# $PET\_QualityCheck.MlAdult\_PET6$

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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: 3.37586mGy

 $\bullet$  Scan time: 5.630 s

Delay: 4.000 sSlice: 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: 50

• Eff. mAs: 50

• kV: 120

Scan time: 5.630 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: AC CT CT 5.0
- Slice: 5.00
- 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

## 3 Pause

## 4 PET QC

#### 4.1 Routine

- Isotope: Ge-68
- Pharm.: Solid Solution
- Inj. Dose: 93.15 MegaBequerels
- Scan mode: Sinogram
- Scan range: Do not match CT FOV
- No. of beds: 1
- Scan duration/bed: 2e+008 Counts

#### 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: Norm Review

• Recon range (bed): -1 to -1

• Output image type: Corrected

• Recon method: Iterative

Iterations: 2 Subsets: 8

• Image size: 128

• Zoom: 1

Filter: GaussianFWHM (mm): 2.5

• Offset X: 0 mm

• Offset Y: 0 mm

• Attenuation correction: On (1)

• Scatter correction: On

 $\bullet\,$  Match CT slice location: Off

• Save intermediate data: Off