# WB\_Flourid\_PET\_5.MlAdult\_PET5

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## 1 Topogram

## 1.1 Routine

• mA: 35

• kV: 120

• Topogram length: 34 mm

• Tube position: Top

#### 1.2 Scan

• mA: 35

• kV: 120

• Delay: 4s

• Topogram length: 34 mm

• Direction: Caudocranial

• Tube position: Top

• API: None

• Kernel: 80s shar

• Window: Topogram Body

## 2 Lav CT knogle

## 2.1 Routine

• Eff. mAs: 40

• kV: 120

• CARE Dose4D: Off

• CareDoseType: CareDoseAEC

• CTDlvol: 2.70069mGy

• Scan time: 38.240 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: 50

• Eff. mAs: 40

• kV: 120

 $\bullet$  Scan time: 38.240 s

• Rotation time: 0.500 s

• Delay: 4.000 s

• Slice: 3.00 mm

• Pitch: 0.8000000

• Direction: Craniocaudal

## 2.3 Recons

#### 2.3.1 Recon 1

• Series description: AC CT

• 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: 574

## 3 Pause

## 4 PET knogle

### 4.1 Routine

• Isotope: F-18

• Pharm.: Fluoride

• Inj. Dose: 200 MegaBequerels

 $\bullet\,$  Inj. date (date/month - year): 21/08 - 2012

• Inj. time: 14:00:00

• Scan mode: Sinogram

• Scan range: Match CT FOV

• No. of beds: 11

• Scan duration/bed: 120 Seconds

### 4.2 Scan

• Autoload: On

• Rebinner LUT: Off

• Scan output: Sinogram

• Sinogram mode: Trues

• Input trigger signal: None

• LLD (keV): 435

• ULD (keV): 650

## 4.3 Recons

#### 4.3.1 Recon 1

• Series description: PET knogleskanning Image check

• Recon range (bed): Automatic/All/dunno

• Output image type: Corrected

• Recon method: Iterative

• Iterations: 4

• Subsets: 8

• Image size: 168

• Zoom: 1

• Filter: Gaussian

• FWHM (mm): 4

• Offset X: 0 mm

• Offset Y: 0 mm

• Attenuation correction: On (1)

• Scatter correction: On

• Match CT slice location: On

• Save intermediate data: Off

#### 4.3.2 Recon 2

- Series description: PET knogleskanning True X 2 mm
- Recon range (bed): Automatic/All/dunno
- 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: On (1)
- Scatter correction: On
- Match CT slice location: On
- Save intermediate data: Off

#### 4.3.3 Recon 3

- Series description: no AC
- Recon range (bed): Automatic/All/dunno
- Output image type: Uncorrected
- Recon method: Iterative
- Iterations: 3
- Subsets: 21
- Image size: 336

- Zoom: 1
- Filter: Gaussian
- FWHM (mm): 4
- Offset X: 0 mm
- Offset Y: 0 mm
- Attenuation correction: On (1)
- Scatter correction: On
- Match CT slice location: On
- Save intermediate data: Off

#### 4.3.4 Recon 4

- Series description: PET WB UDEN Trie X
- Recon range (bed): Automatic/All/dunno
- Output image type: Corrected
- Recon method: Iterative
- Iterations: 3
- Subsets: 21
- Image size: 336
- Zoom: 1
- Filter: Gaussian
- FWHM (mm): 4
- Offset X: 0 mm
- Offset Y: 0 mm
- Attenuation correction: On (1)
- Scatter correction: On
- Match CT slice location: On
- Save intermediate data: Off