$Hjerne_FDG_PET_CT_5.MlAdult_PET5$

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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: 20s standar

• Window: Topogram Head

2 Diagn.CT Brain

2.1 Routine

• Eff. mAs: 380

• kV: 120

• CARE Dose4D: Off

• CareDoseType: CareDoseAEC

• CTDlvol: 52.2276mGy

• Scan time: 8.010 s

• Delay: 4.000 s

• Slice: 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: 380

• kV: 120

 \bullet Scan time: 8.010 s

• Rotation time: 1.000 s

• Delay: 4.000 s

• Slice: 3.00 mm

• Pitch: 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: 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

2.3.2 Recon 2

• Series description: Diagn. CT Brain Suppl.zoom

• Slice: 3.00

• Kernel: H31s medium smooth +

• Window: Cerebrum

• Extended FoV: Off

• FoV: 220

• 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: F-18

• Pharm.: FDG

• Inj. Dose: 227 MegaBequerels

• Scan mode: Sinogram

• Scan range: Match CT FOV

• No. of beds: 1

• Scan duration/bed: 10 Minutes

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 Recons

4.3.1 Recon 1

• Series description: PET Brain corrected 3 mm zoom 3

• Recon range (bed): -1 to -1

• Output image type: Corrected

- Recon method: Iterative3D
- Iterations: 6
- Subsets: 16
- Image size: 336
- Zoom: 3
- Filter: Gaussian
- FWHM (mm): 3
- Offset X: 0 mm
- Offset Y: 0 mm
- Attenuation correction: On (1)
- Scatter correction: On
- Match CT slice location: On
- Save intermediate data: Off

4.3.2 Recon 2

- Series description: PET Brain neurostat
- Recon range (bed): -1 to -1
- Output image type: Corrected
- Recon method: Iterative
- Iterations: 4
- Subsets: 8
- Image size: 128
- Zoom: 3
- Filter: Gaussian
- FWHM (mm): 8
- Offset X: 0 mm

• Offset Y: 0 mm

• Attenuation correction: On (1)

• Scatter correction: On

• Match CT slice location: On

• Save intermediate data: Off