$PETCT_AC_WholeBody_XL.MlAdult_PET5$

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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: None

• Kernel: 80s shar

• Window: Topogram Body

2 CT WB

2.1 Routine

• Eff. mAs: 50

• kV: 120

• CARE Dose4D: Off

• CareDoseType: CareDoseAEC

• CTDlvol: 3.37586mGy

• Scan time: 18.670 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: 18.670 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: tmp CT WB RTD

• 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: 11.500

• No. of images: 72

2.3.2 Recon 2

• Series description: AC CT

• 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: 274

3 Pause

4 PET WB

4.1 Routine

• Isotope: F-18

• Pharm.: FDG

• Inj. Dose: 10 Millicuries

• Scan mode: Sinogram

• Scan range: Match CT FOV

• No. of beds: 5

• Scan duration/bed: 2 Minutes

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 WB

• Recon range (bed): -1 to -1

• Output image type: Corrected

- Recon method: Iterative
- Iterations: 2
- Subsets: 8
- Image size: 168
- Zoom: 1
- Filter: Gaussian
- FWHM (mm): 5
- 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 WB Uncorrected
- Recon range (bed): -1 to -1
- Output image type: Uncorrected
- Recon method: Iterative
- Iterations: 2
- Subsets: 8
- Image size: 168
- Zoom: 1
- Filter: Gaussian
- FWHM (mm): 5
- Offset X: 0 mm

• Offset Y: 0 mm

• Attenuation correction: On (1)

• Scatter correction: Off

• Match CT slice location: On

• Save intermediate data: Off