$PETCT_WholeBody_XXL.Adult_PET4$

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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 CT WB

2.1 Routine

- Eff. mAs: 210
- kV: 120
- CARE Dose4D: On
- CareDoseType: CareDoseAEC
- \bullet CTDlvol: 15.288mGy

• Scan time: 24.310 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: 210

• Eff. mAs: 210

• kV: 120

 \bullet Scan time: 24.310 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 CT WB 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: 182

2.3.2 Recon 2

• Series description: AC CT WB

• 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: 242

2.3.3 Recon 3

• Series description: CT WB 3.0 B30f

• Slice: 3

• Kernel: B30f

• 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: 2.000

• No. of images: 363

2.3.4 Recon 4

• Series description: CT LUNG 3.0 B70f

• Slice: 3

• Kernel: B70f

• Window: Lung

• Extended FoV: Off

• FoV: 300

• 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: 363

3 Pause

4 PET WB

4.1 Routine

• Isotope: F-18

• Pharm.: Fluorodeoxyglucose

 $\bullet\,$ Inj. Dose: 370000000 Bequerels

• Scan mode: Sinogram

 $\bullet\,$ Scan range: Do not match CT FOV

 $\bullet\,$ No. of beds: Not given. Check recon range

• Scan duration/bed: 3.000000 Minutes

4.2 Scan

• Autoload: On

• Rebinner LUT: Off

• Scan output: Sinogram

• Sinogram mode: false

• Input trigger signal: None

- LLD (keV): n/a
- ULD (keV): n/a

4.3 Recons

4.3.1 Recon 1

- Series description: PET WB
- Recon range (bed): 1 to 5
- Output image type: Corrected
- Recon method: Iterative3D
- Iterations: 2
- Subsets: 24
- Image size: 200
- Zoom: 1
- Filter: Gaussian
- FWHM (mm): 6.000000
- Offset X: 0 mm
- Offset Y: 0 mm
- Attenuation correction: Off ()
- 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 5
- Output image type: Uncorrected
- Recon method: Iterative3D
- Iterations: 2
- Subsets: 24
- Image size: 200
- Zoom: 1
- Filter: Gaussian
- FWHM (mm): 6.000000
- Offset X: 0 mm
- Offset Y: 0 mm
- Attenuation correction: Off ()
- Scatter correction: On
- Match CT slice location: On
- Save intermediate data: Off