mirroring: None

• Extended CT scale: Standard

• Recon job: Axial

• Recon Axis: Axial

• Image order: Craniocaudal

• Recon increment: 3.000

• No. of images: 274

## 13 Pause

## 14 PET WB

## 14.1 Routine

• Isotope: F-18

• Pharm.: FDG

• Inj. Dose: 200 MegaBequerels

• Scan mode: Sinogram

• Scan range: Match CT FOV

• No. of beds: 1

• Scan duration/bed: 4 Minutes

## 14.2 Scan

• Autoload: On

• Rebinner LUT: Off

• Scan output: Sinogram

• Sinogram mode: Trues

• Input trigger signal: None

• LLD (keV): 435

• ULD (keV): 650

### 14.3 Recons

#### 14.3.1 Recon 1

• Series description: PET WB Image check

• Recon range (bed): -1 to -1

• Output image type: Corrected

• Recon method: Iterative3D

• Iterations: 4

• Subsets: 8

• Image size: 168

• Zoom: 1

• Filter: Gaussian

• FWHM (mm): 4

• Offset X: 0 mm

• Offset Y: 0 mm

• Attenuation correction: Off (1)

• Scatter correction: On

• Match CT slice location: On

• Save intermediate data: Off

#### 14.3.2 Recon 2

- $\bullet$  Series description: PET WB True X 2 mm
- Recon range (bed): -1 to -1
- Output image type: Corrected
- Recon method: TrueX
- Iterations: 3
- Subsets: 21
- Image size: 336
- Zoom: 1
- Filter: Gaussian
- FWHM (mm): 2
- Offset X: 0 mm
- Offset Y: 0 mm
- Attenuation correction: Off (1)
- Scatter correction: On
- Match CT slice location: On
- Save intermediate data: Off

## 15 Pause

## 16 PET WB

- Isotope: F-18
- Pharm.: FDG
- Inj. Dose: 200 MegaBequerels
- Scan mode: Sinogram

- Scan range: Match CT FOV
- No. of beds: 1
- Scan duration/bed: 4 Minutes

#### 16.2 Scan

- Autoload: On
- Rebinner LUT: Off
- Scan output: Sinogram
- Sinogram mode: Trues
- Input trigger signal: None
- LLD (keV): 435
- ULD (keV): 650

### 16.3 Recons

#### 16.3.1 Recon 1

- Series description: PET WB Image check
- Recon range (bed): -1 to -1
- Output image type: Corrected
- Recon method: Iterative3D
- Iterations: 4
- Subsets: 8
- Image size: 168
- Zoom: 1
- Filter: Gaussian
- FWHM (mm): 4
- Offset X: 0 mm

- Offset Y: 0 mm
- Attenuation correction: Off (1)
- Scatter correction: On
- Match CT slice location: On
- Save intermediate data: Off

#### 16.3.2 Recon 2

- Series description: PET WB True X 2 mm
- Recon range (bed): -1 to -1
- Output image type: Corrected
- Recon method: TrueX
- Iterations: 3
- Subsets: 21
- Image size: 336
- Zoom: 1
- Filter: Gaussian
- FWHM (mm): 2
- Offset X: 0 mm
- Offset Y: 0 mm
- Attenuation correction: Off (1)
- Scatter correction: On
- Match CT slice location: On
- Save intermediate data: Off