$test_WB_FDG_PET_CT_IV_Lunge_6.MlAdult_PET6$

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1 Topogram

1.1 Routine

- mA: 35
- kV: 120
- Topogram length: 0 mm
- Tube position: Top

1.2 Scan

- mA: 35
- kV: 120
- Delay: 5s
- Topogram length: 0 mm
- Direction: Craniocaudal
- Tube position: Top
- API: API 31
- Kernel: 80s shar
- Window: Topogram Body

2 CT BANAN

2.1 Routine

- Eff. mAs: 100
- kV: 120
- CARE Dose4D: On
- CareDoseType: CareDoseAEC
- CTDlvol: 7.6543mGy

• Scan time: 7.840 s

• Delay: 8.000 s

• Slice: 5.00 mm

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

• Tilt: 0.0 grader

2.2 Scan

• Quality ref. mAs: 100

• Eff. mAs: 100

• kV: 120

 \bullet Scan time: 7.840 s

• Rotation time: 0.500 s

 \bullet Delay: 8.000 s

• Slice: 5.00 mm

• Pitch: 1.4000000

• Direction: Caudocranial

2.3 Recons

2.3.1 Recon 1

• Series description: CT BANAN B18f very smooth

• Slice: 5.00

• Kernel: B18f very smooth

• Window: Mediastinum

• Extended FoV: Off

• FoV: 315

• 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: 127

2.3.2 Recon 2

• Series description: CT BANAN 2.0 B70f

• Slice: 2.00

• Kernel: B70f very sharp

• Window: Lung

• Extended FoV: Off

• FoV: 315

• Center X: 0

• Center Y: 0

• Mirroring: None

• Extended CT scale: Standard

• Recon job: Axial

• Recon Axis: Axial

• Image order: Craniocaudal

• Recon increment: 1.500

• No. of images: 253

3 Pause

4 CT WB

4.1 Routine

• Eff. mAs: 225

• kV: 120

• CARE Dose4D: On

• CareDoseType: CareDoseAEC

• CTDlvol: 15.1475mGy

• Scan time: 12.680 s

• Delay: 60.000 s

• Slice: 3.00 mm

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

• Tilt: 0.0 grader

4.2 Scan

• Quality ref. mAs: 225

• Eff. mAs: 225

• kV: 120

• Scan time: 12.680 s

 \bullet Rotation time: 0.500 s

• Delay: 60.000 s

• Slice: 3.00 mm

• Pitch: 1.2000000

• Direction: Craniocaudal

4.3 Recons

4.3.1 Recon 1

- Series description: AC CT
- Slice: 3.00
- Kernel: B18f 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

4.3.2 Recon 2

- Series description: CT WB 3.0 B30f
- Slice: 3.00
- Kernel: B30f medium smooth
- 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: 3.000

• No. of images: 274

4.3.3 Recon 3

• Series description: CT WB 3.0 COR

• Slice: 3.00

• Kernel: B30f medium smooth

• Window: Abdomen

• Extended FoV: Off

• FoV: 821

• Center X: 0

• Center Y: 0

• Mirroring: None

• Extended CT scale: Standard

• Recon job: 3D

• Recon Axis: Coronal

• Image order: AP

• Recon increment: 3.000

• No. of images: 167

4.3.4 Recon 4

• Series description: CT WB 3.0 SAG

• Slice: 3.00

• Kernel: B30f medium smooth

• Window: Abdomen

• Extended FoV: Off

• FoV: 821

• Center X: 0

• Center Y: 0

• Mirroring: None

• Extended CT scale: Standard

• Recon job: 3D

• Recon Axis: Sagittal

• Image order: LeftToRight

• Recon increment: 3.000

• No. of images: 274

5 Pause

6 PET WB

6.1 Routine

• Isotope: F-18

• Pharm.: FDG

• Inj. Dose: 1 MegaBequerels

• Scan mode: Sinogram

• Scan range: Match CT FOV

- No. of beds: 1
- Scan duration/bed: 150 Seconds

6.2 Scan

- Autoload: On
- Rebinner LUT: Off
- Scan output: Sinogram
- Sinogram mode: Trues
- Input trigger signal: None
- LLD (keV): 435
- ULD (keV): 650

6.3 Recons

6.3.1 Recon 1

- Series description: PET WB Image check
- Recon range (bed): -1 to -1
- Output image type: Corrected
- Recon method: Iterative3D
- Iterations: 4
- Subsets: 8
- Image size: 168
- Zoom: 1
- Filter: Gaussian
- FWHM (mm): 4
- Offset X: 0 mm
- Offset Y: 0 mm

- Attenuation correction: Off (1)
- Scatter correction: On
- Match CT slice location: On
- Save intermediate data: Off

6.3.2 Recon 2

- Series description: PET WB True X 2 mm
- Recon range (bed): -1 to -1
- Output image type: Corrected
- Recon method: TrueX
- Iterations: 3
- Subsets: 21
- Image size: 336
- Zoom: 1
- Filter: Gaussian
- FWHM (mm): 2
- Offset X: 0 mm
- Offset Y: 0 mm
- Attenuation correction: Off (1)
- Scatter correction: On
- Match CT slice location: On
- Save intermediate data: Off