$PETCT_GantryOffset.Adult$

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1 Topogram

1.1 Routine

- mA: 36
- kV: 120
- Topogram length: 512 mm
- Tube position: Top

1.2 Scan

- mA: 36
- kV: 120
- Delay: 4s
- Topogram length: 512 mm
- Direction: Craniocaudal
- Tube position: Top
- API: API
- Kernel: 80
- Window: Topogram

2 CT

2.1 Routine

- Eff. mAs: 50
- kV: 120
- CARE Dose4D: Off
- CareDoseType: CareDoseAEC
- CTDlvol: 3.64mGy

• Scan time: 7.980 s

• Delay: 2.000 s

• Slice: 2 mm

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

• Tilt: 0.0 grader

2.2 Scan

• Quality ref. mAs: 120

• Eff. mAs: 50

• kV: 120

 \bullet Scan time: 7.980 s

• Rotation time: 0.500 s

• Delay: 2.000 s

• Slice: 2 mm

• Pitch: 0.80

• Direction: Craniocaudal

2.3 Recons

2.3.1 Recon 1

• Series description: tmp CT RTD

• Slice: 4

• Kernel: B19f

• Window: Abdomen

• Extended FoV: false

• 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: 4.000

• No. of images: 56

2.3.2 Recon 2

• Series description: CT 2.0 B19f

• Slice: 2

• Kernel: B19f

• Window: Abdomen

• Extended FoV: false

• 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: 2.000

• No. of images: 111

3 Pause

4 PET Offset

4.1 Routine

- Isotope: Ge-68
- Pharm.: Crossed Rods
- Inj. Dose: 59200000.000000 Bequerels
- Inj. date (date/month year): 24/06 2008
- Inj. time: 14:12:43
- Scan mode: Sinogram
- Scan range: Do not match CT FOV
- No. of beds: Not given. Check recon range
- Scan duration/bed: 10000000.000000 Counts

4.2 Scan

- Autoload: true
- Rebinner LUT: N/A
- Scan output: Sinogram
- Sinogram mode: false
- Input trigger signal: None
- LLD (keV): N/A
- \bullet ULD (keV): N/A

4.3 Recons

4.3.1 Recon 1

- Series description: PET Gantry Offset
- Recon range (bed): Automatic/All/dunno
- Output image type: Uncorrected
- Recon method: Iterative3D
- Iterations: 4
- Subsets: 8
- Image size: 400
- Zoom: 1
- Filter: Gaussian
- FWHM (mm): 5
- Offset X: 0 mm
- Offset Y: 0 mm
- Attenuation correction: On (1)
- Scatter correction: true
- Match CT slice location: false
- Save intermediate data: false