$TUMOR_HYPOXI_HUND_DAG_4.MlAdult$

27. september 2012

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1 Topogram

1.1 Routine

- mA: 35
- kV: 120
- Topogram length: 512 mm
- Tube position: Top

1.2 Scan

- mA: 35
- kV: 120
- Delay: 4s
- Topogram length: 512 mm
- Direction: Craniocaudal
- Tube position: Top
- API: None
- Kernel: T80s sharp
- Window: Topogram Body

2 CT WB

2.1 Routine

- Eff. mAs: 170
- kV: 4
- CARE Dose4D: Off
- CTDlvol: 11.4448mGy
- \bullet 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: 170

• Eff. mAs: 170

• kV: 120

• Scan time: 4.000 s

 \bullet 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

 $\bullet\,$ 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: 74

2.3.2 Recon 2

- Series description: CT TUMOR 3.0 B30f
- 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: 74

3 Pause

4 PET WB

4.1 Routine

- Isotope: Cu-64
- Pharm.:
- Inj. Dose: 0 MegaBequerels
- $\bullet\,$ Inj. date (date/month year): 22/04 2008
- Inj. time: 12:03:00
- Scan mode: Sinogram
- Scan range: Match CT FOV
- No. of beds: 1
- Scan duration/bed: 20 PtMinute

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 Recon

4.3.1 Recon 1

- Series description: PET TUMOR Corrected 180 min
- Recon range (bed): ups..
- 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 Uncorrected 180 min
- Recon range (bed): ups..
- Output image type: Uncorrected
- 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: Off

• Match CT slice location: On

• Save intermediate data: Off