PET_AC_WB_FDG.MlChild_PET5

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1 Topogram

1.1 Routine

- mA: 50
- kV: 80
- Topogram length: 0 mm
- Tube position: Top

1.2 Scan

- mA: 50
- kV: 80
- Delay: 4s
- Topogram length: 0 mm
- Direction: Craniocaudal
- Tube position: Top
- API: None
- Kernel: 20s standar
- Window: Topogram Body

2 CT lavdosis

2.1 Routine

- Eff. mAs: 30
- kV: 80
- CARE Dose4D: Off
- CareDoseType: CareDoseAEC
- \bullet CTDlvol: 0.540268mGy

• Scan time: 11.000 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: 25

• Eff. mAs: 30

• kV: 80

 \bullet Scan time: 11.000 s

• Rotation time: 0.500 s

• Delay: 4.000 s

• Slice: 3.00 mm

• Pitch: 1.4000000

• Direction: Craniocaudal

2.3 Recons

2.3.1 Recon 1

• Series description: AC CT lavdosis

• Slice: 3.00

• Kernel: B19f PET very smooth

• Window: Baby 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: 274

3 Pause

4 PET WB

4.1 Routine

• Isotope: F-18

• Pharm.: FDG

• Inj. Dose: 1 MegaBequerels

• Scan mode: Sinogram

• Scan range: Match CT FOV

• No. of beds: 5

• 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): 425
- ULD (keV): 650

4.3 Recons

4.3.1 Recon 1

- Series description: PET WB corrected
- 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

- \bullet Series description: PET WB True X
- 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: Off
- Match CT slice location: On
- Save intermediate data: Off