# $PETCT\_GantryOffset.MlAdult\_PET5$

7. marts 2013

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## 1 Topogram

#### 1.1 Routine

mA: 35kV: 120

• Topogram length: 512 mm

• Tube position: Top

#### 1.2 Scan

mA: 35kV: 120Delay: 4s

Topogram length: 512 mmDirection: Craniocaudal

• Tube position: Top

API: NoneKernel: 80s shar

• Window: Topogram Body

## 2 CT

#### 2.1 Routine

• Eff. mAs: 50

• kV: 120

• CARE Dose4D: Off

• CareDoseType: CareDoseAEC

CTDlvol: 3.37586mGyScan time: 4.390 s

Delay: 4.000 sSlice: 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: 4.390 sRotation time: 0.500 s

Delay: 4.000 sSlice: 5.00 mmPitch: 0.8000000

• Direction: Craniocaudal

#### 2.3 Recons

#### 2.3.1 Recon 1

• Series description: CT 5.0

• Slice: 5.00

• Kernel: B19f PET very smooth

Window: AbdomenExtended FoV: On

FoV: 700Center X: 0Center Y: 0Mirroring: None

• Extended CT scale: Standard

Recon job: AxialRecon Axis: Axial

Image order: CraniocaudalRecon increment: 3.000

• No. of images: 55

## 3 Pause

## 4 PET

#### 4.1 Routine

• Isotope: Ge-68

Pharm.: Crossed RodsInj. Dose: 1.67 MillicuriesScan mode: Sinogram

• Scan range: Do not match CT FOV

• No. of beds: 1

• Scan duration/bed: 1e+007 Counts

#### 4.2 Scan

• Autoload: On

Rebinner LUT: OffScan output: SinogramSinogram mode: Trues

• Input trigger signal: None

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

## 4.3 Recons

#### 4.3.1 Recon 1

• Series description: PET Gantry Offset

• Recon range (bed): -1 to -1

• Output image type: Uncorrected

• Recon method: Iterative

Iterations: 2 Subsets: 8

• Image size: 256

• Zoom: 1

Filter: GaussianFWHM (mm): 2.5Offset X: 0 mmOffset Y: 0 mm

• Attenuation correction: On (1)

• Scatter correction: Off

Match CT slice location: OffSave intermediate data: Off