$Hjerne_PiB_PET_Dyn_LM_6.MlAdult_PET6$

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: None

• Kernel: 20s standar

• Window: Topogram Head

2 AC CT Brain

2.1 Routine

• Eff. mAs: 30

• kV: 120

• CARE Dose4D: Off

• CareDoseType: CareDoseAEC

CTDlvol: 4.71218mGyScan time: 11.230 s

Delay: 4.000 sSlice: 3.00 mm

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

• Tilt: 0.0 grader

2.2 Scan

• Quality ref. mAs: 380

• Eff. mAs: 30

• kV: 120

Scan time: 11.230 sRotation time: 1.000 s

Delay: 4.000 sSlice: 3.00 mmPitch: 1.2000000

• Direction: Caudocranial

2.3 Recons

2.3.1 Recon 1

- Series description: AC CT
- Slice: 3.00
- Kernel: H19s PET very smooth
- 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: 3.000
- No. of images: 74

3 Pause

4 PET Brain

4.1 Routine

- Isotope: C-11
- Pharm.: PIB
- Inj. Dose: 1 MegaBequerels
- Scan mode: List mode
- Scan range: Match CT FOV
- No. of beds: 1
- Scan duration/bed: 30 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 FDG Brain corrected

Recon range (bed): -1 to -1
Output image type: NoRecon
Recon method: Iterative3D

Iterations: 6Subsets: 16Image size: 336

• Zoom: 3

Filter: GaussianFWHM (mm): 3Offset X: 0 mmOffset Y: 0 mm

• Attenuation correction: On (1)

• Scatter correction: On

Match CT slice location: OnSave intermediate data: Off