# $PETCT\_AC\_WholeBody\_Ped.Child\_PET4$

11. januar 2013

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# 1 Topogram

# 1.1 Routine

- mA: 50
- kV: 80
- Topogram length: 768 mm
- Tube position: Top

#### 1.2 Scan

- mA: 50
- kV: 80
- Delay: 4s
- Topogram length: 768 mm
- Direction: Craniocaudal
- Tube position: Top
- API: API
- Kernel: 80
- Window: Topogram

# 2 CT WB

## 2.1 Routine

- Eff. mAs: 25
- kV: 80
- CARE Dose4D: Off
- CareDoseType: CareDoseAEC
- $\bullet$  CTDlvol: 0.5125mGy

• Scan time: 9.580 s

• Delay: 2.000 s

• Slice: 5 mm

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

• Tilt: 0.0 grader

## 2.2 Scan

• Quality ref. mAs: 65

• Eff. mAs: 25

• kV: 80

 $\bullet$  Scan time: 9.580 s

• Rotation time: 0.500 s

• Delay: 2.000 s

• Slice: 5 mm

• Pitch: 1.40

• Direction: Craniocaudal

## 2.3 Recons

#### 2.3.1 Recon 1

• Series description: tmp 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: 5

• Kernel: B19f

• Window: Abdomen

• Extended FoV: On

• FoV: 780

• 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: 158

# 3 Pause

# 4 PET WB

## 4.1 Routine

- Isotope: F-18
- Pharm.: FDG
- $\bullet\,$  Inj. Dose: 92500000 Bequerels
- $\bullet$  Inj. date (date/month year): 07/12 2008
- Inj. time: 02:00:33
- Scan mode: Sinogram
- Scan range:
- No. of beds: 0
- Scan duration/bed: 2.000000 Minutes

#### 4.2 Scan

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

#### 4.3 Recons

#### 4.3.1 Recon 1

- Series description: PET WB
- Recon range (bed): 1 to 3
- Output image type: Corrected
- Recon method: Iterative3D
- Iterations: 2
- Subsets: 24
- Image size: 200
- Zoom: 1
- Filter: Gaussian
- FWHM (mm): 5.000000
- 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 Uncorrected
- Recon range (bed): 1 to 3
- Output image type: Uncorrected
- Recon method: Iterative3D
- Iterations: 2
- Subsets: 24

• Image size: 200

• Zoom: 1

• Filter: Gaussian

• FWHM (mm): 5.000000

• Offset X: 0 mm

• Offset Y: 0 mm

• Attenuation correction: Off ()

• Scatter correction: On

• Match CT slice location: On

• Save intermediate data: Off