

HUND_64Cu_Liposom_Dag1.MlAdult

27. september 2012

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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: T80s sharp
- Window: Topogram Body

2 CT WB

2.1 Routine

- Eff. mAs: 170
- kV: 12
- CARE Dose4D: Off
- CTDIvol: 11.4448mGy
- Scan time: 1.2000000 s

- Delay: 4.000 s
- Slice: 3.00 mm
- No. of images: Samme som i foerste recon, slet?(y/n)
- Tilt: 170 grader

2.2 Scan

- Quality ref. mAs: 225
- Eff. mAs: 170
- kV: 120
- Scan time: 12.680 s
- Rotation time: 0.500 s
- Delay: 4.000 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: 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 3.0 B30f medium smooth
- 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 WB

4.1 Routine

- Isotope: Cu-64
- Pharm.: liposome
- Inj. Dose: 1 MegaBequerels
- Inj. date (date/month - year): 08/02 - 2012
- Inj. time: 16:03:00
- Scan mode: List mode
- Scan range: Match CT FOV
- No. of beds: 1
- Scan duration/bed: 40 PtMinute

4.2 Scan

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

4.3 Recon

4.3.1 Recon 1

- Series description: PET WB Image check
- Recon range (bed): ups..
- Output image type: NoRecon
- 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