# $Hjerne\_PIB\_AC\_LM\_4.Adult\_PET4$

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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 Lav CT Brain

## 2.1 Routine

• Eff. mAs: 32

• kV: 120

• CARE Dose4D: Off

• CareDoseType: CareDoseAEC

• CTDlvol: m 1=0.000000 5.69736mGy

• Scan time: 17.652 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: 350

• Eff. mAs: 32

• kV: 120

 $\bullet$  Scan time: 17.652 s

• Rotation time: 0.500 s

• Delay: 2.000 s

• Slice: 3 mm

• Pitch: m 1=-1 0.55

• Direction: Caudocranial

## 2.3 Recons

#### 2.3.1 Recon 1

 $\bullet$  Series description: m 1=12345MA m 2= Lav CT Brain RTD

• Slice: 4

• Kernel: H31f

• 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: 4.000

• No. of images: 56

#### 2.3.2 Recon 2

 $\bullet$  Series description: m 1=45AC CT MA m 2= AC CT

• Slice: 3

• Kernel: H31f

• 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

## 3 Pause

## 4 PET Brain LM

## 4.1 Routine

• Isotope: C-11

• Pharm.: C11-PIB

• Inj. Dose: 185000000 Bequerels

• Scan mode: List mode

• Scan range: Do not match CT FOV

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

• Scan duration/bed: 30.000000 Minutes

## 4.2 Scan

• Autoload: Off

• Rebinner LUT: Off

• Scan output: List mode

• 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: PIB 40-70 min 5mm

• Recon range (bed): 1 to 1

• Output image type: Uncorrected

- Recon method: Iterative3D
- Iterations: 4
- Subsets: 12
- Image size: 400
- Zoom: 1
- Filter: Gaussian
- FWHM (mm): 5
- Offset X: 0 mm
- Offset Y: 0 mm
- Attenuation correction: Off ()
- Scatter correction: Off
- Match CT slice location: Off
- Save intermediate data: Off

#### 4.3.2 Recon 2

- Series description: PIB 40-60 min 5mm
- Recon range (bed): 1 to 1
- Output image type: Uncorrected
- Recon method: Iterative3D
- Iterations: 4
- Subsets: 12
- Image size: 400
- Zoom: 1
- Filter: Gaussian
- FWHM (mm): 5
- Offset X: 0 mm

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

#### 4.3.3 Recon 3

- Series description: PIB 45-60 min 5mm
- Recon range (bed): 1 to 1
- Output image type: Uncorrected
- Recon method: Iterative3D
- Iterations: 4
- Subsets: 12
- Image size: 400
- Zoom: 1
- Filter: Gaussian
- FWHM (mm): 5
- Offset X: 0 mm
- Offset Y: 0 mm
- Attenuation correction: Off ()
- Scatter correction: Off
- Match CT slice location: Off
- Save intermediate data: Off