# $TUMOR\_HUND\_FDG.MlAdult\_PET5$

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## 1 Topogram

## 1.1 Routine

- mA: 35
- kV: 120
- Topogram length: 0 mm
- Tube position: Top

#### 1.2 Scan

- mA: 35
- kV: 120
- Delay: 4s
- Topogram length: 0 mm
- Direction: Craniocaudal
- Tube position: Top
- API: None
- Kernel: 80s shar
- Window: Topogram Body

## 2 CT WB

## 2.1 Routine

- Eff. mAs: 170
- kV: 120
- CARE Dose4D: Off
- CareDoseType: CareDoseAEC
- CTDlvol: 11.4448mGy

• Scan time: 12.680 s

• Delay: 4.000 s

• Slice: 3.00 mm

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

• Tilt: 0.0 grader

## 2.2 Scan

• Quality ref. mAs: 225

• Eff. mAs: 170

• kV: 120

 $\bullet$  Scan time: 12.680 s

• Rotation time: 0.500 s

• Delay: 4.000 s

• Slice: 3.00 mm

• Pitch: 1.2000000

• Direction: Craniocaudal

## 2.3 Recons

#### 2.3.1 Recon 1

• Series description: AC CT

• Slice: 3.00

• Kernel: B19f PET very smooth

• Window: Abdomen

• Extended FoV: On

• FoV: 700

• 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: 274

#### 2.3.2 Recon 2

• Series description: CT WB 3.0 B30f

• Slice: 3.00

• Kernel: B30f medium smooth

• 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: 3.000

• No. of images: 274

## 3 Pause

## 4 PET TUMOR

## 4.1 Routine

• Isotope: F-18

• Pharm.: FDG

• Inj. Dose: 200 MegaBequerels

• 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 TUMOR Image check

• Recon range (bed): -1 to -1

• Output image type: Corrected

- Recon method: Iterative
- Iterations: 4
- Subsets: 8
- Image size: 168
- Zoom: 1
- Filter: Gaussian
- FWHM (mm): 4
- Offset X: 0 mm
- Offset Y: 0 mm
- Attenuation correction: On (1)
- Scatter correction: On
- Match CT slice location: On
- Save intermediate data: Off

#### 4.3.2 Recon 2

- Series description: PET TUMOR True X 2.0 mm
- Recon range (bed): -1 to -1
- Output image type: Corrected
- Recon method: TrueX
- Iterations: 3
- Subsets: 21
- Image size: 336
- Zoom: 1
- Filter: Gaussian
- FWHM (mm): 2
- Offset X: 0 mm

- Offset Y: 0 mm
- Attenuation correction: On (1)
- Scatter correction: Off
- Match CT slice location: On
- Save intermediate data: Off

## 5 Pause

## 6 PET WB

## 6.1 Routine

- Isotope: F-18
- Pharm.: FDG
- Inj. Dose: 200 MegaBequerels
- Scan mode: Sinogram
- Scan range: Match CT FOV
- No. of beds: 1
- Scan duration/bed: 2 Minutes

#### 6.2 Scan

- Autoload: On
- Rebinner LUT: Off
- Scan output: Sinogram
- Sinogram mode: Trues
- Input trigger signal: None
- LLD (keV): 435
- ULD (keV): 650

#### 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: Iterative
- Iterations: 4
- Subsets: 8
- Image size: 128
- Zoom: 1
- Filter: Gaussian
- FWHM (mm): 4
- Offset X: 0 mm
- Offset Y: 0 mm
- Attenuation correction: On (1)
- Scatter correction: On
- Match CT slice location: On
- Save intermediate data: Off

#### 6.3.2 Recon 2

- Series description: PET WB True X 2.0 mm
- Recon range (bed): -1 to -1
- Output image type: Corrected
- Recon method: TrueX
- Iterations: 3
- Subsets: 21

• Image size: 336

• Zoom: 1

• Filter: Gaussian

• FWHM (mm): 2

• Offset X: 0 mm

• Offset Y: 0 mm

• Attenuation correction: On (1)

• Scatter correction: Off

• Match CT slice location: On

• Save intermediate data: Off

## 7 Pause

## 8 DynMulti

## 8.1 Routine

• Eff. mAs: 170

• kV: 120

• CARE Dose4D: Off

• CareDoseType: CareDose

• CTDlvol: 1032.37mGy

• Scan time: 90.100 s

• Delay: 4.000 s

• Slice: 7.20 mm

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

• Tilt: 0.0 grader

## 8.2 Scan

• Quality ref. mAs: 100

• Eff. mAs: 170

• kV: 120

• Scan time: 90.100 s

 $\bullet$  Rotation time: 1.000 s

• Delay: 4.000 s

• Slice: 7.20 mm

• Pitch: 1.0000000

• Direction: Craniocaudal

## 8.3 Recons

#### 8.3.1 Recon 1

• Series description: DynMulti 7.2 B30s

• Slice: 7.20

• Kernel: B30s medium smooth

• Window: Abdomen

• Extended FoV: Off

• FoV: 0

• Center X: -3

• Center Y: 0

• Mirroring: None

• Extended CT scale: Standard

• Recon job: Axial

• Recon Axis: Axial

• Image order: Craniocaudal

• Recon increment: 1.000

• No. of images: 360