# 3\_PET\_AC\_FDG\_Brain\_Neurostat.Adult\_PET4

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## 1 Topogram

## 1.1 Routine

• mA: 35

• kV: 120

• Topogram length: 256 mm

• Tube position: Lateral

#### 1.2 Scan

• mA: 35

• kV: 120

• Delay: 4s

• Topogram length: 256 mm

• Direction: Craniocaudal

• Tube position: Lateral

• API: API

• Kernel: 20

• Window: Topogram

## 2 CT Brain

## 2.1 Routine

• Eff. mAs: 30

• kV: 120

• CARE Dose4D: On

• CareDoseType: CareDoseAEC

• CTDlvol: 5.124mGy

• Scan time: 2.640 s

• Delay: 2.000 s

• Slice: 3 mm

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

• Tilt: 0.0 grader

### 2.2 Scan

• Quality ref. mAs: 30

• Eff. mAs: 30

• kV: 120

 $\bullet$  Scan time: 2.640 s

• Rotation time: 1.000 s

• Delay: 2.000 s

• Slice: 3 mm

• Pitch: 1.50

• Direction: Craniocaudal

### 2.3 Recons

#### 2.3.1 Recon 1

• Series description: tmp CT Brain RTD

• Slice: 5

• Kernel: H19s

• 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: Craniocaudal

• Recon increment: 5.000

• No. of images: 1

#### 2.3.2 Recon 2

• Series description: AC CT

• Slice: 3

• Kernel: H19s

• 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: Craniocaudal

• Recon increment: 3.000

• No. of images: 1

## 3 Pause

## 4 PET Brain

#### 4.1 Routine

- Isotope: F-18
- Pharm.: Fluorodeoxyglucose
- Inj. Dose: 200000000 Bequerels
- Scan mode: Sinogram
- Scan range: Do not match CT FOV
- No. of beds: Not given. Check recon range
- Scan duration/bed: 10.000000 Minutes

#### 4.2 Scan

- Autoload: On
- Rebinner LUT: Off
- Scan output: Sinogram
- Sinogram mode: false
- Input trigger signal: None
- LLD (keV): n/a
- ULD (keV): n/a

#### 4.3 Recons

#### 4.3.1 Recon 1

- Series description: PET Brain Corrected 3 mm zoom 3
- Recon range (bed): 1 to 1
- Output image type: Corrected

• Recon method: Iterative3D

• Iterations: 4

• Subsets: 24

• Image size: 400

• Zoom: 3.000000

• Filter: Gaussian

• FWHM (mm): 3.000000

• Offset X: 0 mm

• Offset Y: 0 mm

• Attenuation correction: Off ()

• Scatter correction: On

• Match CT slice location: On

• Save intermediate data: Off

#### 4.3.2 Recon 2

• Series description: Neurostat

• Recon range (bed): 1 to 1

• Output image type: Corrected

• Recon method: Iterative3D

• Iterations: 4

• Subsets: 8

• Image size: 128

• Zoom: 3.000000

• Filter: Gaussian

• FWHM (mm): 8.000000

• Offset X: 0 mm

• Offset Y: 0 mm

 $\bullet$  Attenuation correction: Off ()

• Scatter correction: On

• Match CT slice location: On

• Save intermediate data: Off