

4_PETCT_AC_WholeBody_Test.Adult

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1 Topogram

1.1 Routine

- mA: 35
- kV: 120
- Topogram length: 1024 mm
- Tube position: Top

1.2 Scan

- mA: 35
- kV: 120
- Delay: 4s
- Topogram length: 1024 mm
- Direction: Craniocaudal
- Tube position: Top
- API: API
- Kernel: 80
- Window: Topogram

2 ReadOnly=1 ResID=921 AC CT Body

2.1 Routine

- Eff. mAs: 225
- kV: 120
- CARE Dose4D: On
- CareDoseType: CareDoseAEC
- CTDIvol: 16.38mGy

- Scan time: 7.290 s
- Delay: 60.000 s
- Slice: 3 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: 7.290 s
- Rotation time: 0.500 s
- Delay: 60.000 s
- Slice: 3 mm
- Pitch: 1.40
- Direction: Craniocaudal

2.3 Recons

2.3.1 Recon 1

- Series description: tmp AC CT Body RTD
- Slice: 4
- Kernel: B19f
- Window: Abdomen
- Extended FoV: false
- 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: 88

2.3.2 Recon 2

- Series description: AC CT
- Slice: 3
- Kernel: B19f
- Window: Abdomen
- Extended FoV: true
- 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: 117

2.3.3 Recon 3

- Series description: CT WB 3.0 B30f
- Slice: 3
- Kernel: B30f
- Window: Abdomen
- Extended FoV: false
- 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: 117

2.3.4 Recon 4

- Series description: CT WB 3.0 cor B30f
- Slice: 3
- Kernel: B30f
- Window: Abdomen
- Extended FoV: false
- FoV: 524
- 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: 76

2.3.5 Recon 5

- Series description: CT WB 3.0 sag B30f
- Slice: 3
- Kernel: B30f
- Window: Abdomen
- Extended FoV: false
- FoV: 500
- 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: 76

3 Pause

4 PET WB

4.1 Routine

- Isotope: F-18
- Pharm.: Fluorodeoxyglucose
- Inj. Dose: 370000000.000000 Bequerels
- Inj. date (date/month - year): 13/09 - 2010
- Inj. time: 15:09:14
- Scan mode: Sinogram
- Scan range: Match CT FOV
- No. of beds: Not given. Check recon range
- Scan duration/bed: 3.000000 Minutes

4.2 Scan

- Autoload: true
- Rebinner LUT: N/A
- Scan output: Sinogram
- Sinogram mode: false
- Input trigger signal: None
- LLD (keV): N/A
- ULD (keV): N/A

4.3 Recons

4.3.1 Recon 1

- Series description: PET WB Corrected HD
- Recon range (bed): 1 to 2
- Output image type: Corrected
- Recon method: TrueX
- Iterations: 2
- Subsets: 12
- Image size: 200
- Zoom: 1
- Filter: Gaussian
- FWHM (mm): 4.000000
- Offset X: 0 mm
- Offset Y: 0 mm
- Attenuation correction: On (1)
- Scatter correction: true
- Match CT slice location: true
- Save intermediate data: false

4.3.2 Recon 2

- Series description: PET WB Corr TrueX+TOF
- Recon range (bed): 1 to 2
- Output image type: Corrected
- Recon method: TrueXTOF
- Iterations: 3
- Subsets: 21

- Image size: 400
- Zoom: 1
- Filter: Gaussian
- FWHM (mm): 2.000000
- Offset X: 0 mm
- Offset Y: 0 mm
- Attenuation correction: On (1)
- Scatter correction: true
- Match CT slice location: true
- Save intermediate data: false