

PETCT_NEMA_2007_IQ_MOD.Adult_PET4

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1 Topogram

1.1 Routine

- mA: 35
- kV: 120
- Topogram length: 1024 mm
- Tube position: Top

1.2 Scan

- mA: 35
- kV: 120
- Delay: 4s
- Topogram length: 1024 mm
- Direction: Craniocaudal
- Tube position: Top
- API: API
- Kernel: 80
- Window: Topogram

2 AC CT NEMA

2.1 Routine

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

- Scan time: 8.050 s
- Delay: 2.000 s
- Slice: 5 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
- Scan time: 8.050 s
- Rotation time: 0.500 s
- Delay: 2.000 s
- Slice: 5 mm
- Pitch: 0.80
- Direction: Craniocaudal

2.3 Recons

2.3.1 Recon 1

- Series description: tmp AC CT NEMA RTD
- Slice: 4
- Kernel: B19f
- 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: 4.000
- No. of images: 56

2.3.2 Recon 2

- Series description: AC CT NEMA
- Slice: 5
- Kernel: B19f
- Window: Abdomen
- Extended FoV: On
- FoV: 780
- 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: 75

3 Pause

4 PET NEMA IQ

4.1 Routine

- Isotope: F-18
- Pharm.: FDG
- Inj. Dose: 370000000 Bequerels
- Scan mode: Sinogram
- Scan range: Do not match CT FOV
- No. of beds: 0
- Scan duration/bed: 540.000000 Seconds

4.2 Scan

- Autoload: On
- Rebinner LUT: Off
- Scan output: Sinogram
- Sinogram mode: PromptsRandoms
- Input trigger signal: None
- LLD (keV): n/a
- ULD (keV): n/a

4.3 Recons

4.3.1 Recon 1

- Series description: PET NEMA IQ
- Recon range (bed): 1 to 1
- Output image type: Corrected

- Recon method: Iterative3D
- Iterations: 3
- Subsets: 24
- Image size: 200
- Zoom: 1
- Filter: Gaussian
- FWHM (mm): 5
- Offset X: 0 mm
- Offset Y: 0 mm
- Attenuation correction: Off ()
- Scatter correction: On
- Match CT slice location: Off
- Save intermediate data: Off