# WB\_FDG\_PET\_CT\_IV\_4.Child\_PET4

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# 1 Topogram

# 1.1 Routine

mA: 30kV: 120

• Topogram length: 1536 mm

• Tube position: Top

# 1.2 Scan

mA: 30kV: 120Delay: 4s

Topogram length: 1536 mmDirection: Craniocaudal

• Tube position: Top

API: APIKernel: 80

• Window: Topogram

# 2 PreMonitoring

# 2.1 Routine

• Eff. mAs: 20

• kV: 120

• CARE Dose4D: Off

• CareDoseType: CareDoseAEC

• CTDlvol: 1.18627mGy

• Scan time: 0.500 s

• Delay: 2.000 s

• Slice: 10 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: 20

• kV: 120

Scan time: 0.500 sRotation time: 0.500 s

• Delay: 2.000 s

Slice: 10 mmPitch: 0.00

• Direction: Craniocaudal

# 2.3 Recons

#### 2.3.1 Recon 1

• Series description: PreMonitoring 10.0 eFoV

Slice: 10Kernel: B19s

Window: AbdomenExtended FoV: On

FoV: 780Center X: 0Center Y: 0Mirroring: None

• Extended CT scale: Standard

Recon job: AxialRecon Axis: Axial

Image order: CraniocaudalRecon increment: 10.000

• No. of images: 1

# 3 Monitoring

### 3.1 Routine

• Eff. mAs: 20

• kV: 120

• CARE Dose4D: Off

• CareDoseType: CareDoseAEC

CTDlvol: 35.5881mGyScan time: 0.500 s

Delay: 10.000 sSlice: 10 mm

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

• Tilt: 0.0 grader

# 3.2 Scan

• Quality ref. mAs: 65

Eff. mAs: 20kV: 120

Scan time: 0.500 sRotation time: 0.500 s

Delay: 10.000 sSlice: 10 mm

• Pitch: 0.00

• Direction: Craniocaudal

# 3.3 Recons

#### 3.3.1 Recon 1

• Series description: Monitoring 10.0 eFoV

Slice: 10Kernel: B19s

Window: AbdomenExtended FoV: On

FoV: 780Center X: 0Center Y: 0

• Mirroring: None

• Extended CT scale: Standard

Recon job: AxialRecon Axis: Axial

Image order: CraniocaudalRecon increment: 10.000

• No. of images: 30

# 4 CT WB

### 4.1 Routine

• Eff. mAs: 225

• kV: 120

• CARE Dose4D: On

• CareDoseType: CareDoseAEC

CTDlvol: 16.4093mGyScan time: 11.011 s

• Delay: 10.000 s

• Slice: 2 mm

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

• Tilt: 0.0 grader

# 4.2 Scan

• Quality ref. mAs: 225

• Eff. mAs: 225

• kV: 120

Scan time: 11.011 sRotation time: 0.500 s

• Delay: 10.000 s

Slice: 2 mmPitch: 1.20

• Direction: Craniocaudal

#### 4.3 Recons

#### 4.3.1 Recon 1

• Series description: CT WB RTD

• Slice: 4

• Kernel: B19f

Window: AbdomenExtended FoV: Off

FoV: 500Center X: 0Center Y: 0

• Mirroring: None

• Extended CT scale: Standard

Recon job: AxialRecon Axis: Axial

Image order: CraniocaudalRecon increment: 4.000

• No. of images: 119

#### 4.3.2 Recon 2

• Series description: AC CT WB

• Slice: 2

• Kernel: B19f

Window: AbdomenExtended FoV: Off

FoV: 500Center X: 0Center Y: 0Mirroring: None

• Extended CT scale: Standard

Recon job: AxialRecon Axis: Axial

• Image order: Craniocaudal

• Recon increment: 2.000

• No. of images: 237

#### 4.3.3 Recon 3

• Series description: CT WB 2.0 B30f

• Slice: 2

• Kernel: B30f

- Window: AbdomenExtended FoV: Off
- FoV: 500Center X: 0
- Center Y: 0
- Mirroring: None
- Extended CT scale: Standard
- Recon job: AxialRecon Axis: Axial
- Image order: Craniocaudal
- Recon increment: 2.000
- No. of images: 237

#### 4.3.4 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

#### 4.3.5 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

- Image order: LeftToRightRecon increment: 2.000
- No. of images: 251

# 5 Pause

# 6 PET WB

# 6.1 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

# 6.2 Scan

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

#### 6.3 Recons

#### 6.3.1 Recon 1

- Series description: PET WB image check
- Recon range (bed): 1 to 1
- 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

#### 6.3.2 Recon 2

- $\bullet$  Series description: PET WB Corr-TrueX + TOF 2 mm
- Recon range (bed): 1 to 1
- Output image type: Corrected
- Recon method: TrueXTOF
- $\bullet$  Iterations: 2
- Subsets: 21
- Image size: 400
- Zoom: 1
- Filter: Gaussian
- FWHM (mm): 2.000000
- $\bullet$  Offset X: 0 mm
- Offset Y: 0 mm
- Attenuation correction: Off ()
- Scatter correction: On
- Match CT slice location: On
- Save intermediate data: Off