PETCT_AC_WholeBody_Ped.MlChild_PET5

27. februar 2013

Indhold

1 Topogram

1.1 Routine

- mA: 50
- kV: 80
- Topogram length: 768 mm
- Tube position: Top

1.2 Scan

- mA: 50
- kV: 80
- Delay: 4s
- Topogram length: 768 mm
- Direction: Craniocaudal
- Tube position: Top
- API: None
- Kernel: 80s shar
- Window: Topogram Body

2 CT WB

2.1 Routine

- Eff. mAs: 25
- kV: 120
- CARE Dose4D: Off
- CareDoseType: CareDoseAEC
- CTDlvol: 1.685mGy

- Scan time: 6.830 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: 50

- Eff. mAs: 25

- kV: 120

- Scan time: 6.830 s

- Rotation time: 0.500 s

- Delay: 4.000 s

- Slice: 5.00 mm

- Pitch: 1.5000000

- Direction: Craniocaudal

2.3   Recons

2.3.1   Recon 1

- Series description: tmp CT WB RTD

- Slice: 5.00

- Kernel: B19f PET very smooth

- Window: Abdomen

- 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: 21.600

- No. of images: 26

2.3.2   Recon 2

- Series description: AC CT

- Slice: 5.00

- Kernel: B19f PET very smooth

- Window: Abdomen

- 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: 174

3 Pause

4 PET WB

4.1 Routine

- Isotope: F-18

- Pharm.: FDG

- Inj. Dose: 2.5 Millicuries

- Scan mode: Sinogram

- Scan range: Match CT FOV

- No. of beds: 3

- Scan duration/bed: 2 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 WB

- Recon range (bed): -1 to -1

- Output image type: Corrected

- Recon method: Iterative

- Iterations: 4

- Subsets: 8

- Image size: 168

- Zoom: 1

- Filter: Gaussian

- FWHM (mm): 5

- 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 WB Uncorrected

- Recon range (bed): -1 to -1

- Output image type: Uncorrected

- Recon method: Iterative

- Iterations: 4

- Subsets: 8

- Image size: 168

- Zoom: 1

- Filter: Gaussian

- FWHM (mm): 5

- Offset X: 0 mm

- Offset Y: 0 mm

- Attenuation correction: On (1)

- Scatter correction: On

- Match CT slice location: On

- Save intermediate data: Off