

WB_FDG_PET_CT_IV_5.MlAdult

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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 (-1 betyder None, ellers skal tallet oversaettes (stoerre arbejde): -1
- Kernel: T80s sharp
- Window: Topogram Body

2 CT WB

2.1 Routine

- Eff. mAs: 225
- kV: 14
- CARE Dose4D: On
- CTDIvol: 15.1475mGy
- Scan time: 1.2000000 s

- Delay: 60.000 s
- Slice: 3.00 mm
- No. of images: Samme som i foerste recon, slet?(y/n)
- Tilt: 225 grader

2.2 Scan

- Quality ref. mAs: 225
- Eff. mAs: 225
- kV: 120
- Scan time: 14.840 s
- Rotation time: 0.500 s
- Delay: 60.000 s
- Slice: 0.6 mm
- Pitch: 1.2000000
- Direction: Craniocaudal

2.3 Recon

2.3.1 Recon 1

- Series description: AC CT
- Slice: 3.00
- Kernel: "B19f PET very smooth"
- Window: "Abdomen"
- Extended FoV(1/0): 1
- FoV: 700
- Center X: 0
- Center Y: 0

- Mirroring: None
- Extended CT scale: Standard
- Recon job: AxialRJ
- Recon Axis: Axial
- Image order: Craniocaudal
- Recon increment: 3.000
- No. of images: 324

2.3.2 Recon 2

- Series description: CT WB 3.0 B30f
- Slice: 3.00
- Kernel: "B30f medium smooth"
- Window: "Abdomen"
- Extended FoV(1/0): 0
- FoV: 500
- Center X: 0
- Center Y: 0
- Mirroring: None
- Extended CT scale: Standard
- Recon job: AxialRJ
- Recon Axis: Axial
- Image order: Craniocaudal
- Recon increment: 3.000
- No. of images: 324

2.3.3 Recon 3

- Series description: CT WB 3.0 SPO cor
- Slice: 3.00
- Kernel: "B30f medium smooth"
- Window: "Abdomen"
- Extended FoV(1/0): 0
- FoV: 971
- Center X: 0
- Center Y: 0
- Mirroring: None
- Extended CT scale: Standard
- Recon job: VolumeRJ
- 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(1/0): 0
- FoV: 971
- Center X: 0
- Center Y: 0

- Mirroring: None
- Extended CT scale: Standard
- Recon job: VolumeRJ
- Recon Axis: Sagittal
- Image order: LeftToRight
- Recon increment: 3.000
- No. of images: 324

3 Pause

4 PET WB

4.1 Routine

- Isotope: F-18
- Pharm.: FDG
- Inj. Dose: 1(P gaBequerels
- Inj. date (date/month - year): 08/08 - 2012
- Inj. time: 14:17:00
- Scan mode: Sinogram
- Scan range: Match CT FOV
- No. of beds: 6
- Scan duration/bed: 3(P nutes

4.2 Scan

- Autoload: On
- Rebinner LUT: 0
- Scan output: Sinogram

- Sinogram mode: Trues
- Input trigger signal: None
- LLD (keV): 435
- ULD (keV): 650

4.3 Recon

4.3.1 Recon 1

- Series description: ET WB Image check
- Recon range (bed): ups..
- Output image type: Corrected
- Recon method: Iterative
- Iterations: 4
- Subsets: 8
- Image size: 168
- Zoom: 1
- Filter: Gaussian
- FWHM (mm): 4
- Offset X (mm): 0
- Offset Y (mm): 0
- Attenuation correction: (1,1)
- Scatter correction: 1
- Match CT slice location: 1
- Save intermediate data: 0

4.3.2 Recon 2

- Series description: ET WB Corr True X 2 m
- Recon range (bed): ups..
- Output image type: Corrected
- Recon method: TrueX
- Iterations: 3
- Subsets: 21
- Image size: 336
- Zoom: 1
- Filter: Gaussian
- FWHM (mm): 2
- Offset X (mm): 0
- Offset Y (mm): 0
- Attenuation correction: (1,1)
- Scatter correction: 1
- Match CT slice location: 1
- Save intermediate data: 0