$TUMOR_HUND_FLT.MlAdult_PET5$

30. maj 2013

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1 Topogram

1.1 Routine

mA: 35kV: 120

 \bullet 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 TUMOR

2.1 Routine

• Eff. mAs: 170

• kV: 120

• CARE Dose4D: Off

 \bullet CareDoseAEC

CTDlvol: 11.4448mGyScan time: 4.000 s

Delay: 4.000 sSlice: 3.00 mm

• No. of images: Samme som i foerste recon, slet?(y/n)

 \bullet Tilt: 0.0 grader

2.2 Scan

• Quality ref. mAs: 170

• Eff. mAs: 170

• kV: 120

Scan time: 4.000 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: 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: 74

2.3.2 Recon 2

- Series description: CT TUMOR 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: 74

3 Pause

4 PET Statisk

4.1 Routine

• Isotope: F-18

- Pharm.: FLT
- Inj. Dose: 199 MegaBequerels
- Scan mode: Sinogram
- Scan range: Match CT FOV
- No. of beds: 1
- Scan duration/bed: 10 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 TUMOR Image Check
- Recon range (bed): -1 to -1
- Output image type: Corrected
- Recon method: Iterative
- Iterations: 4
- Subsets: 8
- Image size: 256
- Zoom: 1
- Filter: Gaussian
- FWHM (mm): 3
- 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 TUMOR True X
- 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: Off
- Match CT slice location: On
- Save intermediate data: Off

5 Pause

6 DynMulti

6.1 Routine

- Eff. mAs: 170
- kV: 120
- CARE Dose4D: Off
- CareDoseType: CareDose
- CTDlvol: 1032.37mGy
- Scan time: 90.100 s
- Delay: 4.000 s
- Slice: 7.20 mm
- No. of images: Samme som i foerste recon, slet?(y/n)
- Tilt: 0.0 grader

6.2 Scan

- Quality ref. mAs: 100
- Eff. mAs: 170
- kV: 120
- Scan time: 90.100 s
- Rotation time: 1.000 s
- Delay: 4.000 s
- Slice: 7.20 mm
- Pitch: 1.0000000
- Direction: Craniocaudal

6.3 Recons

6.3.1 Recon 1

- Series description: DynMulti 7.2 B30s
- Slice: 7.20

• Kernel: B30s medium smooth

Window: AbdomenExtended FoV: Off

• FoV: 0

Center X: -3Center Y: 0

• Mirroring: None

• Extended CT scale: Standard

Recon job: AxialRecon Axis: Axial

Image order: CraniocaudalRecon increment: 1.000No. of images: 360