$Terapi_WB_FDG_PET_CT_4.Child_PET4$

27. februar 2013

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1 Topogram

1.1 Routine

- mA: 30
- kV: 120
- Topogram length: 1536 mm
- Tube position: Bottom

1.2 Scan

- mA: 30
- kV: 120
- Delay: 4s
- Topogram length: 1536 mm
- Direction: Craniocaudal
- Tube position: Bottom
- API: API
- Kernel: 80
- Window: Topogram

2 CT WB

2.1 Routine

- Eff. mAs: 225
- kV: 120
- CARE Dose4D: On
- CareDoseType: CareDoseAEC
- CTDlvol: 16.4093mGy

• Scan time: 11.011 s

• Delay: 6.000 s

• Slice: 2 mm

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

• Tilt: 0.0 grader

2.2 Scan

• Quality ref. mAs: 225

• Eff. mAs: 225

• kV: 120

 \bullet Scan time: 11.011 s

• Rotation time: 0.500 s

• Delay: 6.000 s

• Slice: 2 mm

• Pitch: 1.20

• Direction: Craniocaudal

2.3 Recons

2.3.1 Recon 1

• Series description: CT WB RTD

• Slice: 4

• Kernel: B19f

• Window: Abdomen

• Extended FoV: Off

• FoV: 500

• Center X: 0

• Center Y: 0

• Mirroring: None

• Extended CT scale: Standard

• Recon job: Axial

• Recon Axis: Axial

• Image order: Craniocaudal

• Recon increment: 4.000

• No. of images: 119

2.3.2 Recon 2

• Series description: AC CT WB

• Slice: 2

• Kernel: B19f

• Window: Abdomen

• Extended FoV: Off

• FoV: 500

• Center X: 0

• Center Y: 0

• Mirroring: None

• Extended CT scale: Standard

• Recon job: Axial

• Recon Axis: Axial

• Image order: Craniocaudal

• Recon increment: 2.000

• No. of images: 237

2.3.3 Recon 3

• Series description: CT WB 2.0 B30f

• Slice: 2

• Kernel: B30f

• Window: Abdomen

• Extended FoV: Off

• FoV: 500

• Center X: 0

• Center Y: 0

• Mirroring: None

• Extended CT scale: Standard

• Recon job: Axial

• Recon Axis: Axial

• Image order: Craniocaudal

• Recon increment: 2.000

• No. of images: 237

2.3.4 Recon 4

• Series description: CT WB 2.0 cor B30f

• Slice: 2

• Kernel: B30f

• Window: Abdomen

• Extended FoV: Off

• FoV: 500

• Center X: 0

• Center Y: 0

- Mirroring: None
- Extended CT scale: Standard
- Recon job: 3D
- Recon Axis: Coronal
- Image order: AP
- Recon increment: 2.000
- No. of images: 251

2.3.5 Recon 5

- Series description: CT WB 2.0 sag B30f
- Slice: 2
- Kernel: B30f
- Window: Abdomen
- Extended FoV: Off
- FoV: 500
- Center X: 0
- Center Y: 0
- Mirroring: None
- Extended CT scale: Standard
- Recon job: 3D
- Recon Axis: Sagittal
- ullet Image order: LeftToRight
- Recon increment: 2.000
- No. of images: 251

3 Pause

4 PET WB

4.1 Routine

- Isotope: F-18
- Pharm.: Fluorodeoxyglucose
- Inj. Dose: 0 Bequerels
- Scan mode: Sinogram
- Scan range: Do not match CT FOV
- No. of beds: Not given. Check recon range
- Scan duration/bed: 2.000000 Minutes

4.2 Scan

- Autoload: On
- Rebinner LUT: Off
- Scan output: Sinogram
- Sinogram mode: false
- Input trigger signal: None
- LLD (keV): n/a
- ULD (keV): n/a

4.3 Recons

4.3.1 Recon 1

- Series description: PET WB image check
- Recon range (bed): 1 to 3
- Output image type: Corrected

• Recon method: TrueX

• Iterations: 2

• Subsets: 12

• Image size: 200

• Zoom: 1

• Filter: Gaussian

• FWHM (mm): 3.000000

• Offset X: 0 mm

• Offset Y: 0 mm

• Attenuation correction: Off ()

• Scatter correction: On

• Match CT slice location: On

• Save intermediate data: Off

4.3.2 Recon 2

• Series description: PET WB Corr-TrueX + TOF 2 mm

• Recon range (bed): 1 to 3

• Output image type: Corrected

• Recon method: TrueXTOF

• Iterations: 2

• Subsets: 21

• Image size: 400

• Zoom: 1

• Filter: Gaussian

• FWHM (mm): 2.000000

• Offset X: 0 mm

• Offset Y: 0 mm

 \bullet Attenuation correction: Off ()

• Scatter correction: On

• Match CT slice location: On

• Save intermediate data: Off