$X_Cal_PET5_FDG.MlAdult_PET5$

30. maj 2013

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1 Topogram

1.1 Routine

mA: 35kV: 120

• Topogram length: 256 mm

• Tube position: Top

1.2 Scan

mA: 35kV: 120Delay: 4s

Topogram length: 256 mmDirection: Craniocaudal

• Tube position: Top

API: NoneKernel: 80s shar

• Window: Topogram Body

2 CT WB

2.1 Routine

• Eff. mAs: 225

• kV: 120

• CARE Dose4D: On

• CareDoseType: CareDoseAEC

CTDlvol: 15.1475mGyScan time: 4.000 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: 225

• Eff. mAs: 225

• kV: 120

Scan time: 4.000 sRotation time: 0.500 s

Delay: 4.000 sSlice: 3.00 mmPitch: 1.2000000

• Direction: Craniocaudal

2.3 Recons

2.3.1 Recon 1

• Series description: AC CT

• Slice: 3.00

• Kernel: B19f PET very smooth

Window: AbdomenExtended 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: 74

3 Pause

4 PET WB

4.1 Routine

Isotope: F-18 Pharm.: FDG

• Inj. Dose: 86 MegaBequerels

• Scan mode: Sinogram

• Scan range: Match CT FOV

• No. of beds: 1

• Scan duration/bed: 5 Minutes

4.2 Scan

• Autoload: On

Rebinner LUT: OffScan output: Sinogram

Sinogram mode: TruesInput trigger signal: None

LLD (keV): 435ULD (keV): 650

4.3 Recons

4.3.1 Recon 1

• Series description: PET WB Corr True X 2 mm

Recon range (bed): -1 to -1Output image type: Corrected

• Recon method: TrueX

Iterations: 3Subsets: 21Image size: 336

• Zoom: 1

Filter: GaussianFWHM (mm): 2Offset X: 0 mmOffset Y: 0 mm

• Attenuation correction: On (1)

• Scatter correction: On

Match CT slice location: OnSave intermediate data: Off