$PETCT_AC_WholeBody_Ped.MlChild_PET5$

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1 Topogram

1.1 Routine

mA: 50kV: 80

• Topogram length: 768 mm

• Tube position: Top

1.2 Scan

mA: 50kV: 80Delay: 4s

Topogram length: 768 mmDirection: Craniocaudal

• Tube position: Top

API: NoneKernel: 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 sSlice: 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 sRotation time: 0.500 s

Delay: 4.000 sSlice: 5.00 mmPitch: 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: AbdomenExtended 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: AbdomenExtended 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: GaussianFWHM (mm): 5Offset X: 0 mmOffset Y: 0 mm

• Attenuation correction: On (1)

• Scatter correction: On

Match CT slice location: On Save intermediate data: Off