

PETCT_GantryOffset.MlAdult_PET5

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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: 80s shar
- Window: Topogram Body

2 CT

2.1 Routine

- Eff. mAs: 50
- kV: 120
- CARE Dose4D: Off
- CareDoseType: CareDoseAEC
- CTDIvol: 3.37586mGy
- Scan time: 4.390 s
- Delay: 4.000 s
- Slice: 5.00 mm
- No. of images: Samme som i foerste recon, slet?(y/n)
- Tilt: 0.0 grader

2.2 Scan

- Quality ref. mAs: 50
- Eff. mAs: 50
- kV: 120
- Scan time: 4.390 s
- Rotation time: 0.500 s
- Delay: 4.000 s
- Slice: 5.00 mm
- Pitch: 0.8000000

- Direction: Craniocaudal

2.3 Recons

2.3.1 Recon 1

- Series description: CT 5.0
- Slice: 5.00
- 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: 55

3 Pause

4 PET

4.1 Routine

- Isotope: Ge-68
- Pharm.: Crossed Rods
- Inj. Dose: 1.67 Millicuries
- Scan mode: Sinogram
- Scan range: Do not match CT FOV
- No. of beds: 1
- Scan duration/bed: 1e+007 Counts

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 Gantry Offset
- Recon range (bed): -1 to -1
- Output image type: Uncorrected
- Recon method: Iterative
- Iterations: 2
- Subsets: 8
- Image size: 256
- Zoom: 1
- Filter: Gaussian
- FWHM (mm): 2.5
- Offset X: 0 mm
- Offset Y: 0 mm
- Attenuation correction: On (1)
- Scatter correction: Off
- Match CT slice location: Off
- Save intermediate data: Off