$21_PET_AC_Brain_LM_ACI.MlAdult_PET6$

30. maj 2013

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1 Topogram

1.1 Routine

mA: 35kV: 120

• Topogram length: 1024 mm

• Tube position: Top

1.2 Scan

mA: 35kV: 120Delay: 4s

Topogram length: 1024 mmDirection: Craniocaudal

• Tube position: Top

API: NoneKernel: 80s shar

• Window: Topogram Body

2 CT Brain

2.1 Routine

• Eff. mAs: 55

• kV: 120

• CARE Dose4D: On

• CareDoseType: CareDoseAEC

CTDlvol: 3.71345mGyScan time: 18.670 s

Delay: 4.000 sSlice: 5.00 mm

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

• Tilt: 0.0 grader

2.2 Scan

• Quality ref. mAs: 55

• Eff. mAs: 55

• kV: 120

Scan time: 18.670 sRotation time: 0.500 s

Delay: 4.000 sSlice: 5.00 mmPitch: 0.8000000

• Direction: Craniocaudal

2.3 Recons

2.3.1 Recon 1

• Series description: AC CT efov 700

• Slice: 5.00

• Kernel: B19f PET very smooth

Window: CerebrumExtended FoV: On

FoV: 700Center X: 0Center Y: 0Mirroring: None

• Extended CT scale: Standard

Recon job: AxialRecon Axis: Axial

Image order: CraniocaudalRecon increment: 3.000

• No. of images: 274

2.3.2 Recon 2

• Series description: AC CT fov 500

• Slice: 5.00

• Kernel: B19f PET very smooth

Window: CerebrumExtended FoV: Off

FoV: 500Center X: 0Center Y: 0

• Mirroring: None

• Extended CT scale: Standard

Recon job: AxialRecon Axis: Axial

Image order: CraniocaudalRecon increment: 3.000

• No. of images: 274

3 Pause

4 PET WB

4.1 Routine

• Isotope: F-18

- Pharm.: Florbetaben
- Inj. Dose: 1 MegaBequerels
- Scan mode: List mode
- Scan range: Match CT FOV
- No. of beds: 1
- Scan duration/bed: 20 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 WB Image check
- Recon range (bed): -1 to -1
- Output image type: NoRecon
- Recon method: Iterative
- Iterations: 4
- Subsets: 14
- Image size: 128
- Zoom: 2
- 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