# $PETCT\_QualityCheck.MlAdult\_PET5$

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# Indhold

1	Topogram 1			
	1.1	Routine		
	1.2	Scan		
2	CT	1		
	2.1	Routine		
	2.2	Scan		
	2.3	Recons		
		2.3.1 Recon 1		
3	Pau	${ m se}$		
4	PET	$\Gamma$ QC		
	4.1	Routine		
	4.2	Scan		
	4.3	Recons		
		4.3.1 Recon 1		

# 1 Topogram

#### 1.1 Routine

- 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

#### 2.1 Routine

- Eff. mAs: 50
- kV: 120
- CARE Dose4D: Off
- CareDoseType: CareDoseAEC
- CTDlvol: 3.37586mGy

• Scan time: 5.630 s

• Delay: 4.000 s

• Slice: 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: 5.630 s

• Rotation time: 0.500 s

• Delay: 4.000 s

• Slice: 5.00 mm

• Pitch: 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: 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: 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: Gaussian
- FWHM (mm): 2.5
- Offset X: 0 mm
- Offset Y: 0 mm
- Attenuation correction: On (1)
- Scatter correction: On
- Match CT slice location: Off
- Save intermediate data: Off