$PET_CT_WB_U_25_kg.MlChild_PET6$

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1 Topogram

1.1 Routine

mA: 50kV: 80

 \bullet Topogram length: 0 mm

• Tube position: Top

1.2 Scan

mA: 50kV: 80Delay: 4s

Topogram length: 0 mmDirection: Craniocaudal

• Tube position: Top

API: NoneKernel: 80s shar

• Window: Topogram Body

2 CT WB

2.1 Routine

• Eff. mAs: 35

• kV: 120

• CARE Dose4D: Off

• CareDoseType: CareDoseAEC

CTDlvol: 2.359mGy
Scan time: 6.830 s
Delay: 20.000 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: 50

• Eff. mAs: 35

• kV: 120

Scan time: 6.830 sRotation time: 0.500 s

Delay: 20.000 sSlice: 3.00 mmPitch: 1.5000000

• 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: 3.000No. of images: 174

3 Pause

4 PET WB

4.1 Routine

Isotope: F-18 Pharm.: FDG

• Inj. Dose: 200 MegaBequerels

• Scan mode: Sinogram

• Scan range: Match CT FOV

• No. of beds: 3

• Scan duration/bed: 3 Minutes

4.2 Scan

• Autoload: On

Rebinner LUT: OffScan output: Sinogram

• Sinogram mode: Trues

• Input trigger signal: None

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

4.3 Recons

4.3.1 Recon 1

- Series description: PET Image check
- 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): 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

4.3.2 Recon 2

- Series description: PET WB True X 2 mm
- Recon range (bed): -1 to -1
- Output image type: Corrected
- Recon method: TrueX
- Iterations: 2
- Subsets: 21
- Image size: 336
- Zoom: 1
- Filter: Gaussian
- FWHM (mm): 2
- Offset X: 0 mm
- Offset Y: 0 mm
- Attenuation correction: On (1)
- Scatter correction: On
- Match CT slice location: On
- Save intermediate data: Off