TUMOR_HYPOXI_HUND_DAG_3.MlAdult_PET5

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1 **Topogram**

Routine 1.1

• 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 TUMOR

2.1 Routine

• Eff. mAs: 170

• kV: 120

• CARE Dose4D: Off

• CareDoseType: CareDoseAEC

• CTDlvol: 11.4448mGy

• Scan time: 4.000 s

• Delay: 4.000 s

• Slice: 3.00 mm

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

• Tilt: 0.0 grader

2.2 Scan

• Quality ref. mAs: 170

• Eff. mAs: 170

• kV: 120

• Scan time: 4.000 s

• Rotation time: 0.500 s

• Delay: 4.000 s • Slice: 3.00 mm • Pitch: 1.2000000 • Direction: Craniocaudal

2.3 Recons

2.3.1 Recon 1

- Series description: AC CT
- Slice: 3.00
- Kernel: B19f PET very smooth
- Window: AbdomenExtended 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: 74

2.3.2 Recon 2

- Series description: CT TUMOR 3.0 B30f
- Slice: 3.00
- Kernel: B30f medium smooth
- Window: AbdomenExtended 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: 74

3 Pause

4 PET STATIC

4.1 Routine

• Isotope: Cu-64

• Pharm.: ATSM

• Inj. Dose: 1e-006 MegaBequerels

• Scan mode: Sinogram

• Scan range: Match CT FOV

• No. of beds: 1

• Scan duration/bed: 20 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 TUMOR Corrected 24 hr

• Recon range (bed): -1 to -1

• Output image type: Corrected

• Recon method: Iterative

• Iterations: 4

• Subsets: 8

• Image size: 256

• Zoom: 1

• Filter: Gaussian

• FWHM (mm): 3

• 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 TUMOR Uncorrected 24 hr

• Recon range (bed): -1 to -1

• Output image type: Uncorrected

• Recon method: Iterative

• Iterations: 4

• Subsets: 8

- Image size: 256
- Zoom: 1
- Filter: Gaussian
- FWHM (mm): 3
- Offset X: 0 mm
- Offset Y: 0 mm
- Attenuation correction: On (1)
- Scatter correction: Off
- Match CT slice location: On
- Save intermediate data: Off

5 Pause

6 DynMulti

6.1 Routine

- Eff. mAs: 170
- kV: 120
- CARE Dose4D: Off
- CareDoseType: CareDose
- CTDlvol: 1032.37mGy
- Scan time: 90.100 s
- Delay: 4.000 s
- Slice: 7.20 mm
- No. of images: Samme som i foerste recon, slet?(y/n)
- Tilt: 0.0 grader

6.2 Scan

- Quality ref. mAs: 100
- Eff. mAs: 170
- kV: 120
- Scan time: 90.100 s
- Rotation time: 1.000 s
- Delay: 4.000 s
- Slice: 7.20 mm
- Pitch: 1.0000000
- Direction: Craniocaudal

6.3 Recons

6.3.1 Recon 1

- Series description: DynMulti 7.2 B30s
- Slice: 7.20

• Kernel: B30s medium smooth

Window: AbdomenExtended FoV: Off

• FoV: 0

Center X: -3Center Y: 0

• Mirroring: None

• Extended CT scale: Standard

Recon job: AxialRecon Axis: Axial

Image order: CraniocaudalRecon increment: 1.000No. of images: 360