# $1\_PET\_CT\_WB\_FDG\_Barn.Child\_PET4$

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## 1 Topogram

## 1.1 Routine

• mA: 30

• kV: 120

• Topogram length: 768 mm

• Tube position: Bottom

#### 1.2 Scan

• mA: 30

• kV: 120

• Delay: 4s

• Topogram length: 768 mm

• Direction: Craniocaudal

• Tube position: Bottom

• API: API

• Kernel: 80

• Window: Topogram

## 2 CT WB

## 2.1 Routine

• Eff. mAs: 40

• kV: 120

• CARE Dose4D: On

• CareDoseType: CareDoseAEC

• CTDlvol: 2.912mGy

• Scan time: 11.070 s

• Delay: 6.000 s

• Slice: 2 mm

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

• Tilt: 0.0 grader

## 2.2 Scan

• Quality ref. mAs: 40

• Eff. mAs: 40

• kV: 120

 $\bullet$  Scan time: 11.070 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: 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: 2

• 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: 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.: FDG

• Inj. Dose: 100000000 Bequerels

• Scan mode: Sinogram

• Scan range: Do not match CT FOV

• 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 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): 3.000000

• Offset X: 0 mm

• Offset Y: 0 mm

 $\bullet$  Attenuation correction: Off ()

• Scatter correction: On

• Match CT slice location: On

• Save intermediate data: Off