$WB_Flourid_PET_5.MlAdult$

1. oktober 2012

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1 Topogram

1.1 Routine

- mA: 35
- kV: 120
- Topogram length: 34 mm
- Tube position: Top

1.2 Scan

- mA: 35
- kV: 120
- Delay: 4s
- Topogram length: 34 mm
- Direction: Caudocranial
- Tube position: Top
- API: None
- Kernel: T80s sharp
- Window: Topogram Body

2 CT WB

2.1 Routine

- Eff. mAs: 40
- kV: 38
- CARE Dose4D: Off
- CTDlvol: 2.70069mGy
- \bullet Scan time: 0.8000000 s

• Delay: 4.000 s

• Slice: 3.00 mm

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

• Tilt: 40 grader

2.2 Scan

• Quality ref. mAs: 50

• Eff. mAs: 40

• kV: 120

• Scan time: 38.240 s

• Rotation time: 0.500 s

• Delay: 4.000 s

• Slice: 0.6 mm

• Pitch: 0.8000000

• Direction: Caudocranial

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: 574

3 Pause

4 PET WB

4.1 Routine

• Isotope: F-18

• Pharm.: Fluoride

• Inj. Dose: 200 MegaBequerels

• Inj. date (date/month - year): 21/08 - 2012

• Inj. time: 14:00:00

• Scan mode: Sinogram

• Scan range: Match CT FOV

• No. of beds: 11

• Scan duration/bed: 120 PtSecond

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 knogleskanning 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: 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 knogleskanning True X 2 mm
- 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: 0 mm
- Offset Y: 0 mm
- Attenuation correction: On (1)
- Scatter correction: On
- Match CT slice location: On
- Save intermediate data: Off

4.3.3 Recon 3

- Series description: no AC
- Recon range (bed): ups..
- Output image type: Uncorrected
- Recon method: Iterative
- Iterations: 3
- Subsets: 21
- Image size: 336

- 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

4.3.4 Recon 4

- Series description: PET WB UDEN Trie X
- Recon range (bed): ups..
- Output image type: Corrected
- Recon method: Iterative
- Iterations: 3
- Subsets: 21
- Image size: 336
- 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