PETCT_Brain.MlAdult_HD

1. oktober 2012

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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: None

• Kernel: T20s standard

• Window: Topogram Head

2 CT WB

2.1 Routine

• Eff. mAs: 380

• kV: 16

• CARE Dose4D: Off

• CTDlvol: 59.4292mGy

 \bullet Scan time: 0.8000000 s

• Delay: 4.000 s

• Slice: 3.00 mm

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

• Tilt: 380 grader

2.2 Scan

• Quality ref. mAs: 380

• Eff. mAs: 380

• kV: 120

• Scan time: 16.010 s

• Rotation time: 1.000 s

• Delay: 4.000 s

• Slice: 0.6 mm

• Pitch: 0.8000000

• Direction: Craniocaudal

2.3 Recon

2.3.1 Recon 1

 $\bullet\,$ Series description: AC CT

• Slice: 3.00

• Kernel: "H19s PET very smooth"

• 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: 1.500
- No. of images: 148

2.3.2 Recon 2

- Series description: CT Brain 3.0 H31s
- Slice: 3.00
- Kernel: "H31s medium smooth +"
- Window: "Cerebrum"
- Extended FoV: Off
- FoV: 250
- Center X: 0
- Center Y: 0
- Mirroring: None
- Extended CT scale: Standard
- Recon job: Axial
- Recon Axis: Axial
- Image order: Caudocranial
- Recon increment: 1.500
- No. of images: 148

3 Pause

4 PET WB

4.1 Routine

• Isotope: F-18

• Pharm.: FDG

• Inj. Dose: 5 Millicuries

 $\bullet\,$ Inj. date (date/month - year): 30/10 - 2006

• Inj. time: 08:50:00

• Scan mode: Sinogram

• Scan range: Match CT FOV

• No. of beds: 1

• Scan duration/bed: 10 PtMinute

4.2 Scan

• Autoload: On

• Rebinner LUT: Off

• Scan output: Sinogram

• Sinogram mode: Trues

• Input trigger signal: None

• LLD (keV): 435

• ULD (keV): 650

4.3 Recon

4.3.1 Recon 1

- Series description: PET Brain
- Recon range (bed): ups..
- Output image type: Corrected
- Recon method: TrueX
- Iterations: 3
- Subsets: 21
- Image size: 336
- 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