HUND_64Cu_Liposom_Dag2.MlAdult_PET5

3. december 2012

Indhold

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

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

- 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 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 HEART

4.1 Routine

- Isotope: Cu-64

- Pharm.: liposome

- Inj. Dose: 1 MegaBequerels

- Inj. date (date/month - year): 08/02 - 2012

- Inj. time: 16:04:00

- Scan mode: Sinogram

- Scan range: Match CT FOV

- No. of beds: 1

- Scan duration/bed: 3 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 HEART True X Day2

- Recon range (bed): Automatic/All/dunno

- 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

5 Pause

6 PET LIVER

6.1 Routine

- Isotope: Cu-64

- Pharm.: liposome

- Inj. Dose: 1 MegaBequerels

- Inj. date (date/month - year): 08/02 - 2012

- Inj. time: 16:04:00

- Scan mode: Sinogram

- Scan range: Match CT FOV

- No. of beds: 1

- Scan duration/bed: 3 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 LIVER True X Day2

- Recon range (bed): Automatic/All/dunno

- 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

7   Pause

8   PET TUMOR

8.1   Routine

- Isotope: Cu-64

- Pharm.: liposome

- Inj. Dose: 1 MegaBequerels

- Inj. date (date/month - year): 08/02 - 2012

- Inj. time: 16:04:00

- Scan mode: Sinogram

- Scan range: Match CT FOV

- No. of beds: 1

- Scan duration/bed: 10 Minutes

8.2 Scan

- Autoload: On

- Rebinner LUT: Off

- Scan output: Sinogram

- Sinogram mode: Trues

- Input trigger signal: None

- LLD (keV): 435

- ULD (keV): 650

8.3 Recons

8.3.1 Recon 1

- Series description: PET Tumor True X Day2

- Recon range (bed): Automatic/All/dunno

- 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

9 Pause

10 PET WB

10.1 Routine

- Isotope: Cu-64
- Pharm.: liposome
- Inj. Dose: 1 MegaBequerels
- Inj. date (date/month - year): 08/02 - 2012
- Inj. time: 16:04:00
- Scan mode: Sinogram
- Scan range: Match CT FOV
- No. of beds: 1
- Scan duration/bed: 2 Minutes

10.2 Scan

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

10.3 Recons

10.3.1 Recon 1

- Series description: PET WB Image Check Day2

- Recon range (bed): Automatic/All/dunno

- Output image type: Corrected

- Recon method: Iterative

- Iterations: 4

- Subsets: 8

- Image size: 128

- 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

10.3.2 Recon 2

- Series description: PET WB True X Day 2

- Recon range (bed): Automatic/All/dunno

- 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