# TUMOR\_HYPOXI\_HUND\_LM.MlAdult\_PET5

30. maj 2013

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## 1 Topogram

#### 1.1 Routine

mA: 35kV: 120

Topogram length: 256 mmTube position: Lateral

#### 1.2 Scan

mA: 35kV: 120Delay: 4s

Topogram length: 256 mmDirection: CraniocaudalTube position: Lateral

• API: None

• Kernel: 20s standar

• Window: Topogram Head

### 2 CT WB

#### 2.1 Routine

• Eff. mAs: 40

• kV: 120

• CARE Dose4D: Off

• CareDoseType: CareDoseAEC

CTDlvol: 5.504mGy
Scan time: 9.310 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: 380

• Eff. mAs: 40

• kV: 120

Scan time: 9.310 sRotation time: 1.000 s

Delay: 4.000 sSlice: 3.00 mmPitch: 1.0000000

• Direction: Craniocaudal

#### 2.3 Recons

#### 2.3.1 Recon 1

• Series description: AC CT

• Slice: 3.00

• Kernel: H19s PET very smooth

Window: CerebrumExtended FoV: Off

FoV: 300Center X: 0Center Y: 0Mirroring: None

• Extended CT scale: Standard

Recon job: AxialRecon Axis: Axial

Image order: CraniocaudalRecon increment: 3.000

• No. of images: 74

## 3 Pause

## 4 PET LM

#### 4.1 Routine

Isotope: Cu-64Pharm.: ATSM

• Inj. Dose: 700 MegaBequerels

• Scan mode: List mode

• Scan range: Match CT FOV

• No. of beds: 1

• Scan duration/bed: 60 Minutes

#### 4.2 Scan

• Autoload: On

• Rebinner LUT: Off

• Scan output: List mode

• Sinogram mode: Trues

• Input trigger signal: None

LLD (keV): 435ULD (keV): 650

### 4.3 Recons

#### 4.3.1 Recon 1

• Series description: PET Brain LM

Recon range (bed): -1 to -1Output image type: NoRecon

• Recon method: Iterative

Iterations: 6 Subsets: 16 Image size: 256

• Zoom: 2

Filter: GaussianFWHM (mm): 2Offset X: 0 mmOffset Y: 0 mm

• Attenuation correction: On (1)

• Scatter correction: On

Match CT slice location: On Save intermediate data: Off