$3_PET_AC_Brain_LM_FDG.Adult_PET4$

30. maj 2013

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1 Topogram

1.1 Routine

mA: 35kV: 120

Topogram length: 256 mmTube position: Lateral

1.2 Scan

mA: 35kV: 120Delay: 4s

Topogram length: 256 mmDirection: CraniocaudalTube position: Lateral

API: APIKernel: 20

• Window: Topogram

2 Lav CT Brain

2.1 Routine

• Eff. mAs: 350

• kV: 120

• CARE Dose4D: On

• CareDoseType: CareDoseAEC

CTDlvol: 60.9301mGyScan time: 17.652 s

Delay: 2.000 sSlice: 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: 350

• kV: 120

Scan time: 17.652 s
Rotation time: 0.500 s

Delay: 2.000 sSlice: 3 mmPitch: 0.55

• Direction: Caudocranial

2.3 Recons

2.3.1 Recon 1

- Series description: Lav CT Brain RTD
- Slice: 4
- Kernel: H31f
- Window: CerebrumExtended 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

- Series description: 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: F-18

- Pharm.: Fluorodeoxyglucose
- $\bullet\,$ 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: 10.000000 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: Brain LM
- 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