PROJ_DOTATATE_PET_CT_6.MlAdult_PET6

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Indhold

1	Topogram 1			
	1.1	Routine	1	
	1.2	Scan	1	
2	CT	$_{ m VB}$	1	
	2.1	Routine	1	
	2.2	Scan	1	
	2.3	Recons	2	
		2.3.1 Recon 1	2	
		2.3.2 Recon 2	2	
		2.3.3 Recon 3	2	
		2.3.4 Recon 4	3	
3	Pau	\mathbf{e}	3	
4	PE	WB	3	
	4.1	Routine	3	
	4.2	Scan	3	
	4.3	Recons	4	
		4.3.1 Recon 1	4	
		4.3.2 Recon 2	4	
		4.3.3 Recon 3	5	
		4 3 4 Recon 4	5	

1 Topogram

1.1 Routine

mA: 35kV: 120

• Topogram length: 0 mm

• Tube position: Top

1.2 Scan

mA: 35kV: 120Delay: 4s

Topogram length: 0 mmDirection: Craniocaudal

• Tube position: Top

API: NoneKernel: 80s shar

• Window: Topogram Body

2 CT WB

2.1 Routine

• Eff. mAs: 225

• kV: 120

• CARE Dose4D: On

• CareDoseType: CareDoseAEC

CTDlvol: 15.1475mGyScan time: 12.680 s

Delay: 4.000 sSlice: 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: 225

• kV: 120

Scan time: 12.680 sRotation time: 0.500 s

Delay: 4.000 sSlice: 3.00 mmPitch: 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: AbdomenExtended FoV: On
- FoV: 700Center X: 0
- Center Y: 0
- Centrer 1.0
- 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: AbdomenExtended 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

2.3.3 Recon 3

- Series description: CT WB 3.0 SPO cor
- Slice: 3.00
- Kernel: B30f medium smooth
- Window: Abdomen
- Extended FoV: Off

- FoV: 821
- Center X: 0
- Center Y: 0
- Mirroring: None
- Extended CT scale: Standard
- Recon job: 3D
- Recon Axis: Coronal
- Image order: AP
- Recon increment: 3.000
- No. of images: 167

2.3.4 Recon 4

- Series description: CT WB 3.0 SPO sag
- Slice: 3.00
- Kernel: B30f medium smooth
- Window: Abdomen
- Extended FoV: Off
- FoV: 821
- Center X: 0
- Center Y: 0
- Mirroring: None
- Extended CT scale: Standard
- Recon job: 3D
- Recon Axis: Sagittal
- Image order: LeftToRight
- Recon increment: 3.000
- No. of images: 274

3 Pause

4 PET WB

4.1 Routine

- Isotope: Cu-64
- Pharm.: DOTATATE
- Inj. Dose: 1 MegaBequerels
- Scan mode: Sinogram
- Scan range: Match CT FOV
- No. of beds: 5
- Scan duration/bed: 180 Seconds

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 WB Image check
- Recon range (bed): -1 to -1
- Output image type: Corrected
- Recon method: Iterative3D
- 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 WB DOTATATE True X 2 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: On
- Match CT slice location: On

• Save intermediate data: Off

4.3.3 Recon 3

- Series description: PET WB UDEN True X
- Recon range (bed): -1 to -1
- Output image type: Corrected
- Recon method: Iterative
- Iterations: 3
- Subsets: 21Image size: 336
- 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

4.3.4 Recon 4

- Series description: no AC
- Recon range (bed): -1 to -1
- Output image type: Uncorrected
- Recon method: Iterative
- Iterations: 3
- Subsets: 21
- Image size: 336
- 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