PETCT_Brain_Ped.MlChild_PET5

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1 Topogram

1.1 Routine

mA: 50kV: 80

Topogram length: 256 mmTube position: Lateral

1.2 Scan

mA: 50kV: 80Delay: 4s

Topogram length: 256 mmDirection: CraniocaudalTube position: Lateral

• API: None

• Kernel: 20s standar

• Window: Topogram Head

2 CT Brain

2.1 Routine

• Eff. mAs: 170

• kV: 120

• CARE Dose4D: Off

• CareDoseType: CareDoseAEC

CTDlvol: 28.798mGyScan time: 24.750 s

Delay: 4.000 sSlice: 3.00 mm

• No. of images: Samme som i foerste recon, slet?(y/n)

• Tilt: 0.0 grader

2.2 Scan

• Quality ref. mAs: 380

• Eff. mAs: 170

• kV: 120

Scan time: 24.750 sRotation time: 1.000 s

Delay: 4.000 sSlice: 3.00 mmPitch: 0.8000000

• Direction: Caudocranial

2.3 Recons

2.3.1 Recon 1

- Series description: AC CT
- Slice: 3.00
- Kernel: H19s PET very smooth
- Window: CerebrumExtended FoV: Off
- FoV: 300
- Center X: 0
- Center Y: 0
- Mirroring: None
- Extended CT scale: Standard
- Recon job: Axial
- Recon Axis: Axial
- Image order: Caudocranial
- Recon increment: 1.500
- No. of images: 148

2.3.2 Recon 2

- Series description: CT Brain 3.0 C30s
- Slice: 3.00
- Kernel: C30s medium smooth
- Window: Baby Brain
- Extended FoV: Off
- FoV: 250
- Center X: 0
- Center Y: 0
- Mirroring: None
- Extended CT scale: Standard
- Recon job: Axial
- Recon Axis: Axial
- Image order: Caudocranial
- Recon increment: 1.500
- No. of images: 148

3 Pause

4 PET Brain

4.1 Routine

• Isotope: F-18

• Pharm.: FDG

• Inj. Dose: 2.5 Millicuries

• Scan mode: Sinogram

• Scan range: Match CT FOV

• No. of beds: 1

• Scan duration/bed: 10 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 Brain

• Recon range (bed): -1 to -1

• Output image type: Corrected

• Recon method: Iterative

• Iterations: 6

• Subsets: 16

• Image size: 336

• Zoom: 2

• 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