20_PET_AC_Brain_LM_Roche.MlAdult_PET6

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1 Topogram

1.1 Routine

- mA: 35
- kV: 120
- Topogram length: 1024 mm
- Tube position: Top

1.2 Scan

- mA: 35
- kV: 120
- Delay: 4s
- Topogram length: 1024 mm
- Direction: Craniocaudal
- Tube position: Top
- API: None
- Kernel: 80s shar
- Window: Topogram Body

2 CT Brain

2.1 Routine

- Eff. mAs: 55
- kV: 120
- CARE Dose4D: On
- CareDoseType: CareDoseAEC
- CTDlvol: 3.71345mGy

• Scan time: 18.670 s

• Delay: 4.000 s

• Slice: 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

 \bullet Scan time: 18.670 s

• Rotation time: 0.500 s

• Delay: 4.000 s

• Slice: 5.00 mm

• Pitch: 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: Cerebrum

• Extended FoV: On

• FoV: 700

• Center X: 0

• Center Y: 0

• Mirroring: None

• Extended CT scale: Standard

• Recon job: Axial

• Recon Axis: Axial

• Image order: Craniocaudal

• Recon 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: Cerebrum

• Extended FoV: Off

• FoV: 500

• Center X: 0

• Center Y: 0

• Mirroring: None

• Extended CT scale: Standard

• Recon job: Axial

• Recon Axis: Axial

• Image order: Craniocaudal

• Recon increment: 3.000

• No. of images: 274

3 Pause

4 PET WB

4.1 Routine

- Isotope: F-18
- Pharm.: Florbetapir
- Inj. Dose: 1 MegaBequerels
- Scan mode: List mode
- Scan range: Match CT FOV
- No. of beds: 1
- Scan duration/bed: 15 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